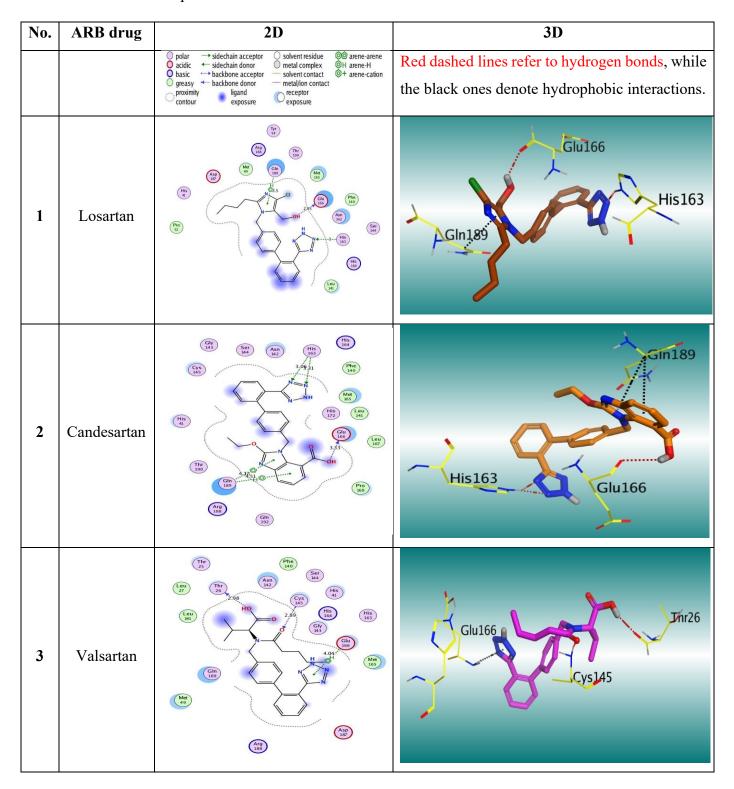
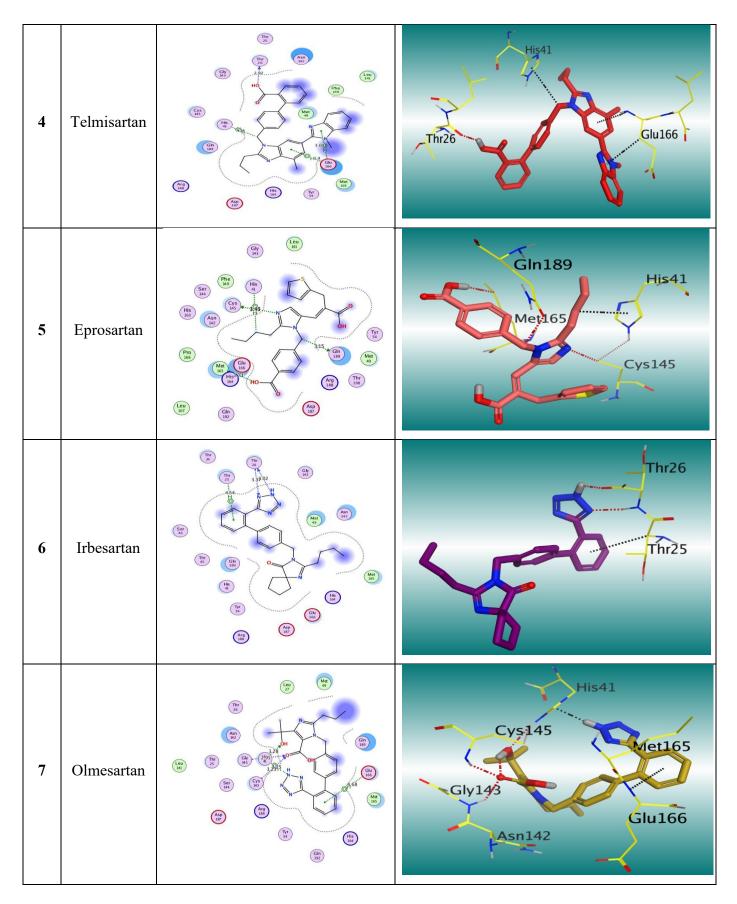
Supplementary data

Fig. 1: 2D and 3D docking representations of ARBs and N3 inhibitor against the binding site of the COVID-19 main protease.





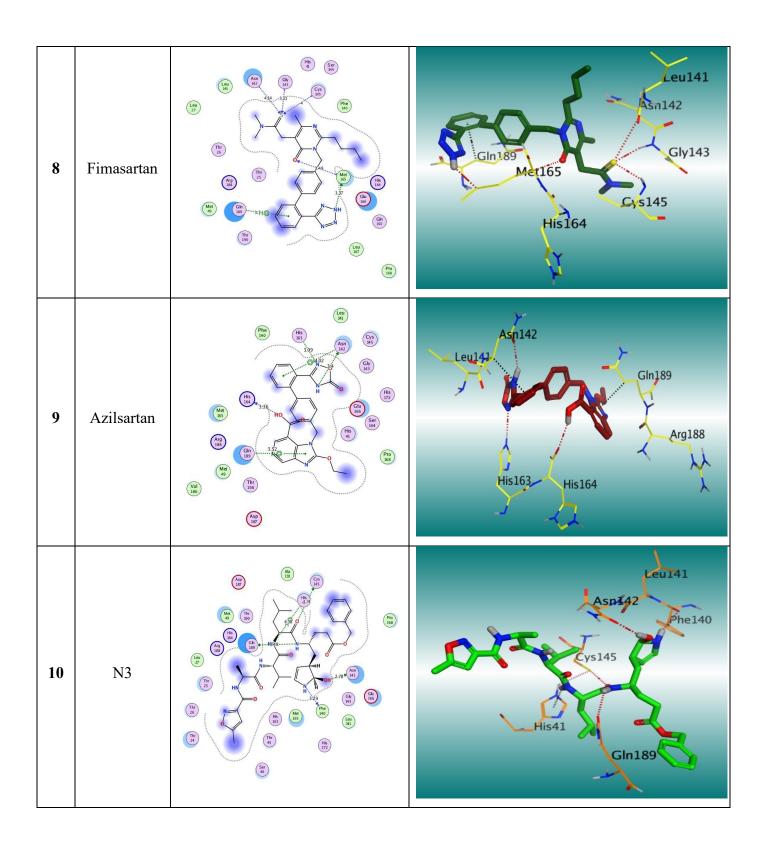
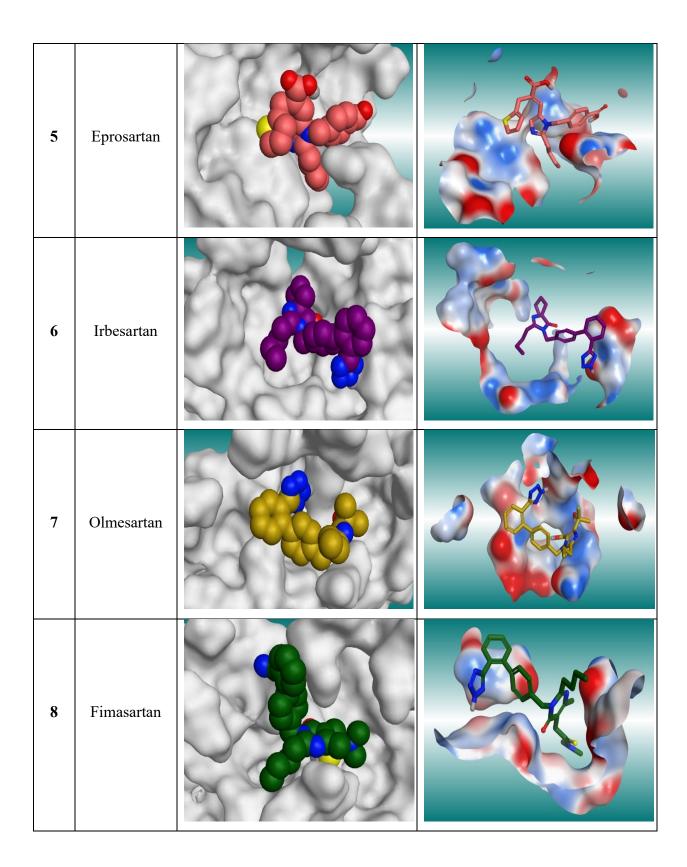


Fig. 2: A) Surface of M^{pro} showing the positioning and fitting of tested compounds, B) surface and maps of the tested compounds compared to N3 inhibitor

No.	ARB drug	A	В
1	Losartan		
2	Candesartan		
3	Valsartan		
4	Telmisartan		



9	Azilsartan	
10	N3	