Supplementary information, Figure S3

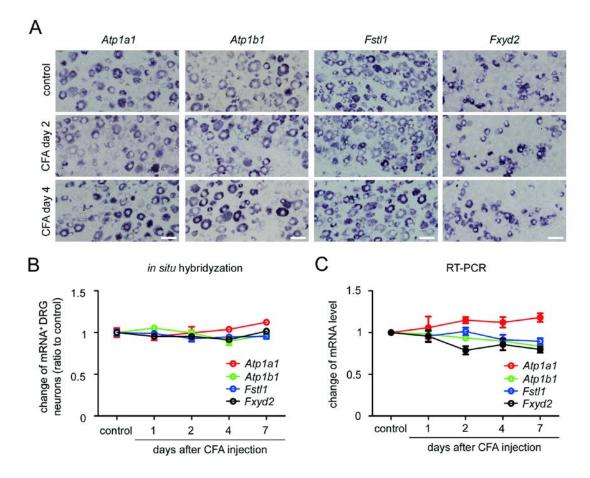


Figure S3 Expression of α1 and β1 subunits of NKA, FSTL1 and FXYD2 in DRGs after peripheral inflammation. (A, B) Representative images and quantitative data for *in situ* hybridization showed that the cellular distribution and levels of mRNA encoding the α1 subunit (*Atp1a1*) and β1 subunit (*Atp1b1*) of NKA, FSTL1 (*Fstl1*) and FXYD2 (*Fxyd2*) were not obviously changed in the DRGs after peripheral inflammation (n = 3). Scale bar, 50 µm. (C) Real-time PCR showed that the mRNA levels of the α1 and β1 subunits of NKA, FSTL1 and FXYD2 in the DRGs were not significantly altered after peripheral inflammation (n = 3).