

Spark - spark_3D

File Edit View Display Run Database Help

New Open Save Add Prot Wizard Constraints Process Reference And Results

Reset Atoms H X F Fields Shape Display Color Labels Clear Mol +ve -ve vdW Hyd Diff 2.0

Main toolbar

Ctrl+N Ctrl+O Ctrl+S

H X F

+ - * Ctrl+G

Results

Rank	Fav	Structure	BIF%	Score
118	☆		61	
119	☆		61	0.941 0.5
120	☆		60	0.941 0.5
121	☆		60	0.941 0.5
122	☆		60	0.941 0.5
123	☆		60	0.941 0.5
124	☆		60	0.941 0.5

Reference And Results Ctrl+A

All References

Protein Ctrl+X

All Results Ctrl+R

Selected Results Ctrl+E

Show Fields as Spheres 0

Show Fields as Tetrahedra 1

Show Fields as Cubes 2

Show Fields as Octahedra 3

Show Fields as Dodecahedra 4

Show Fields as Icosahedra 5

Index Shift+I

Element Shift+E

Type Shift+Y

Charge Shift+C

Formal Charge Ctrl+Shift+C

Field Size Shift+F

Chirality Shift+H

Residue Shift+R

Temperature Shift+T

Reset Labels Shift+N

Lines L

Thin Stick T

Capped Stick C

Ball and Stick B

CPK K

Starter

Right click

Copy...

Delete Selected Results Del

Delete Entire Cluster

Export Selected Results

Mark as Favorite Space

Tag Results

View Log for Selected Results Ctrl+L

View Parent Structure for Selected Result Ctrl+U

Copy Molecule(s) in 2D Ctrl+Alt+C

Copy Molecule(s) in 3D Ctrl+C

Starter weight 1 - 533

Left mouse rotate, right mouse translate, middle mouse/wheel/Alt-left scales. Shift-mouse z-clip.

Shortcut	File operation
F1	Show Manual
Ctrl+Shift+S	Save Project As
Ctrl+T	Open Project Notes Editor
Ctrl+P	Print All Results
Ctrl+Shift+P	Print Selected Results
Alt+F4	Exit

Shortcut	Molecule operation
Ctrl+F	Copy Favorites 3D
Ctrl+Alt+F	Copy Favorites 2D
Ctrl+V	Paste Reference from Clipboard
Ctrl+Space	Clear Favorites
Ctrl+Shift+V	Paste Excluded Volume Molecule

Shortcut	Display operation
Ctrl+Shift+R	Reset Display
Ctrl+Alt+R	Rock Display
Ctrl+Alt+S	Spin Display
F11	Full Screen