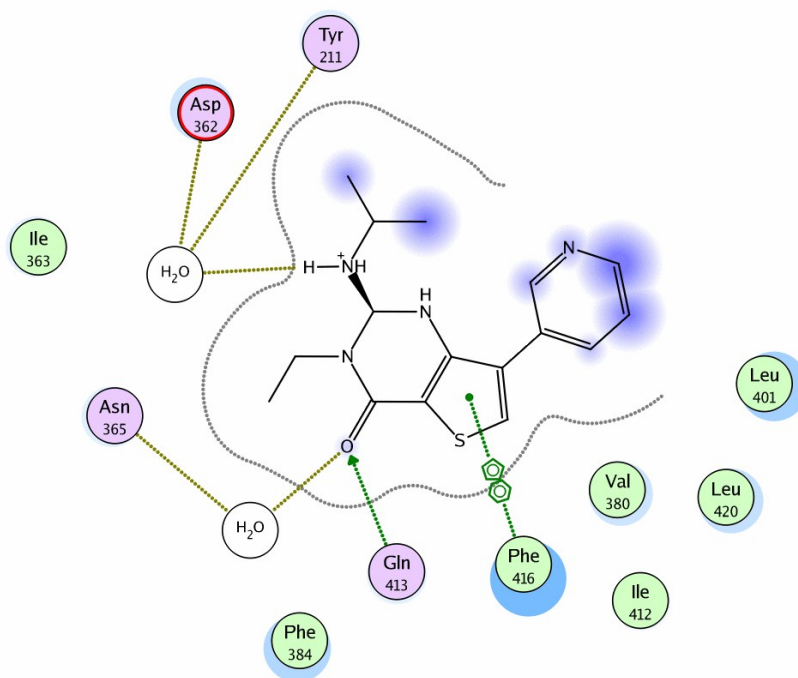**Exhaustive 3D-QSAR analyses as a computational tool to explore the potency and selectivity profiles of thieno[3,2-*d*]pyrimidin-4(3*H*)-one derivatives as PDE7 inhibitors**

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SUPPORTING INFORMATION

S1. Ligplot representation of the binding mode identified at the compound **56**-PDE7A X-ray crystallographic complex.