

Ctrl+N Ctrl+O Ctrl+S

H X F

+ - *

The screenshot displays the Spark software interface. At the top, there is a menu bar (File, Edit, Project, View, Display, Run, Database, Window, Help) and a toolbar with icons for various functions. Below the toolbar is a 'Results' table with columns for Rank, Fav, Structure, BIF%, and a 'Reference And Results' column. The table contains four rows of results. A 3D molecular model is shown in the center, with several context menus overlaid on it. These menus include options for showing fields as different shapes (Spheres, Tetrahedra, Cubes, Octahedra, Dodecahedra, Icosahedra), lines, thin stick, capped stick, ball and stick, and CPK. Other menus include 'Copy...', 'Delete Selected Results', 'Export Selected Results', 'Mark as Favorite', 'Tag Results', 'View Log for Selected Results', 'View Parent Structure for Selected Result', 'Copy Molecule(s) in 2D', 'Copy Molecule(s) in 3D', 'Atom RMB+click', and 'Center on Picked Atoms'. A status bar at the bottom shows 'Starter weight 1 - Untitled' and '0.974 - Fm-c(cn1)c(ccc2)n1c2'. A footer note reads: 'Left mouse rotate x/y, Ctrl left rotate z, right mouse translate, middle mouse/wheel/Alt-left scales. Shift-mouse z-clip.'

Rank	Fav	Structure	BIF%	Reference And Results
1	☆		98	All References Protein All Results Selected Results
2	☆		90	0.976
3	☆		89	0.974
4	☆		89	0.973

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
Ctrl+L	View log for selected molecules

Shortcut	View/ Display operation
Ctrl+Shift+R	Reset Display
Ctrl+Alt+R	Rock Display
Ctrl+Alt+S	Spin Display
F11	Full Screen
F10	Stereo
F4	Capture scene

Right click to access menu
Atom
Molecule in table
Toolbars and window headers
Column title