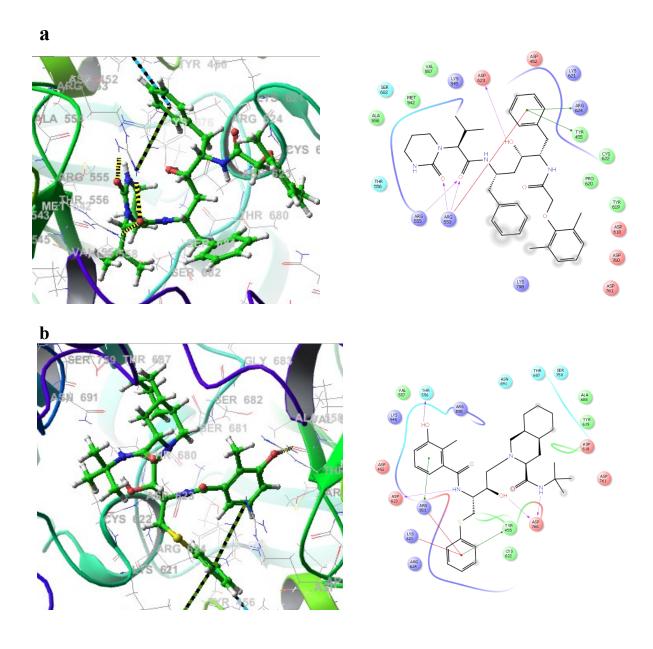
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Potential role of North African plants and their constituents in the mitigation of SARS-

## CoV-2: identifying related therapeutic targets using network pharmacology and molecular docking analyses

## Eman Shawky a, Ahmed A. Nada, Reham S. Ibrahim



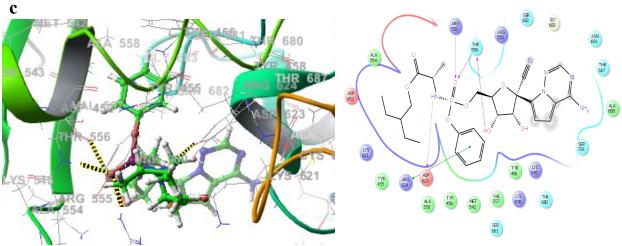
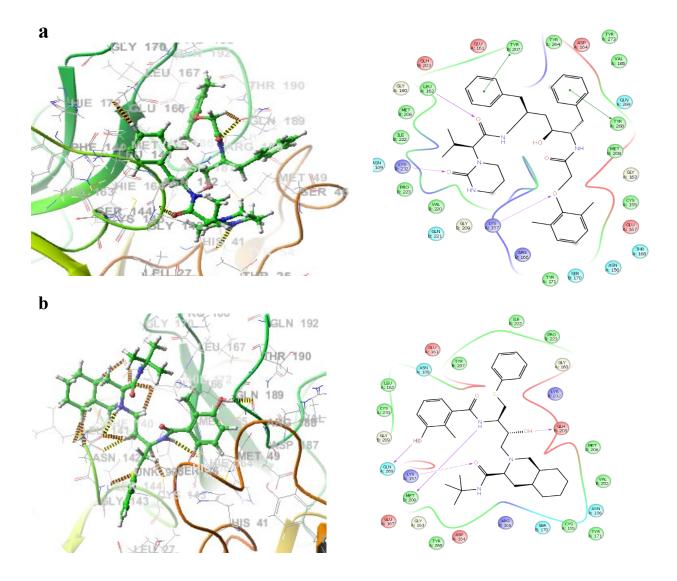


Fig.S1. 3D (to the left) and 2D (to the right) interaction diagrams of a) lopinavir, b) nelfinavir and c) remdesivir with the crystal structure of RdRp (PDB ID: 6M71).



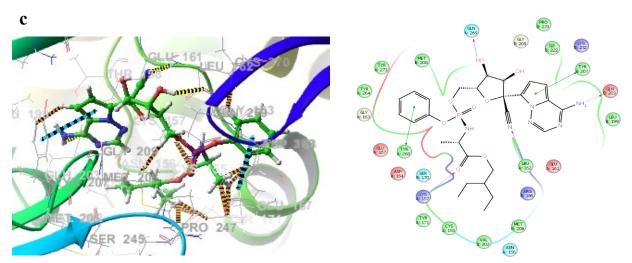
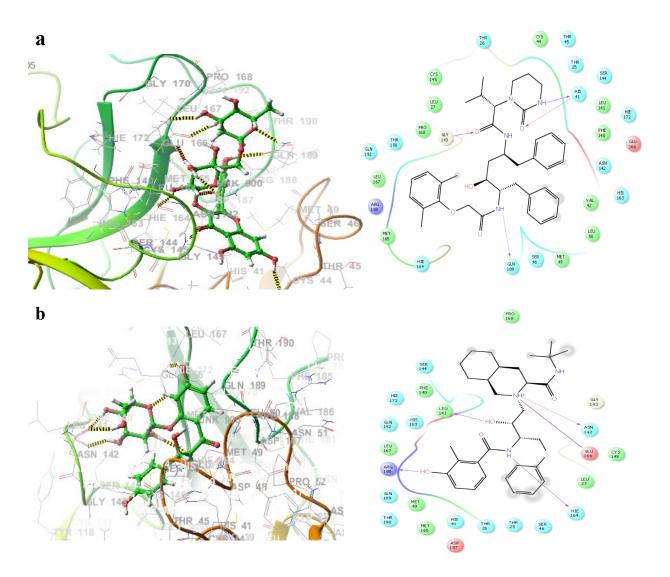


Fig.S2. 3D (to the left) and 2D (to the right) interaction diagrams of a) lopinavir, b) nelfinavir and c) remdesivir with the crystal structure of PLpro (PDB ID: 6w9c).



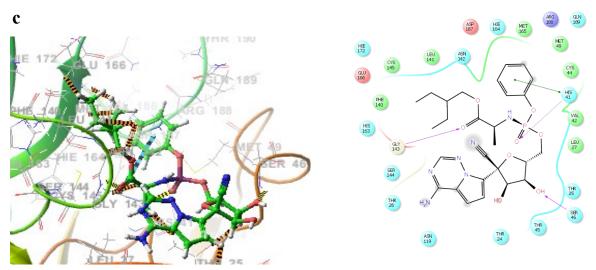


Fig.S3. 3D (to the left) and 2D (to the right) interaction diagrams of a) lopinavir, b) nelfinavir and c) remdesivir with the crystal structure of 3CLpro (PDB ID: 5r7y).