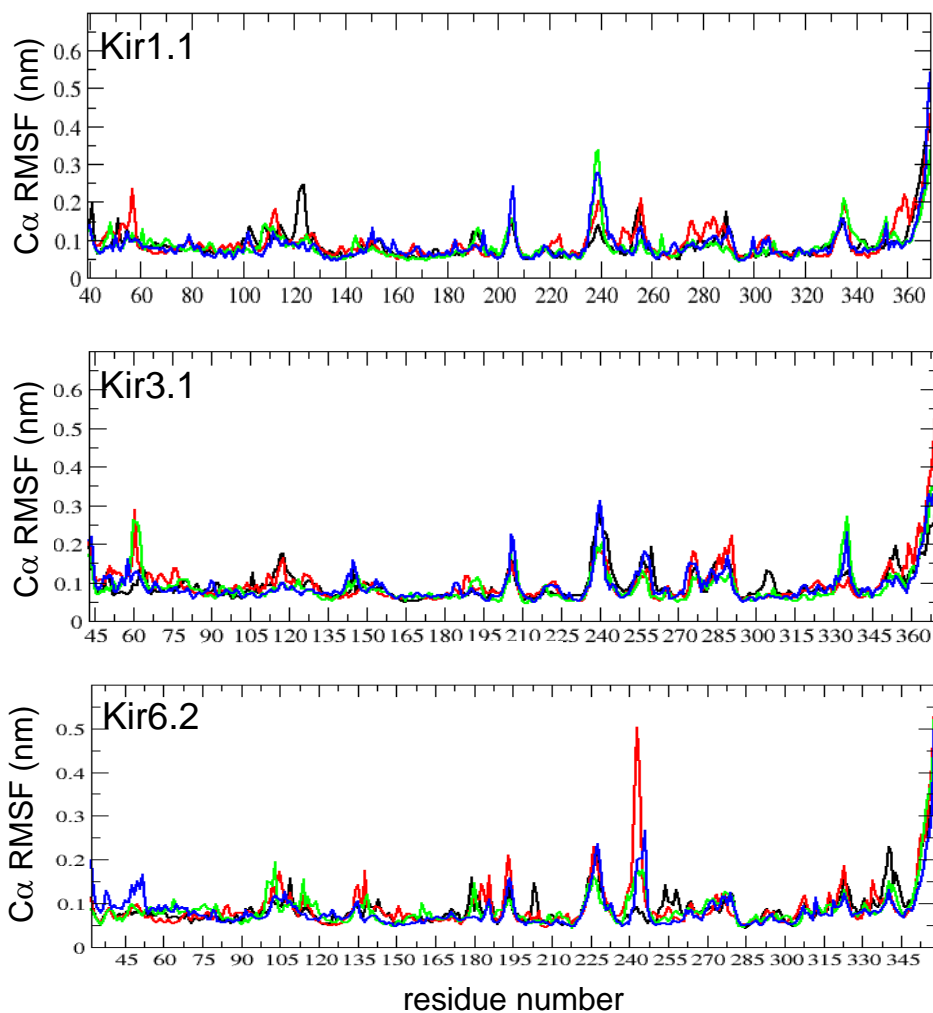


## Supplementary Material

### Molecular Dynamics Simulations of Inwardly Rectifying (Kir) Potassium Channels: A Comparative Study

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*Figure S1*

C $\alpha$  atom root mean square fluctuation (RMSF) as a function of residue number for the three simulations. In each case, the four differently coloured lines correspond to the four subunits of the channel protein.