

Table S4 Hps5I76N protein modeling parameters from SwissModel of N-terminus (residues #1-370).

Residues	Sequence identity	E-score	Q-MEAN4 raw score	Q-MEAN4 Z-score	ANOLEA score at #76	GROMOS score at #76
30 to 104	19.5%	3.2E-24	0.584	-1.27	~ -2.5	~ -160