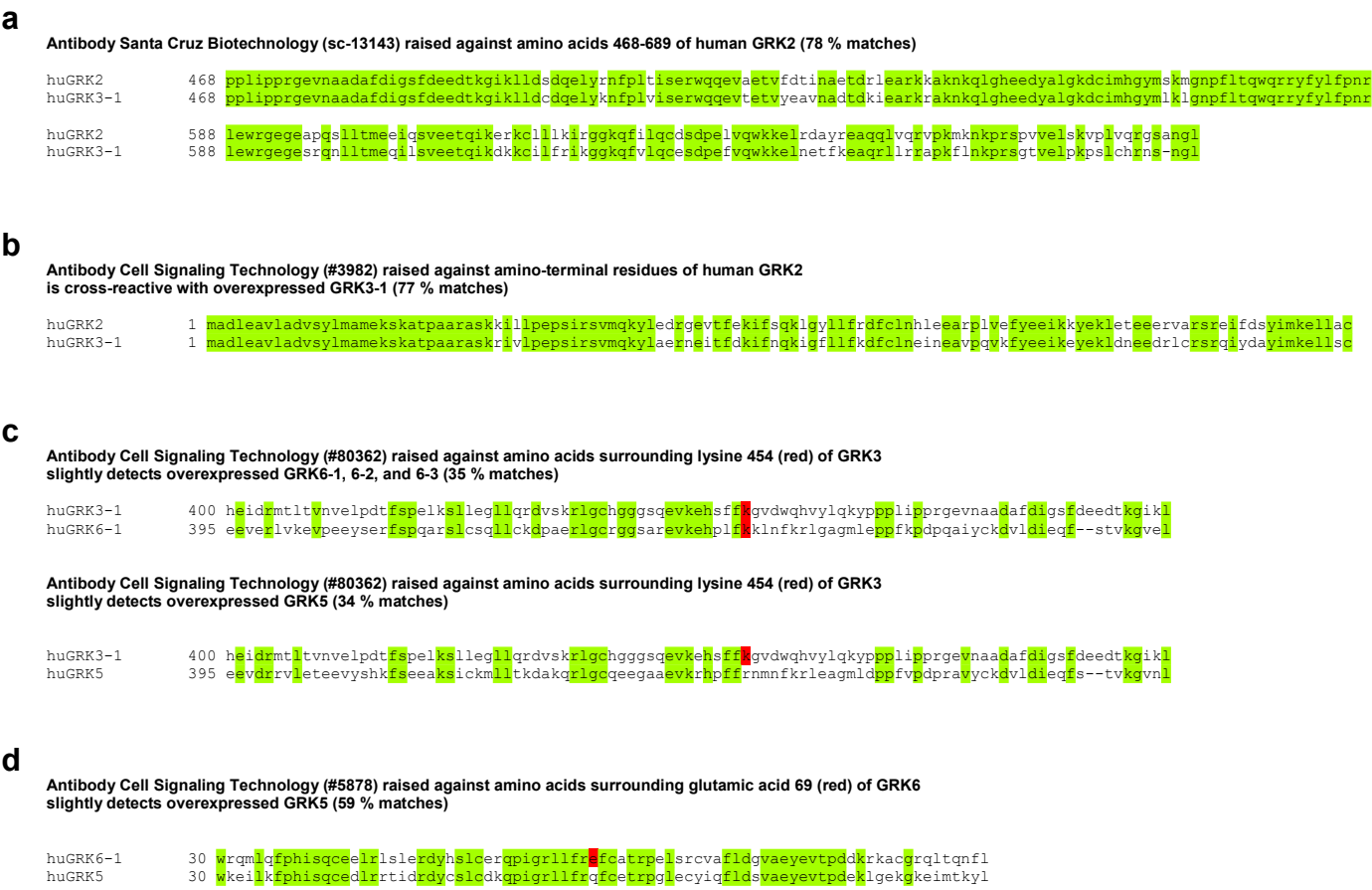
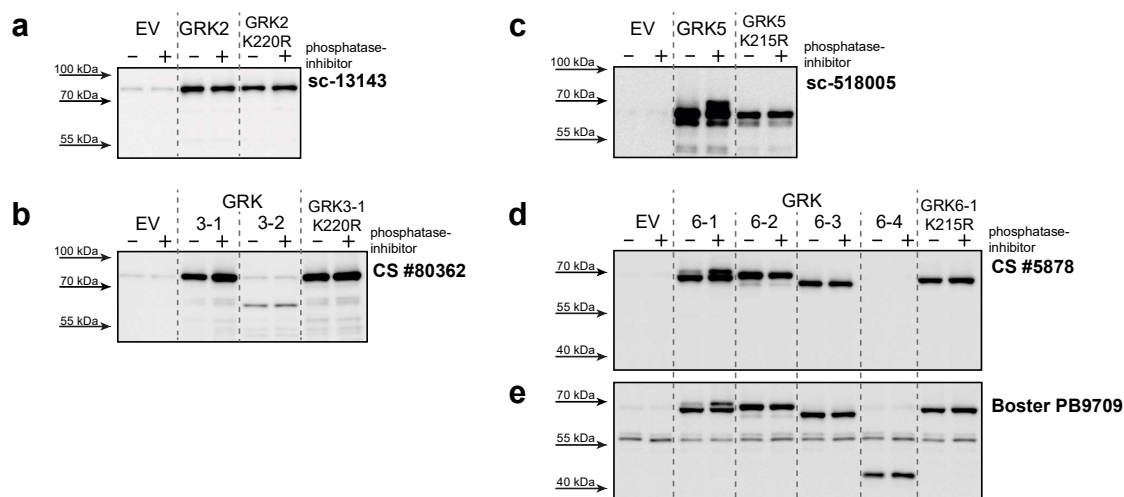


Supplementary Figure S1



Supplementary Figure S1: Alignment of protein sequences used to create designated antibodies and the corresponding sequence of the indicated GRK isoform which are also detected by the antibody. Amino acids identical in both sequences are highlighted in green. Total identity is denoted in per cent. **a** Alignment of human GRK2 and human GRK3-1 protein sequences from amino acid 468 to 689 which are both recognized by antibody sc-13143 (Santa Cruz Biotechnology). **b** Alignment of human GRK2 and human GRK3-1 protein sequences from the first amino acid to amino acid 120 which are both recognized by antibody CS #3982 (Cell Signaling Technology). **c** Alignment of human GRK3-1 with human GRK6-1 and GRK5 protein sequences from amino acid 400 to 499 and 395 to 494, respectively. Antibody CS #80362 (Cell Signaling Technology) was raised against GRK3 but also detects GRK6-1, -2, -3 and GRK5. **d** Alignment of human GRK6-1 and human GRK5 protein sequences from amino acid 30 to 499 which are both recognized by antibody CS #5878 (Cell Signaling Technology).

Supplementary Figure S2



Supplementary Figure S2:

HEK293 cells were transfected with either empty vector (EV), GRK2 (**a**), 3 (**b**), 5 (**c**), and 6 isoforms (**d**) or their corresponding kinase-dead variants GRK2-K220R (**a**), GRK3-1-K220R (**b**), GRK5-K215R (**c**), and GRK6-1-K215R (**d**). Lysates were prepared in presence (+) or absence (-) of phosphatase inhibitors and EDTA. Western blot analysis of the lysates using the indicated antibodies was performed.

Supplementary Figure S3

GRK2

a Antibody Santa Cruz Biotechnology (sc-13143) raised against amino acids 468-689 of human GRK2

huGRK2	468	pplipprgeevnaadafdigsfdeedtkgikllsdsgelyrnfpltiserwqgevaetvfdtinaetdrlearkkaknkqlgheedyalgkdcimhgyskmgnpfltqwqrryfyfnpnr	
muGRK2-1	468	pplipprgeevnaadafdigsfdeedtkgikllsdsgelyrnfpltiserwqgevaetvfdtinaetdrlearkkaknkqlgheedyalgkdcivhgyskmgnpfltqwqrryfyfnpnr	
ratGRK2	468	pplipprgeevnaadafdigsfdeedtkgikllsdsgelyrnfpltiserwqgevaetvfdtinaetdrlearkkaknkqlgheedyalgkdcimhgyskmgnpfltqwqrryfyfnpnr	
hamGRK2-x2	468	pplipprgeevnaadafdigsfdeedtkgikllsdsgelyrnfpltiserwqgevaetvfdtinaetdrlearkkaknkqlgheedyalgkdcimhgyskmgnpfltqwqrryfyfnpnr	
monGRK2	468	pplipprgeevnaadafdigsfdeedtkgikllsdsgelyrnfpltiserwqgevaetvfdtinaetdrlearkkaknkqlgheedyalgkdcimhgyskmgnpfltqwqrryfyfnpnr	
huGRK2	588	lewrgegeapqsltmeeiqsveetqikerkc1llkirgkqfvlqcdsdpe1vgwkkelrdayreaqqlvqrvpkmknkprspvvelskvplvqrgsangl	98 % identity to huGRK2
muGRK2-1	588	lewrgegeapqsltmeeiqsveetqikerkc1llkirgkqfvlqcdsdpe1vgwkkelrdayreaqqlvqrvpkmknkprspvvelskvplvqrgsangl	98 % identity to huGRK2
ratGRK2	588	lewrgegeapqsltmeeiqsveetqikerkc1llkirgkqfvlqcdsdpe1vgwkkelrdayreaqqlvqrvpkmknkprspvvelskvplvqrgsangl	99 % identity to huGRK2
hamGRK2-x2	588	lewrgegeapqsltmeeiqsveetqikerkc1llkirgkqfvlqcdsdpe1vgwkkelrdayreaqqlvqrvpkmknkprspvvelskvplvqrgsangl	99 % identity to huGRK2
monGRK2	588	lewrgegeapqsltmeeiqsveetqikerkc1llkirgkqfvlqcdsdpe1vgwkkelrdayreaqqlvqrvpkmknkprspvvelskvplvqrgsangl	99 % identity to huGRK2

b Antibody Cell Signaling Technology (#3982) raised against amino-terminal residues of human GRK2

huGRK2	1	madleavladvsylmamekskatpaaraskkillpepsirsvmqyledrgvtfekifsqklgyllfrdfc1nhleearplvefyeeikkyekleteeervrsreifdsyimkellac	
muGRK2-1	1	madleavladvsylmamekskatpaaraskkillpepsirsvmqyledrgvtfekifsqklgyllfrdfc1nhleearplvefyeeikkyekleteeervrsreifdsyimkellac	98 %
ratGRK2	1	madleavladvsylmamekskatpaaraskkillpepsirsvmqyledrgvtfekifsqklgyllfrdfc1nhleearplvefyeeikkyekleteeervrsreifdsyimkellac	96 %
hamGRK2-x2	1	madleavladvsylmamekskatpaaraskkillpepsirsvmqyledrgvtfekifsqklgyllfrdfc1nhleearplvefyeeikkyekleteeervrsreifdsyimkellac	97 %
monGRK2	1	madleavladvsylmamekskatpaaraskkillpepsirsvmqyledrgvtfekifsqklgyllfrdfc1nhleearplvefyeeikkyekleteeervrsreifdsyimkellac	100 %

Supplementary Figure S3: Alignment of human, mouse, rat, hamster, and monkey GRK2 protein sequences from amino acid 468 to 689 **(a)** and 1 to 120 **(b)**. Differing amino acids are highlighted in yellow. Total identity is denoted in per cent.

Supplementary Figure S4

GRK3

Antibody Cell Signaling Technology (#80362) raised against amino acids surrounding lysine 454 (red)

huGRK3-1	400	heidrmtltvnvelpdtfspelksllegllqrdvskrlgchgggsqevkehsff	gvdwqhvy	lqkypplipprgevn	aa	afdigsfdeedtkgiklld	
muGRK3-1	400	heidrmtltvnvlpd	fspel	sllegllqrdvs	rlgcggg	arelkehiff	gldwqhvy
ratGRK3	400	heidrmtltvnvlpd	fspel	sllegllqrdvs	rlgcggg	arelkehiff	gldwqhvy
hamGRK3-x3	400	heidrmtltvnvlpd	fspel	sllegll	rdvs	rlgcggg	qevkehsff
monGRK3-x1	400	heidrmtltvnvelpdtfspelksllegllqrdvskrlgchgggsqevkehsff	gvdwqhvy	lqkypplipprgevn	aa	afdigsfdeedtkgiklld	

Supplementary Figure S4: Alignment of human, mouse, rat, hamster, and monkey GRK3 protein sequences from amino acid 400 to 100. Differing amino acids are highlighted in yellow. Total identity is denoted in per cent.

Supplementary Figure S5

GRK5

Antibody Santa Cruz Biotechnology (sc-518005) raised against amino acids 94-157 of human GRK5

huGRK5	94	pdeklgekgkeimtkyltpkspvfiaqvggdlvsqteekllqkpkelfsacagsvheylrgep	
muGRK5	94	pdehlgaakgkeimtkyltpkspvfiaqvggdlvsqteekllqspckelfsacagsvhdylkgdp	89 %
ratGRK5	94	pdehlgaakgkeimtkylspkspvfiaqvggdlvsqteekllqspckelfsacagsvhdylkgdp	87 %
hamGRK5-3	94	pdehlgekgkeiitkyltpkspvfiaqvggdlvsqteakllqkpkelfsacagsvhdylkgdp	89 %
monGRK5	94	pdeklgekgkeimtkyltpkspvfiaqvggdlvssteekllqkpkelfsacagsvheylrgep	95 %

Supplementary Figure S5: Alignment of human, mouse, rat, hamster, and monkey GRK5 protein sequences from amino acid 94 to 157. Differing amino acids are highlighted in yellow. Total identity is denoted in per cent.

Supplementary Figure S6

GRK6

a Antibody Cell Signaling Technology (#5878) raised against amino acids surrounding glutamic acid 69 (red) for GRK6

huGRK6-1	30	wrqmlqfphisqceelrlslerdyhslcerqpigrllfrfc	atrpelsrcvafldgvaeyevtpddkrkacgrqltgnfl	
muGRK6-3	30	wrqmlqfphisqceelrlslerdyhslcerqpigrllfrfc	atrpeltrctafldgvseyevtpdekrkacgrlmgnfl	92 %
ratGRK6-3	30	wrqmlqfphisqceelrlslerdyhslcerqpigrllfrfc	atrpeltrctafldgvaeyevtpdekrkacgrlmgnfl	93 %
hamGRK6-x1	30	wrqmlqfphisqceelrlslerdyhslcerqpigrllfrfc	atrpeltrctafldgvaeyevtpdekrkacgrlmgnfl	93 %
monGRK6-x1	30	wrqmlqfphisqceelrlslerdyhslcerqpigrllfrfc	atrpelsrcvafldgvaeyevtpdekrkacgrlmgnfl	96 %

b Antibody Boster Biological Technology (PB9709) raised against amino acids 382-417 of human GRK6

huGRK6-1	382	qspfqqrkkkkikreeverlvkevpeeyserfspqar	
muGRK6-3	382	qspfqqrkkkkikreeverlvkevpeeytdrfsqar	88 %
ratGRK6-3	382	qspfqqrkkkkikreeverlvkevpeeytdrfsqar	91 %
hamGRK6-x1	382	qspfqqrkkkkikreeverlvkevpeeysehfspqar	97 %
monGRK6-x1	382	qspfqqrkkkkikreeverlvkevpeeyserfspqar	100 %

Supplementary Figure S6: Alignment of human, mouse, rat, hamster, and monkey GRK6 protein sequences from amino acid 30 to 110 **(a)** and 382 to 417 **(b)**. Differing amino acids are highlighted in yellow. Total identity is denoted in per cent.

Supplementary Table S1

Last edited on 19th of October 2021
Overview of mRNA and protein accession identifiers of GRK isoforms in different mammals retrieved from <https://www.ncbi.nlm.nih.gov>. In case of human, mouse, and rat, the computationally predicted sequences reported are not shown. In case of hamster and monkey, all sequences shown are computationally predicted. Of note, some proteins within one species and GRK isoform are identical despite different accession numbers.

human (homo sapiens)			mouse (mus musculus)			rat (rattus norvegicus)			hamster (cricetus griseus)			monkey (chlorocebus sabaeus)						
	mRNA accession	protein accession	# AA	mRNA accession	protein accession	# AA	mRNA accession	protein accession	# AA	mRNA accession	protein accession	# AA	mRNA accession	protein accession	# AA			
huGRK2	NM_001619.5	NP_001610.2	689	muGRK2-1	NM_001290818.1	NP_001277747.1	689	ratGRK2	NM_0127780.2	NP_036908.2	689	hamGRK2-x1	XM_02740884.2	XP_027264485.1	699			
				muGRK2-2	NM_130863.2	NP_570933.1	647						XM_007650600.4	XP_007648790.2	699			
													XM_027408885.2	XP_027264486.1	689			
													XM_003500943.5	XP_003500991.1	689			
												hamGRK2-x2						
huGRK3-1	NM_005160.4	NP_005151.2	688	muGRK3-1	NM_177078.4	NP_796052.2	688	ratGRK3	NM_012897.3	NP_037029.2	688	hamGRK3-x1	XM_007647887.3	XP_007646077.1	401			
	huGRK3-2	NM_001362778.2	NP_001349707.1		575	hamGRK3-x2	XM_007647223.3		XP_007645413.3	245								
				muGRK3-3	NM_001285806.1	NP_001272735.1	581					hamGRK3-x3	XM_027415948.2	XP_027271749.1	688			
huGRK5	NM_005308.3	NP_005299.1	590	muGRK5	NM_018869.3	NP_061357.3	590	ratGRK5	NM_030829.2	NP_110456.2	590	hamGRK5-x1	XM_035436880.1	XP_035292771.1	601			
													hamGRK5-x2	XM_035441693.1	XP_035297584.1	601		
													hamGRK5-x3	XM_027406842.2	XP_027262643.1	590		
													hamGRK5-x4	XM_035441694.1	XP_035297585.1	569		
												hamGRK5-x5	XM_027406846.2	XP_027262647.1	426			
huGRK6-1	NM_001004106.3	NP_001004106.1	576	muGRK6-1	NM_001038018.5	NP_001033107.1	589	ratGRK6-1	NM_031657.4	NP_113845.2	589	hamGRK6-x1	XM_007623236.4	XP_007621426.2	576			
	huGRK6-2	NM_002082.4	NP_002073.2		589	muGRK6-2	NM_011938.5		NP_036068.2	560	ratGRK6-2	NM_001112713.1	NP_00106184.1	560	hamGRK6-x2	XM_007623241.4	XP_007621431.2	581
	huGRK6-3	NM_001004105.3	NP_001004105.1		560	muGRK6-3	NM_001106182.1		NP_00106182.1	576	ratGRK6-3	NM_001112712.2	NP_00106183.1	576	hamGRK6-x3	XM_007623249.4	XP_007621439.2	481
	huGRK6-4	NM_001364164.2	NP_001351093.1		366	muGRK6-4	NM_001286063.2		NP_001272992.1	557				hamGRK6-x4	XM_027409791.2	XP_027265592.1	576	
				muGRK6-5	NM_001286064.2	NP_001272993.1	584				hamGRK6-x5	XM_027409792.2	XP_027265593.1	560				
				muGRK6-6	NM_001286065.1	NP_001272994.1	542				hamGRK6-x6	XM_027409793.2	XP_027265594.1	484				
				muGRK6-7	NM_001286066.2	NP_001272995.1	295				hamGRK6-x7	XM_027409794.2	XP_027265595.1	481				
				muGRK6-8	NM_001377076.1	NP_001364005.1	571				hamGRK6-x8	XM_027409795.2	XP_027265596.1	324				
				muGRK6-9	NM_001377077.1	NP_001364006.1	311											