

Drug similarity and structure-based screening of medicinal compounds to target Macrodmain-I from SARS-CoV-2 to rescue the host immune system: A molecular dynamics study

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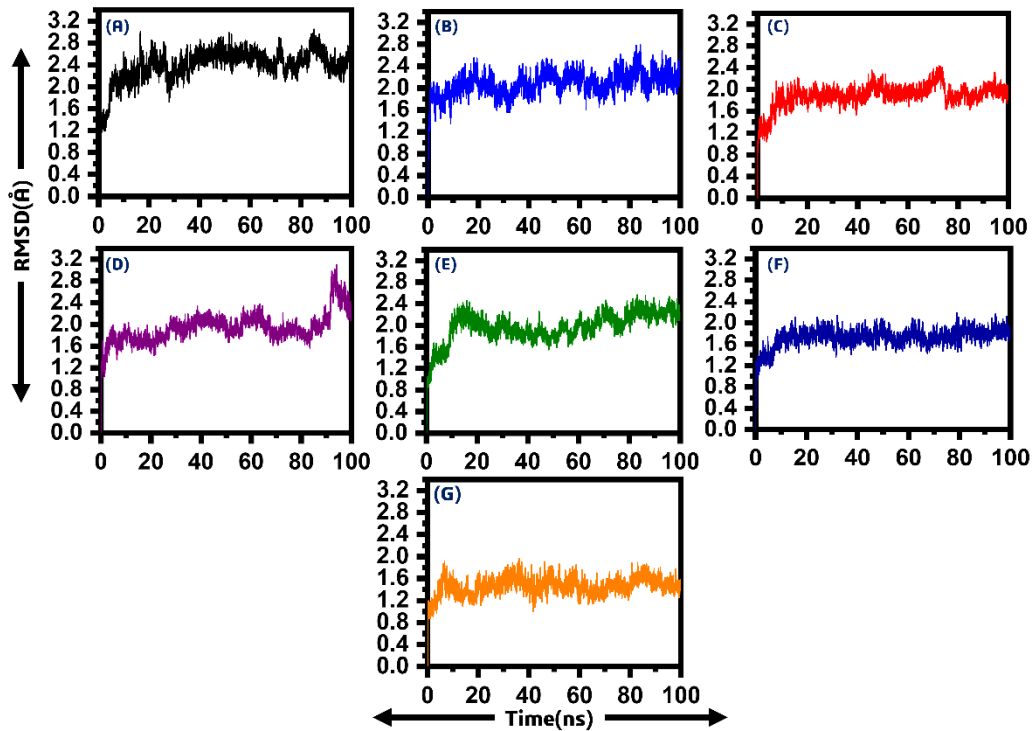
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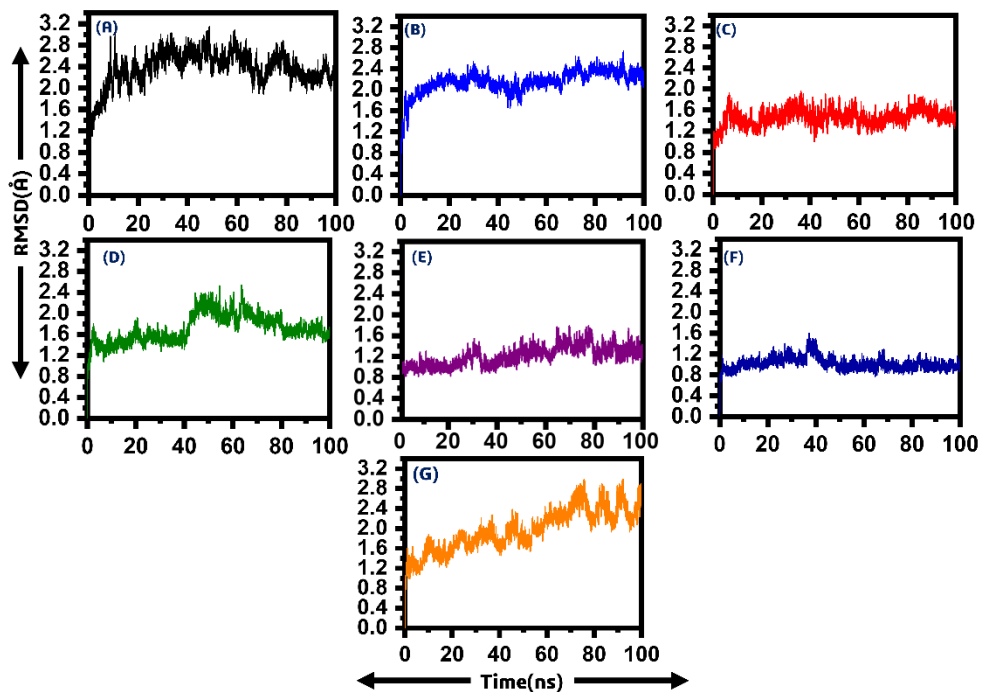
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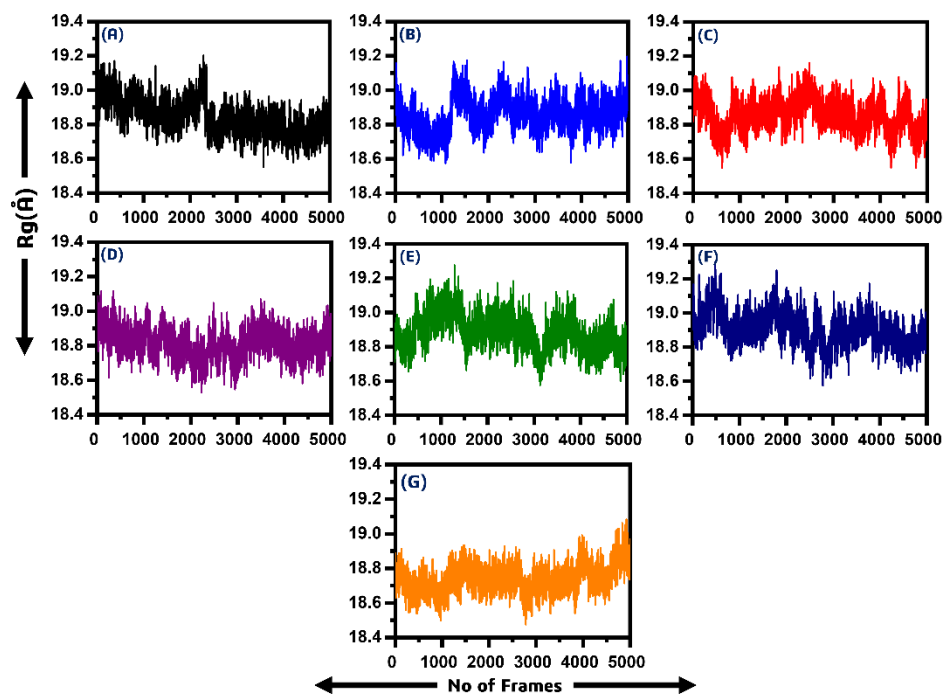
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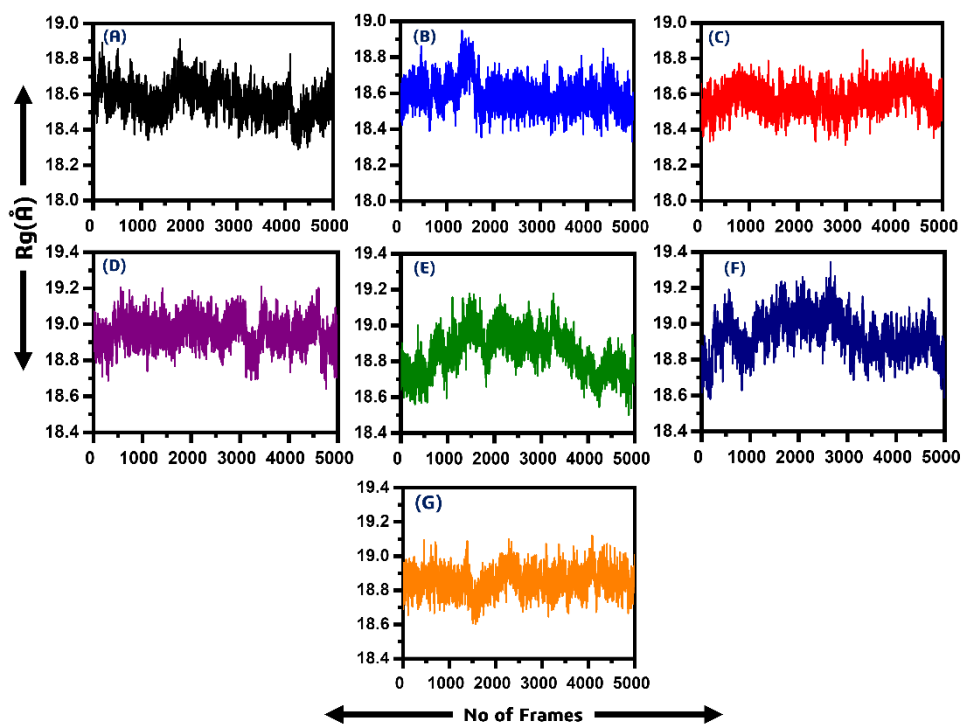
Supplementary Figure S1: RMSDs of all the complexes (replicate 2).



Supplementary Figure S2: RMSDs of all the complexes (replicate 3).



Supplementary Figure S3: R_g of all the complexes (replicate 2).



Supplementary Figure S4: R_g of all the complexes (replicate 3).

Supplementary Table S1: MMGBSA of all the complexes (replicate 2).

Complexes	MM-GBSA (kcal/mol)				
	Δ_{vdW}	Δ_{elec}	Δ_{ps}	Δ_{SASA}	ΔG_{Total}
<i>P1,P5-Di(guanosine-5') pentaphosphate ammonium salt</i> (ChemSpider ID: 4484327)	-85.142	-11.247	34.462	-5.111	-69.326
<i>(R)-RETRO-THIORPHAN</i> (ChemSpider ID: 394363)	-78.716	-22.569	32.669	-4.100	-67.877
<i>P(1),P(6)-bis(5'-adenosyl)hexaphosphate</i> (ChemSpider ID: 110267)	-66.324	-23.426	30.185	-5.212	-68.587
<i>(Rib5)ppA</i> (ChemSpider ID: 30975)	-60.011	-14.178	20.211	-2.287	-68.010
<i>NA7</i> (ChemSpider ID: 393480)	-63.852	-11.981	23.251	-3.252	-69.314
<i>dephospho-CoA</i> (ChemSpider ID: 24785028)	-67.259	-17.548	18.874	-4.276	-63.258
<i>ADPr</i>	-64.147	-17.326	19.253	-5.158	-65.923

Supplementary Table S2: MMGBSA of all the complexes (replicate 3).

Complexes	MM-GBSA (kcal/mol)				
	Δ_{vdW}	Δ_{elec}	Δ_{ps}	Δ_{SASA}	ΔG_{Total}
<i>P1,P5-Di(guanosine-5') pentaphosphate ammonium salt</i> (ChemSpider ID: 4484327)	-84.385	-11.475	29.221	-4.278	-70.531
<i>(R)-RETRO-THIORPHAN</i> (ChemSpider ID: 394363)	-83.224	-17.581	30.272	-4.658	-68.012
<i>P(1),P(6)-bis(5'-adenosyl)hexaphosphate</i> (ChemSpider ID: 110267)	-62.158	-21.759	32.135	-4.522	-65.948
<i>(Rib5)ppA</i> (ChemSpider ID: 30975)	-63.240	-13.586	23.586	-2.485	-68.003
<i>NA7</i> (ChemSpider ID: 393480)	-63.369	-14.485	23.254	-3.758	-70.699
<i>dephospho-CoA</i> (ChemSpider ID: 24785028)	-69.582	-16.125	21.485	-3.478	-66.741
<i>ADPr</i>	-61.186	-13.895	22.241	-5.352	-64.964