

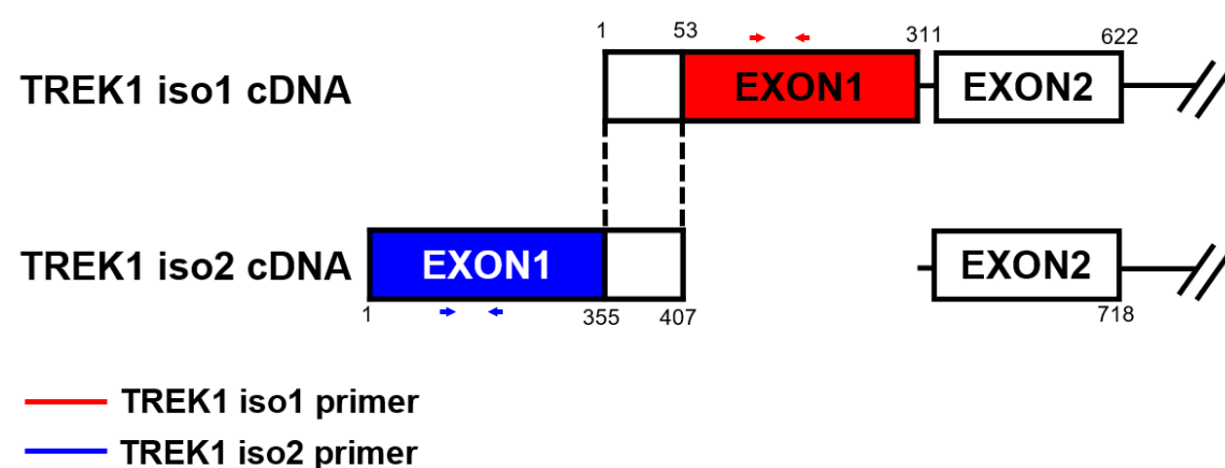
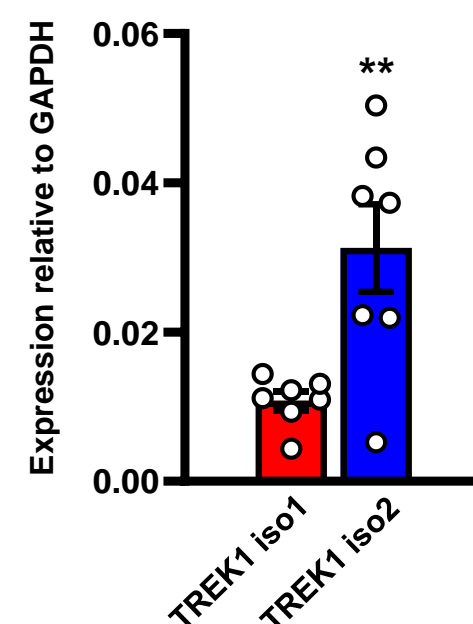
A**B**

Figure S1. Relative expression of TREK1 isoforms in Astrocytes. (A) Schematic diagram of the cDNA for each TREK1 isoform. Primers for qRT-PCR, indicated by small arrows, were designed in exon1 of each isoform. (B) Bar graphs are qRT-PCR results using each primer. Normalized with GAPDH(n=7). All values are mean \pm SEM (** $p < 0.01$). qRT-PCR, quantitative reverse transcription polymerase chain reaction; SEM, standard error of the mean; TWIK1, tandem of pore domains in a weak inward rectifying K⁺ channel; TREK1, TWIK-related K⁺ channel 1; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

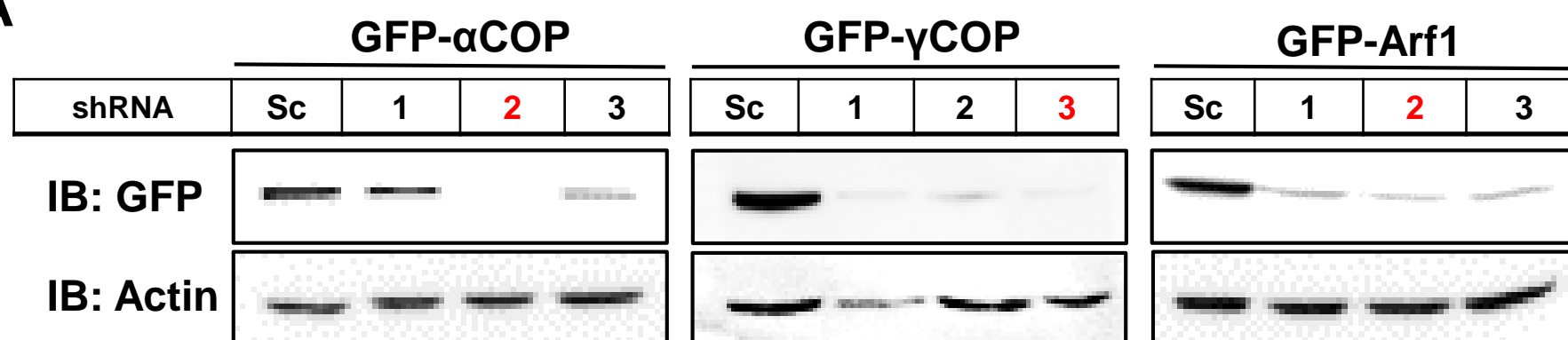
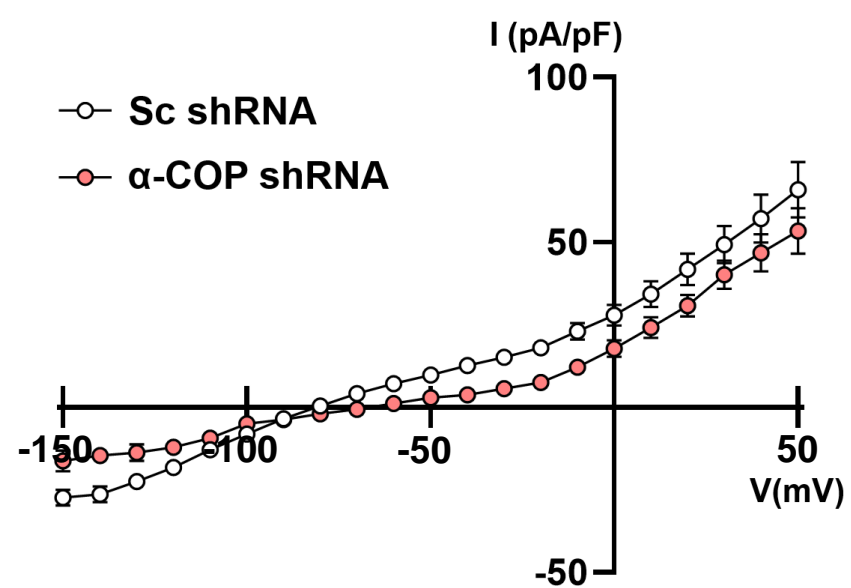
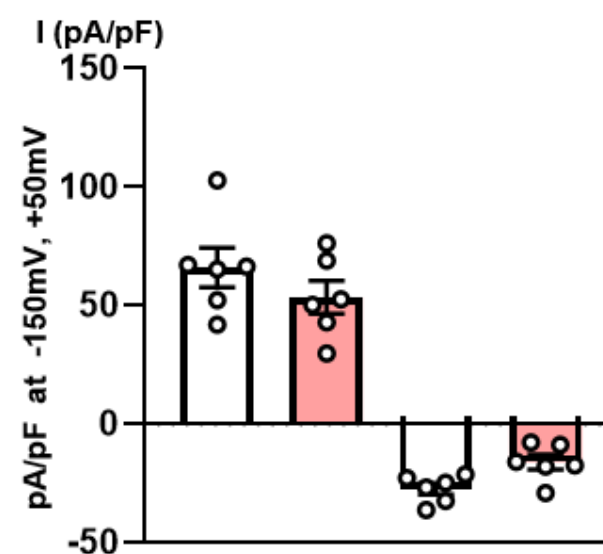
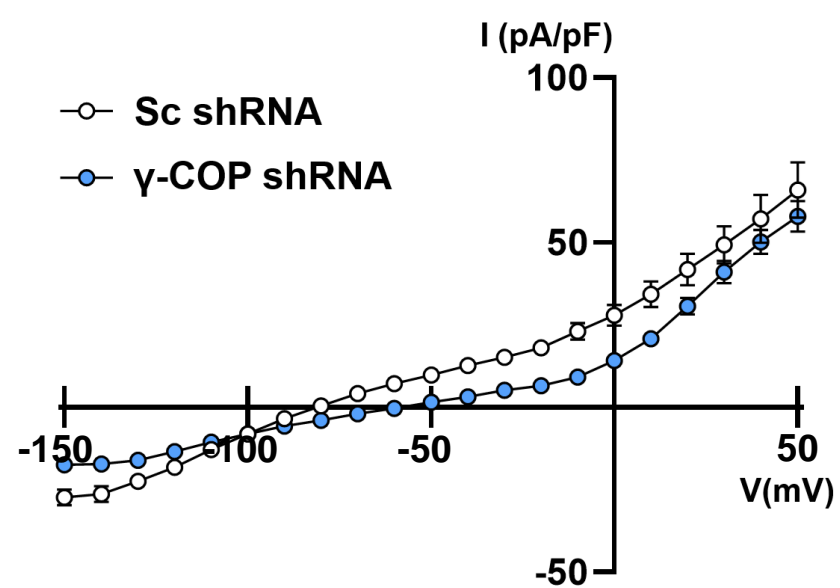
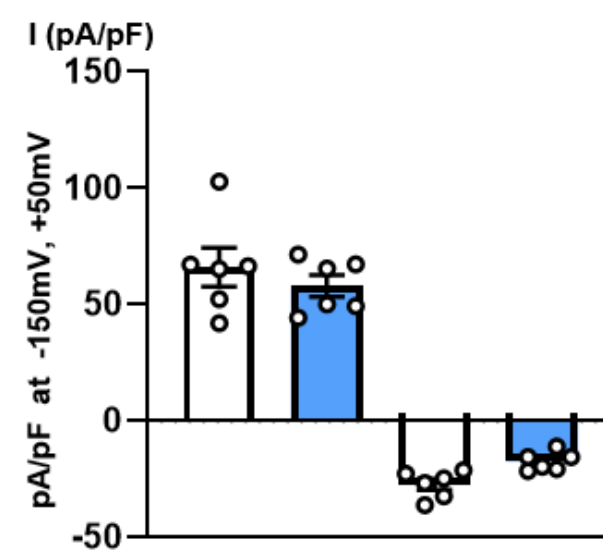
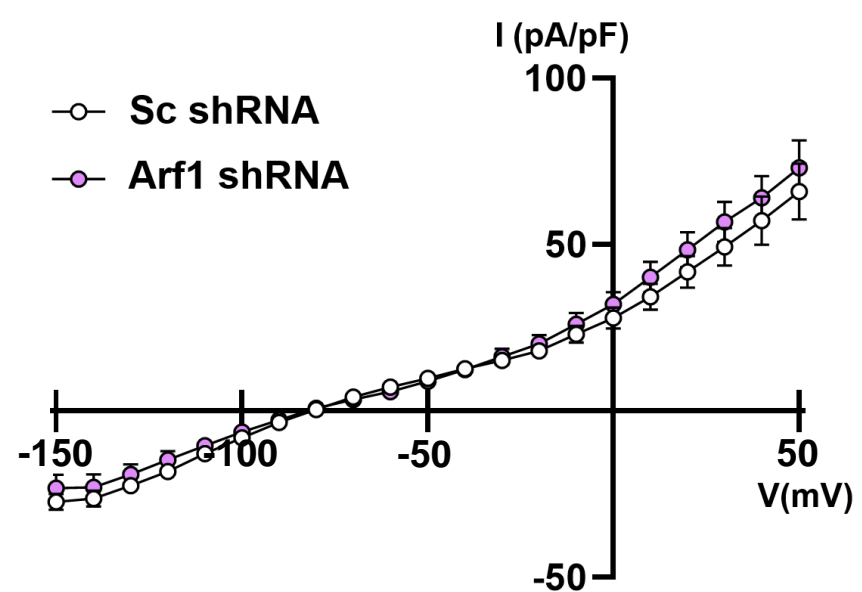
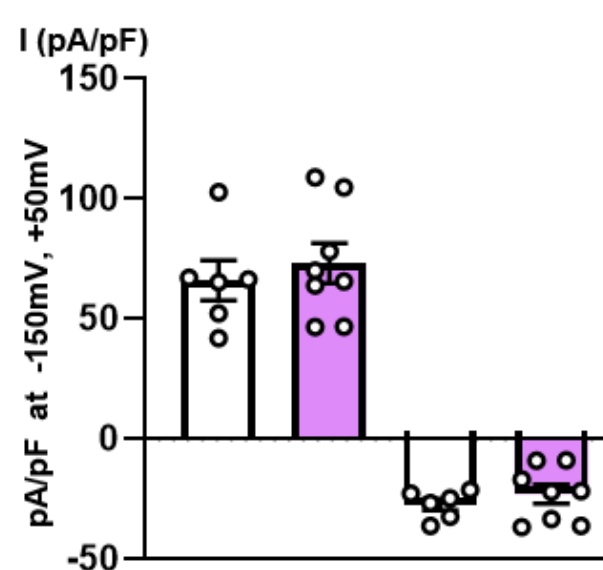
A**B****C****D****E****F****G**

Figure S2. Other COP1 subunits do not affect the K⁺ current in astrocytes. (A) α -COP, γ -COP, Arf1 shRNA validation test in HEK293T cells. GFP-tagged α -COP, γ -COP, and Arf1 were co-transfected with each of three specific shRNAs. The most efficient shRNA (red) was used in the experiment. (B), (D), (F) Average current-voltage (I-V) relationship of Sc shRNA (white), α -COP shRNA (light orange), γ -COP shRNA (light sky), and Arf1 shRNA (light violet) transfected astrocytes. The voltage ramp-induced whole-cell current traces were recorded from each cell. (C), (E), (G) Bar graph at -150 mV and +50 mV in (B), (D), (F). All values are mean \pm SEM (***) $p < 0.001$. SEM, standard error of the mean; COP, coat protein; COP1, coat protein complex 1; Arf1, ADP-ribosylation factor 1; TWIK1, tandem of pore domains in a weak inward rectifying K⁺ channel; TREK1, TWIK-related K⁺ channel 1; GFP, green fluorescent protein; Co-IP, co-immunoprecipitation; Sc, scrambled; HEK, human embryonic kidney.