



Supplementary figure 1 *KCNK18* transgenic mouse models. **a** Scheme of $K_{2p}18.1$ function disrupted in full-knockout ($Kcnk18^{-/-}$) mice. Calcineurin binds to the NFAT-like PQIVID motif in the intracellular loop of $K_{2p}18.1$ and activates the channel by dephosphorylation of S-264 and the S-276 cluster. Phosphorylation of S-264 by PKA (protein kinase A) and S-276 by MARK (mammalian MAP/microtubule affinity-regulating kinase) restores the resting (inhibited) state of $K_{2p}18.1$. Docking of 14-3-3 adaptor protein to phosphorylated S-264 strengthens inhibition. **b** Scheme of G339R point mutation with abolished ion current through $K_{2p}18.1$ but intact intracellular signalling. **c** Scheme of S276A point mutation with facilitated ion current through $K_{2p}18.1$. **d** Scheme of T cell receptor (TCR) and IL-2 dependent intracellular signalling cascades leading to FoxP3 expression. **e** Scheme of $K_{2p}18.1$ -mediated t_{reg} development. T cell receptor (TCR) activation by antigens presented by thymic antigen-presenting cells (tAPC) drives fate decisions of multipotent CD4 single-positive thymocytes (CD4-SP) (step 1). TCR activation induces NF- κ B signaling and thereby $K_{2p}18.1$ upregulation in t_{reg} progenitors (t_{reg}^P) (step 2). $K_{2p}18.1$ -mediated K⁺ efflux hyperpolarizes the membrane potential (V_m) providing the driving force for prolonged high intracellular Ca²⁺ levels, which in turn activate $K_{2p}18.1$ -mediated ion-conductance and NF- κ B signaling forming a positive feedback loop. High intracellular Ca²⁺ levels further promote the translocation of NFAT and NF- κ B related transcription factors into the nucleus (step 3). Those transcription factors induce and stabilize FoxP3 expression and thereby the development of mature t_{reg} (step 4).