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BUNGAP

Screening for nutraceutical - drug interactions toward the non-covalent interactions of their solid binary systems (Case study on Piperine)

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Introduction

The revealed immunomodulating, antioxidant, chemopreventive and anticancer activity

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of the piperine, mayor alkaloid in fruits of the black pepper (*Piper nigrum* Linn.) and the long pepper (*Piper Longum* Linn.) which for many centuries are broadly used both as spice and as remedy in culinary and traditional medicine, respectively spurs our interest on the research of enhancing its low bioavailability. Referring to the ternary amide structure of piperine which is formed between piperidine in chair conformation and piperic acid (5-(3,4-methylenedioxyphenyl)-2,4-pentadienoic acid), its side chain with conjugated double bonds impacts to the appearance of piperine in four possible geometric isomers [1,2]. We envisage that structural flexibility of piperine is favourable for forming amide-amide type of non-covalent H-bond interactions with drugs (e.g. secondary amide moiety in molecule of perindopril – ACE inhibitor, tertiary amide in prazosin- α -adrenoceptor antagonists, primary amide in carbamazepine – antiepileptic drug and etc.) and for forming amide-catvoxylate and amide-hydroxyl H-bonding interaction with drugs and natural compounds (e.g. ascorbic acid with enediole structure and curcumine with β -diketo and hydroxyl benzoic moieties in its structure).









Gorgani L., et al., (2017) Compr Rev Food Sci Food Saf; 16: 124-40