

DARPin/scFV	Position	Peptide	Affinity (nM)	Peptide hit (Human proteome)
Cetuximab	135	ILLTQSPVI	51.93	No Hits
Cetuximab	138	TQSPVILSV	86.36	No Hits
E01	19	ILMANGADV	38.9	100% to GABPB1
E01	97	YIGDTPLHL	82.91	100% to NFKB2
Ec1	116	VLLKNVADV	64.38	No Hits
Ec1	83	VLLKYGADV	96.29	89% to GABPB1
G3	19	ILMANGADV	38.9	100% to GABPB1
G3-Ec1-E01	344	ILMANGADV	38.9	100% to GABPB1
G3-Ec1-E01	18	ILMANGADV	38.9	100% to GABPB1
G3-Ec1-E01	261	VLLKNVADV	64.38	89% to GABPB1
G3-Ec1-E01	422	YIGDTPLHL	82.91	100% to NFKB2
G3-Ec1-E01	228	VLLKYGADV	96.29	89% to GABPB1