

Table S5. Geometric properties of the refined shark A0 homology model compared to structures of human ADORs.

	A0	A2a (2ydv)	A1 (6d9h)
Ramachandran favored (%)	92.74	97.43	96.58
Ramachandran allowed (%)	7.26	2.57	3.42
Ramachandran outliers (%)	0.00	0.00	0.00
Rotamer outliers (%)	0.36	1.92	0.00
C-beta deviations	0	0	0
MolProbity score	2.24	1.61	1.42