

0900Q2 SH2B3 1 362 425		1	YFWHF	GLIS	AKAAQVLVGGT	GSBGFVLRQSETR	GEVLTINFGQR	AKHLRSLINEB	GCQRV	QHLNFSI FDLHEHVRPIL	ESG	GSBVDVLSY	100
014492 SH2B2 1 415 515		1	YFWHF	GLIS	AKAAQVLVGGT	DAGBFLVRQSETR	GEVLTINFGQR	AKHLRSLINEB	GCQRV	QHLNFSI FDLHEHVRPIL	ECG	ACADVLSY	100
014492 SH2B2 1 415 515		1	YFWHF	GLIS	AKAAQVLVGGT	RNGLQVLRQSETR	GEVLTINFGQR	AKHLRSLINEB	GCQRV	QHLNFSI FDLHEHVRPIL	ESG	GSADILSY	100
205 SH2 1 119 421 CRK		1	SSMYW	GRIS	AKAAQVLVGGT	RHGVFLVRQSETR	GDVLSVNSNR	VSHY	INSNGRPPVPSA	QPPGVSFSLRQIQDQFDSIPALFEYKIHLYDT		TLTLEP	107
014492 SH2B2 1 415 515		1	SSMYW	GRIS	AKAAQVLVGGT	RHGVFLVRQSETR	GDVLSVNSNR	VSHY	INSNGRPPVPSA	QPPGVSFSLRQIQDQFDSIPALFEYKIHLYDT		TLTLEP	107
P46108 CRK 1 11 118		1	SSMYW	GRIS	AKAAQVLVGGT	RHGVFLVRQSETR	GDVLSVNSNR	VSHY	INSNGRPPVPSA	QPPGVSFSLRQIQDQFDSIPALFEYKIHLYDT		TLTLEP	107
21W SH2 1 119 207 CRKL		1	SAMYM	GPVS	RQEAQTRLQCG	RHGMFLVRQSDTCE	GDVLSVNSNR	VSHY	INSNLNPN	RFKI	QDQFDSIPALFEYKIHLYDT	TLTLEP	90
P46109 CRKL 1 12 102		1	SAMYM	GPVS	RQEAQTRLQCG	RHGMFLVRQSDTCE	GDVLSVNSNR	VSHY	INSNLNPN	RFKI	QDQFDSIPALFEYKIHLYDT	TLTLEP	90
014492 SH2B3 1 439 589		1	QHWFF	GRIS	REESRLIRQCG	VDGLFLVRQSGNR	KAPVLTICHQK	KNQFLIPCED	DQUTY	FSLDGNTFTSBLQVDFVQLNKGVL		PKCKIRH	98
014492 SH2B3 1 439 589		1	QHWFF	GRIS	REESRLIRQCG	VDGLFLVRQSGNR	KAPVLTICHQK	KNQFLIPCED	DQUTY	FSLDGNTFTSBLQVDFVQLNKGVL		PKCKIRH	98
1M4 SH2 1 119 1139 GRB7		1	QLMFH	GRIS	REESRLIRQCG	VDGLFLVRQSGNR	KAPVLTICHQK	VKHYLLFSESE	EGRLY	FSMDGQRTFTTLQVFEHQLNRGL		PCLLRHC	98
214K SH2 1 119 1139 GRB7		1	QLMFH	GRIS	REESRLIRQCG	VDGLFLVRQSGNR	KAPVLTICHQK	VKHYLLFSESE	EGRLY	FSMDGQRTFTTLQVFEHQLNRGL		PCLLRHC	98
Q14451 GRB7 1 429 527		1	QLMFH	GRIS	REESRLIRQCG	VDGLFLVRQSGNR	KAPVLTICHQK	VKHYLLFSESE	EGRLY	FSMDGQRTFTTLQVFEHQLNRGL		PCLLRHC	98
P42679 MARK 1 120 211		1	MPWFH	GRIS	QKAAVLOLQPG	EDGFLVLRSSARH	GDVLSVNSNR	VHRYVLRMGDS	HLTID	EAVFCNLNMHVEHYSKRGAI		CTKILVRP	93
014492 SH2B3 1 439 589		1	MPWFH	GRIS	QKAAVLOLQPG	EDGFLVLRSSARH	GDVLSVNSNR	VHRYVLRMGDS	HLTID	EAVFCNLNMHVEHYSKRGAI		CTKILVRP	93
P29350 PTFN6 1 108 213		1	ERYWH	GHMS	GGQAEILLQAKG	EPWTLVLRSSLSQ	GDVLSVLSQPKAGPGLR	VTHIKVMCEGG	RYTVG	GLETFDSLTDLVEHFKKTGIE	ASG	AFVYLQRP	105
2YK SH2 2 119 8 PTFN6		1	YH	GHMS	GGQAEILLQAKG	EPWTLVLRSSLSQ	GDVLSVLSQPKAGPGLR	VTHIKVMCEGG	RYTVG	GLETFDSLTDLVEHFKKTGIE	ASG	AFVYLQRP	102
2M7 SH2 2 119 40 PTFN6		1	YH	GHMS	GGQAEILLQAKG	EPWTLVLRSSLSQ	GDVLSVLSQPKAGPGLR	VTHIKVMCEGG	RYTVG	GLETFDSLTDLVEHFKKTGIE	ASG	AFVYLQRP	102
P59350 SH2 2 119 40 PTFN6		1	ERYWH	GHMS	GGQAEILLQAKG	EPWTLVLRSSLSQ	GDVLSVLSQPKAGPGLR	VTHIKVMCEGG	RYTVG	GLETFDSLTDLVEHFKKTGIE	ASG	AFVYLQRP	105
5X34 SH2 1 119 772 PTFN11		1	ERWFH	GHLS	GKEAKLLITEK	KHGSFLVRSSGSH	GDVLSVRTGDDGSGNDKGS	KVTHVMRCQEL	KYDVG	GERFDSLTDLVEHYKKNMVE	TLG	TVLQKLP	106
6R50 SH2 2 119 248 PTFN11		1	ERWFH	GHLS	GKEAKLLITEK	KHGSFLVRSSGSH	GDVLSVRTGDDGSGNDKGS	KVTHVMRCQEL	KYDVG	GERFDSLTDLVEHYKKNMVE	TLG	TVLQKLP	106
5D6 SH2 2 119 378 PTFN11		1	ERWFH	GHLS	GKEAKLLITEK	KHGSFLVRSSGSH	GDVLSVRTGDDGSGNDKGS	KVTHVMRCQEL	KYDVG	GERFDSLTDLVEHYKKNMVE	TLG	TVLQKLP	106
Q06124 PTFN11 2 110 216		1	ERWFH	GHLS	GKEAKLLITEK	KHGSFLVRSSGSH	GDVLSVRTGDDGSGNDKGS	KVTHVMRCQEL	KYDVG	GERFDSLTDLVEHYKKNMVE	TLG	TVLQKLP	106
P10000 SH2 1 119 108 RASAI		1	KIWFH	GRIS	KQAYNLMTVG	QVSSFLVRPSDNT	GVLSYLFRTNNE	LQRFKISFTPN	NQPM	GCGRYNSIGDIIDHYRKEVQE		CYLLKEP	92
8DQ0 SH2 1 119 108 RASAI		1	KIWFH	GRIS	KQAYNLMTVG	QVSSFLVRPSDNT	GVLSYLFRTNNE	LQRFKISFTPN	NQPM	GCGRYNSIGDIIDHYRKEVQE		CYLLKEP	92
8DQ0 SH2 1 119 108 RASAI		1	KIWFH	GRIS	KQAYNLMTVG	QVSSFLVRPSDNT	GVLSYLFRTNNE	LQRFKISFTPN	NQPM	GCGRYNSIGDIIDHYRKEVQE		CYLLKEP	92
P20936 RASAI 2 349 441		1	KIWFH	GRIS	KQAYNLMTVG	QVSSFLVRPSDNT	GVLSYLFRTNNE	LQRFKISFTPN	NQPM	GCGRYNSIGDIIDHYRKEVQE		CYLLKEP	92
1P08 SH2 1 119 142 SHC1		1	EPWFH	GRIS	REAEAILQL	NGDPLVRSSSTT	GVLTISQSGQ	PKHLLIDPEG	VVRT	KDHRFSVSHLSIYHMDNLPI	</		