

Table S2: Prediction of ubiquitination site in ADAM33 protein using BDM-PUB server

Peptide	Position	Score	Threshold
ELLLELEKNHRLLAP	83	0.63	0.3
PWPPRGSKDFSTHEI	159	1.41	0.3
MEQLLTWKGTCGHRD	176	0.65	0.3
GHRDPGNKAGMTSLP	187	0.96	0.3
REARRTRKYLELYIV	210	1.05	0.3
QLRAFFRKGGGACLS	399	1.49	0.3
CCVRCLLKPAALCR	469	1.56	0.3
GRDALCGKLQCQGGK	575	1.66	0.3
KLQCQGGKPSLLAPH	582	0.98	0.3
CQSRRCRKNAFQELQ	644	1.65	0.3
WAPPFCDKPGFGGSM	682	0.68	0.3
DPACSGPKDGPHRDH	749	1.71	0.3
EPSSHPEKPLPAVSP	791	0.75	0.3