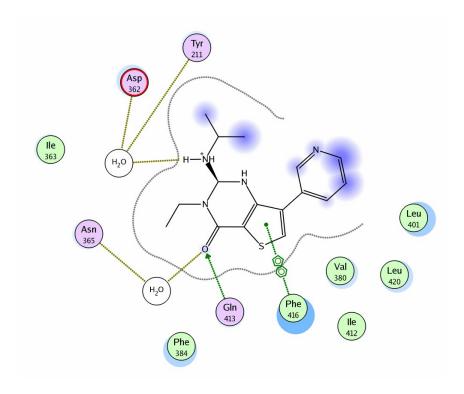
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Exhaustive 3D-QSAR analyses as a computational tool to explore the potency and selectivity profiles of thieno[3,2-d]pyrimidin-4(3H)-one derivatives as PDE7 inhibitors

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SUPPORTING INFORMATION



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