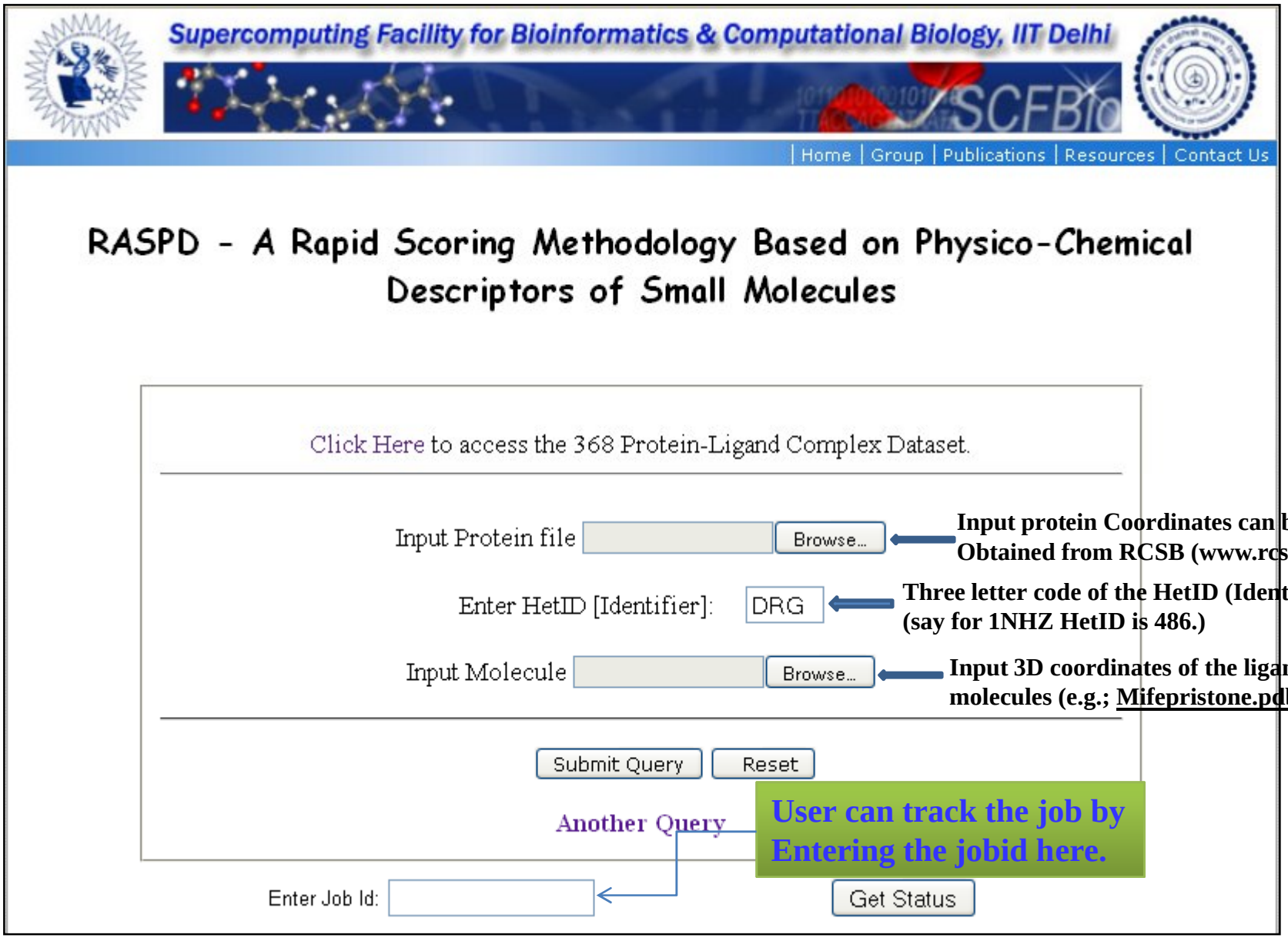
Method D: http://www.scfbio-iitd.res.in/software/drugdesign/raspdmolecule.jsp



Input protein Coordinates can be Obtained from RCSB (www.rcsb.org)

Three letter code of the HetID (Identifier) (say for 1NHZ HetID is 486.)

Input 3D coordinates of the ligand molecules (e.g.; Mifepristone.pdb).

User can track the job by Entering the jobid here.

Result of the Method D is obtained upon clicking View Result option
The input Molecule file can be in any of the following formats (*.pdb, *.mol, *.xyz,*.sdf).

Thank You