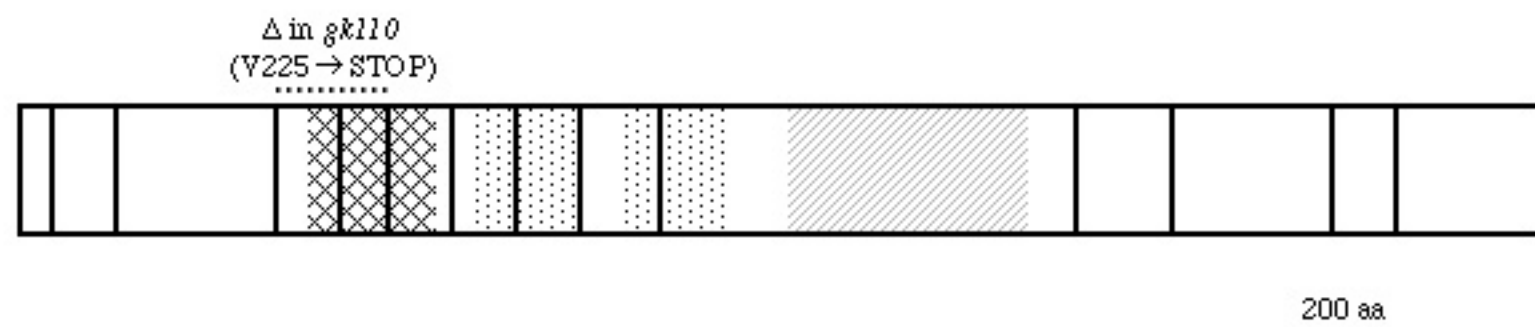


Supplementary Figure 1



Supplementary Figure 1. Structure of DGK-3. Vertical lines indicate exon-intron boundaries. Residues deleted in *gk110* resulting in a protein truncated at V225 are indicated by a dashed overline. Functional domains of DGK-3 as predicted by homology are indicated by patterned boxes. Crosshatched box - residues encoding the predicted calcium-binding EF hands; stippled boxes - cysteine-rich domains predicted to bind DAG and phorbol ester; hatched box - catalytic domain.