

Supplementary Table and Figure

Table S1. Structural statistics for the rSsp1a ensemble

Experimental restraints

Interproton distance restraints	395
Intraresidue	106
Sequential	126
Medium range (i-j < 5)	57
Long range (i-j ≥ 5)	106
Dihedral-angle restraints	32
Hydrogen bonds	16

R.m.s. deviations from mean coordinate structure (Å)

Backbone atoms (residue 1-17)	0.76 ± 0.27
All heavy atoms (residue 1-17)	1.38 ± 0.22

Ramachandran Statistics

% in most favoured region	79%
% in additionally allowed region	21%

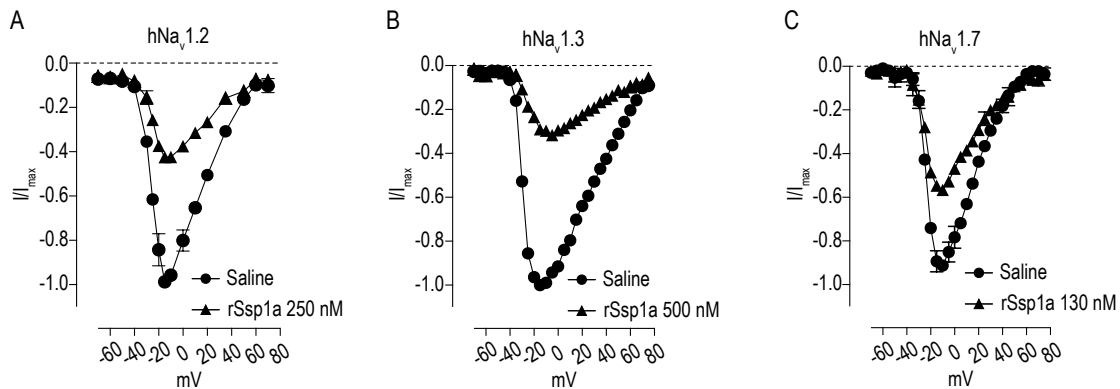


Figure S1. Effect of rSsp1a on current-voltage relationship of hNav1.2, hNav1.4 and hNav1.7. The ~IC₅₀ rSsp1a reduced the peak current without significantly affecting the activation threshold, activation voltage and reversal potential. Each data point in the curve is mean \pm SEM with $n = 3-4$.