



**Figure S11.** Ramachandran plots of *C. albicans* and *S. cerevisiae* enolase 1 structures. The CaEno1 protein structure has 99.6% of all residues in favored (98.2%) and allowed (1.4%) regions. The ScEno1 (PDB ID 2al1) protein structure has 99.7% of all modeled residues in favored (98.1%) and allowed (1.6%) regions.