

Title: *Study of an inhibitory effect of plant polyphenolic compounds against digestive enzymes using bench-working experimental evidence predicted by molecular docking and dynamics*

Author Names: Kaushal Vyas , Supraja Prabaker , Dhamodharan Prabhu , Meenakumari Sakthivelu , Sundararaj Rajamanikandan , Palaniyandi Velusamy , Chia-Hung Su , Subash C.B. Gopinath , Raman Pachaiappan

Abstract: The substantial nutritional content and diversified [biological activity](#) of plant-based [nutraceuticals](#) are due to polyphenolic chemicals. These chemicals are important and well-studied plant [secondary metabolites](#). Their protein interactions are extensively studied. This relationship is crucial for the logical development of functional food and for enhancing the availability and usefulness of polyphenols. This study highlights the influence of protein types and polyphenols on the interaction, where the chemical bindings predominantly consist of hydrophobic interactions and [hydrogen bonds](#). The interaction between [polyphenolic compounds](#) (PCs) and [digestive enzymes](#) concerning their inhibitory activity has not been fully studied. Therefore, we have examined the interaction of four digestive [enzymes](#) (α -amylase, [pepsin](#), trypsin, and α -chymotrypsin) with four PCs (curcumin, [diosmin](#), [morin](#), and 2',3',4'-trihydroxychalcone) through *in silico* and *in vitro* approaches. *In vitro* plate assays, [enzyme kinetics](#), spectroscopic assays, [molecular docking](#), and simulations were performed. We observed all these PCs have significant docking scores and preferable interaction with the active site of the [digestive enzymes](#), resulting in the reduction of [enzyme activity](#). The enzyme-substrate binding mechanism was determined using the [Lineweaver Burk plot](#), indicating that the inhibition occurred competitively. Among four PCs [diosmin](#) and [morin](#) has the highest interaction energy over digestive [enzymes](#) with IC_{50} value of 1.13 ± 0.0047 and 1.086 ± 0.0131 μ M. Kinetic studies show that selected PCs inhibited [pepsin](#), trypsin, and [chymotrypsin](#) competitively and inhibited [amylase](#) in a non-competitive manner, especially by 2',3',4'-trihydroxychalcone. This study offers insights into the mechanisms by which the selected PCs inhibit the enzymes and has the potential to enhance the application of [curcumin](#), [diosmin](#), morin, and 2',3',4'-trihydroxychalcone as natural inhibitors of digestive enzymes.

Keywords: Polyphenolic compounds (PCs) Plant-based nutraceuticals Protein–polyphenol interactions Digestive enzymes Enzyme inhibition In silico analysis Molecular docking

<https://www.sciencedirect.com/science/article/abs/pii/S0141813024000254>

<https://doi.org/10.1016/j.ijbiomac.2024.129222>