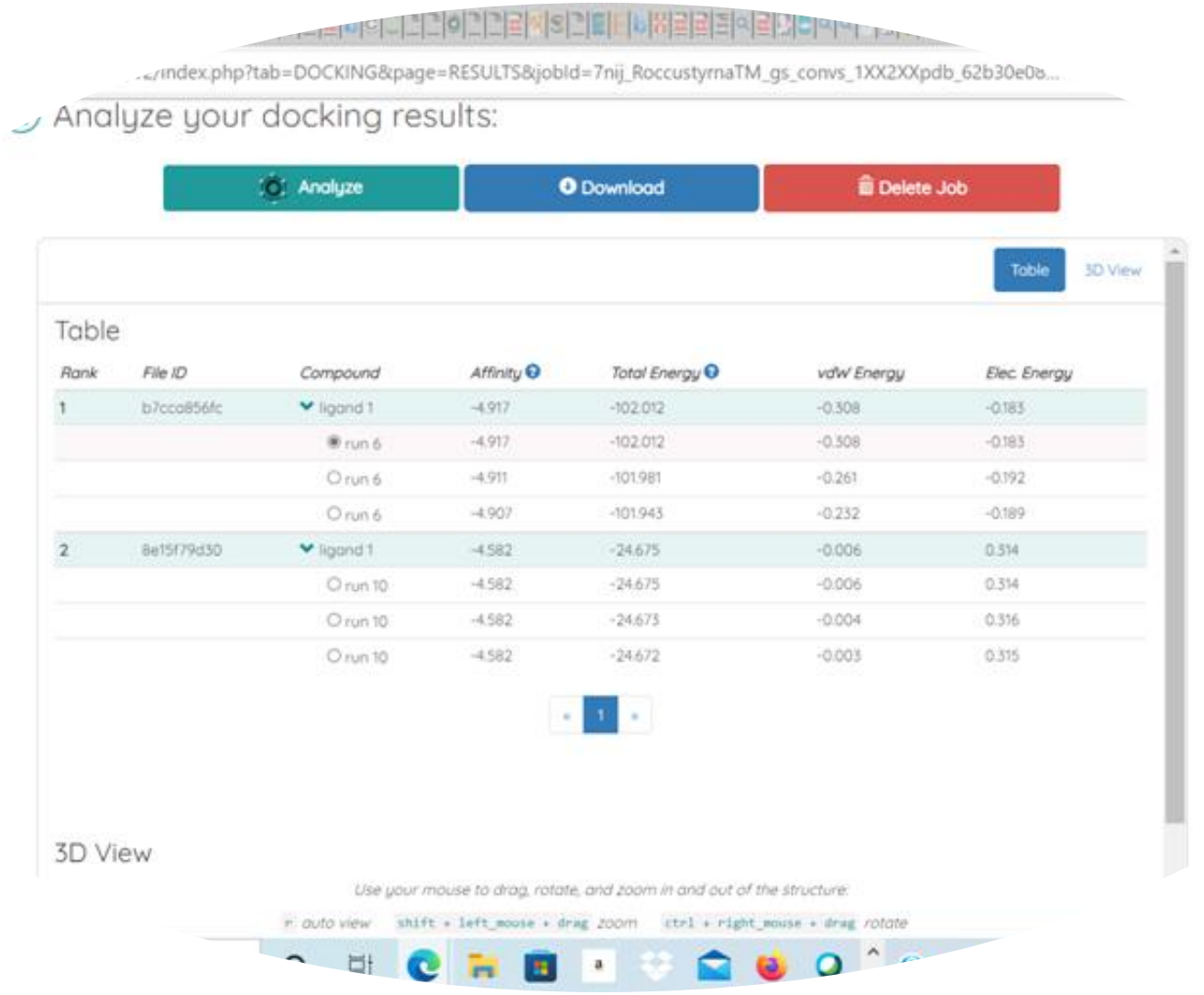
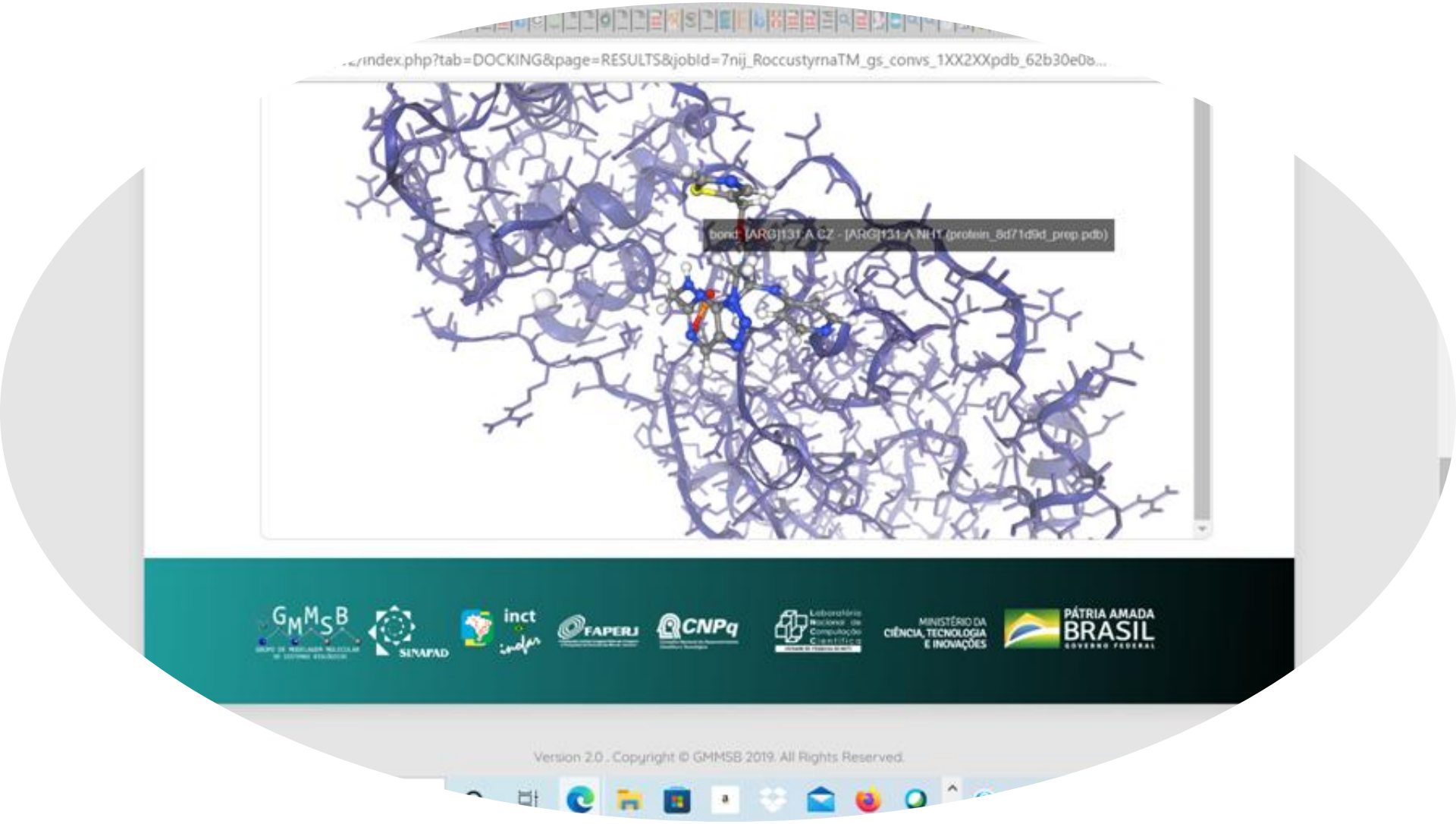

index.php?tab=DOCKING&page=RESULTS&jobId=7nij_RoccustyrnaTM_gs_convs_1XX2XXpdb_62b30e0a...
Analyze your docking results:
Analyze
Download
Delete Job
Table
3D View
Table
Rank File ID Compound Affinity Total Energy vdW Energy Elec. Energy
1 b7cca856fc ligand 1 -4.917 -102.012 -0.308 -0.183
run 6 -4.917 -102.012 -0.308 -0.183
run 6 -4.911 -101.981 -0.261 -0.192
run 6 -4.907 -101.943 -0.232 -0.189
2 8e15f79d30 ligand 1 -4.582 -24.675 -0.006 0.314
run 10 -4.582 -24.675 -0.006 0.314
run 10 -4.582 -24.673 -0.004 0.316
run 10 -4.582 -24.672 -0.003 0.315
1
3D View
Use your mouse to drag, rotate, and zoom in and out of the structure.
r: auto view shift + left_mouse + drag zoom ctrl + right_mouse + drag rotate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bond: [ARG]131:A:CZ - [ARG]131:A:NH1 (protein_8d71d9d_prep.pdb)
GMMSB
SINAPAD
inct
FAPERJ
CNPq
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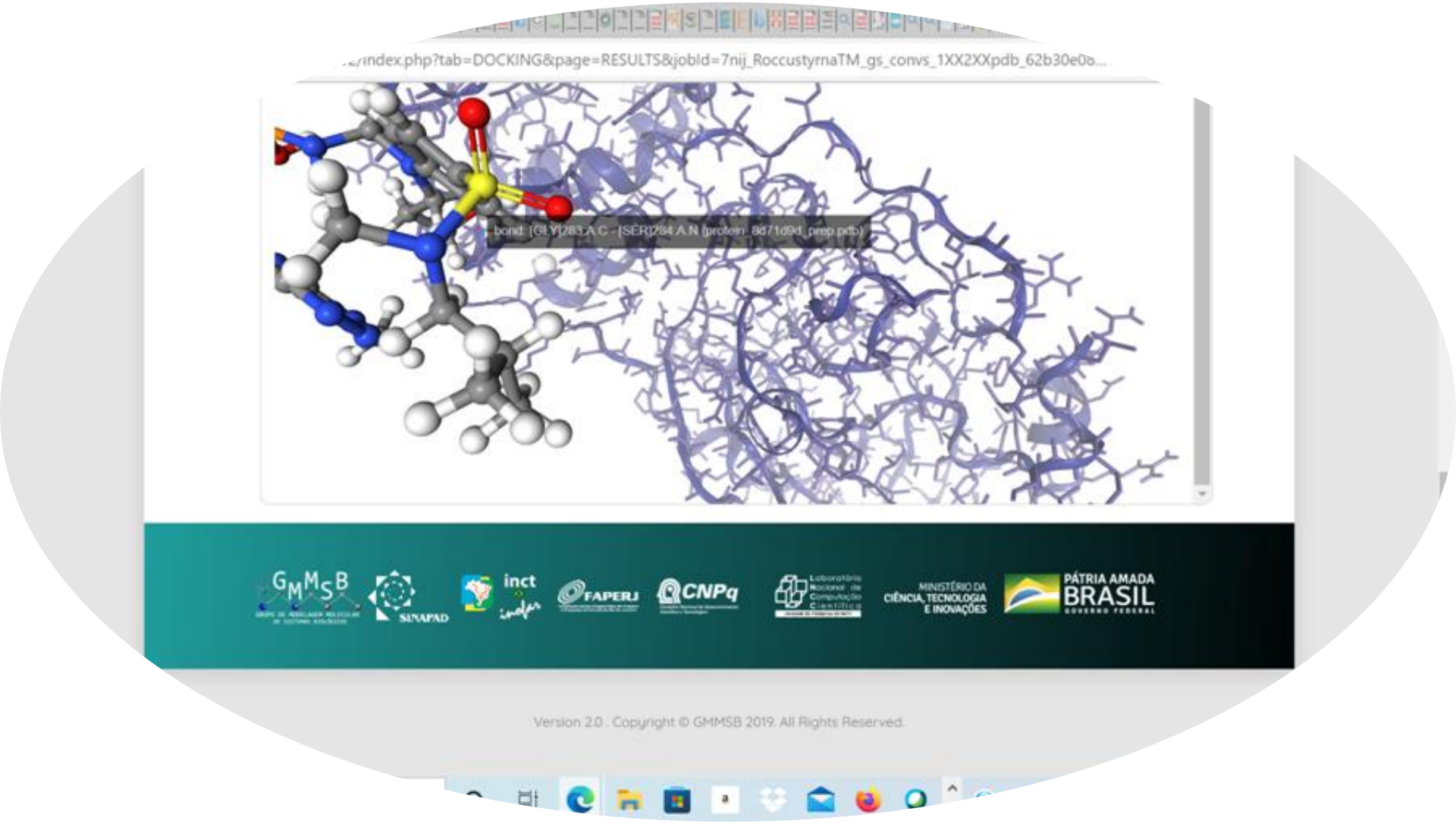
index.php?tab=DOCKING&page=RESULTS&jobId=7nij_RoccustyrnaTM_gs_convs_1XX2XXpdb_62b30e0a...
bond: [GLY:283:A:C - [SER]284:A:N (protein: 8d71d9d_prep.pdb)
GMMSB
SINAPAD
inct
FAPERJ
CNPq
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