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COMPUTATIONAL MODELING OF THE TAUTOMERIC INTERCONVERSIONS OF THE QUERCETIN MOLECULE

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Quercetin molecule (3, 3', 4', 5, 7-pentahydroxyflvanone, $C_{15}H_{10}O_7$) is an important flavonoid compound, which is found in many foods and plants, and is known to act as a natural drug molecule with a wide range of treatment properties, like an anti-oxidant, anti-toxic etc. Notably, that in the composition of the biologically-active compounds in natural conditions quercetin shows quite effective treatment properties, while in the extracted state this molecule shows more effective mechanism of action, however already with toxicological manifestations.

So, at the analysis of the pharmacological properties of the quercetin molecule it should be taking into account its possibility to acquire different tautomeric forms, ranging greatly by their energies.

In this study we revealed and investigated the possible pathways of the tautomeric transformations of the quercetin molecule, which potentially influence on its pharmacological and clinical effectivity (Scheme 1). Also, it was revealed that single proton transfer can induce cascade tautomeric reactions in the quercetin molecule.

These transformations of the quercetin molecule are accompanied by the rearrangement (breakage and formation) of the intramolecular H-bonds, changes of their geometry and dipole moment (value and direction).

Obtained results can be useful for the formulation of the general rules of the formation and importance of tautomeric states, which can be applied to different heterocyclic molecules with potential biological or pharmaceutical activity.

