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Cover picture: Ramachandran plot, a plot of the torsional angles—phi ($\phi$) and psi ($\psi$)—of the residues contained in a protein, was used to evaluate the reliability of simulated TRPM2 NUDT9-H structure by homology modeling, which provided supporting evidence for Yu’s work in identification of ADPR binding sites in the TRPM2 channel (see Research Article by Yu et al., 219–235).