

Supplementary Material

Fig. S1 Hydrophobicity/hydrophilicity and signal peptide prediction of the InaQ domains using bioinformatics tools. (A) Hydrophobicity analysis of the InaQ-N domain. (B) Hydrophilicity analysis of the InaQ-C domain. (C) Signal peptide prediction of InaQ-N. C-score is the “cleavage site” score that should only be significantly high at the cleavage site. S-score is the value of every single amino acid position. Y-score is a derivative of the C-score combined with the S-score.

