

人甲状腺旁素 1 型受体的结构特征

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Structural Characterization of Human Parathyroid Hormone 1 Receptor

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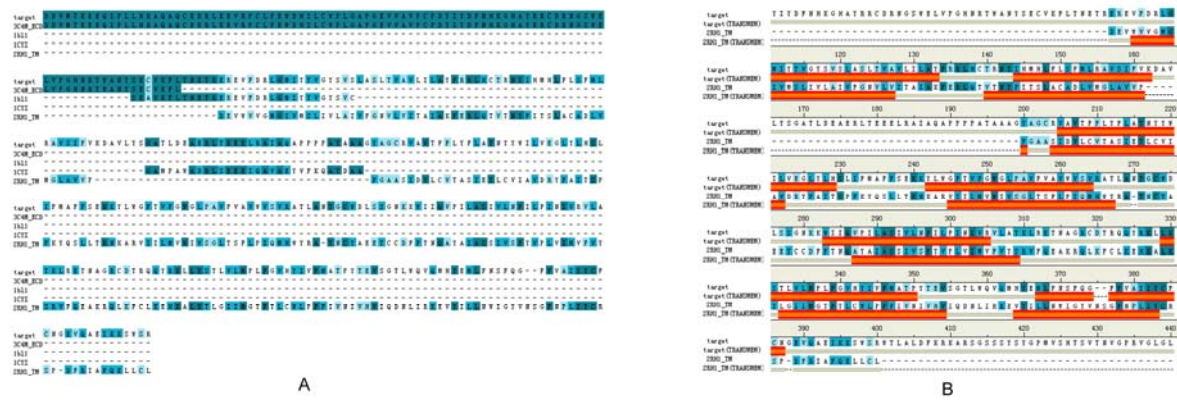


Fig.S1 Alignment of the final model with all templates (A) and the result of seven TMs predicted by DS2.5 (B)

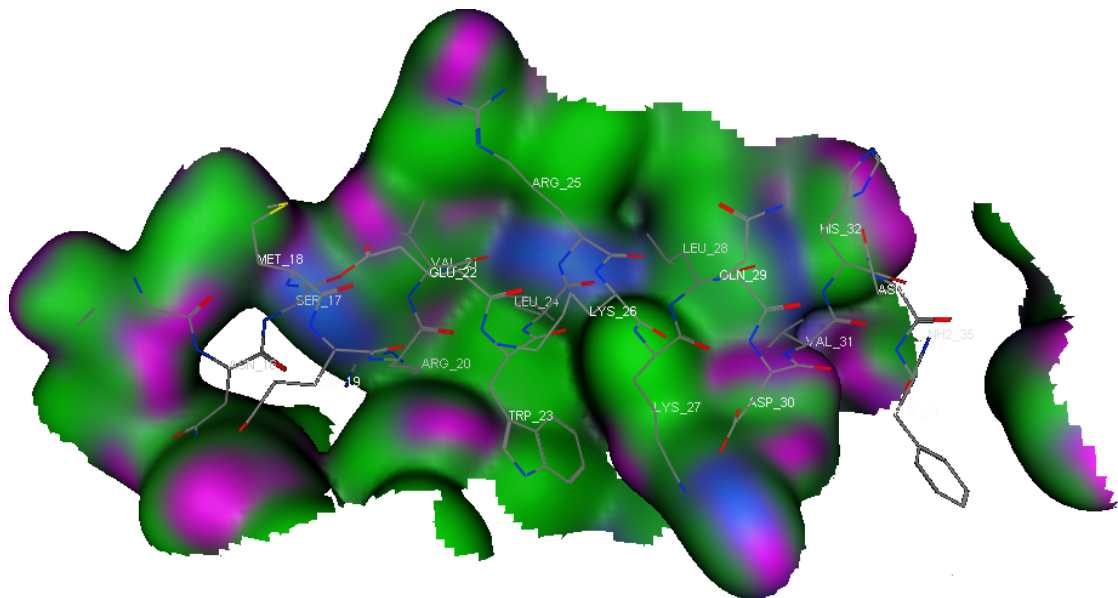


Fig.S2 The interaction diagram of PTH (15-34) binding to PTH1R. Hydrophobic regions are in green, while mild polar and H-bonding regions in blue and purple respectively.

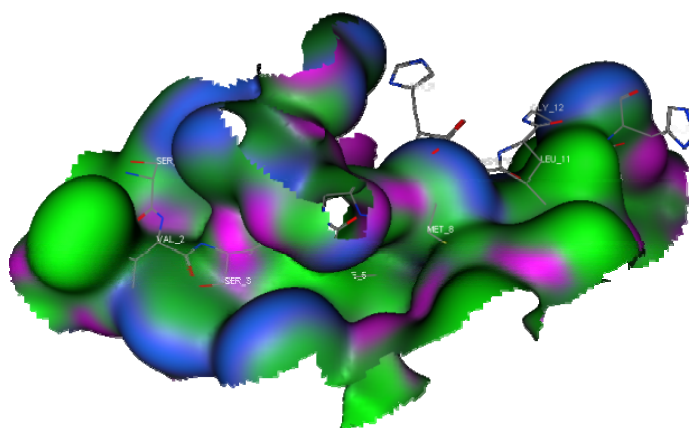


Fig.S3 The hydrophobic interaction diagram of PTH (1-14) binding to PTH1R. Hydrophobic regions are in green, while mild polar and H-bonding regions in blue and purple respectively.