Preliminary study on screening the intermolecular interactions of organic cation drugs from BSC Class III (case study Metformin)

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Among the orally administered drugs, about 40% share the properties of organic cations (protonated bases) or neutral bases at physiological pH, which indicates one important point for studying their transport mechanisms [1]. Antineoplastic platinum compounds [2,3], the histamine H₂ receptor antagonist cimetidine [4], the antiviral drugs (acyclovir, gancyclovir, lamivudine and zalcitabine) [5-7], the antidiabetic drug metformin [8,9], and the antiarrhythmic drug quinidine [10], are the identified to be transported by the organic cation transporters OCT1, OCT2 and OCT3 (membrane transporters) [2]. The case study of drug model (DM) metformin (MET), that according to Biopharmaceutical Classification System (BCS) belongs to the class III drugs (high solubility, low permeability) [11], emphasizes the importance of non-covalent interactions of this dication drug with range of ligands selected from the GRAS (Generally Recognized as Safe by FDA for food additives list) [12].

MET (*N*,*N*-dimethylbiguanide) is the only approved hypoglycemic drug of the biguanide class used in oral therapy of type 2 diabetes, marketed as hydrochloride, embonat (pamoate) and *p*-chlorophenoxy acetate salt [13]. Because of the biguanide π -conjugated system, MET in solution can exist in three resonance-stabilized forms, *i.e.* as neutral molecule (MET), monoprotonated (METH⁺) or diprotonated (METH²⁺) cation, with dissociation constants in water typical of biguanides:

Metformin = L; [HL]/[L][H] $pK_{a1}(N-H^+) \sim 12.40$; [H₂L]/[HL][H] $pK_{a2}(N-H^+) = 2.96$ (NIST database)

The high basicity of pK_{a1} and the difference between the pK_a values qualify MET as organosuperbase and determine the stability of its monoprotonated form METH⁺ within a wide range of pH [14]. Patterns of charge assisted H-bonding between MET and ligands are expected to be useful in predicting its absorption and transportation. Comparable study on predicting ligands for OTC1 by docking, applying High-Throughput Screening (HTS) and *in silico* modeling [15], offer an opportunity for growing the molecular crystals of MET with drugs, that are uses in combo- therapy with MET, as well with excipients and food ingredients, thus by generating the crystallographic, thermodynamic and spectroscopy data MET-ligand interactions, contributes the development of delivery systems with enhanced permeability of BSC class III drug models.

References

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