



Figure S1. ATP5G1(1-67)-RFP overexpression translocates K-RasG12V to mitochondria.

Baby hamster kidney cells co-expressing mGFP-K-RasG12V with mCherry-CAAX, an endomembrane marker or ATP5H1(1-67)-RFP, a mitochondrial marker were maintained in complete growth media (DMEM + 10% FBS + 2mL L-Glut) and fixed with 4% PFA and imaged by a confocal microscope. Scale bar $10\mu m$.



Figure S2. PtdSer supplementation restores the PM localization of K-Ras and LactC2. MDCK cells stably co-expressing ATP5G1(1-67)-RFP with mGFP-LactC2 (A) or -K-RasG12V (B) were supplemented with 10 μ M exogenous PtdSer or PtdCho in growth medium without glucose (DMDM without glucose/pyruvate + 10% FBS + 2mM L-glutamine + 1mM sodium pyruvate) and further incubated for indicated time points. Cells were fixed with 4% PFA and imaged by a confocal microscope. Inserted values represent a mean estimate \pm S.E.M of the fraction of ATP5G1(1-67)-RFP colocalized with mGFP-LactC2 or -K-RasG12V calculated by Manders coefficient from three

independent experiments. Shown are representative images of mGFP-LactC2 and -K-RasG12V. Scale bar 10 μ m. (C)MDCK cells stably expressing mGFP-K-RasG12V or -LactC2 were infected with lentivirus expressing PtdSer decarboxylase (PISD) and maintained in complete growth medium (DMEM + 10% FBS + 2mM L-glutamine). Cells were fixed with 4% PFA and imaged by a confocal microscope. Representative images are shown from three independent experiments. Scale bars 10 μ m.



Figure S3. PI4K inhibitors reduce PI4P contents at the Golgi. MDCK cells stably expressing mGFP-tagged PI4P markers, FAPP1-PH or P4M-SidM were treated with 10 μ M PIK-93 or 50nM phenylarsine oxide for 48h in complete growth medium (DMEM + 10% FBS + 2mM L-glut). Cells were fixed with 4% PFA and imaged by a confocal microscope. Scale bar 10 μ m.



Figure S4. The Golgi-Pl4K inhibition mislocalizes K-RasG12V and PtdSer to mitochondria and endomembrane, respectively. MDCK cells stably expressing (A) mGFP-K-RasG12V or (B) -LactC2 were treated with vehicle (DMSO), 1μ M PIK-93 or 50nM phenylarsine oxide for 48h, and further stained with 100nM MitoTracker Deep Red for 1h. Cells were fixed with 4% PFA and

Figure S4

imaged by a confocal microscope. Inserted values are a mean estimate \pm S.E.M of the fraction of MitoTracker colocalized with mGFP-K-RasG12V or -LactC2 that were calculated by Manders coefficient from three independent experiments. Selected regions indicated by the white squares are shown at a higher magnification. mGFP-K-RasG12V and -LactC2 colocalized or not colocalized with MitoTracker are indicated by closed and open arrowheads, respectilvey. Scale bar 10 $\mu m.$

Figure S5

A. cDNA sequence

Empty ATGGGAGACATGGTGGTGGAACCTGCCCCCTGAAGCCAACTTCTGAGCCCACTACTGGCCTGCCAGGGA sgRNA2 ATGGGAGACATGGTGGTGGAACCTGCCCCCTGAAGCCAACTTCTGAGCCCACTACTGGCCTGCCAGGGA sgRNA6 ATGGGAGACATGGTGGTGGAACCTGCCCCCTGAAGCCAACTTCTGAGCCCACTACTGGCCTGCCAGGGA Empty ATAATGGGGGGTTCCTTGCTAGGCGTCATCA-CAGAGGGGGGTTGGGGAACTGTCAGTGATTGACCGTGAGG sgRNA2 ATAATGGGGGTTCCTTGCTAGGCGTCATCAACAGAGGGGGGTTGGGGAACTGTCAGTGATTGACCGTGAGG sgRBA6 ATAATGGGGGTTCCTTGCTAGGCGTCATCA-CAGAGGGGGTTGGGGAACTGTCAGTGATTGACCGTGAGG Empty TGGCCCAGAAGGCCTGCCAGGAGGTGCTGGAGCAAGTCAAGCTTTTGCATGGAGGCGTGGCCATATCTAG sqRNA2 TGGCCCANAAGGCCTGCCAGGAGGTGCTGGAGCAAGTCAAGCTTTTGCATGGAGGCGTGGCCATATCTAG sgRBA6 TGGCCCAGAAGGCCTGCCAGGAGGTGCTGGAGCAAGTCAAGCTTTTGCATGGAGGCGTGGCCATATCTAG Empty CACAGACACCTCCCTGGAGCTGGTCAATGGGGATGTTCCGGACAGTGCTATCCGTTG-CCTGGATGATCCA sqRNA2 CACAGACACCTCCCTGGAGCTGGTCAATGGGGATGTTCCGGACAGTGCTATCCGTTG-CCTGGATGATCCA sgRBA6 CACAGACACCTCCCTGGAGCTGGTCAATGGGGGATGTTCCGGACAGTGCTATCCGTTGCCCTGGATGATCCA Empty CCTACCCAGATAAGGGAGGAGGAAGATGAAATGGGGGGCCACTGTGGCCTCGGGCACAGCCAAGGGAGCAAG sgRNA2 CCTACCCAGATAAGGGAGGAGGAAGATGAAATGGGGGCCACTGTGGCCTCGGGCACAGCCAAGGGAGCAAG sgRNA6 CCTACCCAGATAAGGGAGGAGGAAGATGAAATGGGGGCCACTGTGGCCTCGGGCACAGCCAAGGGAGCAAG B. Protein sequence Empty MGDMVVEPAPLKPTSEPTTGLPGNNGGSLLGVITEGVGELSVIDREVAQKACQEVLEQVKLLHGGVAISS sqRNA2 MGDMVVEPAPLKPTSEPTTGLPGNNGGSLLGVINRGGWGTVSDsgRNA6 MGDMVVEPAPLKPTSEPTTGLPGNNGGSLLGVITEGVGELSVIDREVAQKACQEVLEQVKLLHGGVAISS

Empty TDTSLELVNGDVPDSAIRCLDDPPTQIREEEDEMGATVASGTAKGARRRQNNSAKQSWLLRLFESKLFD sgRNA2

sgRNA6 TDTSLELVNGDVPDSAIRCPG-

Figure S5. CRISPR/Cas9-mediated sgRNAs targeting canine PI4KB generates a frameshift mutation, resulting in premature stop codon. Nucleotides in red are inserted nucleotide induced by CRISPR/Cas9 system.