Suppo

Fig.1. Structural-based alignment of α$_{1A}$-AR (PDB ID: 3SN6). The sequence shaded in cyan represents sequence similarity

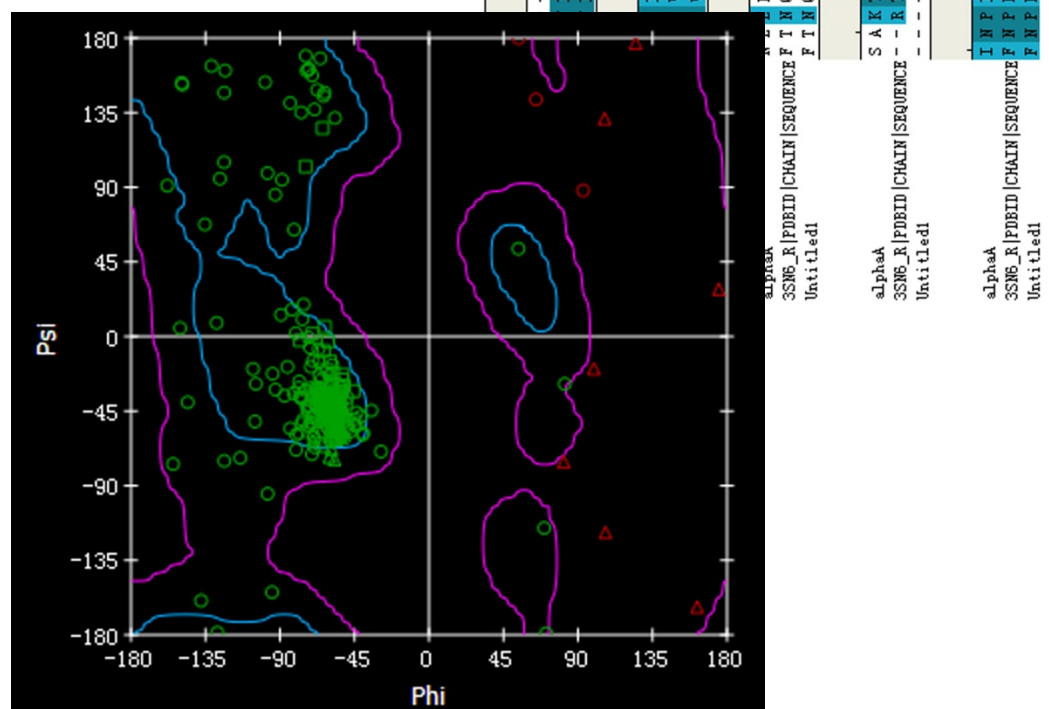Fig.2. Ramachandran plot of the α$_{1A}$-AR model calculated by PROCHECK