

Toggle 3D rendering on / off.

Access all windows such as Layers Infos; Control Panel; Ramachandran plot; Alignments etc.

Centers the screen on the active object(s).

Centers the view on a picked atom.

Lets you select/add/display all amino acids within a given radius of a picked atom.

Identifies an atom you pick on screen.

Measures the distance between two atoms.

Rotates objects. (Keep this activated to use the mouse commands below)

Zooms camera in / out. (OR hold R+L mouse buttons)

Translates camera. (OR hold R mouse button)

Displays the pdb-file in text format.

Switch between displaying the full backbone or just the C_{alpha} trace.

Deselect for a given pdb-file to move/rotate all other molecules.

Selects which pdb-files that are visible.

Sets the active pdb-file.

L-click to select/deselect individual amino acids; R-click to select/deselect entire column.

Column labels; L-click to toggle just the selected amino acids.

L-click to select all groups in that chain.

L-click to select all amino acids in that stretch of 2ndary structure.

Main toggle for showing or hiding an amino acid or group.

Toggle show side chain on / off.

Toggle show label on / off.

Toggle ribbons on / off.

Change colour of amino acid or group.

Changes what the colour commands will act upon: back bone; side chain etc.

DeepView / Swiss PDB viewer v3.7

?Layer	vis	mov	axis	CA	O	H	HondHdst	side	HOH	cyc	Sel
lgbr	v	v									0
1H2H_IgCb1	v										16

group	show	side	labl	ribn	col	b	s
VAL18	v						
LYS19	v						
VAL20	v						
SER21	v						
CYS22	v						
GLN23	v						
ALA24	v						
SER25	v						
GLY26	v						
TYR27	v						
ARG28	v						
PHE29	v						
SER30	v						
ASN31	v						
PHE32	v						
VAL33	v						
ILE34	v						
HIS35	v						
TRP36	v						
VAL37	v						
ARG38	v						
GLN39	v						
ALA40	v						
PRO41	v						
GLY42	v						
GLN43	v						
ARG44	v						
PHE45	v						
GLU46	v						
TRP47	v						
MET48	v						
GLY49	v						
TRP50	v						
ILE51	v						
ASN52	v						
PRO52	v						
TYR53	v						
ASN54	v						
GLY55	v						
ASN56	v						
LYS57	v						
GLU58	v						
PHE59	v						
SER60	v						
ALA61	v						
LYS62	v						
PHE63	v						
GLN64	v						
ASP65	v						
ARG66	v						
VAL67	v						
THR68	v						
PHE69	v						
THR70	v						
ALA71	v						
ASP72	v						
THR73	v						
SER74	v						
ALA75	v						
ASN76	v						
THR77	v						
ALA78	v						
TYR79	v						
MET80	v						
GLU81	v						
LEU82	v						