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Badania *in silico* aktywności antyoksydacyjnej związków fitochemicznych

In silico studies of the antioxidant activity of the phytochemical compounds

Rozprawa doktorska w oparciu o monotematyczny cykl publikacji w dziedzinie nauk medycznych i nauk o zdrowiu w dyscyplinie nauki farmaceutyczne

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STRESZCZENIE

Zmiany oksydacyjne w obrębie biocząsteczek organizmu stanowią istotny czynnik indukujący zaburzenia homeostazy organizmu mogący objawiać się w obrazie klinicznym pacjenta rozwojem szeregu poważnych chorób, takich jak miażdżyca, cukrzyca, Alzheimer, Parkinson czy nowotwory. Fizjologiczne systemy redoks przeciwdziałają temu zjawisku, jednak utrzymanie równowagi między pro– i anty– oksydantami jest w obecnych czasach utrudnione biorąc pod uwagę stale rosnącą ekspozycję na utleniacze. Istotnym wydaje się suplementacja organizmu w składniki zdolne zmiatać wolne rodniki, chelatować jony metali oraz oddziaływać z odpowiednimi enzymami, aby przeciwdziałać narastającemu stresowi oksydacyjnemu.

Większość warzyw, owoców oraz produktów pochodzenia roślinnego zawiera w składzie substancje fenolowe. Liczne badania *in vitro* oraz *in vivo* wykazały prozdrowotny efekt tej klasy związków fitochemicznych wynikający z ich potencjału antyoksydacyjnego. Aktywność, jaką przeciwutleniacz wykazuje w organizmie jest jednak wypadkową wielu czynników które modulują reaktywność czy właściwości fizykochemiczne, pośród których największe znaczenie ma struktura chemiczna.

Metody chemii obliczeniowej stanowią aktualnie nieoceniony sposób badań, jeśli chodzi o analizę zjawisk niekoniecznie mierzalnych eksperymentalnie. W ramach doktoratu skupiono się na zastosowaniu mechaniki kwantowej do badań nad wybranymi związkami z grup fenolokwasów oraz flawonoidów. Dokonano analizy wpływu struktury na aktywność antyoksydacyjną wskazując na istotę zjawiska rezonansu, wiązań wodorowych oraz topologii grup hydroksylowych. Ustalono też indeksy reaktywności dające wstępny wgląd w spodziewaną aktywnością związków. Na przykładzie apigeniny z powodzeniem zastosowano protokół badań umożliwiający projekcję wyników teoretycznych na wymierne wartości, tzn. kinetykę reakcji zmiatania wolnych rodników oraz kompleksowania jonów metali. Dokonano także głębokiego przeglądu literaturowego dotyczącego aktualnie stosowanych technik w badaniach obliczeniowych nad antyoksydantami.

SUMMARY

Oxidative changes within the biomolecules are well-recognized inducers of homeostasis' disorders that can manifest in the patient's clinical profile with the development of several serious diseases such as atherosclerosis, diabetes, Alzheimer's, Parkinson's and cancer. Physiological redox systems counteract this phenomenon but maintaining a balance between pro- and anti-oxidants is difficult these days, given the ever-increasing exposure to oxidants. Hence, supplementing the body with ingredients capable of scavenging free radicals, chelating metal ions and interacting with appropriate enzymes to counteract the present oxidative stress seems vital.

Most vegetables, fruits and products of plant origin contain phenolic substances. Numerous *in vitro* and *in vivo* studies have demonstrated the health–promoting effect of this class of phytochemicals stemming from their antioxidant potential. However, the activity that an antioxidant exhibits in the body is the outcome of many factors that modulate reactivity or physicochemical properties, among which chemical structure is the most important.

Computational chemistry methods are currently an invaluable when it comes to analyzing phenomena that are not necessarily measurable experimentally. This doctoral thesis focused on the application of quantum mechanics to the study of selected compounds from the phenolic acid and flavonoid groups. The effect of structure on the antioxidant activity was analyzed, pointing out the essence of the resonance, the hydrogen bonds, and the topology of hydroxyl groups. Reactivity indices were also determined, giving a preliminary view of the expected activity of the substances investigated. Using apigenin as an example, a research protocol was successfully applied to project theoretical results into quantifiable values, i.e., the kinetics of free radical scavenging and metal ion complexation reactions. An in–depth literature review of currently used techniques in computational studies of antioxidants was also conducted.

WYKAZ PRAC NAUKOWYCH WŁĄCZONYCH DO CYKLU PUBLIKACJI

[A] Spiegel, M.*, Andruniów, T., & Sroka, Z. (2020). Flavones' and Flavonols' Antiradical Structure– Activity Relationship–A Quantum Chemical Study. *Antioxidants (Basel, Switzerland)*, 9(6), 461. https://doi.org/10.3390/antiox9060461

> *IF* = 6.313 *MEiN* = 100

[B] Spiegel, M., Kapusta, K.*, Kołodziejczyk, W., Saloni, J., Żbikowska, B., Hill, G. A., & Sroka, Z. (2020). Antioxidant Activity of Selected Phenolic Acids–Ferric Reducing Antioxidant Power Assay and QSAR Analysis of the Structural Features. *Molecules (Basel, Switzerland)*, 25(13), 3088. https://doi.org/10.3390/molecules25133088

IF = 4.412 *MEiN* = 140

[C] Spiegel, M.*, & Sroka, Z. (2023). Quantum-mechanical characteristics of apigenin: Antiradical, metal chelation and inhibitory properties in physiologically relevant media. *Fitoterapia*, 164(October 2022), 105352. https://doi.org/10.1016/j.fitote.2022.105352

> *IF* = 3.204 *MEiN* =100

[D] Spiegel M.* (2022). Current Trends in Computational Quantum Chemistry Studies on Antioxidant Radical Scavenging Activity. *Journal of Chemical Information and Modeling*, 62(11), 2639–2658. https://doi.org/10.1021/acs.jcim.2c00104

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1. Wprowadzenie

Oddychanie tlenowe jest procesem zarówno efektywniejszym, jak również bardziej skomplikowanym niż jego beztlenowy odpowiednik. Chociaż glikoliza stanowi wspólny początek na drodze pozyskiwania energii w postaci adenozyno–5'–trifosforanu (ATP), to zwiększony zysk energetyczny u organizmów aerobowych oparty jest przede wszystkim na etapach następujących po niej. Zbiorczo określone jako cykl Krebsa, generują one zredukowany dinukleotyd nikotynoamidoadeninowy (NADH) i zredukowany dinukleotyd flawinoadeninowy (FADH₂), które w dalszej kolejności ulegają utlenieniu przez enzymy mitochondrialnego łańcucha oddechowego. Utworzony w ten sposób gradient elektrochemiczny, zależny od różnicy stężeń jonów wodorowych między przestrzenią międzybłonową a macierzą, jest wykorzystywany przez syntazę ATP podczas fosforylacji oksydacyjnej. Transport elektronów, zachodzący z udziałem kompleksów I, III i IV oraz ubichinonu, nie jest jednak całkowicie wydajny. W wyniku niekontrolowanego przepływu pewna ich ilość wchodzi w nieenzymatyczne reakcje jednoelektronowe z pobliskimi cząstkami, prowadząc do powstania produktów ubocznych metabolizmu oksydacyjnego, wśród których najczęściej wymienia się rodniki.^[1–3]

Termin rodnik po raz pierwszy został użyty ponad 300 lat temu przez de Morveau w jego pracy z 1782 roku zatytułowanej "*Sur les dénominations chimiques, la nécessité d'en perfectionner le système et les règles pour y parvenir, suivi d'un tableau d'une nomenclature chimique*". Oryginalne znaczenie różniło się natomiast od dzisiejszej definicji sformułowanej przez IUPAC jako "*[...] indywiduum chemiczne [...] posiadające niesparowany elektron.* ".^[4] Ewolucja koncepcji i rozwój teorii rodnikowej bazował na pracach takich naukowców jak Lavoisiera, Guy–Lussaca, Wöhlera, von Liebiga czy Dumasa; Gomberga oraz Schlenka w obrębie rodników organicznych; a także Michaelisa nad rodnikami generowanymi w procesach biochemicznych. Obecnie uznaje się, że rodniki to liczna klasa indywiduów chemicznych o otwartopowłokowej konfiguracji z jednym, dwoma, trzema^[5] lub nawet czterema^[6] niesparowanymi elektronami. W rezultacie rodniki są zazwyczaj (ale nie zawsze^[7]) bardzo aktywne i spontanicznie reagują z sąsiednimi molekułami w celu stabilizacji swojej struktury elektronowej, co w konsekwencji często skutkuje w zmienionych właściwościach chemicznych oraz fizycznych atakowanej cząstki.

Należy podkreślić, że grupa istotnych biologicznie czynników utleniających obejmuje nie tylko rodniki, ale oprócz tego struktury nierodnikowe, takie jak reaktywne formy tlenu (RFT, *ang. reactive oxygen species, ROS*), azotu (RFA, *ang. reactive nitrogen species, RNS*), siarki (RFS, ang. *reactive sulphur species, RSS*), karbonylu (RFK, *ang. reactive carbonyl species, RCS*) i inne.^[8–10] Ich klasyfikacja opiera się na lokalizacji gęstości spinowej, a każda zdolna jest uszkadzać struktury komórkowe.

Łańcuch oddechowy jest głównym źródłem małych rodników tlenowych, a anionorodnik ponadtlenkowy (O₂⁻⁻) powstaje na skutek niekontrolowanej ucieczki elektronu na tlen atmosferyczny.^[1,11,12] Chociaż O₂⁻⁻, ze względu na ujemny ładunek oraz strukturę elektronową niechętnie przyjmuje dodatkowe elektrony, a tym samym wykazuje minimalną aktywność oksydacyjną wobec biomolekuł^[13,14], to reaguje on z innymi rodnikami i może ulec dysmutacji do nadtlenku wodoru (H₂O₂) w obecności dysmutazy ponadtlenkowej^[15] — H₂O₂ poza tym, że sam jest silnym utleniaczem, jest w stanie także reagować z jonami metali przejściowych, takimi jak żelazo, miedź oraz cynk, w przebiegu reakcji Fentona co prowadzi do jego redukcji do rodnika hydroksylowego ('OH). Ponadto O₂⁻⁻ potrafi również łączyć się z cząsteczkami tlenku azotu (NO'), tworząc peroksynitryty, które powodują nitrowanie składników organicznych^[9]

Pomimo że mitochondria produkują reaktywne formy tlenu w sposób nieprzerwany, nie są one jedynym generatorem endogennych oksydantów. Powszechnie wiadomo, że mikrosomy wątroby posiadają enzymatyczny układ oksydazy NADPH, który katalizując utlenianie NADPH przez tlen wytwarza nadtlenek wodoru.^[9] Do innych *"fabryk"* RFT można zaliczyć peroksysomy, cechujące się wysokim potencjałem oksydacyjnym powiązanym z występowaniem oksydazy zdolnej do redukcji tlenu do nadtlenku wodoru, jak też enzymy cytozolowe — przykładowo oksydaza ksantynowa wskutek konwersji hipoksantyny do ksantyny i kwasu moczowego z użyciem NADP⁺ lub O₂ jako akceptorów elektronów.^[16] Ostatecznie biologiczne tiole^[17], hydrochinony, katecholaminy^[18] i flawiny^[19] w podobny sposób mogą ulegać reakcjom redoks, przyczyniając się do akumulacji wewnątrzkomórkowych utleniaczy. Warto

wspomnieć, że produkcja rodników ma miejsce w przebiegu większości szlaków metabolicznych, aczkolwiek nie w każdym przypadku jest to zjawisko niepożądane.

Kiedy rozpoczęto badania biomedyczne w dziedzinie fizjologii procesów redoks, początkowo uważano, że utleniacze wywołują wyłącznie efekty toksyczne i są bez wyjątku związane ze zmianami patologicznymi.^[20] Jednak komórki wytwarzają oraz uwalniają 'OH, NO' i O²⁻ w trakcie normalnych czynności ustroju, a natura tych struktur jest dwojaka. Część z nich odgrywa niezbędną rolę w utrzymaniu homeostazy, partycypując w szeregu istotnych oraz fundamentalnych procesów fizjologicznych. *Ad exemplum*, fagocytarna oksydaza NADPH była pierwszym poznanym systemem, który generuje RFT nie w charakterze produktu ubocznego, ale podstawowej funkcji swojego działania.^[21] NO' został jednoznacznie zidentyfikowany jako endogenny czynnik sygnalizujący regulujący relaksację śródbłonka naczyń.^[22] RFT są też znaczące w podtrzymywaniu stanu zapalnego podczas odpowiedzi immunologicznej, ze szczególnym uwzględnieniem wybuchu tlenowego będącego metodą walki komórek układu odpornościowego z patogenami.^[23] Z tego powodu przypuszcza się, że niewystarczające stężenie utleniaczy może wręcz osłabić organizm, zapobiegając tzw. pozytywnemu stresowi oksydacyjnemu.^[24,25]

Przyczynę wzmożonej produkcji oksydantów *in vivo* trzeba odnajdywać nie tylko we wpływie już przedstawionych wcześniej czynników wewnętrznych, lecz i zewnętrznych.^[25] Do tych należą m.in. wysokotłuszczowa i wysokocukrowa dieta, nadużywanie leków, palenie tytoniu, spożywanie alkoholu oraz ekspozycja na substancje chemiczne, których metabolizm wiąże się z podwyższoną aktywnością enzymów. Stymulujący efekt ma ponadto zanieczyszczenie powietrza oraz promieniowanie UV, które, choć nie powiązane bezpośrednio, wpływają także na siebie wzajemnie. Modelowo, 'OH, NO' i Cl', pochodzące głównie z odpadów, uczestniczą w reakcjach łańcuchowych niszczenia warstwy ozonowej, zwiększając ilość docierającego do Ziemi ultrafioletu.^[26]

Tym niemniej, niezależnie od źródła, niekontrolowane wytwarzanie oksydantów nie jest korzystne, a ich nagromadzenie w ilościach znacznie przekraczających wartości fizjologiczne jest uznanym inicjatorem uszkodzeń wewnątrzkomórkowych. Energia, którą dysponują z racji niesparowanego elektronu na powłoce walencyjnej lub zajmowanego stanu wzbudzonego, jest przekazywana na biocząsteczki, takie jak lipidy, cukry, DNA, białka i aminokwasy, przekształcając ich struktury na skutek peroksydacji, chociażby w efekcie rozerwania szkieletów węglowych.^[10,27] Utlenianie lipidów zaburza ciągłość błony komórkowej i zmieniając ciśnienie osmotyczne w komórce inicjuje proces apoptozy; przemiany składników lipidowych oraz białkowych wchodzących w skład lipoprotein o niskiej gęstości powodują magazynowanie się cholesterolu^[28]; z kolei alteracje DNA, w których najbardziej rozpoznawalną jest transformacja guaniny w 8–okso–7,8–dihydro–guaninę, pociąga za sobą podniesione ryzyko aktywacji onkogenów i wystąpienia zmian nowotworowych, ze względu na spotęgowaną sposobność transwersji G do T.^[29] Wszystko to może skutkować w nieodwracalnych modyfikacjach strukturalnych, a często też innym profilu działania atakowanej cząstki, prowadzących do patologii w funkcjonowaniu komórek jak również tkanek. Niedostatecznie szybko zahamowane, przedłożone wyżej uszkodzenia ulegają nawarstwieniu i z czasem manifestują swoją obecność w obrazie klinicznym pacjenta.

Przy nieustannym narażeniu na oksydanty i stopniowym wyczerpywaniu się biologicznych reduktorów, mechanizmy obronne organizmu przestają być wystarczające. Długotrwałe stężenie utleniaczy, wykraczające poza ramy fizjologiczne, w skrajnych przypadkach daje początek zaburzeniom homeostazy oraz stresu oksydacyjnego, który ma negatywny wpływ na zdrowie, a jego pojawieniu się towarzyszy rozwój poważnych chorób cywilizacyjnych. Prowadzi on do sukcesywnej degeneracji komórek i narządów.^[30] Badania wykazały, że utrzymujący się wysoki poziom wolnych rodników wiąże się z rozwojem przewlekłego stanu zapalnego, który jest związany z ostrymi oraz chronicznymi przypadłościami, takimi jak choroby stawów^[31], dróg żółciowych^[32], cukrzyca^[33], choroby autoimmunologiczne (np. toczeń^[34], stwardnienie rozsiane^[35], choroba Hashimoto^[36]), czy nowotwory^[37]. Mózg jest wyjątkowo wrażliwy, ponieważ jego komórki potrzebują stałego dopływu tlenu do produkcji neuroprzekaźników, które wpływaja na neuroplastyczność. Nadmiar oksydantów uszkadza neurony dopaminergiczne i sprzyja rozwojowi choroby Parkinsona^[38], a także daje o sobie znać w postaci odkładania blaszek amyloidowych — kluczowych markerów choroby Alzheimera.^[39] Dodatkowo utleniacze odgrywają rolę w patofizjologii miażdżycy^[40]. Finalnie stres oksydacyjny może uruchamiać błędne koło, w którym odpowiedź zapalna amplifikuje generowanie rodników i zaostrza istniejący już stan zapalny.^[41]

Organizm, aby utrzymać docelowy poziom oksydantów, wyposażony jest w zintegrowany system obronny, który potrafi regulować ich stężenie. Przynależą do niego nieenzymatyczne przeciwutleniacze — wśród nich wskazać wypada, między innymi, glutation, kwas moczowy oraz kwas liponowy; nadto, jest on wspierany przez egzogenne reduktory, przykładowo karotenoidy (witamina A), kwas askorbinowy (witamina C) i tokoferole (witamina E).^[42] Współdziałają one ze antyoksydantami enzymatycznymi^[43], w szczególności dysmutazą ponadtlenkową, katalazą i peroksydazą glutationową, wspólnie hamując akumulację rodników lub przerywając łańcuch procesów oksydacyjnych, który już się rozpoczął. Ponieważ te antyoksydanty mają odmienne właściwości fizykochemiczne i odpowiednio charakter hydrofilowy, są one rozmieszczone w różnych kompartmentach organizmu zapewniając wszechstronną ochronę. Jednakże w obecnych czasach wydaje się, że bazowanie wyłącznie na zdolnościach ustroju w tym zakresie nie jest wystarczające.

2. Założenia i cel pracy

Swiadomość wpływowej roli stresu oksydacyjnego w etiologii chorób oraz procesu starzenia się nadała priorytet badaniom nad skutecznymi i nietoksycznymi związkami o aktywności antyoksydacyjnej mogących mieć zastosowanie farmakologiczne. Interesujące w tym względzie są substancje fitochemiczne. Za prowadzeniem prac nad nimi przemawia ponad tysiącletnia tradycja chińskiej medycyny, oparta w znacznej mierze na surowcach pochodzenia roślinnego, a wciąż będąca źródłem nowych substancji leczniczych co potwierdziła nagroda Noble z 2015 roku dla profesor Youyou Tu za odkrycie artemizyny jako lekarstwa na malarie.^[44]

Podstawowym celem badań, na których bazuje cykl publikacji niniejszej pracy doktorskiej była ocena, za pomocą chemii obliczeniowej, aktywności antyoksydacyjnej wyselekcjonowanych związków fitochemicznych, jak i potwierdzenie założenie, że metody *in silico* stanowią aktualnie nieocenione narzędzie, jakie można stawiać na równi z eksperymentami. Tym samym istotnym i zamierzonym elementem było zastosowanie różnych technik oraz sposobów interpretacji wyników, w celu podkreślenia wachlarza możliwości jakie metody obliczeniowe niosą za sobą.

2.1. Cele szczegółowe poszczególnych prac

- [A] Analiza jakościowa wpływu elementów strukturalnych na aktywność przeciwrodnikową oraz estymacja specyficznych indeksów reaktywności dla wybranych flawonoli i flawonów.
- [B] Analiza jakościowa oraz ilościowa wpływu elementów strukturalnych fenolokwasów na ich aktywność przeciwutleniającą wobec układu redoks Fe³⁺/Fe²⁺
- [C] Kompletna analiza termochemiczna i kinetyczna aktywności przeciw-OOH oraz zdolności kompleksowania jonów Cu²⁺ i Fe³⁺, hamowania oksydazy ksantynowej, a także estymacja specyficznych indeksów reaktywności apigeniny w modelowych warunkach fizjologicznych
- **[D]** Przegląd piśmiennictwa podsumowujący aktualne trendy w modelowaniu aktywności przeciwrodnikowej antyoksydantów

3. Materiały i metody

3.1. Oprogramowanie

Niskoenergetyczne konformery studiowanych cząsteczek były generowane przy użyciu programu Gabedit^[45], w którym zaimplementowana jest procedura symulowanego wyżarzania z dynamiką molekularną opartą na potencjałach mechaniki molekularnej pola siłowego Amber99^[46] oraz wartościach półempirycznych. Badania kwantowo–mechaniczne przeprowadzono z zastosowaniem pakietu obliczeniowego Gaussian16^[47], a do wizualizacji struktur 3D posłużono się oprogramowaniem Avogadro^[48] lub GaussView^[49].

W pracy **[B]** analizę topologii wykonano w programie Multiwfn (wersja 3.7)^[50], a program PaDEL– Descriptor^[51] został spożytkowany na potrzeby obliczenia deskryptorów oraz wybranych "*chemicznych* odcisków palca" (ang. fingerprints) dostępnych w bazie PubChem^[52]. Modele QSAR (ang. quantitative structure–activity relationship) zbudowano wykorzystując oprogramowanie QSARINS (wersja 2.2.4).^[53] Wyniki dokowania enzymatycznego przedstawione w publikacji **[C]** otrzymano z użyciem programu AutoDock VINA^[54].

Rysunki oraz grafiki opracowano korzystając z programów MarvinSketch, GIMP, UCSF Chimera^[55] oraz LigPlot+^[56].

Dostęp do programów, zasobów komputerowych oraz przestrzeni dyskowej, a tym samym możliwość realizacji badań, uzyskano z tytułu grantów obliczeniowych we Wrocławskim Centrum Sieciowo– Superkomputerowym (WCSS), grant nr 527; Poznańskim Centrum Superkomputerowo Sieciowym (PCSS), grant nr 467; Centrum Informatycznym Trójmiejskiej Akademickiej Sieci Komputerowej (CI–TASK); oraz Cyfronecie, wchodzącym w skład Infrastruktury PLGrid, grant *plgantioxidants*.

3.2. Metody mechaniki molekularnej oraz kwantowej

3.2.1. Informacje ogólne

Na etapie wyszukiwania lokalnego minimum wdrożono protokół próbkowania konformacyjnego o czasie nagrzewania 1.0 ps, po którym następowała 1.0 ps równoważenia układu. Resztę parametrów pozostawiono domyślnie.

Badania kwantowo–mechaniczne bazowały na teorii funkcjonałów gęstości i jako najwyższe poziomy wykorzystano: B3LYP^[57,58]/6–31+G(d,p)^[59–62]/IEF–PCM^[63] **[A]**, B3LYP/cc–pVDZ^[64]/CPCM^[65] **[B]**, M05–2X^[66]/6–311+G(d,p)^[59,67]/SMD^[68] **[C]**. W przypadku pracy **[C]**, część obejmującą eksplikację zdolności chelatowania jonów Fe³⁺ oraz Cu²⁺ oparta była na funkcjonale M05^[69], ponieważ w przeciwieństwie do M05–2X, został on sparametryzowany także na układach z metalami.^[70] W celu zmniejszenia zużycia zasobów obliczeniowych i ominięcia konieczności uwzględnienia efektów relatywistycznych dla elektronów rdzenia, chmurę elektronową jonów metali opisano przy użyciu pseudopotencjałów Stuttgart–Dresden^[71] i opowiadającej im bazie funkcyjnej^[72].

Jakkolwiek nie wspomniano *explicite*, tak w każdych obliczeniach zastosowano bardzo wąskie odcięcie optymalizacji geometrii (*Opt=VeryTight*) oraz ultradrobną siatkę całkującą (*Int=Ultrafine*). Cząstki otwartopowłokowe — rodniki i jony — były traktowane metodami nierestrykcyjnymi, odpowiednio UB3LYP oraz UM05–2X(UM05). Ponadto, mimo że Gaussian dokonuje anihilacji spinu podczas procedury samouzgodnienia pola, niekiedy jest ona zawodna, zaś zanieczyszczone spinem funkcje falowe nie są wiarygodne i prowadzą do błędów w geometrii, energiach i analizie populacji. Dla takich systemów sprawdzano wartości zanieczyszczenia spinowego, zgodnie z założeniem, że $\langle S^2 \rangle = s(s + 1)$, gdzie *s* to połowa liczby niesparowanych elektronów, a $\langle S^2 \rangle$ to wartość spodziewana.^[73]

3.2.2. Stałe dysocjacji oraz procentowe stężenie molowe

Oszacowanie wartości stałych dysocjacji dla kwasowych protonów oraz ustalenie preferowanych ścieżek deprotonacji w roztworze wodnym wykonano w myśl sposobu zaproponowanego przez *Galano et al.*^[74] Ucieka się on do dopasowania różnicy energii Gibbsa między sprzężoną zasadą, a jej odpowiadającym kwasem, obliczonymi na tożsamym poziomie teorii, do stworzonego w oparciu o dane eksperymentalne modelu regresji liniowej przy wykorzystaniu wartości współczynnika kierunkowego i wyrazu wolnego właściwych użytemu funkcjonałowi oraz bazie funkcyjnej. Na podstawie uzyskanych stałych dysocjacji wykreślone zostaje procentowe stężenie molowe danej formy antyoksydanta w całym zakresie pH.

3.2.3. Indeksy reaktywności

Studiowane indeksy reaktywności skupiały się na mechanizmach powszechnie referowanych w pracach teoretycznych^[75,76], a do jakich zaliczyć można:

a) formalny transferu atomu wodoru (*ang. formal hydrogen atom transfer, f-HAT*[‡]), będący najczęściej wymienianą ścieżką reakcji, której antyoksydant ulega w procesie zmiatania rodników. Opiera się on na homolitycznym rozpadzie wiązania pomiędzy wodorem a atomem tlenu aromatycznej grupie hydroksylowej, według schematu:

$$ArOH \rightarrow ArO^{\bullet} + H^{\bullet}$$

Energetyczną wypadkową procesu jest energia dysocjacji wiązania (*ang. bond dissociation energy, BDE*) sformułowana równaniem:

$$BDE = E(ArO^{\bullet}) + E(H^{\bullet}) - E(ArOH^{\bullet})$$

 b) Transferu elektronu—transferu protonu (ang. electron transfer-proton transfer, ETPT), czyli dwuetapowej sekwencji uwalniania elektronu z cząsteczki i dalszej dysocjacji protonu z utworzonego kationorodnika:

> $ArOH \rightarrow ArOH^{+\bullet} + e^{-}$ $ArOH^{+\bullet} \rightarrow ArO^{\bullet} + H^{+}$

Pierwszy etap definiowany jest przez adiabatyczny potencjał jonizacji (ang. adiabatic ionization potential, aIP), a drugi przez energię dysocjacji protonu (ang. proton dissociation energy, PDE):

 $IP = E(ArOH^{+}) + E(e^{-}) - E(ArOH)$ $PDE = E(ArO^{-}) + E(H^{+}) - E(ArOH^{+})$

 c) Sekwencyjnej utraty protonu—transferu elektronu (ang. sequential proton loss-electron transfer, SPLET), w trakcie której dochodzi do dysocjacji protonu z badanego związku, a następnie emisji swobodnego elektronu:

$$ArOH \rightarrow ArO^{-} + H^{+}$$
$$ArO^{-} \rightarrow ArO^{\bullet} + e^{-}$$

Energetyczny zysk procesu stanowi powinowactwo do protonu (*ang. proton affinity, PA*) i energia transferu elektronu (*ang. electron transfer energy, ETE*) określone przez:

 $PA = E(ArO^{-}) + E(H^{+}) - E(ArOH)$ $ET = E(ArO^{*}) + E(e^{-}) - E(ArO^{-})$

Wartości referencyjne dla atomu protonu i elektronu zaczerpnięto z odpowiednich prac wskazanych w manuskryptach.

[‡]Określenie *formalny* ma szczególne znaczenie. Zasadniczo reakcja może być oparta na dwóch różnych mechanizmach: 1) bezpośrednim przeniesieniu atomu wodoru (ang. *hydrogen atom transfer, HAT),* dominującym w rozpuszczalnikach niepolarnych i niejonowych roztworach wodnych, albo 2) transferze elektronu sprzężonym z protonem (ang. *proton–coupled electron transfer, PCET*), preferowanym w reakcjach, w których uczestniczą silnie elektronegatywne reagenty. W przypadku HAT atom wodoru przenoszony jest jako jedna całość w oparciu o trójelektronową delokalizację gęstości elektronowej wzdłuż ścieżki reakcji, z kolei w PCET proton i elektron przekazywane są niezależnie od siebie, a tym samym mogą pochodzić z innych orbitali. Co więcej, PCET obejmuje delokalizację czteroelektronową.^[91]

3.2.4. Pierwszorzędowa aktywność antyoksydacyjna

Ocena aktywności przeciwrodnikowej, określanej w literaturze jako pierwszorzędowa, projektowana na typowe warunki fizjologiczne bazowała na zaproponowanym protokole QM–ORSA^[75–77] (*ang. Quantum Mechanics–Based Test for Overall Free Radical Scavenging Activity*). Składający się z trzech etapów zakłada badania w modelu lipidowym oraz wodzie o pH=7.4 jako środowiskach biologicznie najważniejszych.

Za rekomendowany rodnik, '*R*, uważa się 'OOH ze względu na wyraźnie niższą reaktywność, w porównaniu do 'OH. Doświadczalnie zmierzone stałe kinetyczne reakcji 'OH z Troloxem C $(k \approx 6.9 \times 10^9 \text{ M}^{-1} \text{ s}^{-1})^{[78]}$ czy anionem kwasu linolowego $(k \approx 8.0 \times 10^9 \text{ M}^{-1} \text{ s}^{-1})^{[79]}$ dowodzą, że 'OH reaguje błyskawicznie z biomolekułami w swoim najbliższym otoczeniu, ponad granicą dyfuzji (>10⁹ M⁻¹ s⁻¹) dla warunków standardowych^[80]). Z drugiej strony, wartości eksperymentalne dla 'OOH są mniejsze, odpowiednio < 10³ M⁻¹ s^{-1[81]} oraz (1.18±0.20) × 10³ M⁻¹ s^{-1[82]}, wskazując że antyoksydant ma możliwość przechwycenia go. Co więcej, trzeba podkreślić, że wprawdzie równowaga układu 'OOH $\rightleftharpoons O_2^-$ w pH=7.4 jest przesunięta na korzyść ponadtlenku (p K_a =4.8; stosowne procenty molowe wynoszą wtedy 0.25% i 99.75%)^[81], to O₂⁻⁻ jest dobrym nukleofilem, jak również umiarkowanym reduktorem.^[13,14] Z racji tego, akceptuje się, że uszkodzenia oksydacyjne pochodzą przede wszystkim od formy protonowanej.

Dla form przeciwutleniaczy o istotnych stężeniach (>0.1mol%) wykonywane są obliczenia termochemiczne wiodące do otrzymania zmian energii swobodnych, ΔG° , mechanizmów transferu elektronu (*ang. single electron transfer, SET*), utworzenia adduktu (*ang. radical adduct formation, RAF*) oraz, już wspomnianego, formalnego transferu atomu wodoru.

SET:	$Ar-OH + R \rightarrow Ar-OH^{+} + R$
RAF:	$Ar-OH + R \rightarrow [R-Ar-OH]$
<i>f</i> –HAT:	$Ar-OH + R \rightarrow Ar-O + RH$

Ścieżki uznane za wykonywalne podlegają dalszym badaniom, tzn. odszukaniu stanów przejściowych łączących relatywne minima (reagenty—produkty) na ścieżce minimum energetycznego reakcji. Jakkolwiek protokół QM–ORSA takie przyjmuje wyłącznie te o negatywnej lub zerowej ΔG° , tak wzięcie pod uwagę dodatkowo tych umiarkowanie endoergicznych (< 10.0 kcal/mol) jest uzasadnione. Mimo że pozytywna zmiana energii swobodnej implikuje odwracalność danego procesu, może on być wciąż ważny, jeśli tylko powstałe cząstki reagują szybko dalej, w toku wystarczająco egzoergicznych procesów zapewniających siłę napędową pod postacią przesunięcia równowagi, oraz gdy energie aktywacji są dostatecznie niskie. Zważając na złożoność systemów biologicznych, jest to szczególnie prawdopodobne.

Należy jednak nadmienić, że powyższy warunek nie dotyczy transferu elektronu z tytułu całkowicie odmiennego charakteru zjawiska i wskazana jest analiza każdego wyniku. Do oszacowania bariery takiej reakcji, ΔG_{ET}^{\neq} , stosuje się teorię Marcusa^[83]:

$$\Delta G_{ET}^{\neq} = \frac{\lambda}{4} \left(1 + \frac{\Delta G_{ET}^0}{\lambda} \right)^2$$
$$\lambda = \Delta E_{ET} - \Delta G_{ET}^0$$

gdzie λ to energia reorganizacji wyliczona z wertykalnej (ΔE_{ET}) i adiabatycznej (ΔG_{ET}^0) energii reakcji, opisująca relaksację orbitali.

Ostatnim etapem jest estymacja kinetyki reakcji opierająca się na konwencjonalnej teorii stanów przejściowych (*ang. conventional transition state theory, TST*) z uwzględnieniem poprawki Eckarta dla tunelowania, w ramach 1M roztworu standardowego, za pomocą równania:

$$k^{TST} = \sigma \kappa(T) \frac{k_B}{h} e^{-\Delta G^{\neq}/RT}$$

gdzie k_B i *h* to, odpowiednio, stałe Boltzmanna i Plancka, ΔG^{\neq} to energia aktywacji, *R* oraz *T* oznaczają stałą gazową i temperaturę reakcji, zaś σ opisuje stopień degeneracji ścieżki reakcji, a $\kappa(T)$ to współczynnik transmisji. Poprzez $\kappa(T)$ brana jest pod uwagę średnia Boltzmanna dla klasycznego prawdopodobieństwa zajścia reakcji oraz jej odpowiednika w świecie kwantowym, i jest przeto sugerowany dla reakcji, których reagenty mogą przejść przez barierę potencjału o wysokości większej niż ich energia — do nich zalicza się chociażby mechanizm *f*–HAT, w którym występuje ruch *"lekkiej"* cząstki H⁺, a także niektóre ścieżki RAF.

Ponadto, do oceny stałych szybkości w pobliżu granicy dyfuzji, tak częstych dla reakcji rodnikowych, wykorzystuje się teorię Collinsa–Kimballa^[84] oraz równania Smoluchowskiego^[85] i Stokesa–Einsteina^[86]:

$$k^{app} = \frac{k^D k^{TST}}{k^D + k^{TST}}$$
$$k^D = 4\pi R D_{AB} N_A$$
$$D = \frac{k_B T}{6\pi \eta r_A}$$

gdzie *R* to dystans między reagentami, D_{AB} oznacza ich wzajemny współczynnik dyfuzji ($D_{AB} = D_A + D_B$), N_A to liczba Avogadro, η stanowi lepkość rozpuszczalnika, zaś *r* to promień substancji, uzyskany obliczeniowo dla objętości wewnątrz konturu o gęstości 0.001elektronów/Bohr³ w ramach modelu polaryzowalnego ośrodka zaproponowanego przez Onsagera.^[87]

3.2.5. Drugorzędowa aktywność antyoksydacyjna

Poza bezpośrednią zdolnością zmiatania wolnych rodników większość antyoksydantów polifenolowych jest w stanie chelatować jony metali przejściowych. Doświadczalnie udowodniono, że Fe³⁺ i Cu²⁺ w formie kompleksu nie uczestniczą tak aktywnie w reakcji Fentona jak niezwiązane, w konsekwencji czego ilość wygenerowanych na drodze katalizowanej redukcji nadtlenku wodoru 'OH oraz 'OOH maleje.^[88] Badania teoretyczne biorą za cel przetestowaniu możliwości utworzenia takich układów i ich trwałości na podstawie stałych kinetycznych procesu zachodzącego według ogólnego równania:

$$Me(H_2O)_i^j + mL^n \rightarrow [Me(H_2O)_{i-2m}L_m]^{j-mn},$$

gdzie *j* to ładunek akwakompleksu jonu metalu, a *i* to liczba cząsteczek wody koordynowanych w jego pierwszej warstwie solwatacyjnej. Zmienne *m* i *n* to, kolejno, liczba ligandów budujących kompleks (m=1 dla kompleksów monodentnych, m=2 dla kompleksów bidentnych, etc...) oraz ładunek liganda.

W oparciu o otrzymaną energię swobodną chelatowania w otoczeniu $f(\Delta G_f)$ ustala się stałe równowagi K_f , K_i^{II} oraz K_i^{app} określone równaniami:

$$K_{f} = e^{-\frac{\Delta G_{f}}{RT}}$$
$$K_{i}^{II} = \sum K_{f} \times {}^{m}f_{i}$$
$$K_{i}^{app} = \sum K_{i}^{II}$$

a opisujące, odpowiednio, stałą równowagi kompleksowania w otoczeniu *f*, stałą równowagi poszczególnej formy antyoksydanta biorąc pod uwagę jego ułamek molowy w badanych warunkach oraz pozorną stałą chelatowania studiowanej substancji. Przez *otoczenie* należy rozumieć daną topologię grup hydroksylowych uczestniczących w tworzeniu centrum koordynacyjnego. Warto wspomnieć, że metodyka została rozwinięta o ocenę potencjału pro– i antyoksydacyjnego powstałych kompleksów z uwzględnieniem askorbinianu oraz jonu ponadtlenkowego jako fizjologicznych reduktorów, lecz właściwa publikacja^[89] nie wchodzi w zakres rozprawy doktorskiej.

4. Wyniki, dyskusja i wnioski

4.1. Publikacja [A]

W rezultacie wykonanych badań udało się opisać elementy strukturalne odpowiadające za reaktywność wybranych flawonów i flawonoli, a także wstępnie oszacować ich pierwszorzędową aktywność przeciwrodnikową bazując na termochemii dostępnych mechanizmów działania. W wyborze związków decydujące było zgromadzenie takich o jak największych różnicach w budowie, co pozwoliło uwiarygodnić obserwacje.

Aby scharakteryzować chmure elektronowa, w początkowej fazie prac zastosowano analize NBO (ang. Natural Bond Orbital)^[90] polegająca na transformacji funkcji falowej do postaci zlokalizowanej – graficznie tożsamej strukturze Lewisa — otrzymując bardziej zrozumiały opis interakcji. Nieobsadzone orbitale p_v w następstwie koniugacji i idącej za nią delokalizacji zmniejszają energię systemu. Wzajemna planarność pierścieni układu AC wobec B stanowi bezpośrednią determinantę efektu. Flawonoidy posiadające podstawnik w pozycji C3, takie jak na przykład moryna, tworzą zawadę przestrzenną, uniemożliwiając osiągnięcie zerowego kąta dwuściennego. Z drugiej strony, czyni to wskazane ugrupowanie nadzwyczaj podatnym na usunieciu atomu wodoru, gdyż prowadzi to do spontanicznej rotacji pierścienia B, aż do uzyskania relatywnie płaskiej struktury. Rezonans jest również modulowany obecnością wiązania podwójnego C2=C3, a jego nasycenie drastycznie osłabia silę redukcyjną polifenolu. Trzeba nadmienić, że istota procesu jest zamknięta niemal całkowicie w obrębie dwóch, niezależnych od siebie układów elektronowych AC oraz BC, istniejących na skutek koniugacji krzyżowej z powodu obecności grupy karbonylowej przy C4. Swoboda ruchu elektronów w granicach danego układu szczególnie istotnie wpływa na aktywność, o ile gestość spinowa powstaje na atomie, z którego zdolna jest ulec jak największej delokalizacji. Dla pierścienia B takie zjawisko dotyczy przede wszystkim grupy hydroksylowej C_4 , bowiem gęstość spinowa może wtedy sięgać aż do ugrupowania karbonylowego, zaś sam pierścień, wraz z mostkiem łączącym go z układem C, tworzy trwały układ o-chinonu.

Powyższe spostrzeżenia znalażły odzwierciedlenie w obliczonych indeksach reaktywności. Rzeczywiście w badaniach nad mechanizmem *f*–HAT najkorzystniejszym miejscem abstrakcji wodoru okazało się ugrupowanie hydroksylowe przy C₄, ze względu na delokalizację chmury elektronowej na większość układu flawonoidu, lub C₃, gdy stanowi to warunek *sine qua non* osiągniecia zerowego kąta dwuściennego między pierścieniami, a więc samą możliwość rozszerzenie rezonansu. Nie jest to jednak reguła — wewnątrzcząsteczkowe wiązania wodorowe pochodzące z sąsiadujących grup hydroksylowych niekiedy prowadzą do stabilizacji tak dużej, że jest ona energetycznie korzystniejsza niż podane wcześniej. Można to zaobserwować w przypadku kwercetyny czy mirycetyny. Dla obu, oraz luteoliny i fisetyny, dokonano analizy zjawiska, wskazując zasadnicze różnice we wspomnianym efekcie, uwarunkowane liczbą partycypujących grup hydroksylowych. Tożsamo ciekawą obserwacją jest stwierdzenie, że spośród dwóch potencjalnie współistniejących konformerów, tzn., gdzie grupa hydroksylowa przy C₂. formuje wiązania wodorowe albo z atomem tlenu pierścienia piranu albo z grupą hydroksylową przy C₃, właśnie to drugie – pomimo większego kąta torsyjnego między A i B — jest energetycznie faworyzowane.

W tejże pracy udało się oprócz tego wskazać SPLET jako mechanizm preferowany w środowisku wodnym, zaś – powtarzając raz jeszcze – grupę hydroksylową przy C4[,] jako najbardziej reaktywną nie tylko w przytoczonym procesie, ale też w jednostkowym wyznaczniku, jakim jest BDE. Również wykazano, że flawonole są aktywniejsze od flawonów. Na niewątpliwą uwagę zasługuje także ugrupowanie katecholowe (C3[,]C4[,]), w którym po oderwaniu atomu wodoru z C4[,]–OH, atom wodoru z C3[,]–OH jest niejako zawieszony pomiędzy oboma tymi elementami strukturalnymi faworyzując kolejne abstrakcję atomu wodoru i skutkujące nią uformowanie znacznie stabilniejszego układu katecholu.

4.2. Publikacja [B]

Tematyka pracy też skupiała się na analizie strukturalnej, z tym że fenolokwasów i w kontekście ich siły redukowania jonów żelaza Fe³⁺. Zastosowano w niej modele ilościowe QSAR, skonstruowane na łączonych wynikach eksperymentalnych i teoretycznych. W ramach prac przebadano trzy cechy determinujące mierzalne właściwości — wzajemną pozycję grup hydroksylowych, wpływ metylacji, oraz odległość między grupami fenolowymi a karboksylową.

Zauważono, że substancje o największej aktywności w teście FRAP miały dwie lub więcej grup hydroksylowych położonych w pozycjach *orto* lub *para* względem siebie. Dla kontrastu, najniższą reprezentowały związki z wyłącznie jedną grupą hydroksylową, dwoma w pozycji *meta*, a także metylowe pochodne kwasu benzoesowego. Analogicznie jak w przypadku flawonoidów efekt rezonansu ma ogromne znaczenie. Widać to szczególnie dla układów *para* oraz *orto*, bowiem w rezultacie powstania gęstości spinowej na jednej z grup hydroksylowych tworzących go dochodzi do aktywacji drugiej grupy. Jest to fenomen bliźniaczy z zaobserwowanym wcześniej dla flawonoidów utworzeniem, odpowiednio, chinonu i katecholu.

Do osłabienia potencjału redukcyjnego przykłada się metylacja, powodująca zmniejszenie ilości grup mogących brać udział w procesie. Na dodatek, zaistniała zawada steryczna utrudnia chelatowanie jonów żelaza, zachodzące tuż przed właściwym etapem przekazania elektronu.

W toku prac wyodrębniono dwa czynniki modulujące aktywność poprzez stabilizację rodnika: rezonans oraz wewnątrzcząsteczkowe wiązanie wodorowe. Dzięki delokalizacji, *orto-meta* i *para-meta* hydroksylowane kwasy fenolowe wykazują większą aktywność niż kwasy monohydroksylowe i te, które mają dwie grupy hydroksylowe w pozycji *meta* względem siebie. Oprócz stabilizacji rezonansowej, utworzony w przebiegu reakcji redoks rodnik może być stabilizowany przez międzycząsteczkowe wiązania wodorowe pomiędzy grupami funkcyjnymi a polarnymi rozpuszczalnikami protonowymi, a także wewnątrzcząsteczkowe wiązania wodorowe. Dla badanych związków wyodrębniono dwa rodzaje: (1) obejmujące tylko tleny hydroksylowe; (2) pomiędzy grupami karboksylowymi i hydroksylowej jest mniej wydajnym akceptorem wiązań wodorowych w porównaniu z podwójnie związanym tlenem grupy karboksylowej. Stabilizacja wiązań wodorowych wyjaśniła przyczynę podwyższonej aktywności pochodnych kwasu benzoesowego z podstawioną pozycją *orto*.

W celu przedstawienia otrzymanych danych jakościowych w bardziej praktyczny sposób utworzono model wielorakiej regresji liniowej. Eksperymentalnie zmierzone jednostki aktywności TAU zostały opisane przez odpowiednie deskryptory topologiczne. Wybierając je, zwrócono uwagę na to, że: (1) dwie lub więcej grup hydroksylowych w pozycji *orto* względem siebie pozytywnie wpływa na aktywność antyoksydacyjną ze względu na delokalizację ładunku i wewnątrzcząsteczkowe wiązanie wodorowe OH–OH; (2) pochodne kwasu fenylooctowego są wydajniejsze od innych ze względu na obecność jakichkolwiek podstawników *orto–meta* lub *para–meta*; (3) obecność *orto–meta*–podstawionych pochodnych kwasu benzoesowego, o silnych wewnątrzcząsteczkowych wiązaniach wodorowych pomiędzy *orto*–hydroksylowymi i karboksylowymi grupami powodujących znaczny wzrost wydajności; (4) metylacja pochodnych kwasu cynamonowego nie zmniejsza krytycznie aktywności tych związków. Przy zadowalających parametrach statystycznych i chemicznie uzasadnionych deskryptorach (*p*=0), opracowano model, który przypuszczalnie, może dobrze odzwierciedlać przebieg reakcji i być potencjalnie wykorzystany do przewidywania wydajności kwasów fenolowych o nieznanej aktywność.

4.3. Publikacja [C]

Całkowicie inna ścieżka badań została podjęta w pracy **[C]**, gdzie na przykładzie apigeniny zademonstrowano jak, podążając protokołem ściśle dedykowanym ocenie zdolności antyoksydacyjnych za pomocą metod obliczeniowych, da się uzyskać wyniki mające bezpośrednie przełożenie na wymierne aktywności, tj. kinetykę reakcji. Ponieważ apigenina wchodziła w zakres analiz przedstawionych w publikacji **[A]** to charakter tych stanowi uzupełnienie, a także weryfikację wcześniejszych obserwacji otrzymanych na innym poziomie teorii.

Dla zrozumienia zachowania substancji w roztworze wodnym fundamentalna jest znajomość względnych równowag kwasowo-zasadowych. Różnice między aktywnościami możliwych struktur mogą być diametralne i poważnie wpłynąć na ich potencjał redoks. W badanym przypadku uwidoczniono kolejność dysocjacji apigeniny: C7 $(pK_{a1}=7.40) \rightleftharpoons C_4$ $(pK_{a2}=8.41) \rightleftharpoons C_5 (pK_{a3}=11.61)$. Tym samym ustalono, że formy neutralne oraz mono-aniony dominują w roztworze wodnym (procenty molowe obu wynoszą 48.7%), aczkolwiek di-anion również występuje w ilościach istotnych dla właściwego oszacowania aktywności przy użyciu metod teoretycznych.

Podobnie jak w publikacji **[A]** określono indeksy reaktywności, aby mieć wstępny wgląd w mechanizmy aktywności. Energia rozpadu homolitycznego wiązania O–H grupy hydroksylowej przy C₄[,] okazała się najniższa sugerując tą pozycję jako preferowaną w mechanizmie przekazania atomu wodoru. Jakkolwiek wartości potencjału jonizacji pozostawały wysokie, tak zaobserwowano, że przejście ze środowiska lipidowego do wodnego, oraz dysocjacja, zmniejszają wymagania energetyczne procesu, ułatwiając ewentualny transfer elektronu na rodnik. Porównując wyniki otrzymane w tej pracy dla neutralnej cząsteczki w wodzie z tymi z poprzedniej, ale na poziomie teorii B3LYP/6–31+G(d,p)/IEF–PCM, widać, że pomimo rozdźwięku w wartościach absolutnych, trendy są tożsame. Wyjątkiem jest wyłącznie energetyka etapu transferu elektronu następującego po oddysocjowaniu wodoru z C4[,] oraz C₅; powodem są marginalne różnice między obiema grupami, obecne dla każdego z zastosowanych poziomów teorii. W związku z tym, można uznać, że indeksy reaktywności z pracy **[A]**, uzyskane z użyciem znacznie prostszego poziomu teorii, wiarygodnie — przynajmniej w ramach kryterium jakościowego — odzwierciedlają te obliczone dla M05–2X/6–311+G(d,p)/SMD.

Badania termochemiczne oparte na analizie energetycznej mechanizmów HAT, SET oraz RAF wobec rodnika 'OOH wskazały, że mechanizm tworzenia adduktu nie jest faworyzowany. Identyczny wniosek da się wysnuć odnośnie mechanizmu SET dla neutralnej form w roztworze wodny — nadmienić należy, że ominięcie oceny jego ergiczności w środowisku lipidowym tłumaczy się charakterem procesu, któremu nie sprzyja apolarny rozpuszczalnik, niezdolny dostatecznie solwatować elektron. W analogiczny sposób, na jaki zwrócono uwagę we wcześniejszym akapicie, transfer ze środowiska lipidowego do wodnego oraz procesy deprotonacji obniżają barierę aktywacji zachodzących reakcji.

Wyznaczenie stałych kinetycznych stanowiło końcowy punkt badań pozwalający rzutować wyniki na dane eksperymentalne. W środowisku lipidowym szybkość zmiatania 'OOH przez apigeninę jest stosunkowo niska (całkowita stała kinetyczna, *koverall*, wynosi 5.79×10^{-1} [M⁻¹ s⁻¹]) i zauważalnie mniejsza niż Troloxa czy witaminy C. Mimo że w fazie wodnej stałe kinetyczne są większe i narastają z każdą kolejną deprotonacją, apigenina swoją aktywność przeciwrodnikową nadal nie przewyższa wspomnianych substancji, często uznawanych za referencyjne.

W niniejszej pracy przeprowadzono także preliminarne badania nad zdolnością apigeniny do hamowania generowania rodników hydroksylowych wskutek chelatowania jonów Cu²⁺ i Fe³⁺ oraz inhibicji oksydazy ksantynowej. Otrzymana stała kompleksowania, K^{f}_{app} , jonu żelaza przez apigeninę $(1.09 \times 10^{12} [M^{-1} s^{-1}])$ jest większa o niemal dziewięć rzędów wielkości niż ta uzyskana dla jonu miedzi $(6.74 \times 10^{3} [M^{-1} s^{-1}])$. Takie chelaty wykazują znacząco mniejszą aktywność w procesach redoks, niż ma to miejsce w przypadku ich akwakompleksów. W oparciu o referencyjną strukturę krystaliczna kwercetyny związanej z centrum aktywnym oksydazy ksantynowej wykonano dokowanie apigeniny. Rezultaty dowodzą, że niezależnie od formy jonowej, każda podobnie wiąże się z aminokwasami tego białka. Można więc spodziewać się, iż wygenerowane przez enzym rodniki mogą być błyskawicznie zmiatane przez związany z nim flawonoid.

4.4. Publikacja [D]

W niniejszej pracy przeglądowej dokonano wnikliwej analizy bibliograficznej nad aktualnymi trendami oraz standardami w badaniach nad pierwszorzędową aktywnością antyoksydacyjną. Zwrócono szczególną uwagę na wybór poziomu teorii, zarówno w odniesieniu do jakości uzyskanych wyników, jak i możliwości porównania z innymi. Przedstawiono szereg wskaźników służących analizie struktury elektronowej, takich jak m.in. teoria orbitali granicznych, mapowanie potencjału elektro– oraz wodorodonorowego czy indeksy Fukui, oraz sposobów ich interpretacji w obrębie tematyki antyoksydantów, a także używane aparaty matematyczne. Kwerenda bibliograficzna zgromadziła wiedzę z ponad 250 publikacji, przedstawiła ją w zorganizowany sposób oraz podkreśliła istniejące braki, jak również zasugerowała nowe drogi, jakimi warto zainteresować się przy tego typu badaniach.

5. Piśmiennictwo

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Article

Flavones' and Flavonols' Antiradical Structure–Activity Relationship—A Quantum Chemical Study

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Abstract: Flavonoids are known for their antiradical capacity, and this ability is strongly structuredependent In this research, the activity of flavones and flavonols in a water solvent was studied with the density functional theory methods. These included examination of flavonoids' molecular and radical structures with natural bonding orbitals analysis, spin density analysis and frontier molecular orbitals theory. Calculations of determinants were performed: specific, for the three possible mechanisms of action—hydrogen atom transfer (HAT), electron transfer–proton transfer (ETPT) and sequential proton loss electron transfer (SPLET); and the unspecific—reorganization enthalpy (RE) and hydrogen abstraction enthalpy (HAE). Intramolecular hydrogen bonding, catechol moiety activity and the probability of electron density swap between rings were all established. Hydrogen bonding seems to be much more important than the conjugation effect, because some structures tends to form more intramolecular hydrogen bonds instead of being completely planar. The very first hydrogen abstraction mechanism in a water solvent is SPLET, and the most privileged abstraction site, indicated by HAE, can be associated with the C3 hydroxyl group of flavonols and C4' hydroxyl group of flavones. For the catechol moiety, an intramolecular reorganization to an o-benzoquinone-like structure occurs, and the ETPT is favored as the second abstraction mechanism.

Keywords: flavonoids; polyphenols; antioxidants; quantum chemistry; density functional theory (DFT); structure–activity relationship

1. Introduction

The twofold nature of reactive oxygen and nitrogen species (ROS, RNS) in the organism is broadly reported [1–3]. They not only participate in signal transduction [3,4], but also may lead to the breaking of DNA chains [5], lipid peroxidation [6] and protein decomposition [7]. During oxidative stress, the free radical concentration overwhelms natural antioxidants' capacity, damaging cells and initiating severe diseases such as atherosclerosis [8], neoplasms [9] and Parkinson's [10] or Alzheimer's [11] disease.

Flavonols and flavones belong to a large group of polyphenolic compounds of flavonoids, known for their beneficial activity, deriving from the antiradical potential [12]. With a capacity to scavenge free radicals and a wide distribution in vegetables [13], they play a crucial role as an external source of antioxidants. Therefore, it is important to maintain their recommended intake.

Their antioxidative ability was found to depend greatly on the molecular structure and substitution pattern: availability of hydroxyl groups—their absolute and relative position, as well as their number [14–17]; the stabilizing effect of hydrogen bonds—intramolecular and originating from

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the solvent [18,19]; the electron delocalization across a molecule, which considerably relies on the degree of conjugation [16,20] and hyperconjugation effect [21]; and the substituents effect, especially of methoxy groups [22].

It is believed that A-ring substituents are not directly involved into the scavenging mechanism [17,23]. Therefore, the type of the B-ring substitution is considered as a determinant of flavonoids' antiradical potency—even one hydroxyl group located there guarantees noticeable scavenging potential, especially if it is in position C4'. Similar observations were noted for phenolic acids [16,24,25]. Furthermore, highly active flavonoids usually also possess a catechol moiety the activity of which, found in other classes of polyphenolic compounds, was demonstrated recently [15,26,27]. The C2–C3 double bond extends π -conjugation onto the carbonyl group in the C-ring, so the unsaturated flavonoids' radical scavenging ability is greater than in saturated structures, e.g., flavanones [28–30]. Herein, the catechol moiety is a subject of the investigation in this manuscript as, according to several reports [17,31,32], it can play a significant role in scavenging potential.

Appropriate assessment of flavonoids' activity requires in-depth studies on all possible modes of action: hydrogen atom transfer (HAT), electron transfer—proton transfer (ETPT), and sequential proton loss—electron transfer (SPLET) [33,34]; as well as their defining characteristics, and the mathematical values related to them: bond dissociation enthalpy (BDE)—for HAT; ionization potential (IP) and proton dissociation enthalpy (PDE)—for ETPT; proton affinity (PA) and electron transfer enthalpy (ETE)—for SPLET. Mechanism-independent determinants, reorganization enthalpy (RE) and hydrogen abstraction enthalpy (HAE) were calculated as well, but for more general, kinetic-independent purpose [35].

This study was focused on describing the structure–activity relationship (SAR) and determinants of the flavones' and flavonols' activity, with robust, computational chemistry methods. The investigations include 13 flavonoids differing in a substitution pattern (Table 1). An intramolecular swap reaction was discovered and examined for the catechol moiety in a thermodynamic aspect, and the relevance of the hydrogen bonding in the B-ring was evaluated and the electronic structure deeply explored, involving the spin density distribution, chemical hardness, as well as HOMOs and LUMOs analysis. The obtained results provide the basis for understanding flavonols' and flavones' antioxidative potential and explain the differences between them.

					³ ' 4' B 5'			
	Flavonoid	C2′	C3′	C4′	C5′	C3	C5	C7
Flavones	Acacetin Apigenin Chrysin Chrysoeriol Diosmetin Genkwanin Luteolin		-ОСН ₃ -ОН -ОН	-OCH3 -OH -OH -OCH3 -OH -OH			-OH -OH -OH -OH -OH -OH	-OH -OH -OH -OH -OH -OCH ₃ -OH
Flavonols	Fisetin Galangin Kaempferol Morin Myricetin Quercitin	-OH	-OH -OH -OH	-OH -OH -OH -OH	-OH	-OH -OH -OH -OH -OH	-OH -OH -OH -OH	-OH -OH -OH -OH -OH -OH

Table 1. Flavonoid structures investigated in this paper.

2. Materials and Methods

All energies noted as kcal/mol were converted from atomic units (a.u.) according to the conversion factor, where 1 a.u. equals 627.5 kcal/mol.

2.1. Conformer Geometry Generation

Molecules of the studied flavonoids were generated in Avogadro [36] from their simplified molecular-input line-entry system (SMILES) [37]. The obtained structures were used in the Gabedit10 [38] Amber Molecular Dynamics Conformational Search procedure to obtain the lowest energy conformers. 1.0 ps of heating followed by 1.0 ps of equilibration molecular dynamic protocols were employed. After completion, obtained conformational isomers of each flavonoid were used for quantum chemistry studies.

Each structure was first optimized in a vacuum with Gaussian16 [39] using the HF/3-21G(d) model chemistry method and then with density functional theory (DFT) B3LYP/6-31G(d,p) [40]. Very tight geometry optimization cutoff and an ultrafine integration grid were used for vibrational frequency calculations. All flavones exhibited exactly one imaginary frequency, indicating that a planar conformation is a first-order saddle point on the potential energy surface. For this reason, the C2–C1' bond of each conformer was successively rotated by 60 degrees and saved for further elaboration, until reaching total 300 degrees of rotation from the origin. Obtained geometries were recalculated at the same computational chemistry level of theory as earlier and all real frequency values were confirmed. The lowest energy conformers were selected as the representatives and optimized once more, but in a polarizable continuum model (PCM) of water solvent, using B3LYP/6-31G + G(d,p) method [41,42]. Calculated enthalpies of the lowest energetical flavonoid isomers are shown in Table S1.

It is interesting to point out that based on our results the gas phase equilibrium structures of flavonols are flat, with the notable exception of morin, while flavones reveal strained geometries, in accord with other studies [43]. The planarity of flavonols is lost in water environment contradictory to Todorova et al.'s [44] findings. In fact, according to our B3LYP/6-31+G(d,p) calculations (this trend was also noticed in a larger basis set, such as 6-31+G(d,p) or aug-cc-pVDZ), all flavonols but morin possess a very flat potential energy surface (PES) for dihedral angles, describing the distortion from planarity, spanning a region of PES ranging from 0 degrees to its equilibrium structure value seen in Table 2. The energy difference between the flat and strained compounds is tiny, up to 0.10 kcal/mol, however, the strained structure always has the lower energy and all real vibrational frequencies, in contrast to the one imaginary frequency found for planar structures of galangin, fisetin and myrecetin. It is worth noting that these conclusions hold true for re-optimized equilibrium geometries with an unpruned grid (Grid = 199974) and very tight optimization criteria, as well as re-calculated vibrational frequencies for such high-quality geometries.

2.2. Radical Geometry Generation

The representative geometries from the previous step served as an input for radical calculations. This step includes removing a single hydrogen atom from each hydroxyl group and running computations at UB3LYP/6-31+G(d,p), retaining an implicit water solvent in two ways: with and without geometry optimization. Spin contamination values of open shell DFT results were checked, as they may interfere with the outcome [45,46]. All were in a range of <0.7500, 0.7511> after spin annihilation. Cation-radical, anion-radical and triplet diradical calculations used later in this study were elaborated the same way. The outcomes obtained for the first two forms were in the same range as for radicals, whilst for the latter one they were found in a range of <2.0001, 2.0030>. The ideal values, calculated according to the formula s(s + 1) where s is a half of a number of unpaired electrons, are 0.75 for a radical and 2.0 for a triplet diradical. Since DFT results are in the acceptable range, they could be used for the elaboration. Calculated enthalpies are presented in Table S2. Structural parameters of

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flavonoids, such as dihedral angles around the C2–C1' bond (θ) and corresponding values for relaxed radicals (θ^{\bullet}), are shown in Table 2.

Table 2. C2–C1' dihedral angles of molecules, their corresponding radicals and the difference between them (Δ).

	Flavonoid	θ				θ• C2-C1'			
	riavonoiu	C2-C1'	C2′	C3′	C4′	C5′	C3	C5	C7
	Acacetin	-13.7						12.9	7.9
	neuccun	10.7						$(\Delta = -2.2)$	$(\Delta = -7.2)$
	Anigenin	-15.5			-5.7			-12.5	-8.6
	ripigeinn	1010			$(\Delta = + 9.8)$			$(\Delta = + 3.0)$	$(\Delta = -6.9)$
ŝ	Chrysin	20.6						18.7	16.3
one								$(\Delta = -1.9)$	$(\Delta = -4.3)$
avı	Chrysoeriol	-16.9			-3.7			-18.3	-4.9
E				10.1	$(\Delta = + 13.2)$			$(\Delta = -1.4)$	$(\Delta = + 12.0)$
	Diosmetin	16.7		12.1				17.6	9.9
				$(\Delta = -4.6)$	4 -			$(\Delta = + 0.9)$	$(\Delta = -6.8)$
	Genkwanin	-16.0			-4.5			-14./	
				22.5	$(\Delta = \pm 11.3)$			$(\Delta = \pm 1.3)$	0.2
	Luteolin	15.9		$(\Lambda = \pm 7.6)$	$(\Lambda = -3.7)$			$(\Lambda = \pm 3.3)$	$(\Lambda = -7.6)$
				$(\Delta = +7.0)$	$(\Delta = -5.7)$			$(\Delta = + 0.5)$	$(\Delta = -7.0)$
	Fisetin	8.8		3.3	0.0		0.3		6.2
				$(\Delta = -5.5)$	$(\Delta = -8.8)$		$(\Delta = -8.5)$		$(\Delta = -2.6)$
	Galangin	-15.0					0.0	-3.6	-14.2
ŝ	0				0.0		$(\Delta = + 15.0)$	$(\Delta = + 11.4)$	$(\Delta = + 0.8)$
lor	Kaempferol	-3.5			0.0		0.0	0.1	-3.4
IOV			44.7		$(\Delta = + 3.5)$		$(\Delta = + 3.5)$	$(\Delta = + 3.6)$	$(\Delta = + 0.1)$
Fla	Morin	35.7	44./		32.0		(1 25.7)	(4 1 2)	(1 2 5)
			$(\Delta = + 9.0)$	-10.1	$(\Delta = -3.1)$	-53	$(\Delta = -35.7)$	$(\Delta = -1.5)$	$(\Delta = -2.5)$
	Myricetin	-9.1		$(\Lambda = -1.0)$	$(\Lambda = \pm 6.6)$	$(\Lambda = \pm 3.8)$	$(\Lambda = \pm 0.1)$	$(\Lambda = \pm 2.8)$	$(\Lambda = \pm 1.4)$
				_10.5	0.0	$(\Delta = \pm 0.0)$	0.0	-59	-57
	Quercetin	-8.5		$(\Lambda = -2.0)$	$(\Lambda = +8.5)$		$(\Lambda = +8.5)$	$(\Lambda = +2.6)$	$(\Lambda = +2.8)$
				(11 = 2.0)	(= 10.0)		(= 10.0)	(11 = 12.0)	(11 = 12.0)

2.3. Quantitative Determinants of Antioxidant Potential

The enthalpy values, H(x), used in this section refer to the unrelaxed forms, include thermal correction and were obtained following Hessian calculations by employing Gaussian16 software [39]. Reorganization enthalpy is the only one that makes use of relaxed radicals' results. The values of enthalpy for H⁺, H[•] and e⁻ in a water solvent were taken from another study [47].

2.3.1. Hydrogen Atom Transfer Mechanism

The hydrogen atom transfer is the simplest reaction path an antioxidant can undergo and is based on a homolytic bond dissociation between the hydrogen and the oxygen atom in the hydroxyl residue:

$$ArOH_{relaxed} \rightarrow ArO_{unrelaxed}^{\bullet} + H^{\bullet}$$

A quantitative descriptor of this process can be assigned to the bond dissociation enthalpy [BDE; Equation (1)], defined as the change of the enthalpy after the hydrogen abstraction [Equation (2); see results in Table 3]:

$$BDE = H(ArO^{\bullet}_{unrelaxed}) + H(H^{\bullet}) - H(ArOH_{relaxed})$$
(1)

$$HAT = BDE$$
 (2)

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Bond Dissociation Enthalpy Flavonoid C2′ C3′ C4′ C5′ C3 C5 C7 91.1 102.1 Acacetin 89.3 100.5 91.7 Apigenin Flavones 102.1 Chrysin 92.8 87.0 Chrysoeriol 106.7 91.6 87.8 106.892.0 Diosmetin 88.8 Genkwanin 102.2 88.2 91.8 Luteolin 84.4 106.1 90.3 Fisetin 83.4 85.1 86.7 Galangin 86.9 98.5 91.7 Flavonols Kaempferol 86.7 85.5 98.3 90.3 Morin 93.4 88.8 84.6 99.1 91.4 87.5 80.7 98.3 91.0 Myricetin 84.4 86.1 Quercetin 87.4 81.4 85.5 98.0 90.4

Table 3. Values of bond dissociation enthalpies [kcal/mol].

The numbers in bold typeface indicate the lowest value for a given compound.

2.3.2. Electron Transfer–Proton Transfer Mechanism

Another recently proposed mechanism is a two-step sequence of electron release from the molecule, followed by a proton dissociation from the formed cation-radical:

$$ArOH_{relaxed} \rightarrow ArOH_{unrelaxed}^{\bullet+} + e^{-}$$
$$ArOH_{unrelaxed}^{\bullet+} \rightarrow ArO_{unrelaxed}^{\bullet} + H^{+}$$

The enthalpy of this process is the sum of the adiabatic ionization potential [IP; Equation (3)] and the proton dissociation enthalpy [PDE; Equation (4)], which can be calculated as follows [Equation (5); see results in Table 4]:

$$IP = H(ArOH_{unrelaxed}^{\bullet+}) + H(e^{-}) - H(ArOH_{relaxed})$$
(3)

$$PDE = H\left(ArO_{unrelaxed}^{\bullet}\right) + H\left(H^{+}\right) - H\left(ArOH_{unrelaxed}^{\bullet+}\right)$$
(4)

$$ETPT = IP + PDE \tag{5}$$

Table 4. Values of ionization potentials and proton dissociation enthalpies [kcal/mol].

				Pr	oton Dis	sociati	on Entl	nalpy	
	Flavonoid	Ionization Potential	C2′	C3′	C4′	C5′	C3	C5	C7
	Acacetin	115.9						16.1	5.0
	Apigenin	117.0			2.2			13.4	4.6
sət	Chrysin	120.5						11.4	2.1
IOV	Chrysoeriol	113.4			3.5			23.2	8.1
лa	Diosmetin	113.8		3.9				22.9	8.1
	Genkwanin	117.2			1.4			14.8	
	Luteolin	115.0		3.1	-0.7			20.9	6.6
	Fisetin	108.9		4.4	6.0		7.6		11.3
sl	Galangin	114.3					2.4	14.0	7.2
ouo	Kaempferol	110.2			6.4		5.2	18.0	10.0
avc	Morin	113.3	10.0		5.4		1.3	15.7	8.1
Ε	Myricetin	109.0		8.4	1.5	5.3	6.9	19.2	11.9
	Quercetin	108.9		8.4	2.4		6.5	19.0	11.4

The numbers in bold typeface indicate the lowest value for a given compound.

2.3.3. Sequential Proton Loss-Electron Transfer Mechanism

This path consists of a proton dissociation from the investigated compound and an emission of the free electron afterwards: $4rOH + + \rightarrow 4rO^{-} + H^{+}$

$$ArOH_{relaxed} \rightarrow ArO_{unrelaxed}^{-} + H^{+}$$

 $ArO_{unrelaxed}^{-} \rightarrow ArO_{unrelaxed}^{\bullet} + e^{-}$

The whole reaction is the proton affinity [PA; Equation (6)] enthalpy plus the electron transfer enthalpy [ETE; Equation (7)] [Equation (8); results are presented in Table 5]:

$$PA = H\left(ArO_{unrelaxed}^{-}\right) + H\left(H^{+}\right) - H\left(ArOH_{relaxed}\right)$$
(6)

$$ETE = H\left(ArO_{unrelaxed}^{\bullet}\right) + H(e^{-}) - H\left(ArO_{unrelaxed}^{-}\right)$$
(7)

$$SPLET = PA + ETE \tag{8}$$

Table 5. Values of proton affinity enthalpies and electron transfer enthalpies [kcal/mol].

	Fl		Pr	oton A	ffinity	Enthal	ру			Ele	ctron T	ransfei	Entha	lpy	
	Flavonold	C2′	C3′	C4′	C5′	C3	C5	C7	C2′	C3′	C4′	C5′	C3	C5	C7
	Acacetin						45.0	30.8						87.0	90.2
	Apigenin			31.8			43.3	31.4			87.4			87.1	90.1
Jes	Chrysin						44.3	31.0						87.7	91.7
IOA	Chrysoeriol			33.3			45.0	30.7			83.5			91.6	90.8
Fla	Diosmetin		36.7				45.3	30.7		81.0				91.4	91.1
щ	Genkwanin			31.1			46.0				87.5			86.0	
	Luteolin		35.8	27.4			43.8	31.3		82.3	86.8			92.1	90.3
	Fisetin		30.7	34.1		38.5		30.0		82.6	80.8		78.1		90.2
s	Galangin					35.2	40.8	30.3					81.6	87.6	91.3
ouc	Kaempferol			32.0		35.5	42.0	30.1			84.6		79.8	86.2	90.1
IOVE	Morin	39.2		32.1		28.3	40.6	29.5	84.0		86.6		86.2	88.3	91.8
Ē	Myricetin		36.0	29.1	30.1	34.9	41.2	29.9		81.3	81.5	84.2	81.0	87.0	90.9
	Quercetin		36.7	27.2		35.5	41.5	30.0		80.6	84.0		79.8	86.4	90.3

The numbers in bold typeface indicate the lowest value for a given compound.

2.3.4. Mechanism-Independent Determinants

The non-specific indices—the reorganization enthalpy [RE; Equation (9)] and the hydrogen abstraction enthalpy [HAE; Equation (10)]—were also used for a quantitative analysis of the flavonols' and flavones' antioxidative potential.

The reorganization enthalpy describes the energy change upon shift from the unrelaxed to the relaxed form, hence geometry optimization. Thus, it may be an interesting predictor of conformational changes resulting from the hydrogen abstraction:

$$ArO^{\bullet}_{unrelaxed} \to ArO^{\bullet}_{relaxed}$$
$$RE = H\left(ArO^{\bullet}_{relaxed}\right) - H\left(ArO^{\bullet}_{unrelaxed}\right) \tag{9}$$

The hydrogen abstraction enthalpy is independent of the scavenging mechanism and its kinetics. It serves as a general determinant of the flavonoid activity that includes reorganization enthalpy correction [35]:

$$ArOH_{relaxed} \to ArO_{relaxed}^{\bullet} + H^{\bullet}$$
$$HAE = H(ArO_{relaxed}^{\bullet}) + H(H^{\bullet}) - H(ArOH_{relaxed})$$
(10)

The obtained RE and HAE values are presented in Table 6.

Table 6. Values of reorganization enthalpies and hydrogen abstraction enthalpies [kcal/mol].

	Fl		Rec	organiz	ation I	Enthalp	ies			Hydro	gen Ab	ostracti	on Entl	nalpies	
	Flavonold	C2′	C3′	C4′	C5′	C3	C5	C7	C2′	C3′	C4′	C5′	C3	C5	C7
	Acacetin						-8.7	-5.1						93.5	86.0
(A)	Apigenin			-6.7			-7.1	-5.0			82.6			93.5	86.7
Jes	Chrysin						-8.6	-5.6						93.5	87.2
IOA	Chrysoeriol			-7.7			-2.4	-5.2			79.3			104.3	86.5
Fla	Diosmetin		-7.2				-2.4	-5.4		80.6				104.4	86.6
	Genkwanin			-6.3			-8.5				82.4			93.7	
	Luteolin		-7.0	-8.6			-1.7	-5.0		81.2	75.9			104.4	86.8
	Fisetin		-7.5	-7.8		-9.1		-5.9		75.9	77.3		77.6		84.4
ls	Galangin					-8.6	-8.4	-5.8					78.3	90.1	85.9
QU	Kaempferol			-6.9		-8.7	-8.5	-5.5			79.8		76.8	89.8	84.8
AVC	Morin	-8.4		-6.2		-12.6	-9.1	-5.3	85.0		82.6		72.1	90.0	86.1
Εl	Myricetin		-6.8	-8.3	-7.2	-9.4	-8.5	-6.0		80.7	72.4	77.1	76.7	89.8	85.0
	Quercetin		-7.0	-8.0		-8.8	-8.7	-5.6		80.4	73.4		76.7	89.3	84.9

The numbers in bold typeface indicate the lowest value for a given compound.

3. Results and Discussion

3.1. Molecule and Radical Electronic Structure Investigation

The starting point of the backbone examination was undertaken with a natural population analysis of NBO 3.1 [48] software, implemented in Gaussian16 [39]. In view of the fact that the backbone is identical for flavones and flavonols, the following statements can be ascribed to both of them, with some exceptions indicated in the text.

Natural bonding orbitals (NBO) analysis revealed that every C–C bond is composed of two sp^2 orbitals, so each carbon atom also has one unoccupied p_y orbital. This can lead to the assumption that free electrons, located on oxygen atoms connected with the aromatic ring, will likely interact with orbitals of carbon atoms, in conjugation and hyperconjugation effects.

Thanks to the p_{y} orbitals' conjugation, the system's total energy is lowered. This property is highly dependent on the AC- and B-rings' mutual planarity, and as discussed above, the B-ring does not share the same plane as the AC-complex. The dihedral angle varies (Table 2) depending on intramolecular repulsions or hydrogen bonding caused by C3, C2' or C5' residues. Dimitrić Marković et al. [27], explaining the differences in activity between anthocyanidins, delphinidin and pelargonidin against C3-glycosylated anthocyanin, malvin, suggested that the greater activity of the first two is due to the C3 hydroxyl group, which maintains coplanarity of the B- and C-ring, and hence p_{μ} orbitals' conjugation. Nevertheless, in this study something completely different was observed. Flavonols reach exact planar structure when hydrogen from the C3 hydroxyl group is removed. Moreover, considering reorganization enthalpy as a descriptor of the most favorable abstraction site in terms of geometry change, it is indeed the C3 hydroxyl group. It can be concluded that, upon reaction, the torsion caused by this residue does not have an impact on the B-ring anymore, and rotation is thermodynamically favored. When HAE is investigated, C3 residues are generally favored, unless the investigated flavonol has at least two hydroxyl groups nearby. Then, one of their positions is an abstraction site. That way it can be expected that hydrogen bond stabilization energy is more urgent than p_{y} orbital conjugation. Contrarily, for all flavonols but morin, abstraction from C4' also leads to the planarity of the rings. This can be associated with the formation of radical which, to be transferred onto the C-ring, require double bond formation between the py orbitals of C2 and C1'.

The hyperconjugation effect was investigated considering molecular orbitals (MOs) interactions. One can see in Figure S1 that the oxygen atoms' LUMO phases have a different sign than aromatic carbon atoms, indicating possible interaction. Indeed, Milenković et al. [21], in their study on kaempferol structure, found how greatly the interference of oxygen atoms' free electrons with the antibonding orbital of carbon atoms contributes to structure stabilization, decreasing the system's total energy

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by up to 34 kcal/mol. However, the structure as a whole is not conjugated. The existence of the carbonyl group in the C-ring, due to the cross-conjugation effect, divides the compound into two electron-separated ring complexes—AC and BC [49]. Therefore, electron density cannot flow freely, in a strict orbital manner, from the A-ring to the B-ring or in reverse, as was stated by Jovanović et al. [50]. To check the other possibility, a hydrogen shift between C5 and the carbonyl residue was elaborated (Appendix A) and its results also go against this hypothesis. Even more, this postulate can be refuted with examination of radicals' spin density distribution—upon forming a radical in the A or B-ring, the density is nearly 0 in B and A-ring atoms, respectively (Figure S2).

The C2–C3 saturation limits electrons' delocalization, indirectly decreasing reduction potential. Evidence confirming this thesis can be found by studying the activity results' comparison between flavonols and flavones against flavanols [51] or flavanones [52] in simple assays, e.g., ferric ion-reducing antioxidant power (FRAP) [53], 2,2'-azino-bis (3-ethylbenothiazoline-6-sulfonic acid) diammonium salt (ABTS) [54], and 2,2-diphenyl-1-picrylhydrazyl (DPPH) [55] or isoflavones in an acrylamide reduction test [56].

Removal of the hydrogen atom by an abstractor creates a radical form, for which Lewis resonance structures are shown in Figure 1. One can see that the biggest resonance effect occurs for the C2', C6' and C4' radicals; it is lower for the C3, C5 and C7 radicals, and the lowest for the C3' and C5' radicals.



Figure 1. Lewis structures of radicals with the biggest electron densities of atoms marked with a dot.

Formerly presented HAE values (Table 6) are accepted in this study as a main numerical determinant of antioxidative potential, as they are mechanism-independent. In compliance with them, preferred hydrogen abstraction position patterns can be depicted. Just based on HAE values, flavonoids, indeed, can be divided into two groups: (I) compounds with a C3 hydroxyl group (flavonols); (II) compounds lacking a hydroxyl residue at the C3 position (flavones).

For flavones, the C4' position, if present, is the most active one. The reason for that may be attributed to larger delocalization of the electron density upon radical formation, and the possibility of establishing the hydrogen bond with the adjacent hydroxyl residues at C3' or C5'. Similar observations were derived from studies on the activity of phenolic acids [57] or anthocyanidins [27], where C4'

had the most favorable hydrogen abstraction energy. Elaboration of this process was performed and is demonstrated in a latter section (Section 3.3), where the importance of intramolecular hydrogen bonds is confirmed. The A-ring hydroxyl residues have much higher HAE, up to 25 kcal/mol, so their scavenging potential will be lower; thus, these in the B-ring seem to be the main determinants of antioxidant capacity, as was mentioned in the Introduction section.

However, flavonols with an OH group at C3 and a lone hydroxyl moiety at C4' are more likely to convey hydrogen from the first position instead of the latter one, when HAE is compared, e.g., morin and kaempferol. On the other hand, if there are at least two (e.g., quercetin) or even three (e.g., myricetin), the C4' position will be privileged. This indicates that the p_y conjugation is desired only when the radical formed at the B-ring is not stabilized by at least one intramolecular hydrogen bonding.

3.2. Morin

Interesting properties, noted by Amić et al. [25], are exhibited by morin (Figure 2). In this study, it was found that the preferred molecular isomer is not the more planar one (isomer B), but the one able to form a hydrogen bond between C3–C2' residues (isomer A). The difference in enthalpies between these conformers is ~1.8 kcal/mol, and this value seems to be sufficient to break the tendency to planarity. One can name previously noted steric restrictions disabling complete conjugation of p_y orbitals, but the choice of isomer A can also be explained differently—the C3–C2' hydrogen bond is "retained" even after the radical is formed, from either the C2' or C3 hydroxyl group. In the first case, the hydrogen atom from the C3 hydroxyl group rotates from the carbonyl site to the B-ring site, forming a hydrogen bond with the C2' radical, subsequently forcing rotation up to 45° due to the steric effect. In the second case, when the C3 radical is formed, it is not necessary for the hydrogen atom to move, and only C2–C1' bond rotation is performed. This situation is most likely to occur since no energy is used for hydrogen shift, and planarity is achieved.



Figure 2. Comparison in geometry between morin's A and B isomer.

3.3. Mechanisms of Action in Terms of the Determinants

The very first step of the mechanism is considered as a thermodynamic determinant of a favored pathway.

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3.3.1. Hydrogen Atom Transfer Mechanism

The only investigated mathematical descriptor associated with the HAT mechanism is the bond dissociation enthalpy (Table 3). The lowest is associated with forming the C4' radical, whilst C3' and C3 enthalpies are quite similar. A likely explanation of this could be Mulliken spin density (SD) distribution in the created radicals (Figure S2). The lower the spin density at the radical center, the greater is the delocalization, and the formed radical becomes more stabilized with resonance. For all compounds but morin, the lowest spin density is associated with the C4' radical. However, delocalization cannot be clearly stated as the only determinant of BDE—if it was, the lowest BDE value for fisetin should be at C4' (SD = 0.284), whilst it is C3' (SD = 0.316); for morin, the lowest BDE should be for C2' (SD = 0.303), whilst it is C4' (SD = 0.311). Therefore, it is assumed that some other factors may be crucial to describe the HAT mechanism. One can observe that C4' BDE decreases with the number of adjacent hydroxyl groups—the lowest enthalpy was found for myricetin, where two hydroxyl groups surround C4', while significantly higher values were derived for quercetin, luteolin and fisetin, with only one adjacent hydroxyl group. Conversely, compounds where the C4' hydroxyl group is alone exhibit the highest bond dissociation enthalpies. Thus, it can be stated the HAT definitely depends on intramolecular hydrogen bonding.

3.3.2. Electron Transfer–Proton Transfer Mechanism

The IP values are much bigger than BDE, while PDE is relatively low (Table 4). Since the first step determines the thermodynamically favored reaction pathway, the ETPT mechanism is not likely to be responsible for flavonoid activity. Nevertheless, ionization potential was also examined with frontier molecular orbitals theory (Appendix B).

The lowest energy required for ionization is 108.9 kcal/mol for fisetin and quercetin, while the highest is over 120 kcal/mol for chrysin; these are much greater values than any bond dissociation energies for the same compounds. Even though PDE does not show any regular pattern in the lowest energy centers, it definitely shows the highest—PDE values for the C5 position are the largest. The reason for this may be found in an electronegative repulsion between the carbonyl oxygen and the C5 radical's unpaired electron (the distance between these atoms is about 2.9Å), as well as breakage of the hydrogen bond between these two residues.

3.3.3. Sequential Proton Loss-Electron Transfer Mechanism

The SPLET mechanism is determined with proton affinity. Calculations showed that PA values are much lower than the corresponding values for the reaction path determinants of the HAT (BDE) or ETPT (IP) mechanisms. Therefore, this one appears to represent the most favored mechanism of action for flavonoids in a water (polar) solvent, as could be expected according to the other studies [27,58], including qualitative structure-activity relationship (QSAR) analysis [59-61]. This is rather surprising as, in general, A-ring hydroxyl groups are considered not to possess scavenging potential [17,62], whilst abstraction takes place at the C7 position for most of the investigated compounds. Gibb's free energies of deprotonation, reported by Álvarez-Diduk et al. [63], are in agreement with the obtained results: the first or second pK is linked with hydrogen dissociation from the C7 hydroxyl group. Interestingly, Lin et al. [24], measuring activity of flavonoids with the DPPH assay, noted that upon glycosylation of C7 in luteolin, its antiradical activity noticeably decreased. They attributed this effect to the decreased availability of the free hydroxyl groups. Whilst the lowest proton affinity enthalpy of luteolin is assigned to C4' (27.4 kcal/mol), the second lowest is assigned to C7 (31.3 kcal/mol). An explanation that could be proposed is the activity of a catechol moiety and a change of polarity within the structure, resulting in overall decreased activity of the compound, instead of a strict change of a favored location. Moreover, since SPLET's ETE determinant is adequate for ETPT's IP, and is lower than this, it can be assumed that an electron transfer is preferred for an ion form instead of a molecule.

Based on the discussion conducted earlier for the ETPT mechanism, similar reasons can explain why the highest PA values are found for C5.

3.3.4. Mechanism-Independent Determinants

The performed elaborations considered unrelaxed forms of intermediate compounds. The proposed reorganization enthalpy is a mathematical explanation of a conformer relaxation, decreasing the total enthalpy. Therefore, one can calculate reorganization enthalpy and, based on derived values, as well as chemical structure, identify significant geometry changes. For example, the lowest RE is for the morin C3 hydroxyl group, which was described earlier in the context of hydrogen bonds and planarity. The low value of RE for C5 can be justified by an electronegative interaction between the C5' radical and carbonyl oxygen atom, which is mediated by a molecule by changing its geometry.

Although three mechanisms and their determinants were investigated, no reaction path can be proposed. It greatly depends on the solvent [34,64] and the abstractor molecule [65]. The lowest enthalpy only indicates where the abstraction is most likely to occur, not how often it will happen. For this reason, a hydrogen abstraction enthalpy can be used, and was used for further elaboration. It points out the most active compound or group, without taking into account how the radical state was reached, rejecting kinetic studies but involving reorganization of geometry. Herein, C4' seems to be the most prominent one, and similar conclusions were stated by Dimitrić Marković et al. [64,66] and Sroka et al. [35]. Furthermore, the second most favored position in a SPLET mechanism is also C4', so the importance of reorganization enthalpy correction may be bigger than one would expect.

3.3.5. Antioxidant Capacity Summary

Assuming an activity is inversely proportional to the enthalpy required for the first step of the mechanism to occur, the investigated compounds have been sorted in the descending order (Table 7). As one can see, flavonols are more active than flavones, and structures with a greater number of hydroxyl groups stand out as the most active ones. The pattern of activity does not change greatly, apart from for flavone luteolin, which is in the top three when HAT and SPLET mechanisms are examined, overtaking most of the flavonols.

Activity	HAT	ETPT	SPLET
	Myricetin	Fi di Quadi	Quercetin
	Quercetin	Fisetin, Quercetin	Luteolin
-	Fisetin	Myricetin	Morin
	Luteolin	Kaempferol	Mirycetin
	Morin	Morin	Fisetin
	Kaempferol	Chrysoeriol	Kaempferol
	Galangin	Diosmetin	Galangin
	Chrysoeriol	Galangin	
	Diosmetin	Luteolin	Chrysoeriol, Diosmetin
	Genkanin	Acacetin	Acacetin
\bullet	Apigenin	Apigenin	Chrysine
	Accacetin	Genkwanin	Genkwanin
	Chrysine	Chrysine	Apigenin

Table 7. Relative activity of investigated compounds.

Flavonoids have been sorted in decreasing order of the enthalpy for the first step of given mechanism.

3.4. Intramolecular Hydrogen Bonding

The polyhydroxy structure of flavonoids provides an opportunity to form an intramolecular hydrogen bond if at least two hydroxyl groups are close to each other. Hydrogen bonds are known for increasing stability of both the molecule and the radical, hence decreasing the energy required to form

the radical [19]. Such a situation can actually be found for every flavonol, where the C3 or C5 hydroxyl group interacts with a carbonyl residue or B-ring hydroxyl groups [21].

Fisetin, luteolin and quercetin do possess two hydroxyl moieties, while myricetin has three at the B-ring. This allows intramolecular hydrogen bonds to be formed when hydrogen is abstracted. Generally, the C4' hydroxyl group can interact with either C5' or C3', whilst C5' and C3' can interact only with C4'. The exception among this group is morin, since its C3 group can interact with C2', leading to a twisted structure. Moreover, a hydrogen bond can be formed between carbonyl group and C3 and C5 residues as well. This study focused only on the hydrogen bonding in B-ring groups, since C4' (or C3' for fisetin) was indicated as a favored position in a thermodynamically preferred SPLET mechanism, as well as as a mechanism-independent HAE determinant. In order to measure a hydrogen bonding stabilizing effect on a radical molecule, the difference in the enthalpies between the radical without (ArO_{NHB}) and with the hydrogen bonds (ArO_{HB}) was considered, named here as hydrogen bond enthalpy (*HBE*), and ascribed to the following equation:

$$HBE = H(ArO_{NHB}^{\bullet}) - H(ArO_{HB}^{\bullet})$$
(11)

Since myricetin possesses three hydroxyl groups in the B-ring, in close proximity, two more situations had to be considered. For the C3' radical, the C5' hydroxyl group can be facing the same direction as C4'—named here as a cross hydrogen bond (CHB)—or the opposite one. On the other hand, the C4' radical can be stabilized by two, one or zero hydrogen bonds coming from the C3' and C5' hydroxyl moieties. This leads to the three different situations, as presented in Figure 3, where C4'-radical (DHB) stands for the situation when the hydrogen bond is formed with both C3' and C5' hydroxyl hydrogens. The C3' and C4' radicals assume existence of only one hydrogen bond with a near-situated hydroxyl group.



Figure 3. Differences in hydrogen bond enthalpy.

The results are presented below, while the full list of enthalpies is appended to the Supplementary Materials (Figure S3):

For all compounds except myricetin, one can see that the average value of the hydrogen bond stabilization energy is about 4 kcal/mol, regardless of the radical site. For myricetin, C3' stabilization energy without crossed hydrogen bond is lower by 1.5 kcal/mol than any other C3' HBE, because oxygen electronegative repulsion happens over a short distance (2.718Å), decreasing hydrogen bond stability. However, if all hydrogen bonds are present, this energy difference increases to 3.4 kcal/mol, still being lower than for fisetin, luteolin or quercetin. For the C4' radical, the stabilization energy is lower than the average, because electrons of at least one oxygen interact with the radical at C4' (2.702Å). If there are two hydrogen bonds, the stabilization energy is much greater, reaching up to 7 kcal/mol.

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3.5. Catechol Moiety

Sroka et al. [35] observed an interesting behavior of luteolin, in comparison to structurally similar apigenin. The only difference between them is the existence of a 3,4-diOH catechol moiety in luteolin, and this small dissimilarity resulted in a nearly 100 times greater activity of the compound. Formation of a diradical and its rearrangement to 1,2-benzoquinone is widely stated as an explanation for the antioxidative scavenging potential difference between these already-mentioned structures. Lin et al. [24], in their experimental study, found a large difference of activity between the C3',C4'-dihydroxyl moiety flavonoids and those with a single C4' residue. They have proposed a mechanism explaining this diversity (see Figure 4): (a) a hydrogen abstraction from the most favored position, herein indicated by the HAE value; (b) a hydrogen transfer from the C4' to the C3', if allowed; (c) a second hydrogen abstraction from the C3' hydroxyl group; (d) an intramolecular reorganization and the hydroquinone formation. Within this section, it was investigated in a strict thermochemical scope.



Figure 4. Example of catechol moiety hydrogen swap reaction and reorganization mechanism for C3.

3.5.1. Intramolecular Hydrogen Swap

For all possible B-ring radicals, a synchronous transit-guided quasi-Newton method [67] was conducted to search potential energy surface for imaginary frequencies, corresponding to the presented movement of the hydrogen atom (b). Each compound demonstrated exactly one imaginary value. Next, the optimization to a transition state showed that it is similar to the dioxolane, where the hydrogen atom is suspended between oxygen atoms.

To describe these processes in a thermochemical way, Gibb's free energy was calculated according to the following formula [Equation (\Re)];

$$\Delta_r G^0(298K) = \sum (\varepsilon_0 + G_{corr})_{products} - \sum (\varepsilon_0 + G_{corr})_{reactants}$$
(12)

where:

- $\Delta_r G^0(298K)$ is Gibb's free energy of the reaction, at 298 K (25 °C) and pressure of 1 atmosphere.
- ε₀ is total electronic energy [Hartree].
- *G*_{corr} is thermal free energy [Hartree].

All values necessary for thermochemical calculations and to plot the reactions profile (Figure 5) are presented in Table S3.



Figure 5. Reactions profile of intramolecular hydrogen swap.

According to Table 6, where HAEs are noted, the hydrogen abstraction from fisetin should take place at C3', while for luteolin, myricetin and quercetin, position C4' is preferred. In consequence, corresponding radicals will be created. The Gibb's free energies calculated for these hydrogen swap reactions are respectively -2.5 kcal/mol, 1.9 kcal/mol, 7.5 kcal/mol and 1.9 kcal/mol. The value obtained for fisetin fits with an assumption that the hydrogen from C4' will likely trade for the free electron at C3'. On the other hand, it can be also stated that the reverse process, the hydrogen swap from C3' to C4', is not going to happen.

If hydrogen abstraction takes place at C3' of luteolin, myricetin or quercetin, an intramolecular swap would happen as well. This allows us to state that despite of investigated compound, if the C4' radical can be formed by the movement of hydrogen from C3' or C5', this will occur, the most stable radical will be formed, and the reaction will be favored thermodynamically. This assumption needs to be tested also on different compounds-e.g., phenolic acids or anthocyanins.

3.5.2. Diradical Formation

To investigate a possible mechanism of the second hydrogen abstraction, the distinctive determinants were calculated for diradical unrelaxed structures, in the same way that radicals were (Table 8).

		HAT	ET	РТ	SPLET		
Flavo	noid	BDE	IP	PDE	PA	ETE	
Luteolin		103.6	-282.6	416.1	26.4	107.0	
Fise	tin	91.6	-287.8	409.2	17.1	104.3	
Manufaction	C3'-C4'	90.5	201.0	405.3	19.7	100.6	
Myricetin	C4'-C5'	83.7	-284.9	398.5	15.5	98.1	
Quero	etin	93.8	-285.7	409.4	18.5	105.2	

Table 8. Enthalpies of diradical formation mechanisms [kcal/mol].

Bond dissociation enthalpies are usually larger than for the single radical formation, and only luteolin behaves differently. This is correct according to intuition and a knowledge of chemistry-that creating a structure with two unpaired electrons requires more energy. It is not likely that second hydrogen abstraction occurs this way.

Interesting values were achieved for the first step of the ETPT mechanism, because formation of a cation-radical releases nearly 300 kcal/mol. On the other hand, proton dissociation requires nearly 400 kcal/mol; therefore, total average enthalpy required for this process is about 120 kcal/mol. According to the statements made during the investigation into single radicals, the first step shows the thermodynamically preferred pathway. For this reason, the ETPT mechanism is assumed to be responsible for the diradical formation. It is probable that, in studies with explicit water solvent molecules, where hydrogen bonds are involved, the PDE would decrease too.

The first step of the SPLET mechanism has lower enthalpy values than for a single radical, but the second one has higher values. Proton affinities are lower than corresponding values in the single radical formation, and electron transfer energies are quite larger.

3.5.3. o-Hydroquinone Formation

The reorganization of diradicals into o-hydroquinone was determined the same way as the intramolecular hydrogen swap earlier (thermochemical values are presented in Table S4 and shown in Table 9, where o-HFE stands for o-hydroquinone formation enthalpy from diradicals). The results show that the intramolecular reorganization into o-hydroquinone happens since each Gibb's free energy is negative, especially for fisetin, myricetin C3'–C4' and quercetin. O-hydroquinone products were experimentally found by Maini et al. [68].

Table 9. o-Hydrod	quinone formation	parameters	[kcal/mol]	ŀ
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Flavo	noid	$\Delta_r G^0$	o-HFE	RE	Σ
Lute	olin	-5.0	-5.6	-34.4	-40.0
Fise	etin	-13.2	-14.4	-15.7	30.1
Myricetin	C3'-C4'	-11.9	-11.9	-14.4	-26.3
Myricetin	C4'-C5'	-7.5	-8.2	-13.8	-22.0
Quer	cetin	15.1	-14.4	-15.7	-30.1

The biggest o-hydroquinone formation enthalpy is also denoted for the three stated compounds—all are below -10 kcal/mol. Relaxation of the molecule decreases the system's total enthalpy even more. The reorganization enthalpy of luteolin is nearly -35 kcal/mol. Interestingly, the structure, instead of becoming more planar, increases its dihedral angle to 29.6° .

These values are even greater if o-HFE and RE are summed up. In the end, the whole mechanism of catechol moiety was described.

4. Conclusions

Within this research, flavonols and flavones were analyzed with the B3LYP/6-31+G(d,p) level of computational chemistry theory, and structure–activity relationship dependencies were proposed. First of all, it was noted that free p_y orbitals play a role when the radical is formed. They contribute to the electron delocalization by resonance and hyperconjugation effects, but only in AC- or BC-ring complexes, depending on the hydrogen abstraction site. The reason for this is the cross-conjugation effect of the carbonyl residue. Nevertheless, there is no possibility that hydrogen density would be exchanged between them in any manner—neither directly, nor indirectly via hydrogen atom exchange. Because of this, most flavonols and flavones adapt geometries, for which AC- and B-rings are coplanar, resulting in conjugation enhancement. It was noticed that an intramolecular hydrogen bonding can be even more important, as flavonoids with two or three B-ring hydroxyl groups close together prefer to detach the hydrogen atom from the B-ring instead of the AC-ring, as suggested by the reorganization enhalpy values. Each additional hydrogen bond guarantees greater reduction of the system's total enthalpy due to the stabilization effect. Moreover, if the structure involves a dicatechol moiety, it is likely to form the hydroquinone form via the diradical intermediate state, where ETPT plays a role when the diradical is going to be formed. Thermodynamically, the most favored mechanism of action for

the first hydrogen abstraction in a polar solvent is a C7 SPLET abstraction. On the other hand HAE, by including reorganization enthalpy correction, we see that C3 for flavonols and C4' for flavones (especially if a C3' or C5' hydroxyl group is present) are most favorable.

Supplementary Materials: The following data is available online at http://www.mdpi.com/2076-3921/9/6/461/s1, Figure S1: HOMOs (lower) and LUMOs (upper) visualization with isovalue 0.05, Figure S2: Radicals' Mulliken spin densities with hydrogen summed into heavy atoms, Figure S3: Enthalpies of flavonoids' radicals with and without hydrogen bond stabilization (isomer without H-bond is marked *) (a.u.), Table S1: Enthalpies of the flavonoids' lowest energetic isomer at B3LYP/6-31+G(d,p) level of theory (a.u.), Table S2: Enthalpies of unrelaxed and relaxed radicals of investigated compounds (a.u.), Table S3: Radicals thermochemical values (a.u.), Table S4: Diradicals thermochemical values (a.u.). All geometries are deposited as xyz files in the online repository accessible via link http://dx.doi.org/10.17632/njz3gx3w2d.

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Appendix A. Electron Density Swap

As mentioned earlier, AC- and BC-electron complexes are not in a conjugated system. Thus, electron density is locked in one of them, depending on which the hydroxyl group hydrogen was removed. Looking at Lewis structures (Figure 1) and LUMOs (Figure S1) gives an insight into delocalization, indicating that radicals' electron density is concentrated on carbonyl oxygen, especially for the C4' radical. Amić et al. [29] and Heijnen et al. [17] suggested a mechanism of activity, where the hydrogen atom can swap from the C5 position to the carbonyl in exchange for electron density. It was ascertained whether such a transfer from the one complex to another can occur, based on the following mechanism (Figure A1):



Figure A1. Electron density swap mechanism.

Elaboration of the problem was similar to that presented for a catechol moiety. No imaginary frequency was found. For this reason, a different procedure was used—beside the polarizable continuum model, a water molecule was placed near (≈ 2.000 Å) to the carbonyl oxygen radical and appropriate hydroxyl group. It was thought that it could serve as a hydrogen bridge between carbonyl oxygen. Based on the calculations, the swap mechanism illustrated in Figure A1 cannot be supported. It seems that electron density cannot be moved between A- and C-rings to achieve further stabilization, at least for the model studied here. This finding emphasizes the importance of the B-ring's substitution pattern for flavonoids' antioxidative potential.

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Appendix B. Frontier Molecular Orbitals Theory

Appendix B.1. Highest Occupied (HOMO) and Lowest Unoccupied (LUMO) Molecular Orbitals

HOMO and LUMO, relatively, are the fundamentals of electron-donating and electron-accepting characteristics in the frontier molecular orbitals theory. The higher the HOMO energy and the smaller the energy gap between HOMO and LUMO, the better the reducing agent is [69]. Since flavonoids operate via hydrogen particle donation, which is coupled with the bound electron transfer, an electron affinity [EA; Equation (A1)] was calculated, and a chemical hardness [η ; Equation (A2)] [70] was checked and noted (Table A1). Evaluation of this property was performed according to the following statements:

$$ArOH + e^{-} \rightarrow ArOH^{-}$$
$$EA = H(ArOH^{-}) - H(e^{-}) - H(ArOH)$$
(A1)

$$\eta = \frac{IP - EA}{2} \tag{A2}$$

 E_{HOMO} values do not differ significantly, and the largest can be found for fisetin, kaempferol, myricetin and quercetin. Since chemical hardness tends to be a valuable descriptor of hydrogen donating predisposition [71], flavonols were organized into three classes according to their E_{HOMO} values: low active (L), moderate active (M) and highly active (H). With chemical hardness studies, it can be concluded that flavonoids are hard acids, in the hard and soft acids and bases (HSAB) concept, especially flavonols. Results obtained from these calculations indicate the C7 hydroxyl group as a favorable hydrogen abstraction site. This pattern coincides with IP values from Table 4; similar conclusions were made by Mazzone et al. during hydroxycinnamic acid investigations [72].

Table A1. Energies of HOMO, LUMO and chemical hardness of investigated flavonoids [eV].

							Inl			
I	Flavonoids	E _{HOMO}	E _{LUMO}	C2′	C3′	C4′	C5′	C3	C5	C7
	Acacetin	-6.270 (M)	-2.217						4.572	4.265
	Apigenin	-6.319 (L)	-2.220			4.263			4.512	4.255
Jes	Chrysin	-6.460 (L)	-2.320						4.456	4.170
IOV	Chrysoeriol	-6.135 (M)	-2.221			4.374			4.625	4.319
Fla	Diosmetin	-6.151 (M)	-2.222		4.439				4.624	4.310
_	Genkwanin	-6.329 (L)	-2.195			4.245			4.565	
	Luteolin	-6.215 (M)	-2.240		4.393	4.213			4.565	4.297
	Fisetin	-5.924 (H)	-2.250		4.413	4.487		4.581		4.398
ls	Galangin	-6.172 (M)	-2.373					4.394	4.514	4.288
ouo	Kaempferol	-5.982 (H)	-2.283			4.415		4.491	4.631	4.374
avc	Morin	-6.141 (M)	-2.285	4.505		4.350		4.270	4.534	4.295
Ē	Myricetin	-5.941 (H)	-2.315		4.527	4.377	4.399	4.503	4.638	4.396
	Quercetin	-5.934 (H)	-2.304		4.544	4.340		4.518	4.647	4.400

A visualization of the results (Figure S1) pinpoints the electron density of HOMO on the B-ring, the C2–C3 double bond and the hydroxyl oxygen atoms. Absence of the hydroxyl groups in the B-ring moves density to the A-ring (e.g., chrysin). The C4' hydroxyl group is always occupied with a great amount of electron density, even if the adjacent hydroxyl group is not (e.g., myricetin). This is in agreement with the statement that C4' is the most favored abstraction position. Examination of the LUMO reveals how conjugation between the B-ring and the C2–C3 double bond is formed—one can observe a π -bond swap characteristic for the conjugated system, and creation of C2–C1' and C3–C4 bonds. Apparently, the electron density arises on the chromone and carbonyl group oxygens. Morin is an exception because B- and C-ring p_V orbitals do not overlap, and thus electrons are retained on them.

Appendix B.2. Single Occupied Molecular Orbital (SOMO)

Like HOMOs of molecules, SOMOs can be found when radical structures are described. Being occupied by an unpaired electron, these structures are much more reactive than corresponding neutral particles. Simply, to check whether the particle is nucleophilic or electrophilic, one should compare the radical's E_{SOMO} with the E_{HOMO} and E_{LUMO} of the targeted compound. If the E_{SOMO} is closer to the E_{LUMO} , then the radical will act as nucleophile; otherwise its electrophile properties are demonstrated.

Presented in Table A2, E_{SOMO} is similar to the E_{HOMO} of flavonoids they derive from. Since E_{SOMO} is slightly lower than E_{HOMO} for the corresponding molecule, it may suggest they retain electron-donating properties, and would willingly detach another hydrogen to scavenge another radical. DFT results indicate each SOMO was occupied exactly by one electron.

					ESOMO			
Flavonoids		C2′	C3′	C4′	C5'	C3	C5	C7
Flavones	Acacetin						-6.367	-6.502
	Apigenin			-6.570			-6.456	-6.586
	Chrysin						-6.699	-6.925
	Chrysoeriol			-6.470			-6.205	-6.320
	Diosmetin		-6.471				-6.216	-6.319
	Genkwanin			-6.594			-6.429	
	Luteolin		-6.494	-6.557			-6.343	-6.440
Flavonols	Fisetin		-6.329	-6.290		-6.212		-6.195
	Galangin					-6.463	-6.429	-6.573
	Kaempferol			-6.356		-6.230	-6.176	-6.336
	Morin	-6.237		-6.489		-6.400	-6.305	-6.445
	Myricetin	-6.202	-6.246	-6.265	-6.268		-6.102	-6.258
	Quercetin	-6.189	-6.224	-6.292			-6.122	-6.270

Table A2. Energies of SOMOs of investigated flavonoids' radicals [eV].

According to the obtained results (Table A3), it is clear that luteolin (flavone) SOMO orbitals are quasi-degenerated. The energy gap between them is much lower than for flavonols, where it reaches approximately 0.7–1.2 eV. Moreover, their energies are even lower than for the molecule's HOMO or the radical's SOMO, which the diradical originates from. As a result, SOMO with energy –6.704 eV conforms to other examined diradicals' lowest energetic SOMO.

Table A3. Energies	of SOMOs of investigated flavonoids' dir	adicals	[eV]	ŀ
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i. <u>Materiały dodatkowe</u>





Quercetin



41





Kaempferol C5	Kaempferol C7	Luteolin C3'
OH 0.004	OH _{0.012}	OH ^{0.069}
-0.012 HO	0.287 0 0.031 0	-0.000 HO 0.003 O 0.359
C -0.027	C:-0.070 OH 0.038	C -0.003
0.295 O O -0.015	-0.003 OH O -0.017	-0.000 OH O -0.003
Luteolin C4'	Luteolin C5	Luteolin C7
O _{0.289}	OH-0.000	OH 0.004
0.002 HO 0.005 O OH 0.073	0.001 HO 0.000 O OH 0.000	0.362 O OH -0.000
C -0.019	C	C -0.055
0.002 OH Ö 0.024	0.579 O O 0.356	-0.012 OH Ö _{0.031}
Morin C2'	Morin C3	Morin C4'
UD -0.002 D	0.023 - OH0.021	0.311
0.003 HO	0.007 HO 0.023 O OH 0.042	0.004 HO
C -0.008 0.028	0.022 C	OH 0.033
0.003 OH 0 0.026	0.010 OH 00.053	0.004 OH O 0.032
Morin C5	Morin C7	
HO 0.039 O	0.0250	OH ^{0.031}
-0.012 HO OH_0.001	0.317 OH 0.004	0.005 HO 0.037 O OH -0.003
C-0.032H0.017	C-0.06H 0.022	p.010 C O.298
0.299 0 0.0011	-0.006 OH 0 -0.004	0.008 OH 0 0.042
Mound antim C2/	Marri antin CA	Mauri estin CE
Myricetin C3′	Myricetin C4'	Myricetin C5
Myricetin C3'	Myricetin C4'	Myricetin C5
Myricetin C3' 0.348 j 0.000 HO - 0.002 O - OH ^{0.083} OH ^{0.083}	Myricetin C4'	Myricetin C5
Myricetin C3' 0.348 0 0.000 HO 0.001 OH 0.001 OH	Myricetin C4'	Myricetin C5
Myricetin C3'		Myricetin C5
Myricetin C3' 0.348 0 040 083 0.000 HO 040 083 0.000 OH 000 0HO 0.000 OH <th>Myricetin C4'</th> <th>Myricetin C5</th>	Myricetin C4'	Myricetin C5
Myricetin C3' 0.348 0 040.083 0.000 H0 040.083 0.000 H0 040.083 0.000 H0 000 0 0.001 H0 000 0 0.001 H0 000 0		
$\frac{Myricetin C3'}{\overset{0.348}{0}, 0H^{0.083}}$		
$\frac{Myricetin C3'}{\overset{0.348}{_{(-)}}}$	$\mathbf{Myricetin C4'}_{0.04}$	
$\frac{Myricetin C3'}{\overset{0.346}{0}, 000} + 0^{0.003} + 0^{$	$\mathbf{Myricetin C4'}$	$Myricetin C5$ $\begin{array}{c} & & & & & \\ & & & & \\ & & & & & $
$\mathbf{Myricetin C3'}$ $\overset{0.346}{\longrightarrow} (P^{0.003} C^{0.000} P^{0.003} C^{0.000} C^{0.003} C^{0.000} C^{0.00$	$\mathbf{Myricetin C4'}$ $\overset{0.047}{\longrightarrow} (0.288)$ $\overset{0.004}{\longrightarrow} (0.004) (0.004) (0.004)$ $\overset{0.000}{\longrightarrow} (0.004) (0.004) (0.004)$ $\mathbf{Myricetin C7}$ $\overset{0.000}{\longrightarrow} (0.005) (0.007) (0.001) ($	
	$\mathbf{Myricetin C4'}$ $\overset{0.047}{\longrightarrow} (0.288)$ $\overset{0.004}{\longrightarrow} (0.004) (0.004) (0.004)$ $\mathbf{Myricetin C7}$ $\overset{0.000}{\longrightarrow} (0.004) (0.007) (0.004)$ $\mathbf{Myricetin C7}$ $\overset{0.000}{\longrightarrow} (0.007) (0.007) (0.007)$ $\mathbf{Quercetin C4'}$ $\overset{0.005}{\longrightarrow} (0.007) (0.007) (0.007) (0.007) (0.007)$	
$ \begin{array}{c} \textbf{Myricetin C3'} \\ & & & & & & & & & & & & & & & & & & $		
Hyricetin C3' $(0.000 + 0) + (0.003) + (0.0$		
Hyricetin C3' $(0.000 + 0) + (0.003) + (0.0$	$ \begin{array}{c} \textbf{Myricetin C4'} \\ & & & & & & \\ & & & & & \\ & & & & & & \\ & & & & & \\ & & & & & & \\ & & & & &$	
Hyricetin C3' $(0.000 + 0) + (0.003) + (0.0$	$ \begin{array}{c} \textbf{Myricetin C4'} \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & & \\ & & & & & & \\ & & & & & & & \\ & & & & &$	$\mathbf{Myricetin C5}$ $(\mathbf{Myricetin C5})$ $(\mathbf{Myricetin C5})$ $(\mathbf{Myricetin C5})$ $(\mathbf{Myricetin C5})$ $\mathbf{Quercetin C3}$ $(\mathbf{Myricetin C5})$ $(Myric$
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Figure 2. Radicals' Mulliken spin densities with hydrogen summed into heavy atoms



Figure 3. Enthalpies of flavonoids' radicals with and without hydrogen bond stabilization (isomer without H–bond is marked *) [a.u.]

Tables

Table 1. Enthalpies of the flavonoids' lowest energetic isomer at B3LYP/6–31+G(d,p) level of theory [a.u.]

A		
	Flavonoid	Enthalpy
	Acacetin	-992.861147
	Apigenin	-953.589415
Jes	Chrysin	-878.363148
IOV	Chrysoeriol	-1068.084598
Fla	Diosmetin	-1068.084312
	Genkwanin	-992.861115
	Luteolin	-1028.811831
	Fisetin	-1028.799467
ls	Galangin	-953.582425
ouo	Kaempferol	-1028.808400
avc	Morin	-1104.033327
FI	Myricetin	-1179.253341
	Quercetin	-1104.030864

Table 2. Enthalpies of unrelaxed and relaxed radicals of investigated compounds [a.u.]

	Flavonoids		Unrelaxed radical	Relaxed radical
	Acacotin	C5	-992,199222	-992,213026
	Acacetin	C7	-992,216822	-992,224983
		C4′	-952,947947	-952,958621
	Apigenin	C5	-952,930000	-952,941269
		C7	-952,944139	-952,952039
	<u> </u>	C5	-877,701313	-877,715043
	Chrysin	C7	-877,716141	-877,725026
		C4′	-1067,446832	-1067,459122
sər	Chrysoeriol	C5	-1067,415336	-1067,419226
ION		C7	-1067,439373	-1067,447632
Fla		C3′	-1067,445198	-1067,456637
	Diosmetin	C5	-1067,414907	-1067,418746
		C7	-1067,438585	-1067,447177
	Contranin	C4′	-992,220493	-992,230596
	Genkwanin	C5	-992,199106	-992,212595
		C3′	-1028,172077	-1028,183270
	Teste alla	C4'	-1028,178156	-1028,191782
	Luteolin	C5	-1028,143630	-1028,146283
		C7	-1028,166412	-1028,174353
		C3′	-1028,167323	-1028,179343
ols	Figation	C4′	-1028,164704	-1028,177094
VOL	risetin	C3	-1028,162134	-1028,176690
Fla		C7	-1028,156325	-1028,165763
	Galangin	C3	-952,944720	-952,958463
	<u> </u>			

	C5	-952,926275	-952,939686
	C7	-952,937061	-952,946355
	C4′	-1028,171031	-1028,182027
K (1	C3	-1028,173008	-1028,186897
Kaempieroi	C5	-1028,152508	-1028,166096
	C7	-1028,165319	-1028,174020
	C2′	-1103,385259	-1103,398629
	C4'	-1103,392680	-1103,402521
Morin	C3	-1103,399249	-1103,419264
	C5	-1103,376267	-1103,390761
	C7	-1103,388420	-1103,396868
	C3′	-1178,614708	-1178,625589
	C4'	-1178,625608	-1178,638759
Marricotin	C5′	-1178,619670	-1178,631215
wryffceun	C3	-1178,616992	-1178,631960
	C5	-1178,597496	-1178,611019
	C7	-1178,609123	-1178,618703
	C3′	-1103,392412	-1103,403587
	C4'	-1103,401993	-1103,414770
Quercetin	C3	-1103,395481	-1103,409514
	C5	-1103,375549	-1103,389360
	C7	-1103,387556	-1103,396441

Table 3. Radicals thermochemical values [a.u.]

Flavonoids	Radical	€0+Hcorr	€0+Gcorr
	C3′	-1028.190	-1028.252
Luteolin	TS	-1028.179	-1028.240
	C4′	-1028.192	-1028.255
	C3′	-1028.179	-1028.243
Fisetin	TS	-1028.170	-1028.232
	C4′	-1028.184	-1028.247
	C3'=C5'	-1178.631	-1178.699
Myricetin	TS	-1178.623	-1178.691
	C4′	-1178.643	-1178.711
	C3′	-1103.410	-1103.477
Quercetin	TS	-1103.401	-1103.466
	C4′	-1103.415	-1103.480

Flavor	noids	6 0 ⁺	•Hcorr ⁱ	€0+Gcorr
T	.1:	-10	27.514	-1027.574
Luteo	oun	-1027.523	<u>-1027.578 (r)</u>	-1027.582
Fice	<i>Li</i> .,	-10	27.522	-1027.583
F ISE	un	-1027.545	<u>-1027.570 (r)</u>	-1027.604
	C2' CA'	-11	77.982	-1178.046
Manicotin	05-04	-1178.001	<u>-1178.024 (r)</u>	-1178.065
wiyncenn	CAL CEL	-11	77.993	-1178.057
	C4 -C3	-1178.006	<u>-1178.028 (r)</u>	<u>-1178.069</u>
Ouar	atin	-11	02.753	-1102.815
Quero	eun	-1102.776	-1102.801 (r)	-1102.839

ⁱ (*r*) marks for the relaxed forms of o-hydroquinone.

" Underlined numbers stand for o-hydroquinone unrelaxed form.



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ii. <u>Oświadczenia współautorów</u>

Wrocław, 14.04.2023

mgr farm. Maciej Spiegel Katedra i Zakład Farmakognozji i Leku Roślinnego Uniwersytet Medyczny im. Piastów Śląskich we Wrocławiu

OŚWIADCZENIE WSPÓŁAUTORA CO-AUTHORSHIP STATEMENT

Jako współautor pracy: As a co-author of the research paper:

Spiegel, M., Andruniów, T., & Sroka, Z. (2020). Flavones' and flavonols' antiradical structure– activity relationship—a quantum chemical study. Antioxidants, 9(6), 461.

oświadczam, że mój udział polegał na:

I declare that my contribution consists of: • sformułowaniu i ewolucji nadrzędnych celów i zadań badawczych

- formulation and evolution of overarching research goals and aims
- zastosowaniu technik matematycznych i obliczeniowych do analizy i syntezy danych z badań application of mathematical and computational techniques to analyze and synthesize study data
- prowadzeniu procesu badawczego i dochodzeniowego oraz gromadzeniu danych
- conducting a research and investigation process, and data collection
 rozwoju i zaprojektowaniu metodologii
- development and design of methodology
- weryfikacji ogólnej powtarzalności wyników badań verification of the overall reproducibility of research outputs
- przygotowaniu i stworzeniu opublikowanej pracy, w szczególności wizualizacji i prezentacji danych preparation and creation of the published work, specifically visualization & data presentation
- przygotowaniu i stworzeniu opublikowanej pracy, w szczególności napisaniu jej wstępnego szkicu
 preparation and creation of the published work, specifically writing the initial draft

rel Macies

Podpis współautora Uniwersytet Medyczffy Wertwo (signuture) KATEDRA I ZAKŁAD FARMAKOGNOZJI I LEKU ROŚLINNEGO kierownik prof. dr hab. Zbigniew Signature) (2) Supervisor 3 signature)

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Wanders, 12. 12. 2022 miejscowość, data

(place, date)

tytuł, imię i nazwisko (title, name and surname)

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miejsce zatrudnienia (affiliation)

OŚWIADCZENIE WSPÓŁAUTORA **CO-AUTHORSHIP STATEMENT**

Jako współautor pracy: As a co-author of the research paper:

> Spiegel, M., Andruniów, T., & Sroka, Z. (2020). Flavones' and flavonols' antiradical structure- activity relationship-a quantum chemical study Antioxidants, 9(6), 461.

oświadczam, że mój udział polegał na: I declare that my contribution consists of:

> • opracowaniu i zaprojektowaniu metodologii badań

development and design of the research methodology

nadzorze i odpowiedzialność kierowniczej za planowanie i realizację działań badawczych

oversight and leadership responsibility for the research activity planning and execution

weryfikacji ogólnej odtwarzalności wyników

verification of the overall reproducibility of results

przygotowaniu ostatecznej wersji manuskryptu; w szczególności korekty

preparation of the final version of the manuscript; in particular post-review corrections

Andruniow T. Andruniow

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Wrocław, 18.04.2023 miejscowość, data (place, date)

prof. dr hab. Zbigniew Sroka tytuł, imię i nazwisko (title, name and surname)

Katedra i Zakład Farmakognozji i Leku Roślinnego Uniwersytet Medyczny im. Piastów Śląskich we Wrocławiu miejsce zatrudnienia (affiliation)

OŚWIADCZENIE WSPÓŁAUTORA CO-AUTHORSHIP STATEMENT

Jako współautor pracy: As a co-author of the research paper:

> Spiegel, M., Andruniów, T., & Sroka, Z. (2020). Flavones' and flavonols' antiradical structure– activity relationship—a quantum chemical study Antioxidants, 9(6), 461.

oświadczam, że mój udział polegał na: I declare that my contribution consists of:

- uzyskaniu wsparcia finansowego dla projektu prowadzącego do powstania niniejszej publikacji acquisition of the financial support for the project leading to this publication
- nadzorze i odpowiedzialności kierowniczej za planowanie i realizację działań badawczych oversight and leadership responsibility for the research activity planning and execution
- przygotowaniu ostatecznej wersji manuskryptu; w szczególności korekty po recenzji preparation of the final version of the manuscript; in particular post-review corrections

podpis współautora (co-author's signature)

Uniwersytet Medyczny wo Wroelawiu KATEDRA I ZAKLAD FARMAKOGMOZJU TEKU ROSI UNISEOSignature) kiejownik prof. dr hab. Zbigniew Sroka (2)

II. Publikacja [B]



Article



Antioxidant Activity of Selected Phenolic Acids–Ferric Reducing Antioxidant Power Assay and QSAR Analysis of the Structural Features

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Abstract: Phenolic acids are naturally occurring compounds that are known for their antioxidant and antiradical activity. We present experimental and theoretical studies on the antioxidant potential of the set of 22 phenolic acids with different models of hydroxylation and methoxylation of aromatic rings. Ferric reducing antioxidant power assay was used to evaluate this property. 2,3-dihydroxybenzoic acid was found to be the strongest antioxidant, while mono hydroxylated and methoxylated structures had the lowest activities. A comprehensive structure–activity investigation with density functional theory methods elucidated the influence of compounds topology, resonance stabilization, and intramolecular hydrogen bonding on the exhibited activity. The key factor was found to be a presence of two or more hydroxyl groups being located in *ortho* or *para* position to each other. Finally, the quantitative structure–activity relationship approach was used to build a multiple linear regression model describing the dependence of antioxidant activity on structure of compounds, using features exclusively related to their topology. Coefficients of determination for training set and for the test set equaled 0.9918 and 0.9993 respectively, and Q2 value for leave-one-out was 0.9716. In addition, the presented model was used to predict activities of phenolic acids that haven't been tested here experimentally.

Keywords: phenolic acids; polyphenols; antioxidants; ferric reducing antioxidant power (FRAP) assay; density functional theory (DFT); quantitative structure–activity relationship (QSAR); multiple linear regression (MLR)

1. Introduction

Nowadays, the popularity of the healthy foods has led to the revival of studies on phytochemicals activity. Plant substances, such as fiber regulate gastrointestinal tract function [1] and unsaturated fatty acids are capable of decreasing risk of atherosclerosis [2], and phenolic compounds can prevent oxidative stress [3]. All these compounds are crucial for the proper function of the human body. A number of important research papers concerning plant antioxidants were published in the last ten years, making an investigation of them an interesting branch of 21st century medical research [4–6].

Phenolic acids are a large group of secondary metabolites, originating from shikimic and benzoic acids [7]. They can be found commonly in plants, especially hydroxybenzoic and hydroxycinnamic

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derivatives, which are responsible for organoleptic properties, such as sour and bitter flavor. However, their true medicinal merit is an antioxidant and antiradical activity arising from their chemical structure. It is known that oxidative stress may cause damage to lipid membranes, DNA, and proteins [8], and may further lead to more severe diseases such as diabetes [9], Alzheimer's. disease [10], and Parkinson's disease [11] or neoplasms [12]. Each phenolic acid is made of one aromatic ring with hydroxyl residues and carboxyl residue linked to it. Acting as a donor of a hydrogen atom or single electron, they are capable of neutralizing reactive oxygen species (ROS), reducing transition metals responsible for Fenton's reaction, and overall decreasing oxidative stress. Even though terms "antiradical" and "antioxidant" are often referred to the same property, these activities do not necessarily coincide. Thus, antiradical activity must be clearly distinguished from the antioxidant one [13]. While the first characterizes the ability of compounds to scavenge free radicals (for instance cation radical ABTS^{•+} and a stable radical DPPH[•]), the second one represents the ability to inhibit the process of oxidation. Measurement of antiradical activity most commonly is performed using ABTS or DPPH tests. During the ABTS method, proposed by Re et al. [14], pre-generated dark-green cation radical is reduced by a hydrogen-donating compound, such as phenolic acid. In the DPPH test [15], the radical undergoes a reaction with a reducer, becoming a neutral molecule. Both these assays rely on hydrogen exchange mechanism. As for the antioxidant activity, a wide variety of methods may be used, including biological assays, such as cellular antioxidant activity (CAA) [16] and chemical-based methods (FRAP [17], CUPRAC assays [18], etc.). While biological assays are considered to be more appropriate, they also are more expensive and time-consuming compare to chemical-based methods [16]. With all variety of methods results of efficiency measurements for phenolic acids is found to be slightly controversial in the literature [19,20].

Ferric Reducing Antioxidant Power assay (FRAP) [17] is based on reduction of a colorless Fe³⁺-TPTZ complex into intense blue Fe²⁺-TPTZ once it interacts with a potential antioxidant. At low cost, this method showed to be useful for screening of antioxidant capacities and comparing efficiencies of different compounds. Thus, in this study, we used FRAP method for an investigation of antioxidant activity of selected phenolic acids. Interestingly, the exact mechanism of the antioxidant activity for these compounds, and the influence of the compounds' structure on their activity, is still not fully elucidated and controversial in the literature [21-24]. Computational chemistry proved to be a good support for experimental investigations. A great review of strategies in theoretical antioxidants activity research has been recently published by Galano et al. [25]. Not only hydrogen atom transfer (HAT) mechanism, but also sequential proton loss-electron transfer (SPLET) and single electron transfer-proton transfer (SET-PT) are widely studied with density functional theory (DFT) methods to elaborate the most probable mechanism of action of antioxidants [21,26,27]. Obtained results clearly showed that SPLET is the most favorable mechanism of action in a polar solvent, whilst HAT dominates in benzene. Presented computational results complement the experimental studies by explaining rationale used in the experiment. That indicates the importance of computational chemistry methods as a supporting tool in every modern-age research.

Quantitative structure–activity relationship is widely used to find a relationship between structural features of compounds and their activities [28–30]. Finding proper descriptors used to develop a quantitative structure–activity relationship (QSAR) model is the very first step one should consider. For example, to build a model for wine polyphenols, Rastija et al. [31] used lipophilicity, Balaban index, Balaban-type index, and 3D GETAWAY descriptors. Gupta et al. [32] focused on MOLMAP descriptors selected by genetic algorithms. Filipović et al. [33] in their studies on free radical scavenging potency of 21 phenolic acids and simple phenolics proposed three models based on the number of vicinal hydroxyl groups, bond dissociation energy, proton affinity and electron transfer enthalpy. QSAR concept also was used by Chen et al. [21] in their studies on thermodynamic properties as descriptors for prediction of DPPH radical scavenging assay. These models provided a good insight into the nature of antiradical and antioxidant activity, though they require an involvement of quantum chemical calculations or other software for descriptor generation. The aim of this study is to perform

a comprehensive investigation of antioxidants nature for the set of phenolic acids with different models of hydroxylation, and to develop a QSAR model for prediction of these properties based on a topology of tested compounds. FRAP assay has been used to measure antioxidant potential of selected phenolic acids. Purely topological descriptors used in this paper are easy to generate and give an understanding on how structural features of studied compounds influence an activity.

2. Results and Discussion

2.1. Experimental Results

Antioxidant properties of phenolic acids and their structures are presented in Table 1. The strongest activity reducing ion Fe³⁺ to Fe²⁺ was noted for 2,3-dihydroxybenzoic acid. The number of units per µmol of compound ($TAU_{Fe/µmol}$) was equal to 202 ± 10.6. A slightly lower activity was observed for 3,4-dihydroxybenylacetic (149 ± 10.0) and 2,5-dihydroxybenzoic acid (128 ± 6.3), while 3-hydroxybenzoic acid possessed the poorest efficiency among tested phenolic acids.

Table 1. Structures of investigated phenolic acids.



* Averaged TAU_{Fe/µmol} values with the maximal errors.

The statistical significance of differences between samples was analyzed using HSD Tukey test. Compounds have been divided into more active (1–11) and less active (12–22) set. Statistical significance was tested on level p < 0.05. Among the first group for nearly every pair except 8–9 and 10–11 (Table S1, Figure S1) the given criterion was achieved, whereas among the second group pairs 18–19, 18–20, 18–21, 18–22, 19–20, 19–21, 19–22, 20–21, 20–22, 21–22 didn't met it (Table S2, Figure S2).

2.2. SAR Investigation

The results of measured antioxidant activities were used as a subject of structure–activity investigation. In order to explain the structure–activity relationship, three key factors must be investigated separately: mutual position of hydroxyl groups, their methylation, and the distance between phenyl and carboxylic group. These factors have been numerously mentioned in literature [21,22,34], however, the full clarification of their influence has not been reported.

2.3. Mutual Positions of Hydroxyl Groups and Resonance Stabilization

All tested phenolic acids can be clustered into two groups, one including compounds with high activities (1–11), and the other with compounds possessing extremely low activities (12–22) (Figure 1). Clustering of phenolic acids has shown a response dependence on the relative positions of the hydroxyl groups in the ring. Compounds, that contain only one hydroxyl group, (13, 15–18, 20, 22) have exhibited very low efficiency. Phenolic acids with two hydroxyls substituted in the *meta* position in relation to each other, such as 3,5-dihydroxybenzoic (19) and 2,4-dihydroxybenzoic acid (14) have also shown a poor activity, compare to ones with two hydroxyls on adjacent carbon atoms.



Figure 1. Phenolic acids clustered by their activity.

Since FRAP assay must be performed in low pH values to maintain iron solubility, the ionization potential is also low, which drives hydrogen atom transfer. FRAP assay is based on an electron transfer mechanism with formation of aryloxyl radical [35]:

$$Fe(TPTZ)_2^{3+} + ArOH \rightarrow Fe(TPTZ)_2^{2+} + ArO^{\bullet} + H^+$$

In this case, once aryloxyl radical is formed, its stability determines the efficiency of phenolic acid as an antioxidant compound. The fact that 2,4-dihydroxybenzoic acid (14) is much less efficient than 2,3-dihydroxybenzoic acid (1) can be explained by resonance structures of its radicals (Figure 2). In both cases radical is stabilized by delocalization over conjugated aromatic rings system. When the second hydroxyl group present in *ortho-* or *para-* position related to the first one, it allows second oxygen atom to participate in delocalization (Figure 2a). One can see that it leads to the activation of the second hydroxyl group. Hence, the reaction can proceed further reducing one more Fe³⁺-TPTZ complex with subsequent oxidation to 5,6-dioxo-1,3-cyclohexadiene-1-carboxylic acid. Highly active 2,5-dihydroxybenzoic acid (3) has the same resonance structures with hydroxyl groups being in *para-*position to each other. Meanwhile, the second hydroxyl of compound 14 (Figure 2b) does not participate in a delocalization, thus 2,4-dihydroxybenzoic acid exhibits significantly lower antioxidant activity. To summarize, in order to achieve high antioxidant and antiradical activity it is critical that two or more hydroxyl groups are located either in vicinal positions or on opposite sides of the ring (in *ortho* or *para* position to each other).



Figure 2. Resonance structures of phenolic acids: (a)—2,3-dihydroxybenzoic acid (1), and (b)—2,4-dihydroxybenzoic acid (14).

Low efficiency of two other compounds from the second cluster (4-hydroxy-3-methoxybenzoic acid (12) and 3,4-dimethoxybenzoic acid (21)) cannot be justified by mutual position of hydroxyl groups as similar in structure 3,4-dihydroxybenzoic acid (10) possesses relatively high efficiency as an antioxidant compound. Thus, methylation of hydroxyl groups must be considered separately.

2.4. The Influence of Methylation

The results of FRAP tests showed a tendency of the methylated compounds to have lower activity than their non-methylated counterparts (5 < 4, 21 < 10, 11 < 9, 8 < 2). Methylation yields a decrease of active electron- and hydrogen-donating groups, which consequently leads to reduced efficiency of the compound as an antioxidant. Interestingly, methylation of the cinnamic acid derivatives has no significant influence on the antioxidant activity, since 4-hydroxy-3-methoxycinnamic acid (11) demonstrates slightly lower activity than 3,4-dihydroxycinnamic one (9). With a decreasing distance between carboxylic group and a ring the influence of methylation is increasing, for example: 3,4-dihydroxyphenylacetic acid (2) is almost two times more efficient then its partially methylated counterpart (8), while the ratio between activities of 3,4-dihydroxybenzoic acid (10), 4-hydroxy-3-methoxybenzoic acid (12), and 3,4-dimethoxybenzoic one (21) amounts 598:26:1. It seems that the farther the carboxylic group is from the methoxylated ring, the more efficient the phenolic acid containing methylated hydroxyl groups. This can be explained by the importance of the inductive effect of carboxylic group. The 4-hydroxy-3,5-dimethoxybenzoic acid (5) is the only exception from the presented trend, it exhibits just slightly lower activity when compared to non-methylated counterparts (4).

2.5. Carboxylic Group Influence and H-Bonding

Antioxidant activity decreases with an increase of the carboxylic group electron-withdrawing effect on a radical delocalization. In most cases, cinnamic acid derivatives have demonstrated improved efficiency over their counterparts derived from benzoic acid. Phenolic acids substituted with hydroxyls in para-meta position have their activity decreasing in the following order phenylacetic acid > cinnamic acid > benzoic acid (2 > 9 > 10, and 8 > 11 > 12). This trend suggests that carboxylic group has the biggest influence on the total antioxidant activity. This influence occurs not through inductive effect (-I) (the distance of carboxylic group from the ring), but through the mesomeric effect (-M) (the resonance with the ring). Carboxylic group of phenylacetic acid is not participating in resonance and can influence only through -I effect, while carboxyl group of a benzoic acid shows the strongest -I and -M effects. Interestingly, phenolic acids that are substituted with only one hydroxyl in para position have demonstrated a decrease of activity presented in a row cinnamic acid > phenylacetic acid > benzoic acid (15 > 17 > 20), that may indicate the importance of inductive effect for low efficient compounds. Though, the difference in their efficiency is negligible. Meanwhile, an antioxidant activity of ortho-substituted compounds such as 2,5-dihydroxybenzoic acid (3) and 2-hydroxybenzoic acid (13) measured by FRAP have shown to be elevated when compared to its phenylacetic counterparts (2,5-dihydroxyphenylacetic (7) and 2-hydroxyphenylacetic (16) acids). In addition to the resonance stabilization (Figure 2), a radical can be stabilized by intermolecular hydrogen bonds between functional groups and polar protic solvents, as well as intramolecular hydrogen bonds [24]. There are two possible types of intramolecular hydrogen bonding for compounds tested here. One involves only hydroxyl oxygens, and the other is a hydrogen bond between carboxylic and hydroxyl groups.

One of the methods used for approximation of the hydrogen bonding energy is based on the calculation of potential energy density has been implemented in Multiwfn program package [36]. Energies (E_{HB}) for all the studied compounds, where hydrogen bonding is possible, were calculated and collected in Table 2, along with distances between critical point (*CP*) and hydrogen atom (H_{HB}), the distance between hydrogen (H_{HB}) and acceptor oxygen atom (O_{ac}) as well as the angle ($\angle O_{ac}$ -CP- H_{HB}).

Compound ID:	IUPAC Name	E _{HB} , (kcal/mol)	CP- H _{HB} Distance, (Å)	O _{ac} - H _{HB} Distance, (Å)	∠O _{ac} -CP-H _{HB} Angle, (°)
1	2,3-dihydroxybenzoic	-12.91/-5.56	0.572/0.90	1.69/2.13	172.84/160.38
2	3,4-dihydroxyphenylacetic	-5.47	0.890	2.12	163.15
3	2,5-dihydroxybenzoic	-12.45	0.580	1.71	173.16
4	3,4,5-trihydroxybenzoic	-3.61/-3.69	0.882/0.880	2.18/2.18	161.43/162.80
5	4-hydroxy-3,5-dimethoxybenzoic	-6.32	0.836	2.06	167.15
6	4-hydroxy-3,5-dimethoxycinnamic	-6.37	0.833	2.06	167.30
7	2,5-dihydroxyphenylacetic	-11.41	0.601	1.78	176.03
8	4-hydroxy-3-methoxyphenylacetic	-6.16	0.842	2.07	167.46
9	3,4-dihydroxycinnamic	-5.42	0.902	2.13	161.51
10	3,4-dihydroxybenzoic	-5.58	0.882	2.11	163.67
11	4-hydroxy-3-methoxycinnamic	-6.30	0.835	2.06	167.47
12	4-hydroxy-3-methoxybenzoic	-6.26	0.838	2.06	167.31
13	2-hydroxybenzoic	-12.98	0.572	1.69	173.31
14	2,4-dihydroxybenzoic	-15.77	0.542	1.64	173.62

 Table 2. Hydrogen bonding energies and geometrical parameters calculated using Multiwfn program package.

The highest hydrogen bond energy has been found for bonds between hydrogen of hydroxyl group and double-bonded oxygen of benzoic acid's carboxylic group (compound 1, 3, 13, 14). The lowest hydrogen bond energy is found to be between hydroxyl groups. According to the presented results, the oxygen of hydroxyl group is a less efficient H-bond acceptor when compared to double-bonded oxygen of carboxyl group. Taking into account this difference in H-bond strength, one can explain why for benzoic acid derivatives efficiency is decreasing in the row: *ortho-meta* (2,3-*pattern*) (1) > *ortho-meta* (2,5-*pattern*) (3) > *meta-meta-para* (4) > *meta-para* (10), while the same tendency does not work for phenylacetic and cinnamic acid derivatives. Similar to

2-hydroxybenzoic acid, 2,5-dihydroxyphenylacetic (7) also can be stabilized by a H-bond between carboxylic group and *ortho*-hydroxyl. However, surprisingly high H-bond energy (–11.41 kcal/mol) in the case of 2,5-dihydroxyphenylacetic one (7) does not justify its low efficiency compare to 3,4-dihydroxyphenylacetic acid (2), where energy of hydrogen bond between two hydroxyl groups is –5.47 kcal/mol. Moreover, due to the steric hindrance in case of the phenylacetic acid derivatives, H-bonding between carboxylic group and *ortho*-hydroxyl should not be that favorable as the one between two hydroxyl groups.

To verify results obtained by the method proposed in Multiwfn, an additional method was employed. Density functional theory was used to calculate enthalpies of H-bond formation. Formation of hydrogen bond in 2-hydroxybenzoic acid (13) results in release of 7.59 kcal/mol of energy (Figure 3a). The energy yield of H-bond formation in case of 3,4-dihydroxybenzoic acid (10) is significantly smaller (1.92 kcal/mol) (Figure 3b). These results are in good agreement with previously obtained data with Multiwfn method. Though, in the case of compound 7 (Figure 3c), H-bonding is not that energetically favorable (1.87 kcal/mol released) due to the steric tension occurring in this molecule. This finding is contrary to results calculated using Multiwfn method.



Figure 3. Enthalpies of hydrogen bond formation. (a)—2-hydroxybenzoic acid (13);
 (b)—3,4-dihydroxybenzoic acid (10); (c)—2,5-dihydroxyphenylacetic acid (7).

The 2,3-dihydroxybenzoic acid (1) (Figure 4a) may form two intramolecular hydrogen bonds: one involving only hydroxyl oxygen atoms and the other involving oxygen of carboxylic group (total of -18.47 kcal/mol, as is shown in Table 2). Similarly, two H-bonds stabilize a 3,4,5-trihydroxybenzoic acid (compound 4, total of -7.30 kcal/mol) (Figure 4b). Though, in this case, the carboxylic group does not participate in H-bonding. The following compounds 2,5-dihydroxybenzoic acid (3) and 3,4-dihydroxybenzoic acid (10) (Figure 4c,d) are stabilized by just one intramolecular H-bond, and only in case of 3 strong interactions with oxygen of carboxylic group takes place (-12.45 kcal/mol, compared to -5.58 kcal/mol for compound 10). It can be seen, that our study explained the trends in effectiveness. Hence, it is assumed that the model of phenolic acids' hydroxylation by two hydroxyl groups, one of each is situated next to carboxyl group (in *ortho* position) may be efficient only for benzoic acid derivatives.

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Figure 4. Intramolecular hydrogen bond stabilization of a molecule and its radical: (a)—2,3-dihydroxybenzoic acid (1); (b)—3,4,5-trihydroxybenzoic acid (4); (c)—2,5-dihydroxybenzoic acid (3); (d)—3,4-dihydroxybenzoic acid (10).

2.6. QSAR Model

Original model, Multiple linear regression model, for the prediction of the antioxidant $TAU_{Fe/\mu mol}$ activity measured by FRAP assay had one outlier (4-hydroxy-3,5-dimethoxybenzoic acid). As it was earlier discussed this compound did not follow the general trend illustrated in SAR investigation. Thus, this point was excluded from the training set and model was rebuilt. Elimination of the 4-hydroxy-3,5-dimethoxybenzoic acid had no influence on selected descriptors, though it influenced their coefficients and statistical parameters. Finalized model based on four topological descriptors is represented by Equation (1):

$TAU_{Fe/\mu mol} = -0.507 + 62.03 \cdot HOccOH(count) + 74.83 \cdot Ph_{ortho(para)-meta} + 134.49 \cdot B_{ortho-meta} + 41.804 \cdot C_{methyl}(count)$ (1)

Descriptors used for MLR model have a clear chemical meaning and were in correspondence with the SAR investigation. First and foremost, the presence of HOccOH fragment is crucial. As it was indicated above, two or more hydroxyl groups in *ortho* position to each other positively impact an antioxidant activity due to charge delocalization and OH-OH intramolecular H-bonding. Not only information about mutual position of substituents, but also its methylation is decoded by this descriptor. Only non-methylated groups positively impact on efficiency. The evidence of phenylacetic acid derivatives being more efficient than others is supported by *Phortho(para)-meta* descriptor, that indicates the presence of any *ortho-meta* or *para-meta* substitutes phenylacetic acids (compounds **2**, **7**, **8**). *Bortho-meta* is a specific descriptor that points out on a presence of *ortho-meta*-substituted benzoic acid derivatives (compounds **1** and **3**). The presence of strong intramolecular H-bonds between *ortho* hydroxyl and carboxylic groups results in significant increase of efficiency. Finally, *C_{methyl}(count*) descriptor is correcting total activity calculated by the proposed model whilst taking into account the fact that methylation of cinnamic acid derivatives does not critically decrease an efficiency of these compounds.

Finalized model illustrates a good agreement with presented experimental results (Figure 5a) with correlation coefficient for the training set R = 0.9959. Interestingly, the mutual position of hydroxyl groups appeared to be an essential descriptor in several models for prediction of antioxidant activity proposed earlier. For instance, the simultaneous presence of the 3',4'-dihydroxy structure at the B-ring or the adjustment of the hydroxyl group at the C-3 atom of selected flavonoid compounds was used among the other descriptors for prediction of inhibitory activity of Lipids peroxidation [37]. Using exclusively this descriptor Rasulev et al. developed a model with correlation coefficient R = 0.813. Meanwhile by adding Petijean shape index, dipole moment, and a number of Glycoside-like

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fragments as descriptors it was possible to increase a correlation coefficient to R = 0.938. The number of vicinal hydroxyl groups was used as a descriptor for MLR models in [33] developed for prediction of antiradical activity measured by ABTS assay and evaluated as vitamin C equivalent (VCEAC) for the set of 21 hydroxybenzoic acids and simple phenolic compounds. Models developed based on only this topological descriptor resulted in correlation coefficient R = 0.915. Meanwhile, by combining the number of vicinal hydroxyl groups with a bond dissociation enthalpy proton affinity, or with a proton affinity and an electron-transfer enthalpy as descriptors the accuracy of developed model was improved by R = 0.957 and R = 0.962, respectively. Model developed by us based purely on the topological descriptors achieved slightly better statistical parameters (Table 3) then earlier models.



Figure 5. (a)—correlation plot between experimental and predicted values of $TAU_{Fe/\mu mol}$ antiradical activity; (b)—applicability domain of developed model. Yellow dots represent compounds selected as the training set, while the blue dots represent test set compounds.

Table 3. Statistical parameters for developed model.

	R	R ²	R ² adj	RMSE	MAE	Q^2_{loo}	Q^2_{lmo}
training set cross-validation	0.9959	0.9918	0.9893	5.5211 10.3056	4.0678 7.4203	0.9716	0.9592
external validation	0.9996	0.9993	0.9973	1.5429	1.2627		

An applicability domain represents the response and chemical structure space in which the model makes predictions with a given reliability. All compounds from the second cluster (12–22) appeared to be on the border of a predicted applicability domain as well as compound 1 (Figure 5b). The location of these compounds close to an outlier region is determined by their extreme response values (extremely low for 12–22, and extremely high for 1). Nevertheless, all 21 phenolic acids were found inside an applicability domain.

The number of statistical parameters that are used to validate developed models increase over years, as described in review by Gramatica et al. [38]. In this study, different validation criteria of robustness and predictivity have been used. The coefficient of determination R^2 (the square of the sample correlation coefficient (R) between the experimental endpoint and predicted one), adjusted coefficient of determination R^2_{adj} (in case when slope is set to zero), root mean square error *RMSE* (square root of the average of squared differences between predicted and experimental endpoint observation), mean absolute error *MAE* (the average magnitude of the errors in a set of predictions, without considering their direction) for both training and test sets were calculated. Additionally, specific coefficients of determination Q^2_{LOO} and Q^2_{LMO} (for leave-one-out and leave-many-out cross-validation,

respectively) were calculated and presented in Table 3. Though, it must be noted, that even inadequate models with chance correlation may pass statistical analysis with sufficient values of parameters in case when it is built for data sets that include a small number of samples. When working with small data sets one must rely not only on pure statistics but mostly on a chemical knowledge about mechanism of activity of interest, and structure–activity relationship in order to select an adequate chemically justified model.

With satisfactory statistical parameters and chemically justified descriptors (with *p*-values for all the descriptors equal to 0), developed here model, presumably, can well reflect the course of the reaction, and potentially be used for efficiency prediction of phenolic acids which activities are unknown. Based on both SAR and QSAR studies one can assume that 3,4,5-trihydroxyphenilacetic and 2,3,4-trihydroxybenzoic acids should be efficient as antioxidant compounds (Table 4), which is found to be in good agreement with the literature [39].

Table 4. Structures of phenolic acids and their antioxidant activities predicted by proposed quantitative structure–activity relationship (QSAR) model.



3. Materials and Methods

3.1. Apparatus

Hitachi U 5100 spectrophotometer (Japan, Tokyo) connected with computer for controlling of measurement and data analyzing, temperature stabilizer, and glass cuvette with 1 cm optical path was used.

3.2. Reagents

3.2.1. Phenolic Acids

Extrasynthese, Genay, France: 3,4-dihydroxyphenylacetic; 4-hydroxyphenylacetic; 4-hydroxy-3-methoxyphenylacetic; 2,5-dihydroxyphenylacetic; 4-hydroxycinnamic; 4-hydroxy-3,5-dimethoxycinnamic; 4-hydroxy-3-methoxybenzoic; 3,4,5-trihydroxybenzoic; 2-hydroxycinnamic; 3,4-dihydroxybenzoic.

Koch-Light Laboratories, Haverhill, United Kingdom: 3-hydroxycinnamic; 2,5-dihydroxybenzoic; 2,3-dihydroxybenzoic; 3,5-dihydroxybenzoic; 2,4-dihydroxybenzoic.

Fluka Chemie AG, Buchs, Switzerland: 3-hydroxybenzoic; 3,4-dimethoxybenzoic; 4-hydroxy-3,5-dimethoxybenzoic.

Fluka Chimica, Milano, Italy: 3,4-dihydroxycinnamic.

Sigma-Aldrich, St. Louis, MO, USA: 4-hydroxybenzoic; 2-hydroxybenzoic.

3.2.2. Other Reagents

Merck, Darmstadt, Germany: methanol, gradient grade.

Chempur, Piekary Slaskie, Poland: methanol, pure for analysis; hydrochloric acid 35%, pure for analysis.

Sigma-Aldrich, St. Louis, MO, USA: sodium acetate trihydrate; iron (III) chloride; 2,2-diphenyl-1-picrylhydrazyl radical (*DPPH*•); 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulfonic acid) diammonium salt (*ABTS*•+); potassium persulfate ($K_2S_2O_8$).

Fluka, Buchs, Switzerland: 2,4,6-tris(2-pyridyl)-(S)-triazine (TPTZ).

3.3. Methods

3.3.1. Measurement of Reducing Activity of Phenolic Acids with FRAP Method

The reducing activity of phenolic acids was measured with the Benzie and Strain method [17]. FRAP reagent was freshly prepared as follows: 12.5 mL of 0.3 mol/L acetate buffer (CH3COOH:CH3COONa), pH 3.6 was mixed with the same volume of methanol (Merck, Darmstadt, Germany). 2.5 mL of 10 mmol/L TPTZ in 0.04 mol/L HCl and 2.5 mL of 0.02 mol/L FeCl3·6H2O were added to such a solution.

225 μ L of 50% solution of methanol in water and 75 μ L of investigated phenolic compound solution (concentration chosen individually for each compound) were added to the 2.25 mL of FRAP reagent. The absorbance measurement at 593 nm was made at the beginning of the reaction and after 1 min at 37 °C. Measurements were repeated 5 times, for each phenolic acid. The results of experiments were presented as a reaction rate expressed as number of total antioxidant units per μ mol of substance (*TAU*_{*Fe*/*µmol*). One unit is the amount of substance that reduces one μ mol of Fe³⁺ to Fe²⁺ at 593 nm during 1 min at 37 °C. Measurements were done in a glass cuvette with 1 cm optical path. The number of units per 1 µmol of phenolic acid was calculated with the following Equation (2):}

$$TAU_{Fe/\mu mol} = \frac{1.513 * A_{F0} - A_{F1}}{c}$$
(2)

where $TAU_{Fe/\mu mol}$ —is the number of units per μ mol of phenolic acid; A_{F0} —is the absorbance of the solution at the beginning of the reaction; A_{F1} —is the absorbance of the solution after 1 min of the reaction; c—is the concentration of phenolic acid in the reaction mixture [mM].

The absolute error was calculated with the total differential method and the statistical significance of differences between samples was estimated using ANOVA with Tukey's test, using Statistica version 13.3.

3.3.2. DFT Calculations

Density functional theory was used to perform thermodynamic calculations in order to support hypothesis proposed in structure–activity relationship (*SAR*) investigations. Molecular structures of the studied compounds have been visualized using GaussView application [40] and optimized using B3LYP/cc-pVDZ level of theory with Gaussian16 [41] in an implicit water solvent, using conductor-like polarizable continuum model (*CPCM*) solvation method. Topology analysis of hydrogen-bonding was performed using Multiwfn 3.7 [36]. Hydrogen bond energy was calculated using Equation (3) [42]:

$$E_{HB} = \frac{V(r_{bcp})}{2} \tag{3}$$

where $V(r_{bcp})$ —is a potential energy density V(r) at corresponding bond critical point (*BCP*).

Additionally, relative intramolecular hydrogen bond enthalpy (ΔH) was calculated by comparing the sum of electronic and thermal enthalpies of the conformer with intramolecular hydrogen bonds

and of the lowest-energy conformer without hydrogen bonds, following the procedure demonstrated by Korth et al. [43].

3.3.3. Topological Descriptors

The choice of descriptors, which might be relevant to the activity of interest, is a problematic task. In this study, our aim was to build a QSAR model that is based exclusively on topology of phenolic acids. Optimized structures of studied compounds were saved in MDL MOL (*.mol*) format. PaDEL-Descriptor software [44] was used for the calculation of 2D topological descriptors and fingerprints. Calculated 2D descriptors included number of atoms and atoms of a certain element type, aromatic atoms, aromatic bonds, acidic and basic groups, hydrogen bond acceptors and donors, bonds of certain bond order, as well as topological descriptors characterizing the carbon connectivity, topological descriptors combining distance and adjacency information, etc. List of fingerprints calculated included Estate [45], Pubchem, Substructure [46], Klekota-Roth [47] and 2D atom pairs' fingerprints. Additional specific sets of descriptors have been developed manually based on *ortho-para-meta-* substitution pattern of hydroxyl groups, effect of carboxyl group and methylation. A total of 12,355 descriptors were used.

3.3.4. QSAR Model Development and Validation

QSARINS v2.2.4 [48,49] software (QSAR Research Unit in Environmental Chemistry and Ecotoxicology Department of Theoretical and Applied Sciences (DiSTA), University of Insubria, Varese, Italy) was used for data preparation and model development. The entire data set was preprocessed using a filter to eliminate constant (>80%) and codependent (>95%) descriptors. A splitting procedure plays a critical role for small data sets, since assigning an insufficient number of compounds into the validation set may result in the developed model being overtrained, while too many compounds in validation set may lead to loss of information for proper model development. Thus, only three compounds were selected randomly for validation set. Multiple linear regression (*MLR*) QSAR models were developed using the genetic algorithm (*GA*) method of variable subset selection. QSARINS software was also used for an applicability domain calculation by the leverage from the diagonal values of the Hat matrix.

4. Conclusions

Antioxidant activity for a set of phenolic acids was measured by FRAP assay. SAR investigation showed that mono hydroxylated compounds and compounds with two hydroxyl groups in *meta* position to each other exhibited the lowest efficiency as antioxidants, while compounds with two or more hydroxyl groups in *ortho* or *para* position to each other illustrated the highest antioxidant properties. Methylated phenolic acids derivatives were shown to be less efficient compare to their nonmethylated counterparts. Two stabilization factors were elucidated: resonance stabilization of radical and intramolecular hydrogen bonding. Due to resonance stabilization, *ortho-meta* and *para-meta* hydroxylated phenolic acids have an improved activity over mono hydroxylated ones and those having two hydroxyl groups in *meta* position to each other. Hydrogen bonding stabilization explained the reason behind the elevated activity of benzoic acid derivatives with substituted *ortho* position. Proposed hypotheses have been validated by quantum chemical calculations.

Multiple Linear Regression model for the prediction of antioxidant activity measured by FRAP assay was built based on four topological descriptors that include the presence of HOccOH fragment, any *ortho-meta* or *para-meta* substitutes phenylacetic acids, *ortho-meta*-substituted benzoic acid derivatives, and the number of methylated fragments of cinnamic acid derivatives. All these descriptors were in correspondence with SAR investigation. Having just one outlier (4-hydroxy-3,5-dimethoxybenzoic acid), the developed model has satisfactory statistical parameters and can be used for activity prediction of new, not yet tested, phenolic acids based on their structural features.

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Supplementary Materials: The Supporting Information is available online at Structures of investigated compounds in a .xyz format are accessible at the http://dx.doi.org/10.17632/77xhsdzk3y.

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Sample Availability: Samples of the compounds are not available.



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i. <u>Oświadczenia współautorów</u>

Wrocław, 14.04.2023

mgr farm. Maciej Spiegel Katedra i Zakład Farmakognozji i Leku Roślinnego Uniwersytet Medyczny im. Piastów Śląskich we Wrocławiu

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Jako współautor pracy: As a co-author of the research paper:

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oświadczam, że mój udział polegał na:

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Quantum-mechanical characteristics of apigenin: Antiradical, metal chelation and inhibitory properties in physiologically relevant media

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ABSTRACT

Density functional theory was used to examine the antioxidant activity of apigenin. All protonated species that are present in a non-negligible population at physiological pH were considered in the study. The ability to scavenge the hydroperoxide radical was evaluated in lipid and aqueous environments. The capacity to halt the Fenton reaction by chelating Fe(III) and Cu(II) ions was also investigated, as was the ability to inhibit xanthine oxidase. The results indicate that these activities may be particularly important in describing the beneficial effects of apigenin, especially because of its lower anti-OOH potential than Trolox or vitamin C. The findings underscore the significant role of dianion in the antiradical and chelating properties, despite its presence in much lower molar fractions than other ions.

1. Introduction

The physiological activity of radicals is a double-edged sword. On the one hand, they are essential for basic organism functions such as the immune response or intracellular signaling, and insufficient amounts may debilitate the body by preventing positive oxidative stress. [1] On the other, when accumulated in quantities far in excess of physiological concentrations, they are recognized triggers of intracellular damage, consequently leading to oxidative stress, which manifests itself in the development of severe disorders such as Alzheimer's or Parkinson's disease, hypercholesterolemia or cancer. [2] While the internal redox system consisting of enzymes and small molecules protects against this situation [3,4], due to external triggers the overall concentration of free radicals in the body is no longer so easy to maintain. For these reasons, scientific efforts are focused on the search for and development of new substances with antiradical activity, and polyphenols seem to be the most appealing in this regard.

These phytochemicals, found in abundance in daily consumed spices [5], fruits and vegetables [6,7], are the first line of support for the internal antioxidant system. In the experiments, polyphenols have been shown to satisfactorily scavenge artificial radicals such as DPPH[•] [8,9], chelate Fe(II) and Fe(III) ions involved in the Fenton reaction [10], or modulate enzymes important in the cellular red-ox system, such xanthine oxidase. [11] The biosynthesis of all phenolic compounds originates from the shikimate pathway and gives rise to three major

classes of polyphenols: flavonoids, phenolic acids and anthocyanidins. The former are built up on a cyclized C6-C3-C6 heterocyclic system, divided into three rings A, B and C, with the AC system representing the fundamental oxygen-containing scaffold. Based on the substitution pattern — the position of the B ring, the number and position of the -OH groups or the presence/absence of the C2=C3 bond - several subclasses of flavonoids can be distinguished. [12] Regardless of the structure, all the mentioned activities are associated with the presence of one or more phenolic hydroxyl groups linked to the aromatic ring or the delocalized electron cloud system itself. Namely, the reaction of a highly active radical with an antioxidant results in a much less reactive radicalantioxidant product whose electron density is distributed throughout the structure, rather than on a a single atom, resulting in its stabilization. [13] This structure not only gives them the ability to act as a hydrogen atom/electron donor or form adducts with small radical, but also gives them the ability to form stable complexes with the transition state metals. It has been experimentally proven that antioxidant chelated iron or copper does not participate as actively in the Fenton reaction as free one, thereby reducing the generation of hydroxyl and hydroperoxide radicals through this pathway. [14]

Probably the best recognized dietary polyphenol is apigenin. This flavone is on the top five flavonoids present in the daily diet, found ubiquitously in spices, fruits and vegetables such as parsley, spinach, celery seed, green celery heart, dried oregano, oranges, onions, wheat sprouts, tea and more. Typically presented in glycosylated forms - C-

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and O-glucosides, glucuronides, O-methyl ethers, and acetylated de-- it is easily broken down by the microbiota in the small intestine, followed by absorption and good distribution to tissues. [15] The experimental studies shown the beneficial effects of apigenin intake in preventing the development of diabetes, cancer, Alzheimer's diseases, amnesia, depression or insomnia. [16]

This another theoretical study on apigenin may not considered novel. Over the years, different scientific groups have conducted studies to describe this flavonoid. The impact of solvent and deprotonation on the feasibility of hydrogen atom transfer was studied in the work sof of Klein et al. [17,18]. This process has also been related to the experimental data elsewhere [19,20], while structure-activity relationships have been combined into the QSAR model by Zuvela et al. [21]. The data on the electronic structure, structural features and intrinsic reactivity indices are also available, whether from the original works [22-25] or the reviews [12,26] Nevertheless, as can be seen, these works are relatively old. Moreover, they tend to focus on one aspect, such as the bond dissociation process, leaving the other undiscussed. After all, it was not possible for us to find any theoretical investigations on the chelating properties of apigenin or the antiradical activity against specific radicals. This shows that while this may be considered as a trivial topic, it is actually emerging as a still undiscovered field worthy of further exploration, as was partially done by Zheng et al. [27], who studied the effect of varying substituents on the radical scavenging activity.

To fill this unexpected gap, apigenin has been accurately characterized in this article at density functional level of theory on its primary (anti-OOH) and secondary (Cu2+ and Fe3+ complexing power and xanthine oxidase inhibition potential) activities, demonstrating the mechanistic properties behind its beneficial activity

2. Results and discussion

2.1. Acid-base equilibria in water

The chemical structure of apigenin is shown in Fig. 1. The presence of three OH groups indicate that the system may exist simultaneously at three different protonation states.

The acid-base dissociation constants, pK_a , were calculated for the studied system, and were found to be equal to: $pK_{a1} = 7.40$ (C7), $pK_{a2} =$ 8.41 (C4'), and $pK_{a3} = 11.61$ (C5). According to other observations [28-30], unless the C3 position is not occupied, the C7 position is most easily deprotonated, while the C5 position is least susceptible to proton detachment. The latter is an expected behavior coming from particularly strong intramolecular hydrogen bond interactions between carbonyl and hydroxyl residues, which indirectly enable density delocalization, as it was have demonstrated in previous paper. [23]

Examining a graph plotting the molar fraction as a function of pH (Fig. 2), we find that non-dissociated and single anionic forms are present in equal amounts (${}^{M}f_{H_{3}A} = {}^{M}f_{H_{3}A^{-}} = 48.7\%$) in physiological pH. While di-anion species are also present in a non-negligible population $({}^{M}f_{HA^{2-}} = 4.7\%)$ and thus remain relevant to the study of apigenin activity, completely dissociated structure not (${}^{M}f_{A^{3-}} = 0.0\%$).



Fig. 1. Chemical structure and atom numbering of apigenin.

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2.2. Intrinsic reactivity indices

In order to determine the chemical properties of the isolated species and allow comparison with other substances regardless of the radical scavenged, enthalpies of bond dissociation (a), ionization potential (b). proton affinity (c) and proton dissociation (d) were calculated.

- $BDE = H^{\circ}(C-O^{\bullet}) + H^{\circ}(H^{\bullet}) H^{\circ}(C-OH);$
- $IP = H^{\circ}(C\text{-}OH^{\bullet+}) + H^{\circ}(e^{-}) H^{\circ}(C\text{-}OH);$ $ET = H^{\circ}(C\text{-}O^{\bullet}) + H^{\circ}(e^{-}) - H^{\circ}(C\text{-}O^{-});$
- $PA = H^{\circ}(C-O^{\circ}) + H^{\circ}(H^{+}) H^{\circ}(C-OH);$
- $\begin{aligned} PDE &= H^{\circ}(C \cdot O^{\circ}) + H^{\circ}(H^{+}) \cdot H^{\circ}(C \cdot OH^{+}); \\ The results are shown in Table 1. Ionization appears to be more \end{aligned}$

feasible in aqueous solvent (127.4 kcal/mol — 98.5 kcal/mol) than in pentyl ethanoate (142.4 kcal/mol), which can be linked to the ability to solvate the ultimately antioxidant cation radical. At the same time, it can be seen that successive deprotonations clearly reduce the energetics of the process, confirming the increasing destabilization of the species within each dissociation step. Vaganek et al. [18] reported that IP value for neutral species in water is 117.6 kcal/mol (B3LYP/6-311 + G(d,p)/ IEF-PCM), which is about 10 kcal/mol difference with our results. In the work of Zheng et al. [22] with energies estimated at the M06-2X/6-311 + G** level of theory and the SMD solvation model, IP equals 140.1 kcal/mol for species without intramolecular hydrogen bonds and 141.4 kcal/mol for species with them. In their next paper, a value of 139.3 kcal/mol was obtained. Our previous work treating the structureactivity relationships of 13 common flavonoids predicted an IP of 117.0 kcal/mol when B3LYP/6-31 + G(d,p) and PCM water solvent were used. These discrepancies are actually not astonishing - determining IP and EA values appears to be a difficult task and is strongly dependent on functional and basis set used [31].

The of C4' and C7 bond dissociation do not differ between neutral species in water or pentyl ethanoate, but the process is slightly more favored for C5 in the former than the latter (by about 6.0 kcal/mol). Following the results of Chen et al. [22], strong intramolecular hydrogen bonds weaken the antiradical capacity of C5 hydroxyl, to a greater degree in the weakly polarized solvent of pentyl ethanoate, and to a lesser extent in water. After all, dissociation does not seem to affect the values of this index significantly, and the reactivity pattern can be described as C5 < C7 < C4', what goes along with previous studies on flavonoids. [22,32] While only this work provides BDE values for all species present in the physiological environment, there are other data for neutral ones and can be reported for the set (C4', C5, C7) as (82.1, 92.5, 86.4) [17], (88.3, 94.5, 93.7) and (87.8, 89.3, 93.2) [22], (89.3, 100.5, 91.7) [23] and (96.6, 103.3, 102.2) [25], all in kcal/mol.

The pattern of hydrogen atom donating sites obtained in this work also applied to PDE, but not necessarily to PA, whose reactivity trend can be represented as $C5 < C4^{\prime} < C7$ in the case of water. Similar one was reported in other papers. [18,22,23] In order to provide complete and systematic data on the intrinsic properties, proton affinity values were also calculated for pentyl ethanoate, although deprotonation should not occur in a solvent devoid of solvation potential. This is evidenced by about 15-20 kcal/mol higher values of the index compared to those obtained in water. Also observable is a shift in the preferred site of deprotonation - it is the C4' position rather than C7. Though the difference is marginal and amounts to 0.6 kcal/mol, so both positions can be considered equivalent.

The electron transfer enthalpy results suggest that this process is less thermochemically demanding in pentyl ethanoate than in water. As the earlier discussion shows, this behavior is perfectly clear ---the anionic structure of the antioxidant is less stable than the radical one. While this holds not true for neutral species in water, a similar willingness to detach an electron is already noticeable when the first dissociation (at C5) occurs, and is further amplified in the case of dianion. Like the ionization potential, ETE values also vary in the literature, being in the ranges of 76.3-116.3 (C4'), 67.9-114.4 (C5) and 74.2-119.2 (C7) kcal/ mol [17,18,22,23].

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Fig. 2. Molar fractions of apigenin species plotted as a function of pH.

 Table 1

 IP, BDE, PA, and PDE values calculated for non-negligible apigenin species in pentyl ethanoate and water at 298.15 K. All values are in kcal/mol.

Species	Solvent	Site	IP	BDE	PA	ETE	PDE
		C4'		80.5	46.0	91.8	4.8
	PE	C5	142.4	92.3	59.7	89.8	16.5
		C7		85.6	46.6	96.3	9.8
H ₃ A		C4'		80.1	31.0	101.2	4.8
		C5	127.4	86.7	35.4	103.4	11.4
	water	C7		85.2	29.4	107.9	9.9
H ₂ A ⁻		C4'	107.0	81.3	32.6	98.5	23.2
		C5	107.9	87.0	41.4	95.4	28.9
HA^{2-}		C5	98.5	85.7	42.6	92.9	37.0

These brief and preliminary outcomes suggest that the *f*-HAT mechanism might be relying primarily on the C4⁺ hydroxyl group. At the same time, the SET mechanism, which is approximated by IP, appears to be much less favored that the former, however the difference between IP and BDE for HA²⁻ is much smaller, suggesting that their two related mechanisms may actually compete in radical scavenging activity. Although the overall results are not exactly the same with those reported elsewhere, it should recalled that the intrinsic reactivity indices are suggested for comparative purposes only and shall not be taken as determinants of antiradical activity. This can only be achieved by considering the radical being scavenged and examining the kinetic of feasible reactions. Nevertheless, the patterns generally confirm each other, which proves their usefulness for the purpose just stated.

2.3. Thermochemistry and kinetics

The scavenging potential of apigenin and its dissociated forms towards *OOH was determined by calculating the Gibbs free energies of the reaction (ΔG°) and the activation energies (ΔG°) of the *f*-HAT, RAF and SET mechanisms. For the first two pathways, $\Delta G^{\circ} < 10.0$ kcal/mol was a prerequisite for the reaction to be considered favorable and studied further. Although this implies the reversibility of the reaction, endergonic pathways may still be important channels if their products rapidly react further. This would be especially true if these latter steps are sufficiently exergonic to provide the driving force, and if their reaction barriers are low. Given the complexity of biological systems involving many different chemicals, it is likely that this could occur under such conditions. [33–35] Consideration of low endergonic pathways is found also in other works. [36–38] In contrast, such a limitation is not applicable to SET calculations, because the process is described by Marcus theory. All results are summarized in in Table 2. It is also important to emphasize that although the pK_a of the 'OOH/O² pair is 4.8, suggesting that the deprotonated form is present in the highest concentration under physiological conditions (0.25% vs. 99.75%), the reaction of superoxide with non-radicals is spin-forbidden, so this species is of little importance, and the oxidative damage comes primarily from the protonated form [39] Hence all calculations presented in the text are for 'OOH.

The results show that f-HAT represents a feasible mechanism of action with ΔG° ranging from between 2.2 kcal/mol (C4' of $H_2A^-)$ to 9.9 kcal/mol (C5 of H3A) in all cases expect C5 in pentyl ethanoate, for which $\Delta G^{\circ} = 16.1$ kcal/mol. In contrast, the established Gibbs free energies of SET were found to be significantly higher among neutral species (37.8 kcal/mol) and slightly higher for monoanionic species (18.3 kcal/mol). In the case of HA^{2-} , the trend shown is changed, and although this species is present in solution in a small fraction, its kinetic behavior may still be high enough to be consider an important part of the description of the scavenging of hydroperoxide radical by apigenin. Neutral species, regardless of solvent, are completely inactive in electron transfer reactions on *OOH. This is to be expected, given the activation energies, which actually drives the kinetics according to conventional transition state theory -- so it is anticipated that while non-dissociated species act mainly via f-HAT, the antiradical activity of dissociated species might be a combination of both hydrogen atom transfer and electron donation properties. The RAF mechanism was found to be highly endergonic in almost all cases. However, although ΔG° exceeds the imposed threshold of 10.0 kcal/mol, the calculated values of ΔG^{\ddagger} suggest that the reaction might be competitive with HAT at the C5 and C7 position. The above deliberations are supported by the reaction rates shown in Table 3, for which the transition state structures are depicted

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Table 2

Gibbs free energies of reaction (ΔG° , in kcal/mol) and activation (ΔG^{\ddagger} , in kcal/mol) between the considered apigenin species in water and pentyl ethanoate (mared with ^{PE} apex) at pH = 7.4 and 298.15 K.

Mechanism	Site	H_3A^{PE}		H ₃ A		H_2A^-		HA ²⁻	
		ΔG°	ΔG^{\ddagger}						
f-HAT	C4'	4.4	22.7	3.3	23.9	2.2	23.1		
	C5	16.1		9.9	29.6	7.9	27.3	6.6	28.0
	C7	9.5	25.1	8.4	26.5				
RAF	C2	18.6		16.2		13.8		15.0	
	C3	14.3		12.2	22.8 ^a	10.0	21.0 ^a	11.7	18.3 ^a
	C4a	33.4		31.9		26.7		26.2	
	C5	21.9		18.9		19.5		19.1	
	C6	22.0		20.3		14.4		14.5	
	C7	25.0		22.9		28.9		27.7	
	C8	21.5		19.2		15.6		15.5	
	C1'	25.5		22.9		22.8		20.0	
	C2'	21.8		19.6		19.4		21.4	
	C3'	21.1		19.6		20.0.		13.5	
	C4'	17.5		15.6		15.1		18.8	
	C5'	21.4		20.8		20.8		13.8	
	C6'	19.7		17.7		17.6		19.9	
SET				37.8	83.0	18.3	18.8	8.9	12.8

^a See the main text.

Table 3

Rate constants (k, in $M^{-1} s^{-1})^{a}$ and branching ratios (Γ , in %) of the reactions between *****OOH and the considered apigenin species in water and pentyl ethanoate (indicated by ^{PE} apex) at pH = 7.4 and 298.15 K.

Mechanism	H_3A^{PE}		H ₃ A		H_2A^-		HA ²⁻	
	k	Г	k	Г	k	Г	k	Г
HAT-C4'	$5.75 imes 10^{-1}$	99.4	$6.11 imes10^{-1}$	98.1	$2.99 imes 10^{0}$	93.9		
HAT-C5			1.15×10^{-4}	0.0	$6.47 imes 10^{-3}$	0.2	$1.04 imes 10^{-2}$	0.0
HAT-C7	$3.72 imes 10^{-3}$	0.6	$7.58 imes 10^{-3}$	1.2				
RAF-C3			$3.92 imes 10^{-3}$	0.6	$7.93 imes 10^{-2}$	2.5	$7.83 imes10^{0}$	0.0
SET			8.96×10^{-49}	0.0	$1.08 imes 10^{-1}$	3.4	$1.24 imes 10^5$	100.0
k _{total}			$6.23 imes 10^{-1}$		$3.18 imes 10^{0}$		1.25×10^5	
k _{overall}	$5.79 imes 10^{-1}$		$2.97 imes10^{-1}$		$1.52 imes 10^{0}$		$5.78 imes10^3$	
kcorrected			$7.42 imes 10^{-4}$		3.79×10^{-3}		1.44×10^{1}	

^a k_{total} is the sum of all rate constants for a given species; $k_{overall}$ equals k_{total} multiplied by the molar fraction of the species at pH = 7.4; and $k_{corrected}$ equals $k_{overall}$ multiplied by the molar fraction of *OOH under the given conditions.

in Fig. 3.

The overall reaction rate assumes that the hydroperoxide radical is present in significant amounts, much more than the antioxidant, and therefore does not control the reaction kinetics. This is hardly true and can be an error yielding inaccurate results. For this reason, emphasis should be placed on considering the molar fraction of the radical when establishing the reaction rate, particularly if the value is small. Thereby, $k_{\rm corrected}$ was computed to provide a more accurate description of the kinetic process in solution.

The evaluation of kinetic data indicates fairly low reaction rates. In line with previous assumptions, the activity of neutral and monoanionic species is mainly based on hydrogen atom transfer, with the C4' site being the most reactive. Although aqueous solvent and subsequent deprotonations slightly increase the values of the corrected reaction rates, they remain low. However, in accordance with an earlie hypothesis, although present in small amounts, HA²⁻ has a great impact on the proper description of the radical scavenging potential of apigenin. The branching ratio of an individual reaction rate of SET is much higher than any other reported here, and at the same time $k_{corrected}$ is the highest, despite the fact that only 4.66% of this form is present under the conditions studied, corresponding to $1.44 \times 10^1 \text{ M}^{-1} \text{ s}^{-1}$.

To determine the relative anti-'OOH potential, the corrected reaction rates were summed ($\Sigma k_{o}^{PE} = 5.79 \times 10^{-1} M^{-1} s^{-1}$; $\Sigma k_{corrected} = 2.51 \times 10^{1} M^{-1} s^{-1}$) and compared to the reported k_{overall}, refined to account for the molar fraction of hydroperoxide in solution, as was done for apigenin. The data indicate that apigenin is a worse radical scavenger than both Trolox ($\Sigma k_{corrected}^{PE} = 8.50 \times 10^2 M^{-1} s^{-1}$; $\Sigma k_{corrected} = 2.24 \times$ $10^4~M^{-1}~s^{-1})~[33]$ and vitamin C ($\Sigma k_{corrected}^{ep}=4.05\times 10^1~M^{-1}~s^{-1};$ $\Sigma k_{corrected}=6.4\times 10^5~M^{-1}~s^{-1})~[40].$ It would be wise to make comparisons to kaempferol (flavonol) and narigenin (flavanone) for the observable effect of the presence of the 3-OH group and the absence of the C2-C3 double bond, respectively, but data are lacking in this context.

2.4. Chelating properties

Complexation of Cu(II) and Fe(III) ions with consequences in intercepting the Fenton reaction is another likely activity exhibited by most dietary antioxidants. Apigenin has only one coordination site, constituting of C4 carbonyl and C5 hydroxyl groups, and this one has been studied for all the species. Thermodynamic and kinetic data are shown in Table 4, while a visualization of the optimized structures is shown in Fig. 4.

The copper ion was found to be coordinated by O—H and C=O groups, with Cu—O distances of the former equal to: 2.138 Å (H₃A), 2.112 Å (H₂A⁻) and 2.108 Å (HA²⁻); the latter: 1.958 Å (H₃A), 1.951 Å (H₂A⁻) and 1.946 Å (HA²⁻). In these complexes, the Jahn-Teller effect causing tetragonal deformation is evident, with the axial water molecules located at distances: [2.461 Å, 2.300 Å] (H₃A), [2.475 Å, 2.313 Å] (H₂A⁻) and [2.474 Å, 2.318 Å] (HA²⁻) from the Cu cation. [41] On the other hand, Fe(III) complexes persists their octahedral structure, and the distance between Fe and the O—H and C=O oxygens was found to be: [1.971, 1.882 Å] (H₃A), [1.963 Å, 1.873 Å] (H₂A⁻) and [1.960 Å, 1.863 Å] (H₂A⁻).

The outcomes indicate that the most stable complexes are those with



Fig. 3. Transition state structures of viable hydrogen atom transfers. Distances are in Å and angles are in degrees.

Table 4

Gibbs free energies of complexation $(\Delta G^{\circ}_{f_{5}}$ in kcal/mol) and kinetic constants $(K_{f_{5}}K_{i}^{H}$ and $K_{i}^{opp})$ for Cu(II) and Fe(III) ions chelated by C4-C5 coordination site of apigenin species.

Species	$\Delta G^_f$	K _f	K_i^{II}	K_i^{app}
Cu(II) H ₃ A H ₂ A ⁻ HA ²⁻	-2.2 -4.6 -6.9	$\begin{array}{c} 3.79 \times 10^{1} \\ 2.22 \times 10^{3} \\ 1.22 \times 10^{5} \end{array}$	$\begin{array}{c} 1.81 \times 10^{1} \\ 1.06 \times 10^{3} \\ 5.67 \times 10^{3} \end{array}$	$\textbf{6.74}\times \textbf{10}^3$
Fe(III) H ₃ A H ₂ A ⁻ HA ²⁻	-7.0 -12.9 -18.2	$\begin{array}{c} 1.39 \times 10^{5} \\ 2.91 \times 10^{9} \\ 2.34 \times 10^{13} \end{array}$	$\begin{array}{c} 6.61 \times 10^{4} \\ 1.39 \times 10^{9} \\ 1.09 \times 10^{12} \end{array}$	1.09×10^{12}

the most dissociated species. This behavior is expected due to the equalization of the metal's positive charge by the successively increasing unpaired electron density. The subsequent deprotonation results in an increase in stability of about 2 kcal/mol and 5 kcal/mol for Cu(II) and Fe (III), respectively. We expect that chelation would be further enhanced if the C5 group also dissociated due to the removal of the steric hindrance at the chelation site, however, such a structure does not occur under the conditions considered. Once equilibrium constants are established, the aforementioned behavior can be further detailed by notable changes in K_f between $H_3A(3.79 \times 10^1)$, $H_2A^-(2.22 \times 10^3)$ and $HA^{2-}(1.22 \times 10^5)$. Even larger values occur for Fe(III)-complexes, as would be expected from the much larger ΔG°_f values.

Although the HA^{2-} species accounts for only 4.66% of the total forms in aqueous solution, it seems to prevail when describing Fe(III) complexation, constituting virtually all of apigenin II-type antioxidant activity ($K_f = K_i^{qup} = 1.09 \times 10^{12}$). To a lesser extent, this is true for Cu(II), since $K_i^{qup} = 6.74 \times 10^3$ is the sum of the contriution of both HA²⁻ (the larger part) and H₂A⁻ (the smaller part). In any case, summarizing the complexation process, the results indicates that apigenin is much better Fe(III) chelator than Cu(II), although being complexed their contribution to the Fenton reaction is naturally smaller.

2.5. Xanthine oxidase inhibition

Preliminary molecular docking studies were carried out to gain insight into how the analyzed species bind to the identified xanthine oxidase active site. The approach was validated by comparing the position of the newly introduced ligand with that of the cocrystallized quercetin (Fig. 5) and the interactions with the surrounding amino acids (Fig. 6).

The data show that apigenin, in any state, has a similar conformation to the original ligand and interacts with the same amino acids. Arg880 and Thr1010 are the common hydrogen bond sites interacting with the ligands. The position of the AC ring in apigenin species differs only slightly from quercetin, while the only difference between the structures in this scaffold is the presence of C3 hydroxyl in the latter. Given that apigenin's conformation results in the formation of a hydrogen bond between Val1010 and Ala1079, albeit not as pronounced as in the case of quercetin, it can be assumed that the absence of 3-OH favors binding, allowing the formation of the two hydrogen bonds mentioned. The process of deprotonation seems to have little effect on the length of the hydrogen bond, and while the distances decrease, the changes are negligible. The binding pocket appears to be strongly hydrophobic, and almost all atoms of the phytochemicals under study are involved in these interactions. Additionally, 3D visualization makes it possible to



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Fig. 4. Structures of Cu(II) and Fe(III) complexes with apigenin species.

prediction π - π interactions with the aromatic rings of Phe914 and Phe1009.

Table 5 shows the established inhibition constants. While the scavenging capacity towards 'OOH and chelating properties depend on the most dissociated species, even though their molar fraction was much lower, the inhibition capacities do not. This is due to the fact that the binding free energies of the studied species are relatively similar, thanks to almost identical orientation in the binding pocket and interactions.

3. Conclusions

DFT experiments revealed the antiradical and antioxidative properties of apigenin and its dissociated forms in biologically relevant solvents. Given these activities, the deprotonation process was shown to be extremely beneficial. Nonetheless, while even a small fraction of dianion species had an effect in the first two, it is much smaller in the case of inhibitory activity. The results of the study are summarized as follows:

• •OOH is scavenged at a reaction rate constant of $5.79\times 10^{-1}\,M^{-1}\,s^{-1}$ in the lipid phase and 1.44 \times 10¹ $M^{-1}\,s^{-1}$ in the aqueous phase. These values are lower than Trolox or vitamin C.

- The Fe(III) cation is much more easily chelated than the Cu(II) cation. The corresponding apparent equilibrium constants were determined to be 1.09×10^{12} and 6.74×10^3 , respectively.
- Xanthine oxidase inhibition is similar to that of quercetin, and similar amino acids are involved in the interactions with ligands. Apigenin, on the other hand, due to its lack of 3-OH is able to rotate and form two more exclusive hydrogen bonds with Val1011 and Ala1079.

4. Computational details

DFT studies were performed in the Gaussian16 software package. [42] The antiradical behavior was studied according to a wellestablished computational protocol that includes thermochemical and kinetic aspects of the processes undergoing, the QM-ORSA protocol. [43] The meta-hybrid exchange-correlation GGA functional M05-2X [44] combined with the 6-311 + G(d,p) basis set [45,46] was used. The physiologically relevant solvents were modeled by the water and pentyl ethanoate (PE), respectively, within a universal solvation model based on solute electron density (SMD). [47] The same level of theory was used for the frequency computations necessary to ensure the local minimum or transition state character of the geometries found, as well as to account for zero-point energy corrections. The unrestricted procedure was



Fig. 5. The binding mode of H_3A (top-left), H_2A^- (top-right) and HA^{2-} (bottom) in the active site of xanthine oxidase. Quercetin is represented by the slightly transparent structure in pink. Hydrogen bonds are represented by green lines with their corresponding distances. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

used for open-shell systems. Intrinsic reaction coordinate (IRC) calculations were performed to ensure that the transition states found correctly link reactants and products along the path of the reaction minimum energy.

To account for the presence of differently dissociated species in polar solvent, the pK_a values and molar fractions for neutral and charged species were determined using the fitted parameters method [48,49], which is based on experimentally measured dissociated constants of phenolic groups and has proven reliability. Only those whose population is non-negligible (^Mf > 0.1%) at pH = 7.4 were included in the study. [43]

Intrinsic reactivity indices [13] — bond dissociation enthalpy (BDE), ionization potentials (IP), proton affinities (PA), electron transfer enthalpy (ETE), and proton dissociation energies (PDE) describing the thermochemical behavior of the isolated species and providing initial assurance of their activity were estimated within the framework of the adiabatic approximation using previously proposed H^+ and electron solvation enthalpies. [50]

From a mechanistic point of view, there are three main modes of antiradical activitity— hydrogen atom transfer, single electron transfer (*SET*) and radical adduct formation (*RAF*). When discussing the hydrogen atom transfer, however, it is important to remember that the reaction can involve transfer of a proton and an electron as a single or separated entities. Therefore, the reaction is commonly referred as formal hydrogen atom transfer (*f-HAT*). These are represented by the following reaction schemes:

 $f - HAT : HA + R^{\bullet} \rightarrow HA^{\bullet} + RH$

 $SET: HA + R^{\bullet} \rightarrow HA^{+\bullet} + R^{-}$

 $RAF: HA + R^{\bullet} \rightarrow [HA - R]^{\bullet}$

The reaction rate constants, k, for the HAT and RAF mechanisms

were calculated using conventional transition state theory. [51] The SET reaction barriers were determined adopting Marcus theory. [52] Additionally, Collins-Kimball theory was applied to assess rate constants close to the diffusion limit. [53]

The studies on metal chelation by apigenin considered all possible complexation sites and were evaluated in the M05 functional [54], which unlike M05-2X [44], was parametrized on both metals and nonmetals. The ability of apigenin to chelate copper and iron was determined by calculating the Gibbs free energies of complexation (ΔG_f) for reactions with a general scheme:

 $\left[M(H_2O)_6\right]^n + A^m \rightarrow \left[M(H_2O)_4A\right]^{n-m} + 2H_2O$

where *M* stands for the metal (Cu or Fe), *n* for its charge (2+ and 3+, respectively), *A* stands for the antioxidant, and *m* is the charge of the antioxidant species. The apparent equilibrium constants (K_i^{qpp}) were then calculated according to the formulas:

$$K_{f} = e^{-\frac{m_{f}}{RT}}$$
$$K_{i}^{II} = \sum K_{f}^{*m} f_{i}$$
$$K_{i}^{app} = \sum K_{i}^{II}$$

where K_f is the equilibrium constant at the *f* complexation site of a given species, and K_i^{II} equals $\sum K_f$ multiplied by the molar fraction, ${}^{m}f_i$, of the considered *i* species at pH = 7.4 at 298.15 K. Metals were described by Stuttgart-Dresden effective core potentials. [55,56]

The docking procedure was done on the crystal structure of bovine xanthine oxidase in complex with quercetin (PDB: 3NVY) [57]. To ensure the reliability of the selected protein, milk xanthine oxidase (PDB: 2CKJ) was superimposed and sequentially aligned. The root mean-squared deviation was found to be 0.847 Å, while pairwise sequence alignment showed 84.9% identity and 89.5% similarity, with 5.9% gaps.



Fig. 6. Intramolecular interactions in the active site of xanthine oxidase. Green dashed lines represent hydrogen bonds, while red circles indicate equivalent residues engaged in hydrophobic interactions. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

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Table 5

Bindings free energies ΔG_{b}° , inhibition constants (K_i), molar-fraction weighted inhibition constant (K_{b}^{II}) and apparent inhibition constant (K_{b}^{app}) of apigenin species.

Species	$\Delta G^_b$	K _b	K_b^{II}	K_b^{app}
H ₃ A	-9.1	4.76×10^{6}	2.27×10^{6}	5.89×10^6
H_2A^-	-9.3	6.68×10^{6}	$3.18 imes 10^6$	
HA^{2-}	-9.5	9.36×10^6	4.36×10^5	

After structure validation, 3NVY was pre-processed by removing quercetin, adding hydrogens and fixing charges. The docking box was centered on the original ligand position, with the dimension and volume covering the amino acids involved in the quercetin docking, and the apigenin species were docked using the AutoDock VINA [58]. The apparent binding constant (K_{e}^{gap}) was obtained from the binding free energy (ΔG_{b}°) using the formula:

$$K_b = e^{-\frac{2O_b}{RT}}$$

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 $K_b^{II} = K_b *^m f_b$

 $K_b^{app} = \sum K_b^{II}$

where K_b represents the binding constant of a given b species, K_b^{II} is the individual contribution of a given species determined by accounting for its molar fraction, ${}^{m}f_{b}$, in the system at pH = 7.4.

CRediT authorship contribution statement

Maciej Spiegel: Conceptualization, Methodology, Validation, Formal analysis, Investigation, Resources, Data curation, Visualization, Writing - original draft, Writing - review & editing. Zbigniew Sroka: Supervision, Project administration.

Declaration of Competing Interest

The authors declare no conflict of interest.

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i. <u>Oświadczenia współautorów</u>

Wrocław, 14.04.2023

mgr farm. Maciej Spiegel Katedra i Zakład Farmakognozji i Leku Roślinnego Uniwersytet Medyczny im. Piastów Śląskich we Wrocławiu

OŚWIADCZENIE WSPÓŁAUTORA CO-AUTHORSHIP STATEMENT

Jako współautor pracy: As a co-author of the research paper:

Spiegel, M., & Sroka, Z. (2023). Quantum-mechanical characteristics of apigenin: Antiradical, metal chelation and inhibitory properties in physiologically relevant media Fitoterapia, 164(October 2022), 105352.

oświadczam, że mój udział polegał na: I declare that my contribution consists of:

- sformułowaniu i ewolucji nadrzędnych celów i zadań badawczych formulation and evolution of overarching research goals and aims
- rozwoju i zaprojektowaniu metodologii
- development and design of methodologyweryfikacji ogólnej odtwarzalności wyników badań
- verification of the overall reproducibility of research outputs
- zastosowaniu technik matematycznych i obliczeniowych do analizy oraz syntezy danych z badań application of mathematical and computational techniques to analyse and synthesize study data
- przeprowadzeniu procesu badawczego i dochodzeniowego oraz zbieraniu danych
- conducting a research and investigation process & data collection
- zapewnieniu zasobów obliczeniowych
- provision of computing resources
- wykonywaniu działań mających na celu organizację danych badawczych performing activities to maintain research data
- przygotowaniu i stworzeniu opublikowanej pracy, w szczególności wizualizacji i prezentacji danych preparation and creation of the published work, specifically visualization & data presentation.
- przygotowaniu i stworzeniu opublikowanej pracy, w szczególności napisanie jej wstępnego szkicu
 preparation and creation of the published work, specifically writing the initial draft
- przygotowaniu i stworzeniu opublikowanej, w szczególności naniesieniu korekt preparation and creation of the published work, specifically revision

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oświadczam, że mój udział polegał na: I declare that my contribution consists of:

- pozyskaniu wsparcia finansowego na projekt prowadzący do tej publikacji
- acquisition of the financial support for the project leading to this publication
- odpowiedzialność za zarządzanie i koordynację w zakresie planowania i realizacji działań badawczych management and coordination responsibility for the research activity planning and execution

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IV. Publikacja [D]





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Current Trends in Computational Quantum Chemistry Studies on Antioxidant Radical Scavenging Activity

Maciej Spiegel*

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ABSTRACT: The antioxidative nature of chemicals is now routinely studied using computational quantum chemistry. Scientists are constantly proposing new approaches to investigate those methods, and the subject is evolving at a rapid pace. The goal of this review is to collect, consolidate, and present current trends in a clear, methodical, and reference-rich manner. This paper is divided into several sections, each of which corresponds to a different stage of elaborations: preliminary concerns, electronic structure analysis, and general reactivity (thermochemistry and kinetics). The sections are further subdivided based on methodologies used. Concluding remarks and future perspectives are presented based on the remaining elements.



KEYWORDS: Antioxidants, density functional theory, computational chemistry, electronic structure, weak interactions, kinetics, thermochemistry, QM-ORSA

■ INTRODUCTION

Free radicals play a crucial role in the maintenance of homeostasis by participating in a range of physiologically relevant processes such as immune response and intracellular communication. Nonetheless, whatever the case may be, their uncontrolled accumulation is not favorable. The energy they possess as a due to the unpaired electron on the valence shell or excited state cannot be efficiently neutralized by intracellular antioxidant defense system and is instead transferred to the biologically important targets like lipids, carbohydrates, proteins, and DNA strains. In consequence, these structures degenerate, leading to the development of severe malfunctions resulting in illness such as diabetes, atherosclerosis, Alzheimer's and Parkinson's diseases, or tumor growth.^{1,2}

The expanding awareness of the dual nature of radicals has prompted a surge in interest in compounds that can decrease elevated levels of reactive oxygen, nitrogen, and sulfur species, thereby preventing their harmful activity. These substances, known as *antioxidants*, are a heterogeneous group of molecules able to reduce oxidative stress in different ways. They are classified as follows³

- Type I: chain breakers, which interact directly with radicals by creating species that are more stable and less hazardous to cells than the former ones, thus terminating chain reactions and preventing oxidation of biological targets.
- Type II: preventers, for which, however, a unified mechanism of action is not specified, but it does not include interactions with radicals. Among known activities of that type are metal chelation, particularly iron and copper, which participate in the Fenton reaction,



© 2022 The Author. Published by American Chemical Society as well as regulation of enzymes responsible for radical formation or those directly involved in oxidative stress development.^{1,2,4} Compounds capable of regenerating biological antioxidants¹ or absorbing UV radiation⁵ are also included in this category.

• Type III: substances that effectively repair oxidatively damaged biomolecules.¹

However, because most antioxidants exhibit multiple types of activity at the same time, such categorization is often artificial. It is better to do so on the basis of their chemical structure or origin.

 $\rm \bar{P}$ hytochemicals are plant-derived compounds that are plentiful in many herbs and commonly consumed plants such as beetroot, high in betalains;⁶ tea, rich in catechins;⁷ or grapefruit, abundant in flavanons.⁸ This family can be further subdivided into flavonoids,^{9,10,19–28,11,29–37,12–18} phenolic acids,^{38–47} lignans,⁴⁸ aurones,⁴⁹ chalcones,^{49–51} curcuminoids,^{52,53} anthocyanidins,^{54–56} stilbenoids;^{25,57–59} anthraquinones, ^{60–63} glucosinolates, ^{64,65} alkaloids,^{66,67} coumarins,⁶⁸ terpenes and terpenoids,^{68–72} and others^{41,73,82–86,74–81} (Figure 1), all of them being extensively studied with computational quantum chemistry methods, as evidenced by the number of recent scientific findings. Albeit, plants are not the

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only source of antioxidants —substances with promising antiradical activities have also been found among common drugs, $\frac{87-92}{1000}$ biological substances, $\frac{93-102}{10000}$ and their metabo-

drugs,^{87–92} biological substances,^{93–102} and their metabolites.^{59,93,98,100,103–105} All of the listed substances are being heavily modified in an effort to identify derivatives with an improved safety profile and enhanced radical scavenging potential.^{23,36,102,106–114,46,115–124,83,125,867,89,94,96,101} Finally, completely novel structures are proposed and

investigated on the basis of recognized pharmacophores: phenolic units,^{126–132} five-heterocyclic rings,^{133–139} quinoline backbones,^{140–142} and other moieties.^{143–146} Importantly, despite their diversity, chain breakers share similar reactivity patterns and mechanisms, allowing common theoretical approaches to be used to study any of them.

This review outlines currents trends in "Type I" activity research in a straightforward and methodical manner. The first section deals with preliminary concerns, such as selecting the appropriate level of theory, the solvation model, and the initial structures. Following that, the topic of electronic structure examination is discussed in light of the methodologies documented in the literature. The majority of this work is devoted to thermochemistry and kinetics research, which are the most important for comparing computationally produced results with experimental data. Finally, remaining issues are highlighted and perspectives on the subject provided.

Before proceeding, it is important to note that while this review specifically mentions hydroxyl groups any other residue in which a hydrogen atom is bonded to a highly electronegative atom, such as nitrogen in an amino group or sulfur in a thiol group, can also be considered as the one that may participate in antioxidative activity.⁴⁶

PRELIMINARY CONCERNS

The level of theory chosen, which describes the electronic structure of the molecule *as itself*, and the solvation model, which adjusts the system's electron density cloud to minor perturbations caused by solvent molecules, are two key components influencing chemical behavior that must be taken into account from the beginning of the studies.

Functional and Basis Set. It is frequently advantageous to obtain results that precisely resemble experimental data while needing the least amount of computational time. However, this seemingly easy task is burdened with two fundamental issues: (1) With so many functionals and basis sets available, choosing a level of theory satisfying this condition is difficult. (2) The lack of reference data against which theoretical findings may be compared casts doubt on the latter. Although high-end methods such as CCSD(T)/CBS guarantee quality of the outcomes, this solution is inapplicable for routine computations due to the significant uptake of resources.

As shown in Chart 1, there is currently a trend toward the use of density functional theory (DFT) methods, with $\rm B3LYP^{147,148}$

Chart 1. Share of Functionals in Articles Published in the Last Five Years a



^aPlotted on the basis of refs 9–146.

(~47%), M06-2X¹⁴⁹ (~26%), and M05-2X¹⁵⁰ (~17%) being picked the most commonly. Restricting to any of them is advantageous because it provides researcher a plethora of datapoints the comparison with validate the results in a greater degree than it would happen if was done against outcomes obtained by completely differently constructed functional.

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Moreover, other researchers following the given reasoning would more likely cite the paper that use common level of theory since it enables to compare their own theoretical results with those provided by other researchers.

Given their newness, it is visible that these two Minnesota functionals are gradually displacing B3LYP since their release. This could be a result of their superior performance in estimating thermochemistry, kinetic and noncovalent interactions of nonmetal elements, as well as energies of reactions involving free radicals, which they predict extremely close to actual data, as claimed by the developers $^{149-151}$ and continue to be substantiated by independent scientists, either in the course of original research^{39-41,45,77,84,87,129} or in benchmarks.¹⁵²⁻¹⁵⁴ On the other hand, several researchers who used both B3LYP and one of Truhlar's global hybrids noted that while B3LYP tends to underestimate energies there are no substantial differences and the reactivity patterns hold when the same basis set is used.^{60,64}

Although the reasons for selecting certain functional are usually covered in the manuscript with details, the arguments for choosing "this, but not that" basis set are nearly always glossed over. This bad habit have its consequences in the performance of computations. Increasing the number of basis functions is known to increase task processing time, and it has recently been established that those from the Dunning's and Ahlrich's families are particulary vulnerable.¹⁵³ Furthermore, whereas functionals should be kept constant throughout the research, the basis set does not, namely while one can be used for electronic structure investigations, other may be applied for thermochemistry, which allows maneuvering them in order to obtain the best results at the lowest possible cost.

As a result, Chart 2, which depicts the percentage of basis sets used for thermochemistry computations, is much more divided





^aPlotted on the basis of refs 9-146.

than the previous one, with Pople's leading the way. A particularly good observation is that combining any of them with one of the three most commonly employed functionals presented in Chart 1, appears to have a little influence on the results as evidenced in the following examples. The findings of Shammera Ahamed et al.⁷⁹ (B3LYP and M062X combined with either 6-31+G(d,p) or 6-311++G(d,p)) and Mendes et al. (B3LYP, LC-wPBE, M062X, and BMK in 6-311G(d,p) and 6-311+G(d,p) bases) show that despite the augmentation of basis set with another diffuse function or inclusion of next function

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describing valence shell, the values of reactivity indices found in the gas phase change only slightly, regardless of level of theory used. In their other study,²⁶ the authors demonstrated that shifting from 6-311G(d,p) to 6-311+G(d,p) changed the ionization potential values by less than 4 kcal/mol independently of the environment (gas phase, water, methanol, ethanol, The set of the end of the set of conducted in gas and water. On the other hand, the differences in ionization potential and electron transfer energy associated with ion formation in water milieu were found to be significantly smaller in the case of B3LYP/6-311++G(d,p) outcomes when compared to those obtained at B3LYP/6-31+G(d,p) level of theory.

Moreover, although the presence of polarization functions is undeniably important for determining the energy of highly polarized bonds, the role diffusion functions, which are widely regarded as necessary for accurately modeling electron clouds in ionic systems and radical involving pathways, must be addressed more thoroughly. Still, the recent research¹⁵³ has shed new light on that issue for it was discovered that using 6-311G(d,p) for overall antioxidant studies produces the best results in terms of both the accuracy between theroetical outcomes and experimental values, as well as computational resource uptake. Further confirmation on that indulging issue and more detailed studies on the role of the basis set are welcome.

It is worth noting that open shell computations are hampered by the possibility of spin contamination.¹⁵⁵ This is because the resulting wave function is an artificial mix of spin states rather than an eigenfunction of total spin, $\langle S^2 \rangle$. In an ideal system, $\langle S^2 \rangle$ equals 0.75 for singlet and 2.0 for triplet

$$\langle S^2 \rangle = s(s+1) \tag{1}$$

where s denotes the number of unpaired electrons divided by half. Other values are acceptable as long as they deviate by no more than 10%. ^{10,109,152} Greater ones indicate the presence of higher spin states, which may alter the energy or geometry, just like the population analysis outcomes, resulting in biased conclusions; such structures should not be considered for future research. Spin-restricted open shell computations may be a solution in those cases, but they consume more resources than unrestricted ones and may still produce incorrect energies of unpaired electrons due to the absence of dynamical correlation caused by the vanishing of spin polarization.

Solvation Model. The physiological media in which antioxidants play fundamental biological roles are body fluids and the lipid bilayer of cells membrane. As a consequence, most experiments are conducted in water or in a nonpolar environment, that computational elaborations must account for. The effect of a polar solvent is a fundamental tenet of theoretical investigations because it distorts a molecule's electron cloud due to electrostatic polarization interactions, affecting the shape of the potential energy surface and chemical activity.^{3,20,28,130,156} Furthermore, if the reaction pathway involves proton or electron detachment, the solvation is a known to be a driving force that eases the process and makes it more feasible than it would be in a nonpolar medium. ^{50,116} That is why it is also important for properly modeling dissociation related processes as is stressed later in the text.

For the time being, three methods are employed to incorporate solvent effects: implicit (through a homogeneously

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polarizable medium), explicit (by solvent molecules), and a combination of these two. The integral equation formalism variation of the polarizable continuum model (IEFPCM, often referred to as just PCM).¹⁵⁷ conductor-like polarizable continuum model (CPCM),^{158,159} and solvation model based on density (SMD)¹⁶⁰ are examples of continous solvation models that are implemented in a majority of quantum chemistry software. They are also the most frequently used because they are burdened with a much lower computing cost than implicit or combined approach.

However, the fundamental disadvantage of implicit models is complete negation of intermolecular hydrogen bonds, which are often essential for proper simulation of antiradical activ-ity^{93,103,144} especially when abundant as in the case of ity^{93,103,144} especially when abundant as in the case of capsaicin.⁸⁴ A mutual competition between intramolecular and intermolecular hydrogen bonds is also observed as a relevant factor modying hydrogen atom transfer proclivity.47 The modeling of •OH and •OOH radicals, which are of primary interest in antioxidants research, is an excellent example of how the explicit water molecules can influence the results as well. Accordingly to Pérez-González and Galano,¹²⁵ adding the first water molecule considerably changed the rate constants of the •OH/-OH path, but adding more or repeating the procedure for the 'OOH/'OOH path resulted in no significant change. The fact that their charge is concentrated on a single exposed heteroatom¹⁶¹ is important here.

The application of a homogeneously polarizable medium is a particularly viable method of modeling solvents having large structures. For example, representing the enormous membrane lipid chains would be a time-consuming and inefficient effort. Instead, one of the most common approaches found in the literature is to do that implicitly, by using the largest available aprotic solute, for example, pentyl ethanoate. ^{12,100,107,120,132} To account for the cage effect, which manifests as the loss of entropy of any chemical reaction with a molecularity of two or greater, and improve results, the Okuno's corrections¹⁶² and Benson's free volume theory¹⁶³ can be applied (eq 2), as demonstrated. ^{40,41,104} The corrected Gibbs free energy, ΔG_{solw}^{FV} is expressed then in a form of

$$\Delta G_{\text{solv}}^{\text{FV}} = \Delta G_{\text{solv}}^0 - RT\{\ln[n10^{(2n-2)}] - (n-1)\}$$
(2)

where *n* represents the molecularity of the reaction, and ΔG_{sol}^{v0} is a Gibbs free energy in solvent, *R* a gaseous constant, and *T* the temperature¹⁶⁴

Initial Structure. Let's refere a thorough conformational studies^{156,165} that were performed on a collection of quercetin structures with variable planarity and intramolecular hydrogen bond counts. These two structural features are known to account for antioxidative activity of flavonoids, and so considerable differences have been found among them. This is not the only case, for similar geometry—activity relationships have been pinpointed also in other studies.^{60,91,107,134} This emphasizes the role of selecting the appropriate conformer for the study as a critical first step in theoretical elaborations.

Molecular dynamics (MD) simulations are most effective for producing excellent starting structures. Different approaches can be undertaken to run MD calculations, such as using software that has implemented molecular mechanic potentials and thus allows for the direct conformational search procedure^{166–168} or dedicated ones designed for more demanding studies, like GROMACS,¹⁶⁹ Amber,¹⁷⁰ or CHARMM,¹⁷¹ for which the molecule under consideration must first be parametrized. This pubs.acs.org/jcim

Review

can be done with either the proposed protocols⁸⁴ or available webservices the most well recognized of which is probably CharmmGUI,¹⁷² although AutomatedTopologyBuilder (ATB)¹⁷³ should also be mentioned.

CharmmGUI has an advantage over ATB in that it has a userfriendly interface and instantly generates a complete set of files that can be submitted for molecular dynamics, whereas ATB only produces force field, structure, and topology data, leaving the user to prepare the remaining files. CharmmGUI, on the other hand, predicts topology using CGenFF,¹⁷⁴ which elucidates it through bond perception and atom typing, while ATB processes input using DFT or semiempirical techniques depending on the size of the system. After all, it is the user's expertise that determines which path to take.

MD frequently generates a large number of molecules, whilst upcoming quantum chemical studies should only consider the most populated states. One of the first filters used to get rid of the undesired structures is the geometric clustering algorithm¹⁷⁵ which, in simple terms, groups conformers based on their structure or kinetics. If there are multiples of them, optimization at the appropriate level of theory followed by Maxwell– Boltzmann distribution analysis may be a viable approach for removing superfluous structures, particularly those with molar fractions less than 0.1%. This threshold is proposed^{46,101,112} because a small energy difference between conformers could indicate an interconversion process: rotatability of OH groups¹⁶⁵ or side chains,⁸⁴ bonds deformation due to keto– enol tautomery,²¹ shift in E/Z-conformers equilibria,¹³³ and bending of dihedral angles.^{16,20,28,33,50,67,128,165} They can all modulate hydrogen bonds, electrons cloud delocalization, and polarizability, as well as radical accessibility to specific sites of an antioxidant. Furthermore, the molar fraction cannot be too low if the compound is desired to pass biological barriers.⁴⁶

Two equations (eqs 3 and 4) can be used to evaluate the Maxwell–Boltzmann population f of a specific conformer i, in the set of n conformers

$$f_i = \frac{\exp\left(\frac{-G_{i(a_i)}}{RT}\right)}{\sum_{i=1}^n \exp\left(\frac{-G_{i(a_i)}}{RT}\right)} \text{ for } i = 1, 2, \cdots, n$$
(3)

and

$$\sum_{i=1}^{n} f_i = 1 \tag{4}$$

where *R* is the gas constant, and *T* is the temperature. $G_{i(aq)}$, the aqueous phase Gibbs free energy of the *i*th conformer, can be calculated following eq 5

$$G_{i(aq)} = G_i^{\circ} + \Delta G^{latm \to 1M} + \Delta G_i^*$$
⁽⁵⁾

where G_i° denotes a species' gas phase free energy at a given temperature, $\Delta G^{\text{latm} \to 1\text{M}} = 1.89 \text{ kcal/mol}$ and reflects the shift in standard state from 1 atm (superscripted with °) to 1 M (superscripted with *), and ΔG_i^* denotes a species' aqueous solvation free energy.

Deprotonation and **Dissociation Constants**. The primary activity of type I antioxidants is based on their reductant capacity, which is frequently, but not always^{64,68,125} linked to the hydrogen-donation capacity from one of their aromatic hydroxyl groups; thus, a simple conclusion can be drawn that the more of them, the greater shall be participation of hydrogen-related channels in overall radical scavenging, and so its viability.¹¹⁶

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However, because these residues are also weakly acidic, multiple species can coexist in a water environment at the same time. Their molar fraction is governed both internally by their chemical structure⁸² and externally by the pH of an environment.

If radicals are neutralized by mechanisms inaccessible to other forms, a seemingly small amount of one of them may be critical in accurately measuring scavenging activity. At the given pH, for example, it is possible that the antioxidants have already deprotonated all of the hydroxyl groups and thus exhibit only electron-related channels,^{36,85,102,126} whereas the studied radical is efficiently neutralized solely by formal hydrogen atom transfer. Similarly, with each subsequent dissociation, species are less likely to remove another proton, making routes that include its donation, e.g. sequential proton-loss electron transfer or sequential electron transfer-proton transfer, more energetically demanding; on the other hand, electron-donating processes are expected to occur more easily for them.

There are several methods that can be used to estimate dissociation constants and deprotonation pathways. The oldest ones, such as *direct, proton exchange, hybrid cluster continuum,* and *implicit—explicit,* rely on thermodynamic cycles and have previously been exhaustively discussed in the literature^{176,177} and are thus just briefly mentioned here.

In the *direct* approach, an antioxidant may either (a) adhere to the Arrhenius theory and dissociate directly into anion and proton (Scheme 1a), (b) obey Brønsted–Lowry acid–base theory and react with a water molecule to form conjugated pairs of acid and base (Scheme 1b), or (c) react in the same way as the previous one but in a more sophisticated version in which the formed ion is also solvated by an arbitrary number of water molecules (Scheme 1c). Regardless of the picked cycle, ΔG_s^* represents the free energy of deprotonation, which is calculated using eq 6

$$\Delta G_{\rm s}^* = \Delta G_{\rm gas}^{\circ} + \sum_{i=1}^{\rm N \, products} n_i \Delta G_{\rm solv}^*(i) - \sum_{j=1}^{\rm N \, reactants} n_j \Delta G_{\rm solv}^*(j)$$
(6)

Then, pK_a is determined using a mathematical formula (eq 7)

$$pK_{a}(\text{AntoxOH}) = \frac{\Delta G_{s(\text{AntoxOH})}^{*}}{2.303RT}$$
(7)

Despite the fact that one of the major drawbacks for this method is it requires proton or hydronium ion enthalpies, for which theoretical studies do not perfectly match, due to the impact of different solvation models used¹⁷⁸ or methodology,¹⁷⁹ it is still widely used^{27,114,139} owing to its simplicity. The question and recommended values of solvation enthalpies have been extensively elaborated by Marković and coworkers.^{178,180,181}

Another way for determining pH is the *relative method* or *isodesmic method*. It is based on the proton exchange equilibrium between the acid of interest and the conjugated base of the reference acid and is represented by the following general chemical reaction

 $AntoxOH + Ref^- = AntoxO^- + HRef$

The acid–base pair of a reference chemical is defined as HRef/Ref⁻, and pK_a is computed in the same way as in the direct method but with a modified form of the previously supplied equation, given here as eq 8

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$$pK_{a}(AntoxOH) = \frac{\Delta G_{s(AntoxOH)}^{*}}{RT\ln(10)} + pK_{a}(HRef)$$

Although this method has been shown to produce accurate results, 58,69 it requires HRef to be as structurally close to the HA as possible, and also the knowledge of the experimental value for pK_a (HRef).

Among the newer approaches, the *parameters fitting* method^{182,183} stands out not only due to its much simpler computation protocol but also the accuracy reported in the number of studies that have used it.^{36,41,46,73,85,101,112,126} It employs a linear regression model to determine pK_a values for hydroxyl, carboxylic, amino, and thiol groups in a water solvent. Mathematically, it is expressed by eq 9

 $pK_{a}(AntoxOH) = m\Delta G^{*}_{AntoxOH/AntoxO^{-}} + C$ (9)

where $\Delta G^{*}_{\text{AntoxOH/Antox}^-}$ denotes the difference in Gibbs free energy between the antioxidant's conjugated base and the corresponding acid, and *m* and *C* are empirical parameters available for 20 different functionals, each in one of four Pople's basis sets.

The former, however, is restricted solely to the water solvent. The solution is the *empirical conversion method*¹⁸⁴ presented recently. Although it does not allow for the calculation of pK_a values from scratch, the authors claim that it is useful for converting empirically determined dissociation constants in one solvent to any other with a little error.

Because both, the *parameters fitting method* and the *empirical conversion method*, were developed on strong experimental foundations, combining them in critical situations may not be such a bad idea. However, further testing is required to confirm this.

ELECTRONIC STRUCTURE INVESTIGATIONS

Intrinsic Reactivity Indices. A comprehensive study that covers the entire spectrum of antioxidative activity is difficult and time-consuming process. The intrinsic reactivity indices (Table 1) can be evaluated to obtain preliminary data that will guide further steps of the research. Although they do not consider the radical scavenged, the evaluation of preferred reaction paths of isolated species, identification of the most promising ones for a specific goal, and comparisons across antioxidants with similar chemical nature and modes of action is given by them.

Notably, reactivity indices can be calculated vertically or adiabatically, that means with or without orbital relaxation. IP and EA are particularly important in Marcus theory for calculating the activation energy of electron transfer mechanisms and are thus mostly established in this context. The following equations (eq 10 and eq 11) are used to determine their vertical values

$E_{(N-1)}(g_N) - E_N(g_N) = $ vertical IP	(10)			
$E_{(N)}(g_N) - E_{N+1}(g_N) = \text{vertical EA}$	(11)			
The adiabatic ones, on the other hand, are calculated as (eq 12 d eq 13)				

$E_{(N-1)}(g_{N-1}) - E_N(g_N) = \text{adiabatic IP} $ (12)
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$$E_{(N)}(g_N) - E_{N+1}(g_{N+1}) = adiabatic EA$$

and

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(13)

Table 1. Names, Associated]	Reactions, and Explanations of Int	trinsic Reactivity Indices
Name (typical acronym)	Related reaction	Brief description
Ionization potential (IP)	Antox(OH) _n \rightarrow Antox(OH) _n ⁺ + e ⁻	The ability to contribute an electron, which is interpreted as a willingness to oxidize itself. The lower the IP, the greater the likehood of antioxidant protection through electron transfer via electron donation. This is sometimes referred to as ionization energy (IE) in the literature.
Electron affinity (EA)	$\operatorname{Antox}(\operatorname{OH})_n + e^- \rightarrow \operatorname{Antox}(\operatorname{OH})_n^-$	The ability to accept an electron, which can be interpreted as a desire to reduce itself. The lower the EA, the more likely antioxidant protection through electron transfer via electron acceptance.
Bond dissociation enthalpy (BDE)	$Antox(OH)_n \to Antox(OH)_{n-1}O^{\bullet} + H^{\bullet}$	The amount of energy required to break the O–H bond during homolytic fission, which can be interpreted in the context of the radical's stability. The lower the BDE values, the more active the corresponding –OH residue is in the hydrogen atom transfer mechanism and the more stable the radical formed.
Proton affinity (PA)/Proton dissociation enthalpy (PDE)	$\operatorname{Antox}(\operatorname{OH})_n \to \operatorname{Antox}(\operatorname{OH})_{n\cdot 1}\operatorname{O}^- + \operatorname{H}^+$	The amount of energy required to break the bond during heterolytic fission, which can be interpreted as the anion's stability. The lower the PA/ DDE value, the more corresponding—OH residue will be deprotonated. PA is defined as the inverse of the entialpy change in a gas phase reaction between an electrically neutral chemical species and a proton to form the conjugated acid of the latter, whereas PDE is the deprotonation of a radical cation in any medium any more than the conjugated acid of the latter, whereas PDE is the

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ГаЫе	2 Names	Mathematical Formulations	and Descriptions of Indices Rel	lated to Frontier Molecular Orbitals The	eorv
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Name (typical acronym)	Related formula	Brief description
HOMO–LUMO gap (HLG)	HLG = IP - EA	Represents the ease with which the electron in a molecule can be excited from HOMO to LUMO. The lower it is, the easier the electron migrates from one another, and the radical reaction proceeds more quickly because it is more kinetically stable.
Electronegativity (χ)	$\chi = \frac{\mathrm{IP} + \mathrm{EA}}{2} = - \mu$	The general proclivity to attract electrons.
Chemical potential (μ)	$\mu = -\frac{\mathrm{IP} + \mathrm{EA}}{2} = -\chi$	Indicates the direction of charge flow as well as the capacity to contribute or accept it. Electrons will migrate from high to low μ locations in a quantity proportional to changes in μ , with a corresponding stabilizing energy μ . ²
Global hardness (η)	$\eta = \frac{IP - EA}{2}$	Measures the resistance to electron cloud polarization caused by a minor chemical disturbance or a change in electron number.
Global softness (S)	$S = \frac{\eta}{2}$	The ability to accept electrons. It is inversely proportional to chemical hardness.
Electrophilicity (ω)	$\omega = \frac{\mu^2}{2\eta}$	The ability of a system to acquire a partial charge. When two molecules are involved in a chemical reaction, the one with the higher value is considered the acceptor, while the one with the lower value is considered the donator. It is advised to be used to demonstrate the efficacy of electron donation in compounds with extremely low IP values.

Table 3. Names, Mathematical Formulations, and Descriptions of DAM-Related Indices

Name (typical acronym)	Related formula	Brief description
Electrodonating power $(\omega -)$	$\omega^- = \frac{(3\mathrm{IP} + \mathrm{EA})^2}{16(\mathrm{IP} - \mathrm{EA})}$	The ability of a chemical system to provide a fractional amount of charge. The lower the ω -, the more likely it is that the molecule will behave as an electron donor in weak interactions with other species.
Electron-accepting power (ω +)	$\omega^+ = \frac{(\mathrm{IP} + 3\mathrm{EA})^2}{16(\mathrm{IP} - \mathrm{EA})}$	The ability of a chemical system to receive a fractional amount of charge. The greater the ω +, the more likely it is that the molecule will behave as an electron acceptor in weak interactions with other species.
Donor index (R _d)	$R_{\rm d} = \frac{\bar{\omega_{\rm AntoxOH}}}{\bar{\omega_{\rm Na}}}$	
Acceptor index (R _a)	$R_{\rm a} = \frac{\omega_{\rm AntoxOH}^+}{\omega_{\rm F}^+}$	
Relative value of electron acceptance (REA)	$\text{REA} = \frac{\text{EA}_{\text{AntoxOH}}}{\text{EA}_{\text{F}}}$	
Relative value of electron donation (RIE)	$\text{RIE} = \frac{\text{IP}_{\text{AntoxOH}}}{\text{IP}_{\text{Na}}}$	

where $E_{N}, E_{(N-1)}$, and E_{N+1} denote the total energies of the N, N - 1, and N + 1 electron systems, respectively, computed at ground state geometries of $(g_N), (g_{N-1})$ and (g_{N+1}) systems.

The Hammet sigma constant is one of the tools that draws from reactivity indices. It reflects the electron withdrawing or donating capacities of substituents connected to the aromatic moiety and thus can be applied to assess their impact on the intrinsic reactivity indices in a semiquantitative manner.^{22,23,32,127} They can be also used as a features of quantitative structure—activity relationship (QSAR) models, which numerically relate them and other descriptors to experimental data.^{11,24,43} Recently, an extensive paper on QSAR development and validation was published.¹⁸⁵

Frontier Molecular Orbitals. The energy and distribution of frontier molecular orbitals, specifically the highest-occupied molecular orbital (HOMO), a nucleophilic part of the molecule, and the lowest-unoccupied molecular orbital (LUMO), an electrophilic part of the molecule, can be directly linked to antioxidative activity.¹⁸⁶ This is because Janak's theorem^{187,188} states that the energies (ε) of HOMO and LUMO are related to ionization potential (eq 14) and electron affinity (eq 15), respectively, via the following relationship:

$-\varepsilon(HOMO) = \nu IP$ (1	14)	
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$$-\varepsilon(LUMO) = \nu EA$$
 (15)

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eigenvalue is likely to be a good electron acceptor. Analyzing their values allows for the initial elucidation of the antiradical activity. Although the majority of naturally ocuring radicals are electrophillic, and so the interaction between their SOMO and antioxidants' HOMO is of the greatest importance, there also exist nucleophilic ones, typically carbon-centered, which scavenging potential is subjected to the overlap between SOMO of the reactive specie and LUMO of the scavenger. In addition, visualizing HOMO allows for a prediction of which molecular site is more vulnerable to radical attack^{28,50,116}, but presented later Fukui functions are more reliable option.

A molecule with a low HOMO eigenvalue is likely to be a poor

electron donor, whereas a molecule with a low LUMO

These approximations of orbitals egeinvalues as ionization potential or electron affinity, however, ignore electron correlation and are highly dependent on the method and basis set employed^{27,52,123,153}. This poses further problems for they serve as the foundation for a slew of global descriptive parameters related to the electrophilic character of the species (Table 2)¹⁸⁹ meaning their improper values may lead to further errors in the study and invalid conclusions. Given the importance of being as precise as possible, direct computations, are strongly advised. On the other hand, M062X/6-311G(d,p) level of theory has been found to approximate energies of frontier molecular orbitals with an error no greater than 0.5eV from the ones calculated in the direct vertical mannet.¹⁵³

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To ensure the quality of vertical energies obtained, an electron propagator theory (EPT)^{190,191} and a partial third-order quasiparticle theory (P3)¹⁹¹ can be exploited. These theories lay the groundwork for the systematic inclusion of electronic structure. They have been shown to produce lower mean errors than any other open shell techniques when compared to experimental trial results¹⁹² and are now wiedly used.^{46,101,112} For the EPT estimations to be valid, the pole strength (PS) values must be greater than 0.80–0.85.^{193,154}

Electron and Hydrogen Donation Properties. To do quick and simultaneous comparisons of the relative electron-donating and electron-accepting properties of a set of species the donator-acceptor map (DAM) is a right choice.¹⁹⁵ It is based on the assumption that in a simple charge—transfer model the response of a molecule submerged in an idealized environment that can either remove or donate charge can be represented by a quadratic interpolation for the energy as a function of the number of electrons.^{196,197} Therefore, DAM exhibits the antioxidant's tendency in charge-related processes in terms of electron-donating (ω^-) and electron-accepting (ω^+) powers (Table 3).

The substance is classified into one of four distinct zones (Figure 2) based on precalculated donor (R_d) and acceptor (R_a)



Figure 2. Schematic representation of donator-acceptor map.

indices: (1) the excellent antiradical zone (lower right), where it is both a superior electron donor (small R_d) and acceptor (high R_a), (2) the worst antiradical zone (upper left), where it is both a poor electron donor (high R_d) and a poor electron acceptor (small R_a), (3) the good antireductant zone (upper right), where it is a fine electron acceptor (high R_d and R_a) and thus an effective antiradical, and (4) the strong antioxidant area (lower left), where its good electron donor properties manifest (small R_d and R_a).

 R_d and R_a are valued for their electron-donating and electronaccepting powers, respectively, and are defined in relation to the electron-accepting power of fluorine ($\omega_{F^+} = 3.40$) and electrondonating power of sodium ($\omega_{Na} = 3.46$). F⁻ and Na⁺ are used as references due to their high electron-accepting and electronpubs.acs.org/jcim

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donating capacities. $\omega_{\rm F}^+$ and $\omega_{\rm Na}^-$ can be calculated using experimental data.³² If $R_{\rm a} < 1$, the substance is a poorer electron acceptor than F, and if $R_{\rm d} > 1$, it is a poorer electron donor than Na.

Three of the four zones represent the intended antioxidative activity. In the above graph, the electron flow goes from species in the bottom left to species in the top right. When comparing substances, DAM helps to predict which will behave as stronger or weaker oxidizers or reducers,⁸⁵ while also accounting for the effects of the environement.^{62,71} It is good to remember, there is a general trend of increasing electron-donating properties with each subsequent dissociation, so polyanionic species shall generally have lower R_d values than neutral of cationic forms.^{50,91}

Full electron-donor-acceptor map (FEDAM), an improved version of DAM, has been developed to account for the nature of the interacting free radical, which can have such a significant impact that it can even invert the relative importance of the free radical scavenging mechanisms.¹⁹⁸ FEDAM is based on the relative values of electron acceptance (REA) and electron donation (RIE) indices derived from vertical IP and vertical EA, again with Na and F atoms used as references. These are plotted on a map in the same way that DAM is but for both antioxidants and radicals. It has been used to evaluate antioxidant activity of melatonin, as well as its metabolites and derivatives.^{101,105}

The electron- and hydrogen-donating ability map for antioxidants (eH-DAMA)¹⁰¹ is another approach to enhance DAM. It was designed to identify compounds that are good donors in both electron transfer (low ω) and hydrogen atom transfer (low BDE). It is similar to DAM, but the axes changes, so the Y axis corresponds to ω , while the X axis refers to BDE. As a result, the species in the left-bottom region are more likely to work in both directions, making them particularly valuable as radical scavengers, such as new sesamol¹¹² or melatonin derivatives¹⁰¹, or modified p-coumaric acid analogs with neuroprotective activity.⁴⁶ In most cases, parent molecules, oxidants, or reference antioxidants, e.g. Trolox, are included for comparison purposes.^{46,112}

Radical Attack Site. Frontier molecular orbital theory introduces a numerical method for investigating the reactivity of individual sites of molecule in three types of reactions. Fukui functions¹⁹⁹ were proposed to represent the difference in electron density at a given point, $\rho(r)$, as a function of the number of electrons, N, at a given external potential, v(r) (eq 16). This is based on the notion that the optimal path for a reagent to approach the other species is the one with the highest initial fluctuation of the electronic chemical potential, μ .

$$f(r) = \left[\frac{\delta\mu}{\delta\nu(r)}\right]_{N} = \left[\frac{\partial\rho(r)}{\partial N}\right]_{\nu(r)}$$
(16)

Integrating over that equation for individual atoms in a molecule yields condensed Fukui functions (eqs 17-19), which are a more convenient way of predicting the reaction site than visualization of HOMO or LUMO orbitals. In general, the higher the value, the more reactive this position is to the specific type of attack. For an arbitrary atom A, these functions are defined as

$$f_{\rm A}^- = q_{\rm N-1}^{\rm A} - q_{\rm N}^{\rm A}$$
, for electrophilic attack (17)

$$f_A^+ = q_N^A - q_{N+1}^A$$
, for nucleophilic attack (18)

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$$f_{\rm A}^{0} = \frac{[f_{\rm A}^{+} + f_{\rm A}^{-}]}{2} = \frac{[q_{\rm N-1}^{\rm A} + q_{\rm N+1}^{\rm A}]}{2}, \text{ for radical attack}$$
(19)

where $q_{N-1}^{\rm A}, q_{N+1}^{\rm A}$ represent charges of (N-1) and (N+1)systems obtained vertically from the optimized ground state geometry (N) with charge q_N^A . This ensures that the values are solely determined by the atom's electronegativity and electron density location. At the same time, because of that the atomic charge values used to estimate condensed Fukui functions are heavily reliant on population analysis method.^{124,200,201} Given their varied formulations,²⁰² the condensed Fukui functions may exhibit significant mutual discrepancies-even negative values²⁸—but this has already been explained as a result of the small interatomic distances between atoms.^{203,204} So Atoms-in-Molecule tend to overestimate them;²⁰⁴ Natural Population Analysis is heavily influenced by the functional, and basis set chosen;²⁰⁵ and ESP derived charges, such as ChelpG, have nonsmooth geometry dependence.²⁰⁰ Also a solvent may also affect the final result.¹²⁴ Some studies propose stockholder charge partitioning approaches in this purpose, particularly Hirschfeld analysis,^{15,64,123} since it yields values for which condensed Fukui functions correlate well with the expected data. In fact, a "dual descriptor" better illustrates susceptibility for electrophilic–nucleophilic attack,²⁰⁶ but these types of reactions are beyond the scope of antiradical activity studies.

Redox Potentials. As it was already mentioned, primary activity of antioxidants is related to the direct reduction of radicals. Therefore, the viability of this process can be assessed in terms of electrochemical potentials.²⁰⁷ The Born–Haber thermodynamic cycle is one universal method due to the common pattern of reactivity, and a half-reaction describing that process is always denoted by

AntoxOH^{+•}_(aq) + e⁻ \rightarrow AntoxOH_(aq)

for which the standard reduction potential can be calculated using the Nernst equation (eq $20)\,$

$$E_{\rm red(aq)}^{\circ} = -\frac{\Delta G_{\rm red}^{\circ}}{nF}$$
(20)

where ΔG_{red}^* denotes the standard Gibbs free energy of the reduction, and *n* is the number of electrons transferred and *F* the Faraday's constant (23.06 kcal/molV). According to the Born–Haber cycle, ΔG_{red}^* equals

$$\Delta G_{\text{red}}^* = G^*(\text{AntoxOH}^{+\bullet}) - G^*(\text{AntoxOH}) - nG^{\circ}(e^{-})$$
(21)

with $G^{\circ}(e^{-})$ being free energy of one electron in the gas phase (-0.876 kcal/mol at 298 K). When comparing results to those obtained experimentally, the computed values must be reduced by an absolute potential of the reference electrode, for example, a standard iron electrode or a standard hydrogen electrode.

Notably, because certain compounds may be partially deprotonated at a given pH, and anions have a lower capacity to donate electrons than neutral forms, they also influence the overall reduction potential. As a result, the average can be calculated accordingly to eq 22

$$E = E^{\text{AntoxOH}^{+\bullet}|\text{AntoxOH}} + \frac{RT}{F} \ln(f_{\text{AntoxOH}^{+\bullet}}) - \frac{RT}{F} \ln(f_{\text{AntoxOH}})$$
(22)

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where $E^{\text{AntoxOH}+\bullet|\text{AntoxOH}}$ represents the initial reduction potential, and f_{AntoxOH} denotes the population of the oxidized species that are reduced to produce the species with population f_{AntoxOH} .

 f_{AntoxOH} . **Topology Analysis.** Hydrogen bonds and weak interactions are important in stabilizing radicals or transition states rendering reactions to be more feasible, for example, by lowering BDE values.⁵⁰ Bader's Quantum Theory of Atoms in Molecules (QTAIM) can be used to quantify their strength^{208,209} and thus has been widely applied in antioxidants research.^{31,58,62,103,122,130} Despite the analysis's complexicity, specialized tools such as the Multiwfn software²¹⁰ eases it.

The nature of chemical bonds at bond critical points (BCPs) can be mathematically described by electron densities, ρ , and their associated Laplacian, $\nabla^2 \rho$. The theoretical background is explained in detail in Bader's paper;²⁰⁸ here, it will only be mentioned that BCPs of primary importance in such studies correspond to (3, -1) critical points, being the saddle points with a maximum of electron density in two directions of space and a minimum in the third, and that the number of BCPs must obey the Poincare–Hopf rule.²¹¹

The presence of a bond path between two atoms with a BCP in the middle is the first sign of a bond presence.^{208,215} A second criterion for defining it is that the values of $\rho(\text{BCP})$ and $\nabla^2 \rho(\text{BCP})$ are positive, in ranges of 0.002–0.035 and 0.024– 0.139, respectively.^{215–217} $\nabla^2 \rho(\text{BCP})$ can be expanded as the sum of the eigenvalues $\lambda_1, \lambda_2, \lambda_3$, obtained by diagonalizing the Hessian of the electron density and mutually related as $\lambda_1 < \lambda_2 < \lambda_3$. They can be used to calculate the ellipticity parameter, $\varepsilon = \frac{\lambda_1}{\lambda_2} - 1$, which quantifies the amount of charge that accumulates preferentially. A large ε indicates topological instability and, as a result, an easily ruptured bond. The λ_3 specifies how easily the BCP can be moved along the bond path,²⁰⁸ and the higher the value is, the stronger the interaction is. A local formulation of the virial theorem²⁰⁸ relates $\nabla^2 \rho(BCP)$ to electronic topological parameters by eq 23

$$\frac{\nabla^2 \rho(\text{BCP})}{4} = 2G(\text{BCP}) + V(\text{BCP})$$
(23)

where G(BCP) is the Lagrangian kinetic electron density, and V(BCP) represents the potential electron density (also known as the virial field).

Positive $\nabla^2 \rho(\text{BCP})$ values indicate that G(BCP) is greater than V(BCP), implying that charge is being depleted along the bond path, as is typical of closed-shell interactions such as hydrogen bonding, ionic bonds, and van der Waals. Its negative values, on the other hand, indicate an excess potential energy at BCP in the form of internuclear charge concentration, which corresponds to covalent interactions; in this case, an electron density is localized in between two nuclei and is mutually accessible to both of them.²¹²

Similarly, the -G(BCP)/V(BCP) ratio can be used for that purpose, because -G(BCP)/V(BCP) > 1 indicates that the intramolecular bond is closed and noncovalent, while 0.5 < -G(BCP)/V(BCP) < 1 points out that it is shared—for example partially covalent or ionic.^{208,213,214} Espinosa and coworkers^{219,220} demonstrated that interatomic

Espinosa and coworkers^{219,220} demonstrated that interatomic interaction energy can be related to potential electron energy density at BCP using the following expression

$$E_{\rm HB} = \frac{V(\rm BCP)}{2} \tag{24}$$

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The above relationship, according to Rozas et al.,²¹⁸ allow hydrogen bonds to be classified as weak ($E_{\rm HB} < 12.0 \, \rm kcal/mol$), when $\nabla^2 \rho(\rm BCP) > 0$ and $H(\rm BCP) > 0$, medium ($12.0 < E_{\rm HB} < 24.0 \, \rm kcal/mol$) if $\nabla^2 \rho(\rm BCP) > 0$ and $H(\rm BCP) < 0$, or strong ($E_{\rm HB} < 24.0 \, \rm kcal/mol$) when $\nabla^2 \rho(\rm BCP) < 0$ and $H(\rm BCP) > 0$, where $H(\rm BCP)$ due to the density of electrons total energy, $G(\rm BCP) + V(\rm BCP)$.

This is one of the most useful methods for calculating the energy of hydrogen bond interactions. Furthermore, Korth et al.²²¹⁷ demonstrated how to compute the relative intramolecular hydrogen bond enthalpy by comparing the sum of the conformer's electronic and thermal enthalpies with intra-molecular hydrogen bonds.

However, QTAIM analysis must to be used with caution in course of the studies. The molecule's wavefunction, which is used to evaluated the aforementioned interactions, is determined by the functional and basis set used. As evidenced from refs 222 and 223, no relevant relationship between climbing Perdew's Jacob's ladder rungs and BCP densities was reported. The basis set, on the other hand, appears to be of primary concern. It has been demonstrated²²⁴ that small, double- ζ basis sets from Pople or Dunning's families are insufficient to accurately assess the properties of BCP related to multiple and polar bonds, as well as weak hydrogen bond interactions. Instead, at least triple- ζ are recommended, which is plausible in the context of the current trend in their choice in the studies on antioxidants.

Natural Bond Orbitals. One of the primary requirements for an antioxidant to effectively scavenge free radicals is that it becomes stable after the reaction. The spin density distribution throughout the molecule, which is often larger for conjugated systems, can be used to examine that property;^{28,35,0,82} however, natural bond orbital analysis^{225–228} represents a much more detailed investigation into the topic.

Refining the wavefunction into a Lewis-like structure corresponding to lone pairs and bonds gives an opportunity to track charge transfer by examining changes in the electron density at bonds, investigate hybridization of the orbitals and bonding interactions, as well as study delocalization and hyperconjugation effects. ^{10,34,61,67,85,103,123} During the natural bond orbital analysis, the stabilization energy, $E^{(2)}$, is derived for the electron transfer from filled donor orbital, *i*, to an empty acceptor orbital, *j*, and is related by eq 25

$$E^{2} = -q^{i} \frac{(F_{ij})^{2}}{(E_{j} - E_{i})}$$
(25)

where F_{ij} is the off-diagonal Fock matrix element and q^i the orbital occupancy, and E_j and E_i are diagonal elements. The intrepretion is clear —, the greater the E^2 energy, the greater degree of interaction.

NBO analysis, just like QTAIM, also requires careful application. Alhough it is a straightforward and advantageous method, at the same it is heavily reliant on the geometry of the compound, which stems from the partitioning scheme of the electron density matrix and the localized nature of molecular orbitals. A simple, yet excellent, example of this can be found in the paper of Benassi and Fan²²⁹ where the authors reported on how the delocalization energy and orbital occupancy number differ in pyridine across its seven normal modes and small changes along the displacement coordinates. It has been demonstrated that even minor shift can result in significantly

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different $E^{(2)}$ values, which is expected to be amplified in the case of larger antioxidant structures.

REACTIVITY

Thermochemistry. In reality, the antioxidant's ability to scavenge radicals is influenced not only by the antioxidant itself but also by the species with which it interacts. As a result, thermochemical calculations of known pathways (Table 4) produce far more useful data than intrinsic reactivity indices alone; one way to illustrate them is with O'Ferral-Jencks diagrams^{83,86} (Figure 3). However, before proceeding, a foreword is required: hydroxyl radical, •OH, is so reactive that it quickly reacts with almost any molecule in its vicinity at diffusion-limited rates, before an antioxidant can actually reach it. For this reason, it is skipped from computations and radicals with intermediate to low reactivity are chosen instead for assessing the antiradical potential.^{230–232} Peroxyl radicals, ROO[•] (such as $CH_3OO^{•}$ or $OOH^{•}$),^{233,234} are one of these because their half-lives are long enough that they can be intercepted before oxidizing biological targets.^{235,236} Although acid-base equilibrium of hydroperoxide equals 4.8, what means that O_2 - is typically present in physiological conditions, it is not very reactive species and the oxidation damage are primiarly steming from its protonated form. 237 CCl₃OO $^{\bullet}$ can also be considered because it is used in experiments to simulate larger radicals.²³⁸ Another possible reduction target might be the H_2O_2/O_2 pair, which is more difficult to neutralize than other reactive oxygen species.2

The polarity of the environment also has an effect on reaction energetics. To begin with, it should come as no surprise that reactions generating neutral species, such as RAF or HAT, perform better in nonpolar solvents than reactions that produces ions. This is due to the fact that nonpolar media do not provide enough solvation to stabilize charged species through the charge separation, thereby propelling the reaction forward. In consequence, such reactions are unlikely to occur in a significant number, and conventional studies in nonpolar media focus solely on RAF and HAT, with the remaining pathways being completely ignored.^{60,63,106,121,139,143}

Furthermore, the first step in a multistep mechanism is thermodynamically significant and so determining its energetics allows for assessing reaction's feasability, simplifying the analysis by rejecting unfavorable pathways. Because BDE, IP, and PA are the primary indices of the HAT, SET-PT, and SPLET,^{240–243} they can be used for this purpose.

Moreover, because SPLET mechanism is initiated by proton dissociation, which proclivity is controlled by the environment's pH and acid—base equilibrium, the SET and SPLET processes are extremely closely coupled due to the spontanous. If molar fractions are being considered at the outset, the second step of SPLET actually controls an antioxidant's reactivity, and in this case the entire mechanism becomes equivalent to SET; namely, it is identical to the SET reaction for an acid—base species with N - 1 protons. Herein, I will just mention that it is also an electrostatic potential map, which is a useful tool for distinguishing between electrophilic and nucleophilic centers, highlights positively charges hydroxyl hydrogen to be likely involved in proton dissociation mechanisms.^{28,568}

Finally, because IP and EA values govern electron flow between antioxidant and radical, they can be used to estimate the direction of the SET mechanism, providing an early picture of the process. In general, the bare minimum for SET reactions with electrophilic radicals is IP(antioxidant) < EA(radical) and

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Table 4. Naming, Associ	ted Reactions, and Descriptions of the Mos	: Commonly Studied Antioxidative Reaction Pathways
Name (typical acronym)	Related reaction	Description
Radical adduct formation (RAF)	$Antox(OH)_n + X^{\bullet} \rightarrow [Antox(OH)_nX]^{\bullet}$	In a single step, the radical forms an adduct with the antioxidant, spreading the spin density across the newly formed molecule. The preferred reaction site is determined by the degree of the unpaired electron delocalization.
Hydrogen atom transfer (HAT)/Proton coupled elec- tron transfer (PCET)	$Antox(OH)_n + X^\bullet \rightarrow Antox(OH)_{n-1}O^\bullet + HX$	A one-step mechanism in which an O–H bond is homiletically broken and a hydrogen atom is transferred from antioxidant to free radical, resulting in a more stable antioxidant radical. Low BDE values are common in compounds that promote this path. In the electrochemical sense, it is a reduction process. Although the products of HAT and PCET reactions are identical, the former involves the coordinated transfer of a proton and an electron as a single entity, whereas PCET involves the process of two spatted particles, not necessarily from the same sets of orbitals. Formal HAT refers to dominal reacts that have not been defined as HAT or PCET.
Single electron transfer (SET)	$\operatorname{Antox}(\operatorname{OH})_n + X^\bullet \to \operatorname{Antox}(\operatorname{OH})_n^{\bullet+/\bullet-} + X^{-/+}$	Depending on the mutual IP and EA values, a single electron transfer occurs from an antioxidant to a radical to an antioxidant. The deprotonation influences the thermochemical viability of the SET process to some extent.
Sequential electron transfer-pro- ton transfer (SET-PT)		It occurs in two steps: first, a radical cation Antox(OH), ^{$++$} is formed by electron transfer from an antioxidant to a free radical, and then, it deprotonates to form Antox(OH), _{$++(0+)$ species. The first step is described by the IP values.}
1. Electron transfer	1. Antox(OH) _n + X [•] \rightarrow Antox(OH) _n ^{•+} + X ⁻	
2. Proton transfer	2. Antox(OH) _n ^{•+} \rightarrow Antox(OH) _{n-1} O [•] + H ⁺	
Sequential proton loss-electron transfer (SPLET)		The mechanism is divided into two steps: first, an antioxidant is deprotonated (as described by PA), and then, an electron transfer occurs from the deprotonated antioxidant to a free radical (described by IP). Because pK, values influence the amount of deprotonated species in aqueous
1. Proton loss	1. Antox(OH) _n \rightarrow Antox(OH) _{n-1} O ⁻ + H ⁺	solution, knowing their number <i>a priori</i> can assist in determining the relative importance of this process.
2. Electron transfer	2. Antox(OH) _{n-1} O ⁻ + X [•] \rightarrow Antox(OH) _{n-1} O [•] + X ⁻	
Sequential proton loss-hydro- gen atom transfer (SPLHAT)		The mechanism is identical to SPLET, except that instead of an electron, a hydrogen atom is transferred in the second step. As a result, antioxidants containing at least two hydroxyl groups are particularly appealing PA describes the first step, and BDE describes the second.
1. Proton loss	1. Antox(OH) _n \rightarrow Antox(OH) _{n-1} O ⁻ + H ⁺	
2. Hydrogen atom transfer	2. Antox $(OH)_{n-1}O^- + X^\bullet \rightarrow Antox(OH)_{n-2}(O)_2^{\bullet^+} + HX$	

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Figure 3. O'Farell-Jencks diagram of each single step involved in common reaction mechanisms.

opposite holds true for the nucleophilic ones. In case of not so easily recognizable species, a general rule of knowing that electrons flow from the structure of lower IP to the structure of greater EA, makes it possible to predict which molecule will undergo oxidation and which will undergo reduction, and the reverse path can be thereby ignored.

One final point to mention about thermochemical calculations is that they require the Gibbs free energies of electron, proton, and hydrogen. The very last can be estimated directly at the applied level of theory, but the proton and electron cannot, at least not in a straightforward manner, necessitating the use of widely varying reference values.^{178,180,181,244}

Kinetics. An antioxidant is any substance that, even at low concentrations, significantly delays or prevents the oxidation of radical target. Therefore, it must not only react spontaneously with the oxidizing agent, but it must also react faster than the target it is designed to protect.^{245,246} This aspect can only be modeled using kinetic studies, which account for facets skipped by thermochemical studies-tunneling effects, weighted contribution of different mechanisms and different species to total antioxidative potential, or adherence to the Bell-Evans-Polanyi principle³—and is thus critical for accurately assessing antioxidative behavior.

Although spontaneity is an important criterion for chemical reactivity, it is not always enough because an exergonic reaction can occur at either fast or slow rates. When drawing conclusions from thermochemical data without considering kinetics, the Bell–Evans–Polanyi 247,248 principle, which states that the most exergonic processes have the lowest activation energies and are thus kinetically favored, is implicitly assumed to be followed. On the other hand, ignoring a reaction path due to difficulties in locating a transition state may result in a more significant error than accepting the given rule wihout its confirmation, especially if it was discovered to hold true for a structurally similar compounds.¹¹⁸ It has been recently evidenced that Gibbs free energy is actually proportional to the activation energy of hydrogen atom transfer, and hence reaction rate of this mechanism.90,

Endergonic channels do not need to be included in kinetic calculations in general because even if they occur at high rates,

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they are reversible to the point where no products are detected. However, moderately endergonic processes (typically with ΔG° < 10.0 kcal/mol) may still contribute to antioxidant capacity and should be addressed, particularly if their products evolve into other species quickly, providing a driving force, or the reaction barriers are low.¹⁸² This is the case for some RAF reactions,⁸² but it is particularly common for SET, where thermochemical and kinetic data may show opposing trends; in some cases, the mechanism may be associated with positive Gibbs energies and still be relevant, as has been demonstrated in a number of previously reported studies.^{36,46,73,82,154} Highly exergonic SET reactions involving donors with very low IP (such as monoanionic or polyanionic species) may, on the other hand, be found in the inverted region of the Marcus parabola, where that reaction barriers increase as ΔG° decreases what is often to be found for Gibbs free energies much lower than negative of reorganization energy. $^{251-253}$ Despite this is an unexpected behavior, it underlines that particles with extremely low IP are unlikely to be efficient free radical scavengers.^{46,101,125} That is why using ionization potentials may be misleading, and electrondonating power or electrophilicity is far superior because, while it also relies on IP, it does so in a nonlinear fashion, with the shape of this dependency resembling the Marcus parabola.

The activation barrier of a reaction is determined by the energy difference between the transition state and the reactants. However, in the case of electron-related processes, assessing it is not so straightforward. The barrier of electron transfer reaction $(\Delta G_{\rm ET}^{\pm})$ is calculated in a different way, using the Marcus theory^{153–255} (eq 26), which defines it in terms of the reaction adiabatic free energy $(\Delta G_{\rm ET}^0)$ and nuclear reorganization energy (λ)

$$\Delta G_{\rm ET}^{\neq} = \frac{\lambda}{4} \left(1 + \frac{\Delta G_{\rm ET}^0}{\lambda} \right)^2 \tag{26}$$

The reorganization energy is calculated as the difference between the vertical $(\Delta E_{\rm ET})$ and adiabatic free energies of reaction and accounts for the orbitals relaxation

$$\lambda = \Delta E_{\rm ET} - \Delta G_{\rm ET}^0 \tag{27}$$

The reaction rate constants (k) can be calculated using the conventional transition state theory (TST) which is one of the most robust theoretical methodologies for this purpose, requiring only structural, energy, and vibrational frequency information for reactants and transition states, allowing it to be applied to a wide range of chemical processes.^{256–258} Despite its simplicity, it has been shown to reproduce fine the experimentally measured data on free radicals scavenging kinetics.^{82,84,96,129} It is computed usually in the framework of 1 M standard state²⁵⁸ using the Eyring equation (eq 28)

$$k^{\rm TST} = \frac{k_{\rm B}T}{h} {\rm e}^{-\Delta G^{\neq}/RT}$$
(28)

where $k_{\rm B}$ and h are the Boltzmann and Planck constants, respectively. ΔG^{\dagger} is the free energy of activation, calculated as the difference in energies between transition state and reactants, while R and T denote the gas constant and temperature, respectively.

The more sophisticated Eckart approach,²⁵⁹ also known as the zero-tunneling method, employs the Boltzmann average of the ratio of quantum and classical probabilities²⁶⁰ and is suggested

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for processes in which reactants are transformed into products over energy barriers. Such processes include the HAT reaction, which involves the motion of a light particle (here H[•]) that can easily tunnel, as well as some RAF pathways. Tunneling corrections ($\kappa(T)$), also known in the literature as transmission coefficients ($\gamma(T)$), are included, as is reaction path degeneracy (σ). $\kappa(T)$ can significantly modulate reaction rate values between relatively similar reacting species, such as CH₃OO[•] or OOH[•].¹¹⁸ It can be estimated using external software, for example Eyringpy.²⁶¹ The number of identical reaction paths, on the other hand, is reflected by σ and can be calculated by labeling all similar atoms and counting the number of different but equivalent configurations that can be formed by rotating, but not reflecting them. They can also be established using the Pollak and Pechukas²⁶² scheme (Table S). Taking everything into account, eq 29 is obtained

$$k^{\rm TST} = \sigma \kappa(T) \frac{k_{\rm B}T}{h} {\rm e}^{-\Delta G^{\#}/RT}$$
(29)

Table 5. Point Groups and the Reaction Path Degeneracy Values That Correspond to Them

Point group	σ	Point group	σ
C_1	1	D_{3h}	6
Cs	1	D_{5h}	10
C2	2	D _{coh}	2
C_{2v}	2	D _{3d}	6
C_{3v}	3	T _d	12
C _{cov}	1	O _h	24
D _{2h}	4		

As previously stated, radicals with high reactivity often react at diffusion-limited rates ($k \ge 10^8 \, {\rm M}^{-1} \, {\rm s}^{-1}$) with the vast majority of chemical compounds. For an irreversible bimolecular diffusion-controlled reaction, the Collins–Kimball theory,²⁶³ in conjunction with the steady-state Smoluchowski rate constant²⁶⁴ and the Stokes–Einstein approaches,²⁶⁵ must be used to calculate rate constants properly (Table 6). They are frequently applied in case of RAF mechanisms, which is not surprising given that these types of radical attacks usually occur in the absence of energy barriers, making it difficult to localize the transition state.⁸²

The total reaction rate constant values (k^{TOT}) for all acid– base species (i) present at the specified pH multiplied by the corresponding molar fractions (f) allow the overall reaction rate constant (k^{overall}) , which corresponds to the empirically observed reaction rate, to be calculated

$$k^{\text{overall}} = \sum_{i = \{\text{species}\}} f(i)k^{\text{TOT}}(i)$$
(30)

The reaction rate constants for each antioxidative mechanism (j) are then added to calculate the $k^{\rm TOT}$ values for all acid—base species

$$k^{\text{TOT}} = \sum_{j=\{\text{mechanism}\}} k^{\text{mech}}(j)$$
(31)

The k^{mech} (eq 32) is defined as the sum of reaction rate constants (k^{TST} or k^{app} , depending on the kinetic model used, here simply represented by k) from to the same antioxidative mechanism but calculated at different reaction sites (l)

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Table 6. Collins-Kimball Theory, Steady-State Smoluchowski, and Stokes-Einstein Equation Mathematical Formulations

Name	Related formula	Variables
Collins-Kimball theory	$k^{\rm app} = \frac{k^{\rm D} k^{\rm TST}}{k^{\rm D} + k^{\rm TST}}$	$k_{\rm app}:$ apparent rate constant
Steady-state Smoluchowski	$k^{\rm D} = 4\pi R D_{\rm AB} N_{\rm A}$	$k^{\rm D}$: steady-state Smoluchowski $k^{\rm TST}$: thermal rate constant (obtained from TST) R: reaction distance $N_{\rm A}$: Avogadro number $D_{\rm AB}$: mutual diffusion coefficient of the reactants A (free radical) and B (antioxidant)
Stokes–Einstein equation	$D_{\rm AorB} = \frac{k_{\rm B}T}{6\pi\eta r_{\rm AorB}}$	k _B : Boltzmann constant T: temperature η: viscosity of solvent r: radius of solute

$$k^{\text{mech}} = \sum_{l = \{\text{pathway}\}} k(l) \qquad \qquad \text{ter} \\ (32) \qquad \qquad \text{exi}$$

The branching ratio ($\Gamma(l)$, (eq 33) can be used to calculate the percentage contribution of an antioxidative pathway (l) to the total reaction rate (l) using the following formula

$$\Gamma(l) = \frac{k(l)}{k^{\text{overall}}} \tag{33}$$

Furthermore, the relative antioxidative activity of the examined molecule can be calculated by dividing its k^{overall} by the k^{overall} of ref 45 (e.g., Trolox, Tx) as presented in eq 34

$$r^{T} = \frac{k^{\text{overall}}}{k_{\text{Tx}}^{\text{overall}}}$$
(34)

A threshold of $k^{\text{overall}} = 1.2 \times 10^3 \text{ M}^{-1} \text{ s}^{-1}$ was proposed for quantifying antioxidant activity as a value close to the rate constant of the interaction between HOO[•] and polyunsaturated fatty acids^{3,21,126}. Compounds with higher k^{overall} values are thought to be effective antioxidants, while those with lower are thought to be ineffective.

Thermochemistry and Kinetic Ensemble. Galano, Alvarez-Idaboy and coworkers combined the above considerations into the quantum mechanics-based test for overall free radical scavenging activity (QM-ORSA) protocol,^{3,154,266} which is proposed as a feasible tool to assess radical scavenging activity in physiologically relevant solvents. It entails the impact of all existing acid–base forms, identifying RAF, HAT, and SET mechanisms viabiality, and then subjecting them to kinetic analysis. This method has been already used successfully in numerous papers.^{41,45,58,69,82,88,104,118}

CONCLUDING REMARKS AND ADDITIONAL CONSIDERATIONS

Antioxidants were and are extensively studied using computational quantum chemistry. A number of useful tools are developed to provide insight into their electronic structure and reactivity. Importantly, scientists are not limited in their research to already known procedures but continously suggest new ones and apply them in their research. Therefore, each paper proposes some novel techniques in this matter, as well as new conclusions that provide greater insight into the chemistry of the antioxidants that have been already studied or projected from the scratch. As previously stated, because all antioxidants tend to exhibit similar patterns of activity, they can all be examined in the same way, whether we are talking about fullerenes, natural plant products, or entirely newly synthesized structures.

Certain points, however, should be reconsidered or reinterpreted:

- Almost all of the research considered reactive oxygen species, with only a minor focus on radicals containing nitrogen, carbon, and sulfur elements. These are known to be linked to oxidative stress,^{267–269} so research into them would fill the gap.
- When discussing in vivo activity, although solvation effects and hence deprotonation are considered, the importance of metabolites should be emphasized. The gut microbiota are the first to alter antioxidant structure, influencing primarily their absorption profile. They are then metabolized in the liver, where they undergo further changes that result in products with vastly different activity. By focusing solely on generic structures the outcomes are indeed limited to *in vitro* results, casting doubt on the physiological activities as free radicals scavengers, capable of halting the development of oxidative stress and resulting diseases.
- The dual nature of free radicals extends to the products derived from the antioxidants. Castañeda-Arriaga et al.¹⁰² discussed possible prooxidant behavior of such, in which the new forms may still be capable of oxidizing biologically relevant structures. The authors also proposed them to be able to self-regenerate, allowing them to work multiple times. This fresh topic appears to be a relevant aspect that needs to be considered when attempting to prove experimental observations.

DATA AND SOFTWARE AVAILABILITY

Multiwfn (current version 3.7) is a free, open-source software developed by Tian Lu for wave function analysis and visualization. It is available for download at http://sobereva. com/multiwfn/. Eyring.py (current version 2.0) is a Pythonbased program developed by the Merino et al. group for computing rate constants of unimolecular or bimolecular reactions in the gas phase and in solution using transition state theory. For that purpose, it takes into account reaction symmetry, tunneling corrections, Collins–Kimball theory, Marcus theory, and species molar fractions. It is available at

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https://www.theochemmerida.org/eyringpy. GROMACS (current version 2021.5) is a free package, available under a LGPL license, created to perform molecular dynamics of proteins, lipids, nucleic acids as well as nonbiological systems. It is accessible from http://www.gromacs.org. Amber (current version 20) is a paid versatile suite of biomolecular simulation programs. It can be found at https://ambermd.org/index.php. CHARMM is a molecular simulation program that uses enhanced sampling methods and multiscale techniques to study many-particle systems. For academic users, the version "charmm" is free. The software is available for free download at https://academiccharmm.org.

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Notes

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> https://doi.org/10.1021/acs.jcim.2c00104 J. Chem. Inf. Model. 2022, 62, 2639–2658

V. Wykaz publikacji potwierdzony przez Bibliotekę UMW

Wrocław, 12.04.2023 r.

Maciej Spiegel

Wykaz publikacji

1. Publikacje w czasopismach naukowych

1.1 Publikacje w czasopiśmie z IF

Lp.	Opis bibliograficzny	IF	Punkty ministerialne
1	Spiegel Maciej , Andruniów Tadeusz, Sroka Zbigniew: Flavones' and flavonols' antiradical structure-activity relationship - a quantum chemical study, Antioxidants, 2020, vol. 9, nr 6, art.461 [21 s.], DOI:10.3390/antiox9060461	6,313	100
2	Spiegel Maciej, Kapusta Karina, Kołodziejczyk Wojciech, Saloni Julia, Żbikowska Beata, Hill Glake A., Sroka Zbigniew: Antioxidant activity of selected phenolic acids - ferric reducing antioxidant power assay and QSAR analysis of the structural features, Molecules, 2020, vol. 25, nr 13, art.3088 [15 s.], DOI:10.3390/molecules25133088	4,412	140
3	Machalska Ewa, Zając Grzegorz, Wierzba Aleksandra J., Kapitan Josef, Andruniów Tadeusz, Spiegel Maciej , Gryko Dorota, Bour Petr, Barańska Małgorzata: Recognition of the true and false resonance Raman optical activity, Angewandte Chemie-International Edition, 2021, vol. 60, nr 39, s. 21205-21210, DOI:10.1002/anie.202107600	16,823	200
4	Malchrzak Wojciech, Mastalerz-Migas Agnieszka, Sroka Zbigniew, Spiegel Maciej: One year of the COVID-19 pandemic. What do we know and what is yet to come? - the summarising review, International Journal of Public Health, 2021, vol. 66, art.1603975 [10 s.], DOI:10.3389/ijph.2021.1603975	5,1	100
5	Spiegel Maciej , Gamian Andrzej, Sroka Zbigniew: Antiradical activity of beetroot (Beta vulgaris L.) betalains, Molecules, 2021, vol. 26, nr 9, art.2439 [20 s.], DOI:10.3390/molecules26092439	4,927	140
6	Spiegel Maciej , Gamian Andrzej, Sroka Zbigniew: A statistically supported antioxidant activity DFT benchmark - the effects of Hartree-Fock exchange and basis set selection on accuracy and resources uptake, Molecules, 2021, vol. 26, nr 16, art.5058 [22 s.], DOI:10.3390/molecules26165058	4,927	140
7	Spiegel Maciej , Marino Tiziana, Prejanò Mario, Russo Nino: On the scavenging ability of scutellarein against the OOH radical in water and lipid-like environments: a theoretical study, Antioxidants, 2022, vol. 11, nr 2, art.224 [9 s.], DOI:10.3390/antiox11020224	7,675*	100

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Lp.	Opis bibliograficzny	IF	Punkty ministerialne
8	Dudek Anita, Spiegel Maciej , Strugała-Danak Paulina, Gabrielska Janina: Analytical and theoretical studies of antioxidant properties of chosen anthocyanins; a structure-dependent relationships, International Journal of Molecular Sciences, 2022, vol. 23, nr 10, art.5432 [18 s.], DOI:10.3390/ijms23105432	6,208*	140
9	Spiegel Maciej : Current trends in computational quantum chemistry studies on antioxidant radical scavenging activity, Journal of Chemical Information and Modeling, 2022, vol. 62, nr 11, s. 2639- 2658, DOI:10.1021/acs.jcim.2c00104	6,162*	100
10	Strugała-Danak Paulina, Spiegel Maciej , Hurynowicz Kacper, Gabrielska Janina: Interference of malvidin and its mono- and di- glucosides on the membrane - combined in vitro and computational chemistry study, Journal of Functional Foods, 2022, vol. 99, art.105340 [13 s.], DOI:10.1016/j.jff.2022.105340	5,223*	100
11	Spiegel Maciej , Marino Tiziana, Prejanò Mario, Russo Nino: Primary and secondary antioxidant properties of scutellarin and scutellarein in water and lipid-like environments: a theoretical investigation, Journal of Molecular Liquids, 2022, vol. 366, art.120343 [9 s.], DOI:10.1016/j.molliq.2022.120343	6,633*	100
12	Spiegel Maciej, Krzyżek Paweł, Dworniczek Ewa, Adamski Ryszard, Sroka Zbigniew: In silico screening and in vitro assessment of natural products with anti-virulence activity against Helicobacter pylori, Molecules, 2022, vol. 27, nr 1, art.20 [20 s.], DOI:10.3390/molecules27010020	4,927*	140
13	Spiegel Maciej, Marino Tiziana, Prejanò Mario, Russo Nino: Antioxidant and copper-chelating power of new molecules suggested as multiple target agents against Alzheimer's disease. A theoretical comparative study, Physical Chemistry Chemical Physics, 2022, vol. 24, nr 26, s. 16353-16359, DOI:10.1039/d2cp01918c	3,945*	100
14	Spiegel Maciej , Sroka Zbigniew: Natural dihydroisobenzofuran derivatives as a template for promising radical scavengers: theoretical insights into structure - activity relationships, thermochemistry and kinetics, Theoretical Chemistry Accounts, 2022, vol. 141, nr 11, art.61 [14 s.], DOI:10.1007/s00214-022- 02922-5	2,154*	70
15	Selahi Daniel, Spiegel Maciej , Hadzik Jakub, Pitułaj Artur, Michalak Filip, Kubasiewicz-Ross Paweł, Dominiak Marzena: The appliance of A-PRF and CGF in the treatment of impacted mandibular third molar extraction sockets - narrative review, Applied Sciences-Basel, 2023, vol. 13, nr 1, art.165 [12 s.], DOI:10.3390/app13010165	2,838*	100
16	Spiegel Maciej , Sroka Zbigniew: Quantum-mechanical characteristics of apigenin: Antiradical, metal chelation and inhibitory properties in physiologically relevant media, Fitoterapia, 2023, vol. 164, art.105352, DOI:10.1016/j.fitote.2022.105352	3,204*	100

Lp.	Opis bibliograficzny	IF	Punkty ministerialne
17	Spiegel Maciej , Cel Katarzyna, Sroka Zbigniew: The mechanistic insights into the role of pH and solvent on antiradical and prooxidant properties of polyphenols — Nine compounds case study, Food Chemistry, 2023, vol. 407, art.134677 [10 s.], DOI:10.1016/j.foodchem.2022.134677	9,231*	200
18	Spiegel Maciej , Ciardullo Giada, Marino Tiziana, Russo Nino: Computational investigation on the antioxidant activities and on the Mpro SARS-CoV-2 non-covalent inhibition of isorhamnetin, Frontiers in Chemistry, 2023, vol. 11, art.1122880 [13 s.], DOI:10.3389/fchem.2023.1122880	5,545*	100
19	Maciej Spiegel, Carlo Adamo : Tuning the Photophysical Properties of Ru(II) Photosensitizers for PDT by Protonation and Metallation: A DFT Study, Journal of Physical Chemistry A, 2023, [11 s.] https://doi.org/10.1021/acs.jpca.3c00839 [Ahead of Print]	2.944*	100

*IF 2021

2. Abstrakty

Lp.	Opis bibliograficzny
1	Spiegel Maciej , Żbikowska Beata, Sroka Zbigniew, Hładyszowski Jerzy: Quantum chemistry aided examination of antioxidative potential of phenolic acids, W: 4th International Wroclaw Scientific Meetings. Wrocław, 09-10 October 2020, (red.) Julita Kulbacka, Nina Rembiałkowska, Joanna Weżgowiec, Wrocław 2020, Wydawnictwo Naukowe TYGIEL sp. z o.o., s. 220-222, ISBN 978-83-66489-37-0
2	Spiegel Maciej , Andruniów Tadeusz, Sroka Zbigniew: Influence of structural features on the antiradical Activity of flavones and flavonols - a quantum chemical study, W: 4th International Wroclaw Scientific Meetings. Wrocław, 09-10 October 2020, (red.) Julita Kulbacka, Nina Rembiałkowska, Joanna Weżgowiec, Wrocław 2020, Wydawnictwo Naukowe TYGIEL sp. z o.o., s. 222-223, ISBN 978-83-66489-37-0

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Punkty ministerialne : 2270

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12.04.20231.

Hina Lapo duivile

VI. Osiągnięcia naukowo-badawcze

i. Analiza bibliometryczna całkowitego dorobku naukowego

Publikacji w czasopismach naukowych:	19
h-indeks (Web of Science Core Collection):	6
Sumaryczna ilość cytowań (Web of Science Core Collection):	169
Sumaryczny współczynnik wpływu (Impact Factor):	109.191
Sumaryczna punktacja Ministerstwa Edukacji i Nauki:	2270

ii. Konferencje naukowe

- "4th International Wroclaw Scientific Meetings", Wrocław, Polska, 09–10.10.2020:
 - "Quantum Chemistry Aided Examination of Antioxidative Potential of Phenolic Acids", poster. [współautorzy: Beata Żbikowska, Zbigniew Sroka i Jerzy Hładyszowski]
 - "Influence of Structural Features on the Antiradical Activity of Flavones and Flavonols A Quantum Chemical Study", poster. [współautorzy: Tadeusz Andruniów i Zbigniew Sroka]

iii. Projekty krajowe

- Grant PRELUDIUM21 (kierownik projektu), "Zastosowanie sztucznej inteligencji w poszukiwaniu nowych substancji leczniczych pochodzenia naturalnego o mechanizmach działania ukierunkowanych na zmniejszanie stężenia wolnych rodników w organizmie" (na mocy decyzji nr DEC–2022/45/N/NZ7/02264), Narodowe Centrum Nauki, Kraków, 2022
- iv. <u>Wyróżnienia i nagrody</u>
- Laureat "*Studenckiego Programu Stypendialnego*" organizowanego przez Urząd Miejski we Wrocławiu, Edycja 2022 [online: https://www.wroclaw.pl/akademicki-wroclaw/ znamy-nazwiska-doktorantow-z-przyznanymi-stypendiami-w-studenckim-programiestypendialnym]
- Wybranie artykułu **[D]** na okładkę wydania (*Journal of Chemical Information and Modeling*; June 13, 2022; Volume 62, Issue 11; Pages 2631–2900) [online: https://pubs.acs.org/toc/jcisd8/62/11]
- Zaproszenie do przedstawienia tematyki badań w postaci artykułu opublikowanego w Raporcie Rocznym 2022 Poznańskiego Centrum Superkomputerowo–Sieciowego [online: https://www.pcss.pl/files/2022/12/PCSS_Raport_2022.pdf]
- v. <u>Aktywność recenzencka</u>
- Członek loży recenzentów w czasopismach:
 - Frontiers in Bioscience–Landmark (BRI) [online: https://www.imrpress.com/journal/FBL/about/reviewer_board]
 - Symmetry (MDPI)
 [online: https://www.mdpi.com/journal/symmetry/submission reviewers]
 - Wykonano 8 recenzji dla: Theoretical Chemistry Accounts (1), Phytochemistry (1), Free Radicals
- Research (2), Frontiers in Bioscience–Landmark (1), Journal of Molecular Modeling (1), Molecules (1), Journal of Molecular Liquids (1)

vi. <u>Warsztaty, kursy, seminaria</u>

- "What is a Chemical? Part IV: Innovation in Chemical Descriptions", webinar. IUPAC, 17 lutego 2023.
- "Berlin Digital Science for Drug Discovery Meeting", webinar. DigiDrug.NET, 15 lutego 2023.
- "Cambridge Cheminformatics Meeting", webinar. University of Cambridge, 8 lutego 2023.
- "Quantum Computing: What are the Pharma Use Cases?", webinar. Pistoia Alliance, 7 lutego 2023.
- *"5th International Mini–Symposium on Molecular Machine Learning"*, webinar. University of Münster, 19 stycznia 2023.
- "Protein structure prediction What's next after AlphaFold?", webinar. Discngine Labs, 12 stycznia 2023.
- "Computation in Drug Discovery An Insider's View", webinar. CDD, 1 grudnia 2022.
- "*ELLIS Machine Learning for Molecule Discovery Workshop*", webinar. ELLIS Cambridge, 28 listopada 2022.
- "3rd European Symposium on Chemical Bonding (CBOND2022)", webinar. Vrije Universiteit Amsterdam, 20–22 września 2022.
- "DFT calculations (Abinit, Siesta) within an HPC environment", webinar. Interdyscyplinarne Centrum Modelowania Matematycznego i Komputerowego, Uniwersytet Warszawski, 26 maja 2022.
- "Obliczenia metodami chemii kwantowej: programy Gamess i Gaussian przykłady zastosowań w projektowaniu nowych materiałów funkcjonalnych", webinar. Centrum Informatyczne Trójmiejskiej Akademickiej Sieci Komputerowej, 16 maja 2022.
- "*Exploring the Chemical Bonding in Main Group Chemistry*", wykład. Institute of Chemistry for Life and Health Sciences, Chimie ParisTech, Paryż, Francja, kwiecień 2022.
- "*Studying Potential Energies with Machine Learning*", wykład. Institute of Chemistry for Life and Health Sciences, Chimie ParisTech, Paryż, Francja, kwiecień 2022.
- " Od teorii do eksperymentu czyli wpływ reorganizacji wewnątrz- i międzycząsteczkowej na właściwości cząsteczek w mikro- i makroskali", webinar. Wydział Chemiczny, Uniwersytet Wrocławski. 18 lutego 2021.
- "*Applied Pharmaceutical Bioinformatics*", kurs zdalny. Uppsala University, Uppsala, Szwecja, listopad–grudzień 2019.
- "*Pharmaceutical Bioinformatics*", kurs zdalny. Uppsala University, Uppsala, Sweden, wrzesień–listopad 2019.
- "*Visiting Professors Denis Jacquemin*", wykłady i warsztaty. Politechnika Wrocławska, 22–24 października 2019.
- "*Mapping Polypharmacy Chemical Reactions in At–Risk Populations*", zdalny projekt. Hackseq19 – genomics hackathon collective, October 2019. [online: https://github.com/hackseq/hs19–pop.]
- "*Applied Pharmaceutical Structural Bioinformatics*", kurs zdalny. Uppsala University, Uppsala, Sweden, wrzesień-październik 2019.
- "*Electronic Structure Theory Workshop: Advanced Quantum Chemistry in Python*", warsztaty. Politechnika Wrocławska, 23-25 września 2019.

vii. <u>Staże zagraniczne</u>

- "Introduction to and Application of Novel Approaches in Computational Chemistry" (2023–02 — 2023–04) Weizmann Institute of Science, Rehovot, Izrael; mentor: profesor Jan M.L. Martin;
- "Modeling Optical Properties of Metal Complexes for Photodynamic Therapy Using Quantum Chemical Approaches" (2022–03 2022–06) Institute of Chemistry for Life and Health Sciences (I–CLeHS), Chimie ParisTech, Paris Cedex 05, Paryż, Francja; mentor: profesor Carlo Adamo;
- *"Theoretical Study About the Reaction Mechanisms of Antioxidants"* (2021–07 2021–09) Department of Chemistry and Chemical Technology, University of Calabria, Arcavacata di Rende, Cosenza, Włochy; mentor: **profesor Nino Russo**.

viii. <u>Pozostałe aktywności</u>

 Opiekun naukowy pracy magisterskiej mgr farm. Katarzyny Cel, wykonywanej w Katedrze i Zakładzie Farmakognozji i Leku Roślinnego, zatytułowanej "Wpływ czynników środowiskowych na dwojaką aktywność polifenoli – antyoksydanty i potencjalne prooksydanty – badania in silico".