Kerr-(A)Ds, Myers–Perr black-hole Galilean ον gravitational transformations for the anti-COVID-19 RoccuffirnaTM evolutionary drug design.

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March 30, 2022

Abstract

It is thought that all of the rich content in the present-day Universe based on an array of recent observations developed through gravitational amplification of primeval density fluctuations generated in the very early phase of cosmic evolution. In this paper, we strongly combine machine learning characteristics to achieve very high accuracy levels for the in-silico generation of the RoccuffirnaTM small molecule, a ligand targeted the SARS-COV-2 virus main protease (M pro) using Quantum Kerr-(A)dS and Myers–Perry black microBlackHole-Inspired Gravitational for both Euclidean and Lorentzian signatures in Practice. We provide also an extensive toolbox of methods for performing quantum schrodinger inspired docking algorithms, teleportation and other information-theoretic tasks in MathCast programming language, and compared these algorithms by means of mean percentile free energy ranking, in a new recall-based evaluation metric for the in-silico design of the Novel Series of the RoccuffirnTMQMMMCoRoNNARRFr anti-(nCoV-19) ligands. In this paper we in-silico designed new drug leads that target the COVID-19 virus main protease (M pro). M pro, a key CoV enzyme, which plays a pivotal role in mediating viral replication and transcription, and discuss various general results including Galilean transformation to a rigid QMMM heuristic horizon topology, and near-horizon fragmentation symmetry ranging from supergravity theories to enhance the Roccuffirna's gravity to trap the SARS-COV-2 viruses in practice.

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