Table S1: Summary of sequence-based contact prediction methods in literature.

Methods	FC[1]	FOVC[2]	HBRH[3]	PROFCON[4]	VWP[5]	SVMCON[6]
Contact atoms	Heavy atoms	C_β	C_α	C_β	C_α	C_α
Distance cutoff	4.5Å	8 Å	8 Å	8 Å	8 Å	8 Å
Sequence separation cutoff	4	7	4	6	6	6
# of testing proteins	608	173	1033	633	435	48
Acc	0.16	0.21	0.22	0.32	0.37	0.37
Pct	N/A	0.5	0.5	0.5	0.5	1.0

[1] FC (Fariselli and Casadio, 1999)

[2] FOVC(Fariselli, et al., 2001)

[3] HBRH(Hamilton, et al., 2004)

[4] PROFCON(Punta and Rost, 2005)

[5] VWP(Vullo, et al., 2006)

[6] SVMCON(Cheng and Baldi, 2007)

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