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SUPPLEMENTARY DATA

Oligomerization and Ca^{2+} /calmodulin control binding of the ER Ca^{2+} -sensors STIM1 and STIM2 to plasma membrane lipids

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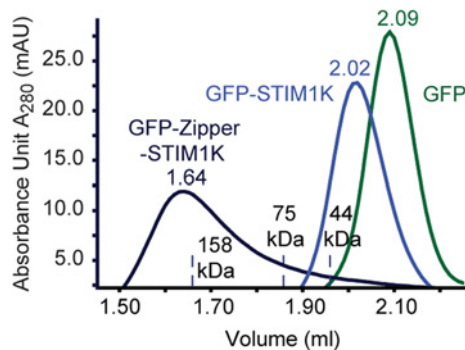


Figure S1 Addition of a leucine zipper to STIM1 K-rich domain leads to tetramerization

Elution profiles of 5 μM GFP (green), GFP-STIM1K (blue) and a GFP-tagged construct with yeast GCN4 leucine zipper fused to STIM1 K-rich domain (GFP-Zipper-STIM1K, black) S200 gel-filtration column run are shown. Positions where markers eluted are indicated.

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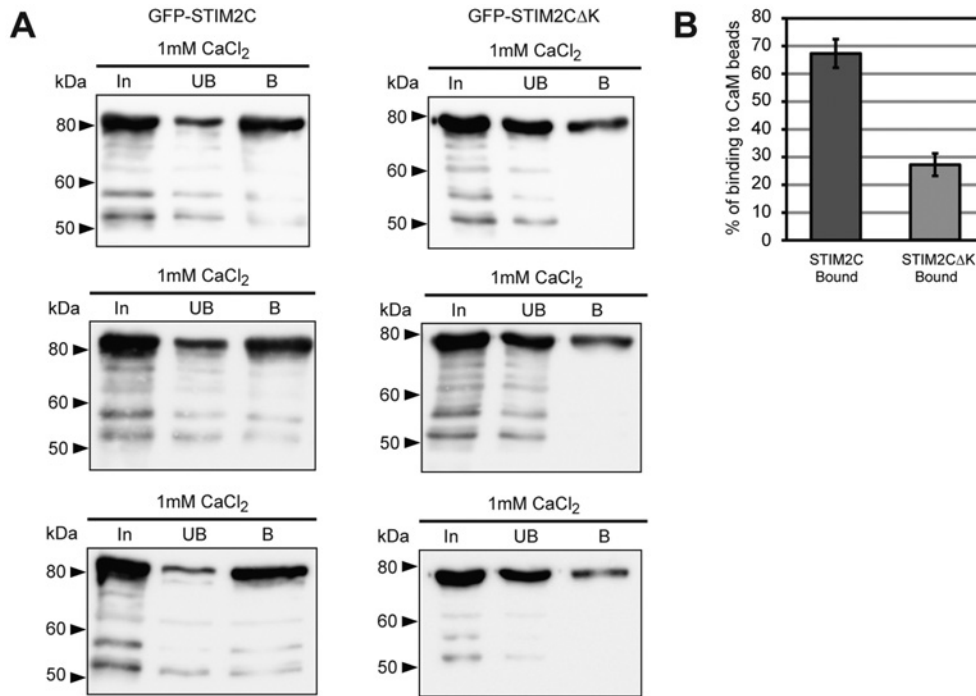


Figure S2 Western blot quantification of GFP-STIM2C and GFP-STIM2CΔK binding to Ca²⁺/calmodulin beads
(A) 1 μM reduced GFP-STIM2C and GFP-STIM2CΔK were incubated with Ca²⁺/CaM beads in presence of 1 mM CaCl₂. Input (In), unbound (UB) and bound (B) material of three separate experiments were detected by a rabbit polyclonal GFP antibody from Santa Cruz Biotechnology, Inc. (1:5000). (B) Quantification of bound GFP-STIM2C and GFP-STIM2CΔK from three experiments. Bars indicate means ± S.D.

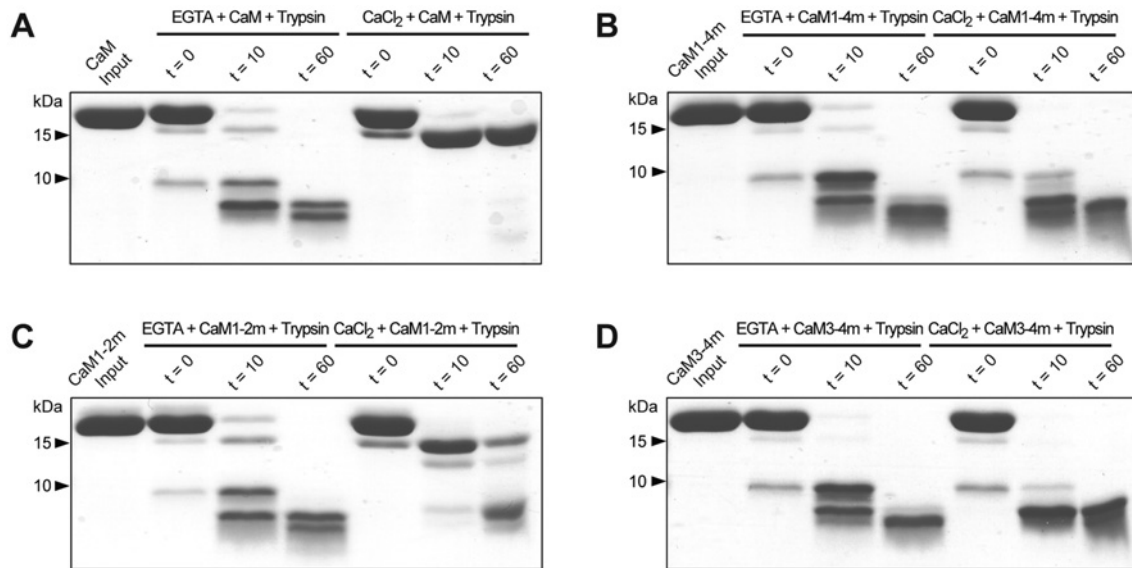


Figure S3 Protease protection assays with purified CaM and CaM mutants

14 μ M CaM (A), CaM1-4m (B), CaM1-2m (C) and CaM3-4m (D) in 1 mM EGTA or 0.2 mM CaCl₂ were incubated with 0.02 μ M trypsin in 20 mM ammonium bicarbonate buffer (pH 8.1) at 20 °C for the indicated times. The proteins were separated by SDS/PAGE and stained with Coomassie Brilliant Blue. The first lane of each gel shows the input of CaM or Ca²⁺-binding mutants of CaM without protease.

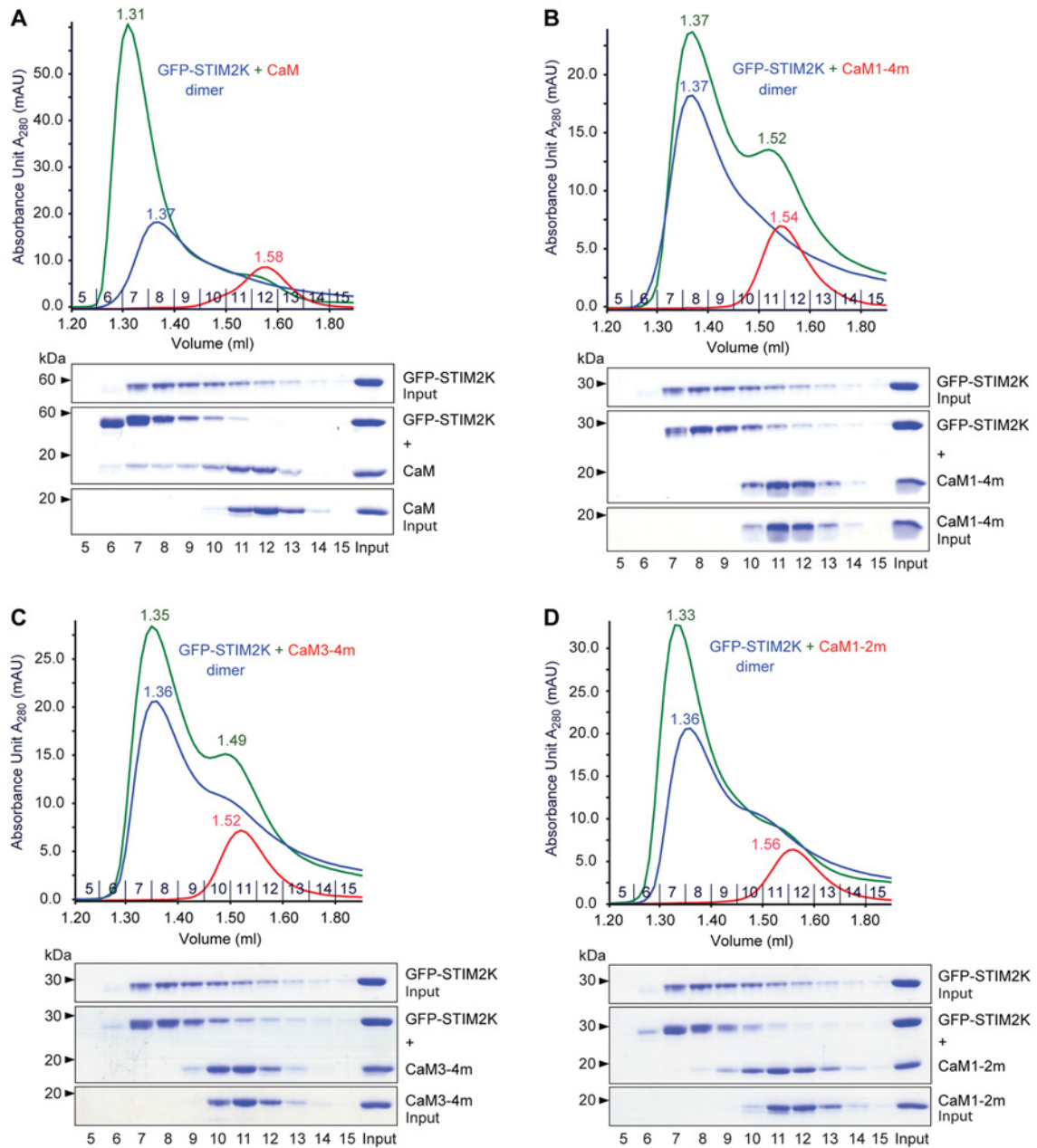


Figure S4 In solution binding of GFP-STIM2K dimer and calmodulin
10 μ M oxidized GFP-STIM2K dimer and 10 μ M purified CaM (A), CaM1-4m (B), CaM3-4m (C) or CaM1-2m (D) in 1 mM CaCl₂ were separated by gel filtration. The elution profiles of GFP-STIM2K dimer (blue), CaM (red) and complexes (green) with indicated volume for elution peaks are shown. Proteins of the indicated eluted fractions separated by SDS/PAGE and Coomassie Brilliant Blue stained are shown below the chromatograms.

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