Figure S1. S4 sequences comparison and concatemerization. (A) Comparison of the amino acids of the S4 segment in Shaker, Neu, and Shaker ILT. A dash means that the amino acid is identical to that of Shaker sequence. The four arginine residues are numbered above the Shaker sequence. The filled circles on the left represent the symbol used to represent the cDNA concatemers constructs. (B) Construction of concatemers. The diagram shows each protomer in a different color and the amino acid sequence of the single amino acid linker. The adjacent amino acids are shown and have the color of the protomer where they belong.