

Supplementary Table1: Summary of the interaction partners and their average protein ratios from at least two pull-down experiments.

Phosphotyrosine	interaction partner	Ratio ± s.d.	Swiss-Prot Number
EGFR			
727	Transforming protein Shc	7.5 ± 1.0	P29353
891	SH3 domain-binding glutamic acid-rich-like protein	13.7 ± 7.6	Q9H299
978	STAT5	9.7 ± 2.9	P516981
	Protein tyrosine phosphatase 2C	3.1 ± 1.4	Q06124
998	STAT5	8.4 ± 3.0	P516981
	Adapter molecule Crk	4.9 ± n.a.	P46108
	Protein tyrosine phosphatase 2C	3.8 ± n.a.	Q06124
	Transforming protein Shc	6.7 ± 1.0	P29353
	Tyrosine protein kinase Csk	3.1 ± n.a.	P41240
1016	Protein tyrosine phosphatase 2C	3.6 ± 1.5	Q06124
1069	Signal transduction protein Cbl-B	19.3 ± 8.5	P22681
1092	Grb2	10.3 ± 2.8	P29354
1110	Grb2	7.7 ± 3.0	P29354
	Transforming protein Shc	6.0 ± 2.8	P29353
1125	Grb2	13.3 ± 7.4	P29354
1138	Grb2	10.5 ± 3.5	P29354
	Transforming protein Shc	5.7 ± 1.9	P29353
1172	Grb2	13.5 ± 7.8	P29354
	Transforming protein Shc	12.6 ± 2.3	P29353
1197	Grb2	9.7 ± 4.8	P29354
	Transforming protein Shc	7.7 ± 3.0	P29353

Phosphotyrosine	interaction partner	Ratio ± s.d.	Swiss-Prot Number
ErbB2			
735	Transforming protein Shc	6.8 ± 1.5	P29353
923	SH3 domain-binding glutamic acid-rich-like protein	5.5 ± 1.2	Q9H299
1005	Transforming protein Shc	5.6 ± 2.1	P29353
1023	Protein tyrosine phosphatase 2C	4.3 ± n.a.	Q06124
1139	Grb2	6.7 ± 3.7	P29354
1196	Transforming protein Shc	7.2 ± 3.1	P29353
	SH3 domain-binding glutamic acid-rich-like protein	4.1 ± n.a.	Q9H299
1222	Transforming protein Shc	5.4 ± 2.3	P29353
1248	Transforming protein Shc	6.7 ± 1.8	P29353

Phosphotyrosine	interaction partner	Ratio ± s.d.	Swiss-Prot Number
ErbB3			
1054	Phosphatidylinositol 3-kinase regulatory beta subunit	11.4 ± 5.4	O00459
1197	Phosphatidylinositol 3-kinase regulatory beta subunit	4.7 ± 2.3	O00459
1199	Grb2	13 ± 4.0	P29354
	Tyrosine protein kinase Csk	6.7 ± 2.3	P41240
1222	Phosphatidylinositol 3-kinase regulatory beta subunit	30.0 ± 4.2	O00459
1260	Phosphatidylinositol 3-kinase regulatory beta subunit	6.5 ± 2.8	O00459
1262	Grb2	8.3 ± 3.1	P29354
1276	Phosphatidylinositol 3-kinase regulatory beta subunit	3.8 ± 1.9	O00459
1289	Phosphatidylinositol 3-kinase regulatory beta subunit	20.0 ± 11.3	O00459
1328	Transforming protein Shc	5.8 ± 2.3	P29353

Phosphotyrosine	interaction partner	Ratio ± s.d.	Swiss-Prot Number
ErbB4			
733	Transforming protein Shc	8.3 ± 2.3	P29353
984	STAT5	9.3 ± 2.8	P516981
	Protein tyrosine phosphatase 2C	3.9 ± 2.4	Q06124
1022	Adapter molecule Crk	12.1 ± 3.2	P46108
1056	Phosphatidylinositol 3-kinase regulatory beta subunit	30 ± n.a.	O00459
1162	Grb2	5.8 ± 2.7	P29354
1188	Grb2	30 ± n.a.	P29354
	Transforming protein Shc	5.3 ± 1.6	P29353
1202	Grb2	6.3 ± 1.0	P29354
1208	Grb2	3.8 ± 2.1	P29354
1221	Grb2	4.8 ± 2.3	P29354
1242	Grb2	8.2 ± 1.7	P29354
	Transforming protein Shc	5.6 ± n.a.	P29353
1268	Grb2	3.5 ± 1.5	P29354
	Nck adapter protein 2	5.0 ± 2.2	O43639
1284	Transforming protein Shc	4.1 ± 1.4	P29353