

## Build menu

- **Experimental Settings...**: The presence of protons depends on pH. Other parameters are also included.
- **Add Block**: Add any residue/base at the end of a structure. This way, a structure can be started from scratch by building the polypeptide chain (or a nucleic acid chain) in a completely extended form ( $\varphi=\psi=180^\circ$ ). 'Add Block' adds also a given sequence to a PDB structure already in memory. Adding bases has not been yet implemented.
- **Mute Block...**: Perform site directed mutagenesis on a selected residue (it is not yet implemented for bases). A single residue must have been selected (Control Click). Conformation of main chain atoms and of side chain  $C\beta$  is preserved.
- **Enter/Add Sequence...**: Type (or paste) in the one-letter code sequence to be built from scratch or added to a structure already in memory.
- **Add Last Block**: ( $\hat{A}CE\sim Y$ ) Repeat the last entered residue/base.
- **Undo Last Block**: ( $\hat{A}CE\sim U$ ) Undo sequentially the structure in memory.