Structural Quality Metrics Across Targets
(green = better, red = worse)

	pAE (median)	CDR RMSD (median, Å)	iPAE (median)	pLDDT (median)	H3 RMSD (median, Å)
B7-H3	8.32	13.28	14.89	0.91	1.28
CD47	7.34	10.88	13.69	0.90	1.10
CEACAM5	9.07	14.69	16.49	0.89	1.63
EGFR	9.01	15.32	16.73	0.91	1.53
EGFRvIII	8.77	12.73	16.16	0.89	1.08
EphA2	8.11	13.43	14.61	0.90	1.40
GPC2	9.28	15.23	16.98	0.88	1.01
HER2-DIV	7.31	21.18	15.23	0.89	1.30
MSLN-C	4.86	20.41	14.18	0.89	1.31
MSLN-N	5.23	8.40	9.87	0.89	1.32

Relative score (lower = better): 0.0 – 1.0