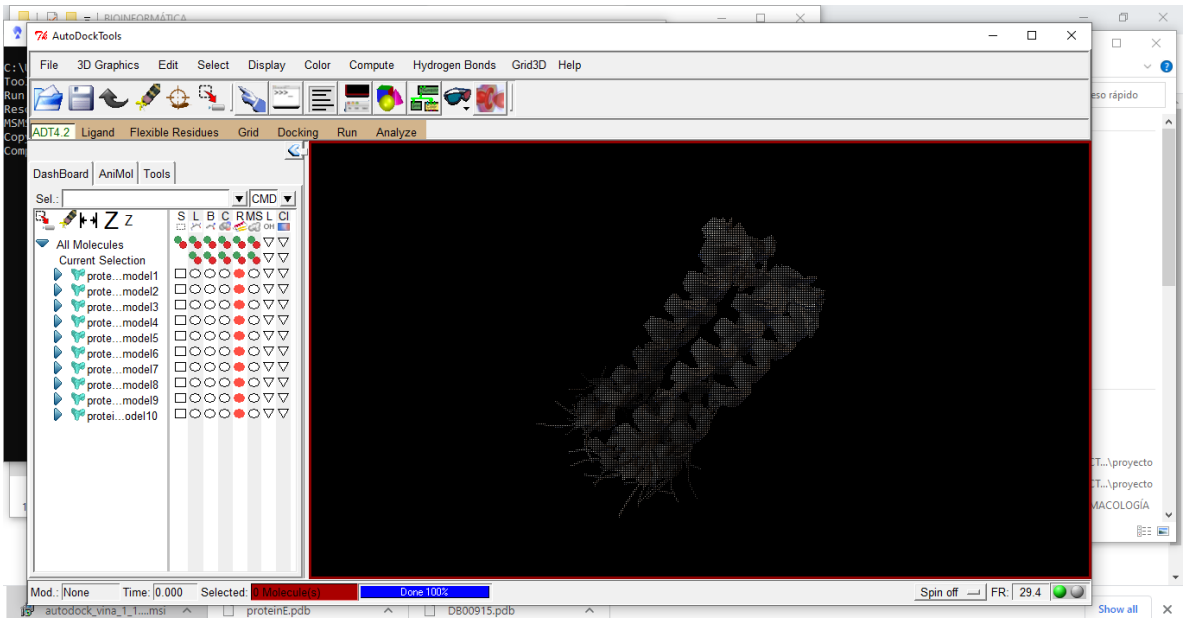
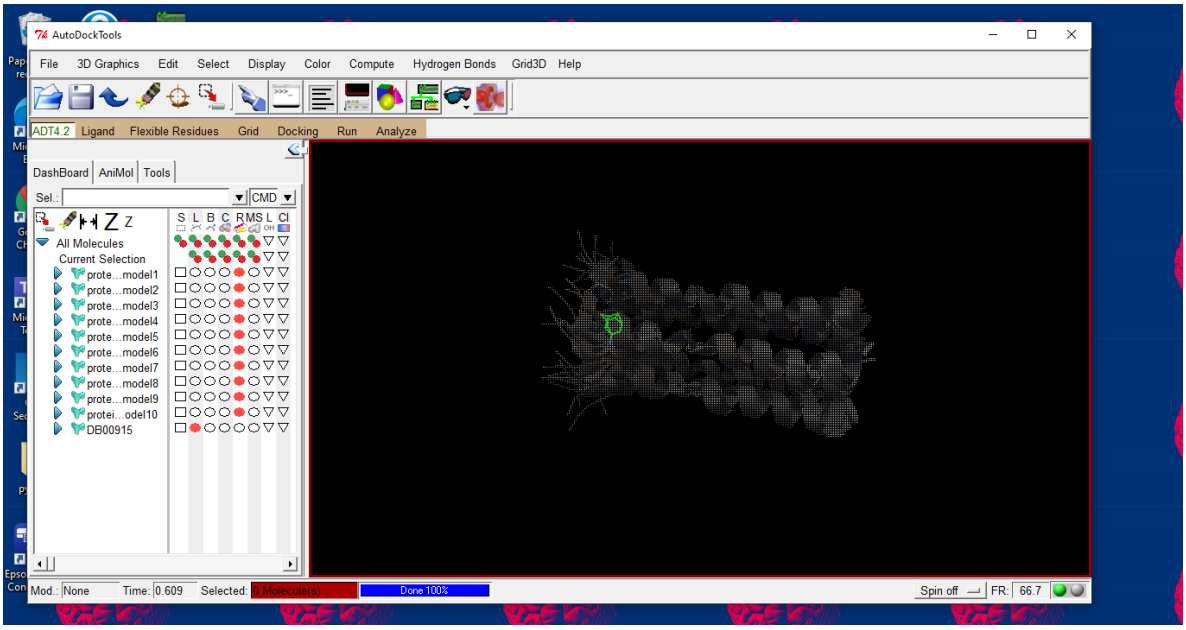
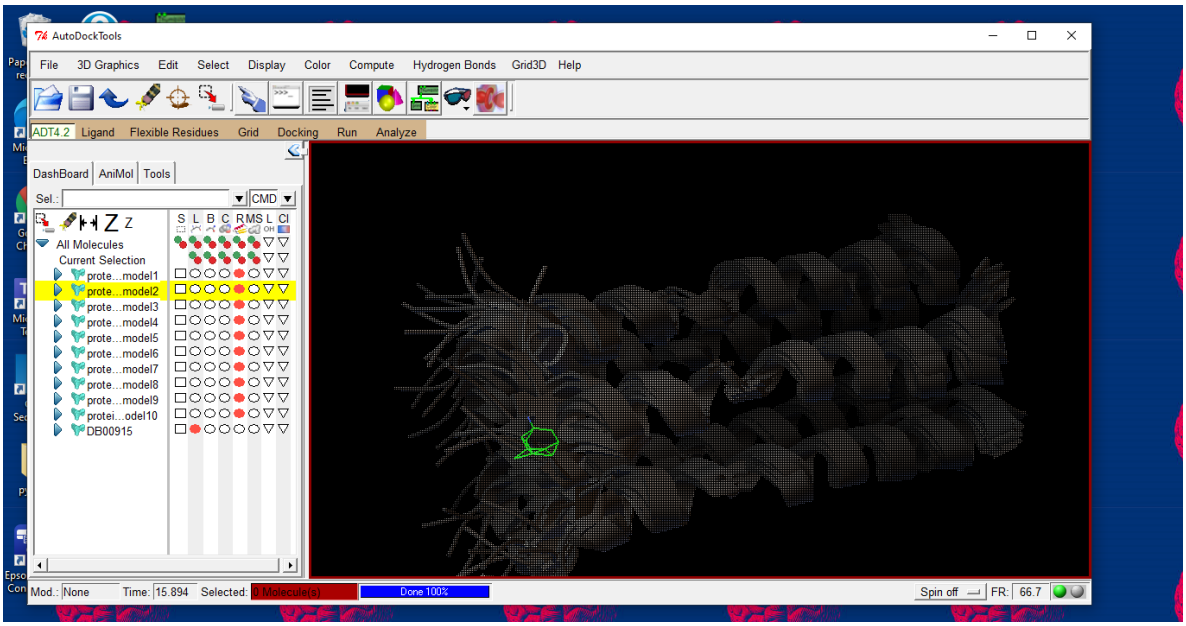
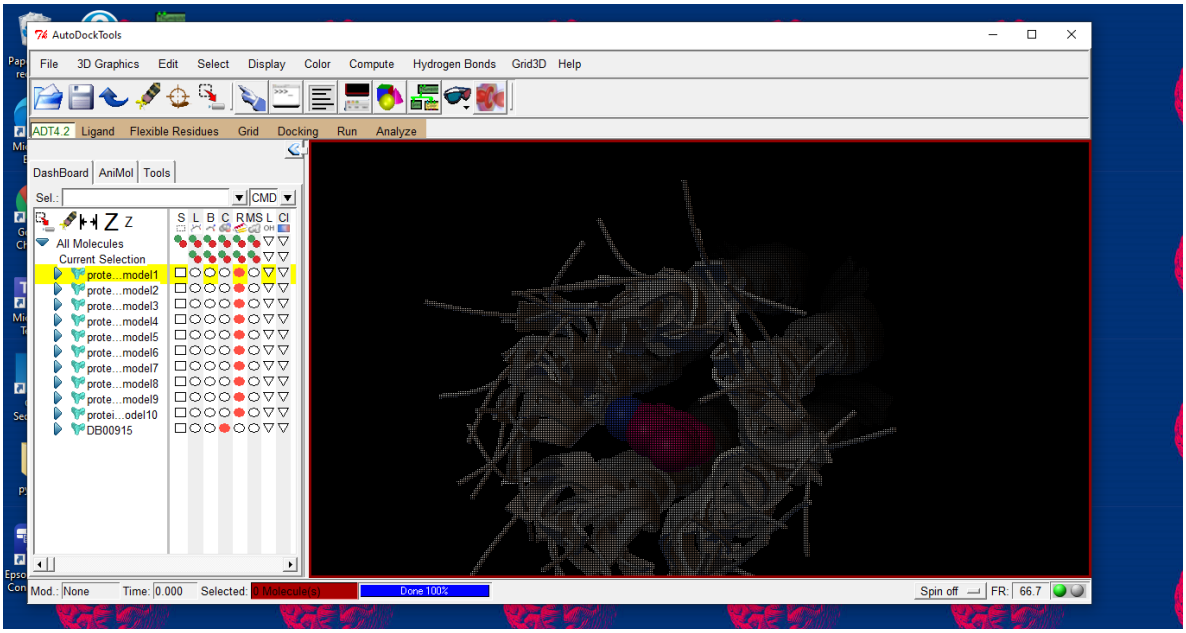
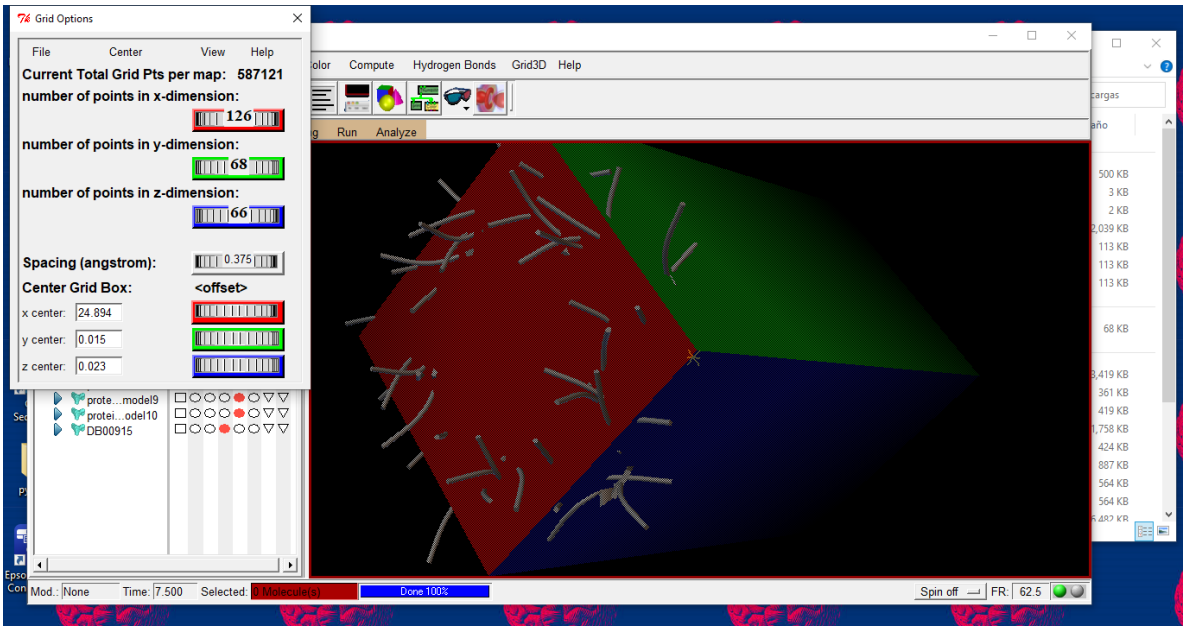
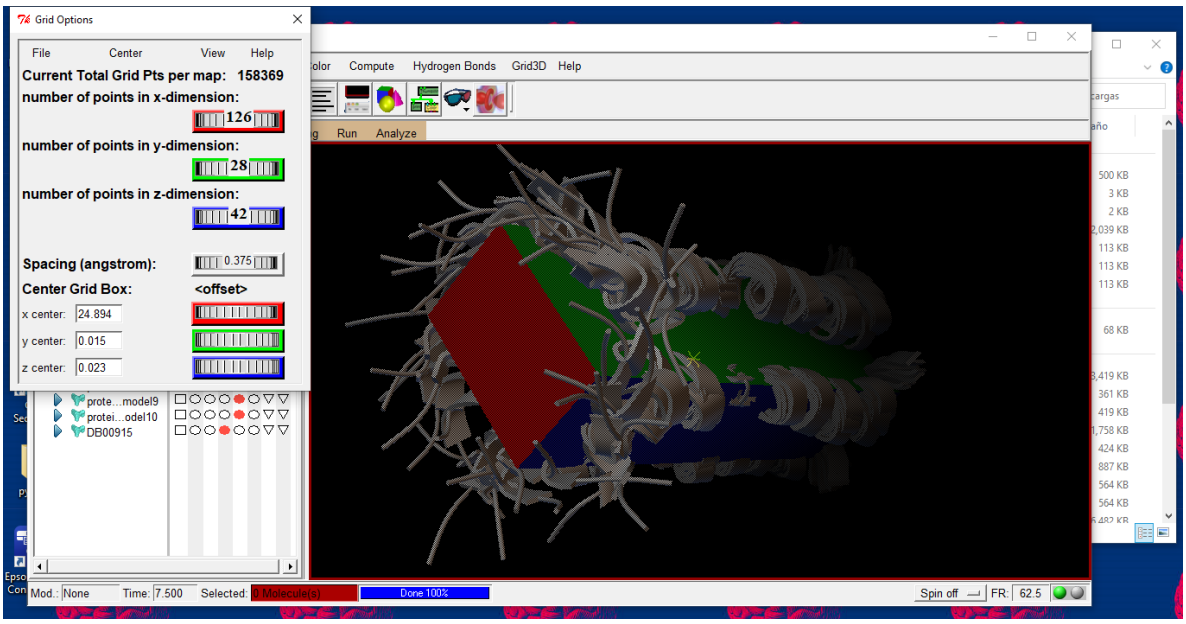

AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
All Molecules
Current Selection
prote...model1
prote...model2
prote...model3
prote...model4
prote...model5
prote...model6
prote...model7
prote...model8
prote...model9
protei...odel10
Mod.: None Time: 0.000 Selected: 0 Molecule(s) Done 100%
Spin off FR: 29.4
autodock_vina_1_1....msi
proteinE.pdb
DB00915.pdb
Show all

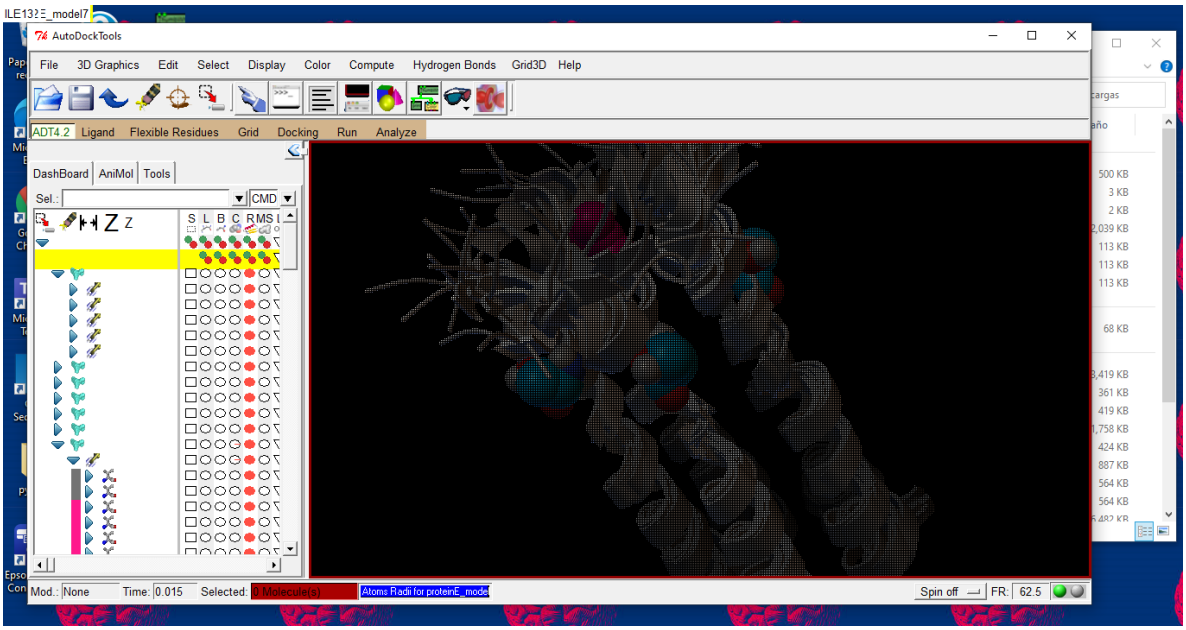

AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
All Molecules
Current Selection
prote...model1
prote...model2
prote...model3
prote...model4
prote...model5
prote...model6
prote...model7
prote...model8
prote...model9
protei...odel10
DB00915
Mod.: None Time: 0.609 Selected: 0 Molecule(s) Done 100%
Spin off FR: 66.7

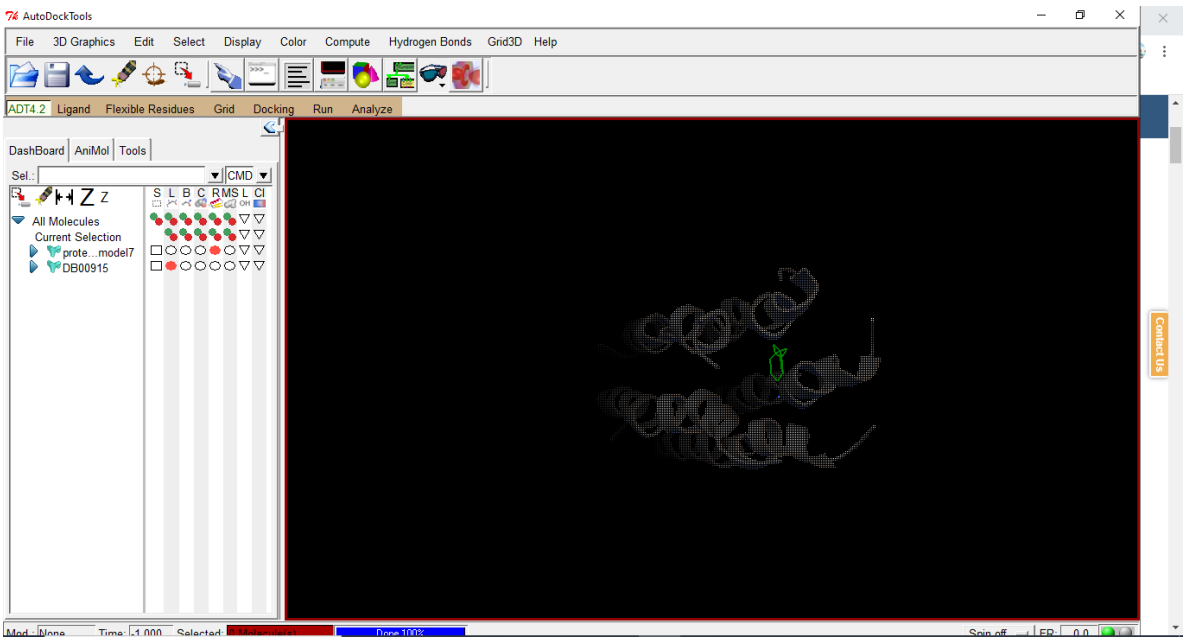

AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
All Molecules
Current Selection
prote...model1
prote...model2
prote...model3
prote...model4
prote...model5
prote...model6
prote...model7
prote...model8
prote...model9
protei...odel10
DB00915
Mod.: None Time: 15.894 Selected: 0 Molecule(s) Done 100%
Spin off FR: 66.7

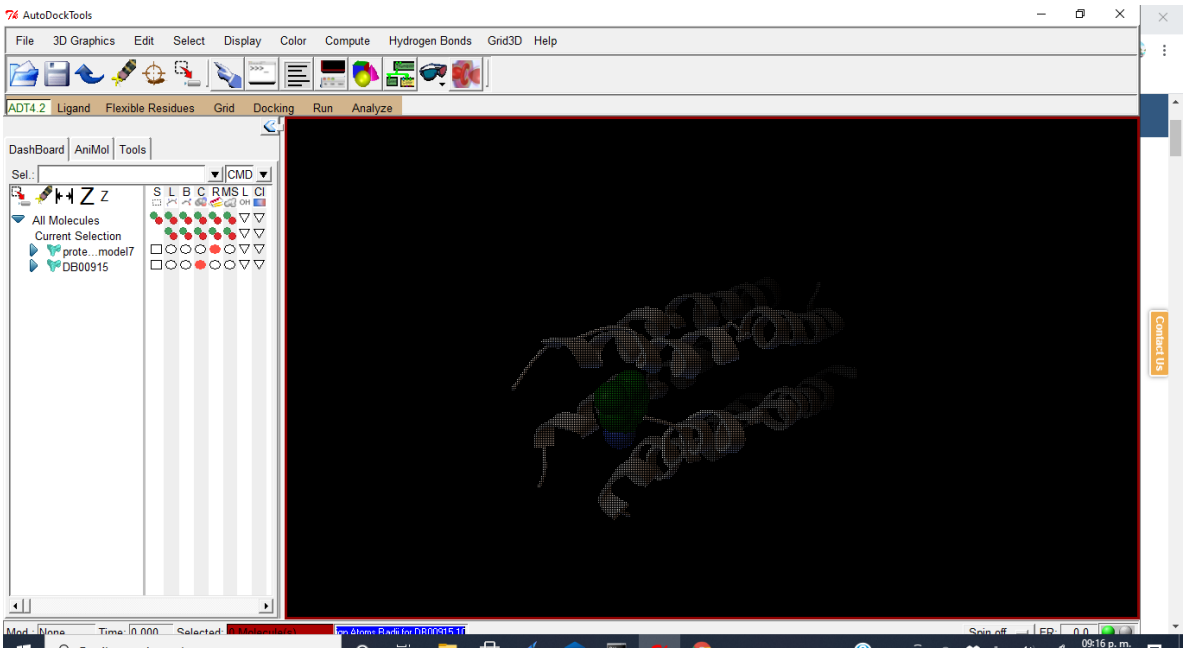

AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
All Molecules
Current Selection
prote...model1
prote...model2
prote...model3
prote...model4
prote...model5
prote...model6
prote...model7
prote...model8
prote...model9
protei...odel10
DB00915
Mod.: None Time: 0.000 Selected: 0 Molecule(s) Done 100%
Spin off FR: 66.7

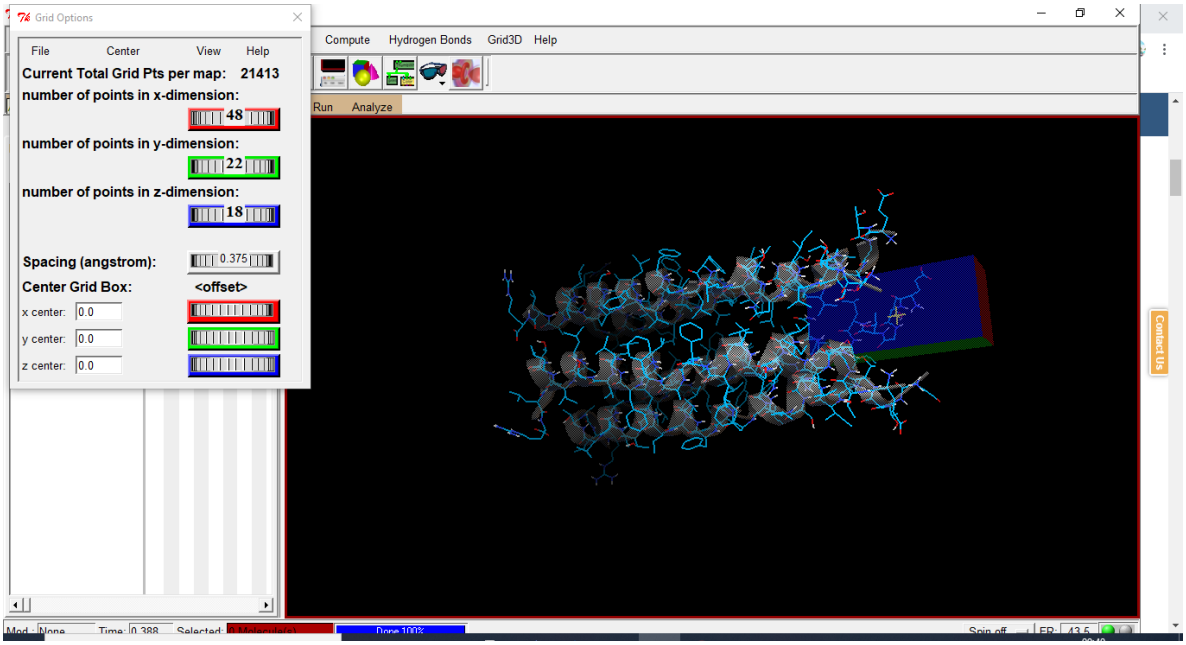

Grid Options
File
Center
View
Help
Current Total Grid Pts per map: 587121
number of points in x-dimension:
126
number of points in y-dimension:
68
number of points in z-dimension:
66
Spacing (angstrom):
0.375
Center Grid Box:
<offset>
x center: 24.894
y center: 0.015
z center: 0.023
prote...model9
protei...odel10
DB00915
Compute
Hydrogen Bonds
Grid3D
Help
Run
Analyze
Mod.: None
Time: 7.500
Selected: 0 Molecule(s)
Done 100%
Spin off
FR: 62.5


Grid Options
File
Center
View
Help
Current Total Grid Pts per map: 158369
number of points in x-dimension:
126
number of points in y-dimension:
28
number of points in z-dimension:
42
Spacing (angstrom):
0.375
Center Grid Box:
<offset>
x center: 24.894
y center: 0.015
z center: 0.023
prote...model9
protei...odel10
DB00915
Compute
Hydrogen Bonds
Grid3D
Help
Run
Analyze
Mod.: None
Time: 7.500
Selected: 0 Molecule(s)
Done 100%
Spin off
FR: 62.5


AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
Mod.: None Time: 0.015 Selected: 0 Molecule(s) Atoms Radii for proteinE_model
Spin off FR: 62.5


AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
All Molecules
Current Selection
prote...model7
DB00915
Contact Us


AutoDockTools
File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help
ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze
DashBoard AniMol Tools
Sel.: CMD
All Molecules
Current Selection
prote...model7
DB00915
Contact Us


Grid Options
File Center View Help
Current Total Grid Pts per map: 21413
number of points in x-dimension:
48
number of points in y-dimension:
22
number of points in z-dimension:
18
Spacing (angstrom):
0.375
Center Grid Box:
<offset>
x center: 0.0
y center: 0.0
z center: 0.0
Compute Hydrogen Bonds Grid3D Help
Run Analyze
Contact Us