A Ubiquitinated IQGAP1 peptides

Coguenee	m/z	Charga		Modification	Mascot
Sequence	observed	Charge	ррп		Score
ILAIGLINEALDEGDAQ <u>K</u> <sup>GG</sup> TLQALQIPAAK	1044.5824	3	0.53	K(GlyGly) 556	71
FLSAIVSSVD <u>K</u> GGIPYGMR	666.3539	3	1.09	K(GlyGly) 1155	81
NLGSIAK <sup>GG</sup> M <sup>Ox</sup> LQHAASNK	604.9838	3	0.83	K(GlyGly) 1230	26
K <sup>GG</sup> LTELGTVDPK	438.9137	3	0.15	K(GlyGly) 1465	34
LTELGTVDP <u>K</u> <sup>GG</sup> NK	476.9280	3	0.15	K(GlyGly) 1475	28
ATFYGEQVDYY <u>K</u> <sup>GG</sup> SYIK	1045.0003	2	0.98	K(GlyGly) 1528	77



**Figure S1. Summary of mass spectrometry data.** (A) Table summarizing mass spectrometry results for each identified peptide. (B-G) Representative mass spectra for ubiquitinated IQGAP1 peptides. (B) 539-567 containing ubiquitinated Lys-556. (C) 1145-1161 containing ubiquitinated Lys-1155. (D) 1224-1239 containing ubiquitinated Lys-1230. (E) 1465-1475 containing ubiquitinated Lys-1465. (F) 1466-1477 containing ubiquitinated Lys-1475. (G) 1517-1532 containing ubiquitinated Lys-1528.



Figure S2. MG132 does not alter the amount of IQGAP1 in cell lysates.

(A) HEK293 cells were incubated with MG132 (+) or DMSO (-) for 4 h. Equal amounts of protein lysate were analyzed by Western blotting using anti-IQGAP1, anti-actin and anti-ubiquitin (Ub) antibodies. (B) The IQGAP1 bands were quantified with Image Studio 2.0 and corrected for the amount of actin in the same sample. Data are expressed as means +/- SD (n=3) and compared using Welch's t-test, with cells incubated with DMSO set as 1. ns: not significant.



**Figure S3.** Lysine mutations in the GRD of IQGAP1 do not disturb interactions between IQGAP1 and CDC42 or RAC1.

(A) Purified His-CDC42(Q61L) was incubated with equal amounts of purified GST-tagged WT IQGAP1 (WT) or IQGAP1 GRD-2K (GRD-2K) or GST alone. Complexes were isolated with glutathione-Sepharose (PD) and bound proteins were analyzed by SDS-PAGE. The gel was cut at ~50kDa; the upper section of the gel was stained with InstantBlue Protein Stain, while the lower section was processed by Western blotting and probed with anti-CDC42 antibodies. (B) RAC1 pulldown was performed as described above for CDC42, except His-RAC1(Q61L) was used and the Western blot was probed with anti-RAC1 antibodies. Data are representative of 2 to 4 experiments. (C) Purified His-CDC42(Q61L) and His-RAC1(Q61L) proteins were resolved by SDS-PAGE and stained with InstantBlue Protein Stain to show purity.

IQGAP1	S KEK RE KLE AY QHL FY LLQ TN PTY LA KL	I FQMPQNK STK FMDSV IF TLY NY ASN QR EEY L	1021
IQGAP2	S KER RK TLE TY QQL FY LLQ TN PLY LA KL	I FQMPQNK STK FMDTV IF TLY NY ASN QR EEY L	934
IQGAP3	SKEKRQKLEAYQHLFYLLQTQPIYLAKL	I FQMPQNKTTKFMEAVIFSLYNYASSRREAYL	1005
IQGAP1	L LRL FK TAL QE EIK SK VDQ I QE IV TG NP	TVI KMVVS FNR GARGQNALRQILAPVVKEIMD	1081
IQGAP2	LLKLFKTALEEEIKSKVDQVQDIVTGNP	TVIKMVVSFNRGARGQNTLRQLLAPVVKEIID	994
IQGAP3	LLQLFKTALQEEIKSKVEQPQDVVTGNP	TVVRLVVRFYRNGRGQSALQEILGKVIQDVLE	1065
IQGAP1	DKSLNIKTDPV DIYKSWVNQMESQTGEA	SKL PYDVT PEQALAHE EVKTR LDSSI RNMRA V	1141
IQGAP2	DKSLIINTN PVEVYKAWVNQLETQTGEA	SKLPYDVTTEQALTYPEVKNKLEASIENLRRV	1054
IQGAP3	DKVL SVHTDPVHLYKNWINQT EAQ TGQR	SHL PYDVT PEQAL SHP EV QRR LD IAL RN LLAM	1125
IQGAP1	T DKF LS AIV SS VD <mark>K</mark> I P YGMRF I AK VL KD	S LH EKF PD AGE DE LLK II GNL LY YRYMN PAI V	1201
IQGAP2	TDKVLNSIISSLDLLPYGLRYIAKVLKN	S IHEKF PDATE DE LLK IVGNL LY YRYMN PAI V	1114
IQGAP3	TDKFLLAITSSVDQIPYGMRYVAKVLKA	T LAEKF PDATD SE VYK VV GNL LY YRF LN PAV V	1185
IQGAP1	APDAFDIIDLSAGGQLTTDQRRNLGSIA	KMLQHAASNKMFLGDNAHLSIINEYLSQSYQK	1261
IQGAP2	APDGFDIIDMTAGGQINSDQRRNLGSVA	KVLQHAASNKLFEGENEHLSSMNNYLSETYQE	1174
IQGAP3	A PDA FD IVAMA AGG AL AAP QR HAL GA VA	Q LL QHA AA GKA FS GQS QH LRV LN DYL EE THL K	1245
IQGAP1	F RRF FQ TAC DV PELQD KFN VD EYS DL VT	L TK PVI YI SIGEI INT HT LLL DH QDA IA PEH N	1321
IQGAP2	F RKY FK EAC NV PE PEE KFN MD KYT DL VT	V SK PVI YI SIE EI IST HS LLL EH QDA IA PEKN	1234
IQGAP3	FRKFIHRACQVPEPEERFAVDEYSDMVA	VAK PMVYI TVGEL VNT HR LLL EH QDC IA PDH Q	1305
IQGAP1	DPIHELLDDLGEVPTIESLIGESS	1345	
IQGAP2	DLLSELLGSLGEVPTVESFLGEGA	1258	
IQGAP3	DPLHELLEDLGELPTIPDLIGESI	1329	

**Figure S4.** Amino acid alignments of the GRD of human IQGAPs. Sequences of the GRD of IQGAP1 (962-1345), IQGAP2 (875-1258) and IQGAP3 (946-1329) were aligned by Clustal W Multiple Alignment. Lys-1155 and Lys-1230 of IQGAP1 are underlined and colored red. **Video S1.** Time-lapse wound healing assay of IQGAP1-null MEFs expressing GFP-tagged WT IQGAP1 or IQGAP1 GRD-2K. The video corresponds to the experiment shown in Fig. 6B.