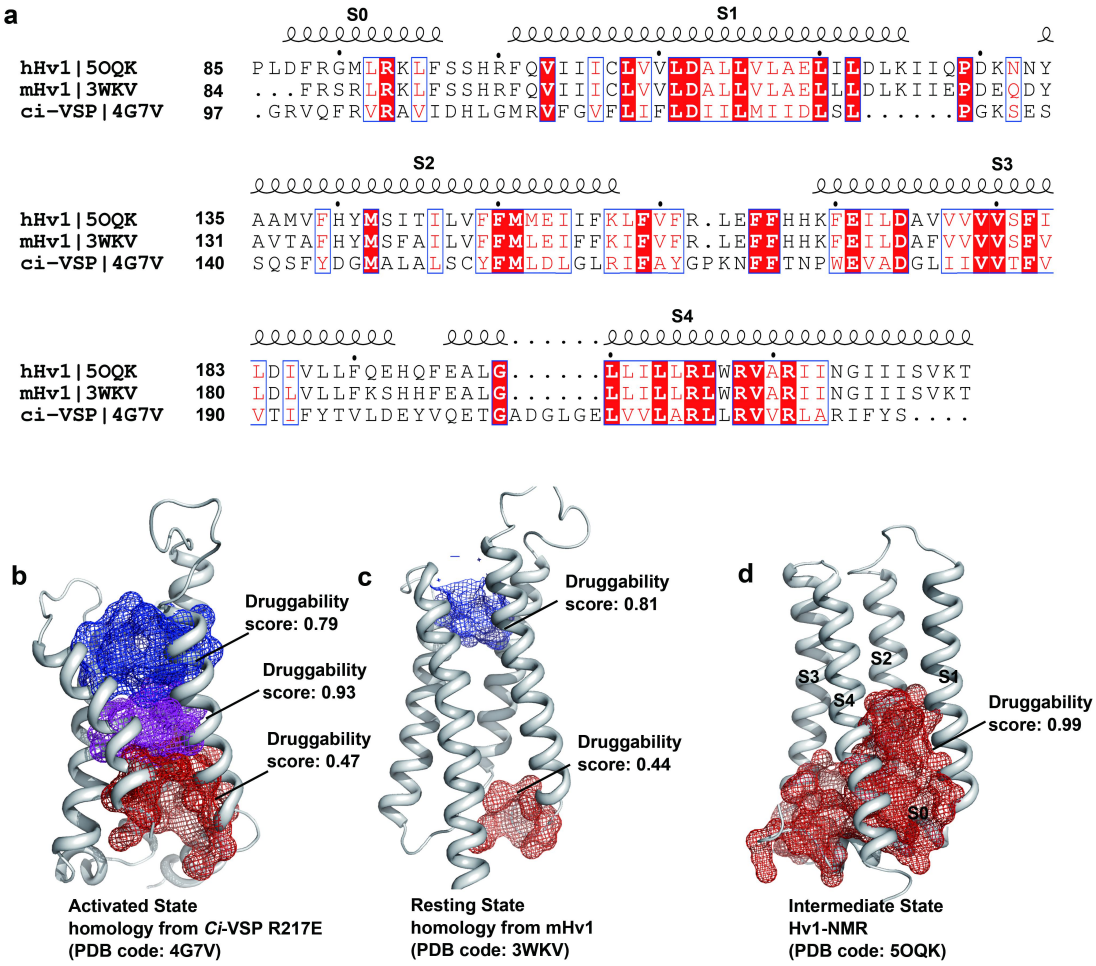

Fig. S4. Multiple alignments and druggable pocket prediction of Hv1 channels. a Alignments among hHv1, mHv1 and ci-VSP used to guide homology modelling. **b-d** The potential druggable pockets identified in activated state (**b**), resting state (**c**) and intermediate state (**d**) of Hv1. The pockets were exhibited in colored mesh. Druggability scores of the pockets were labeled.