The Identification and synthesis of novel allosteric inhibitors Posters

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Introduction

HIV (human immunodeficiency virus) is a lentivirus which can lead to acquired immunodeficiency virus (AIDS). In the last 25 years the number of new infections has reduced from 3.5 million to 2.1 million¹ . The magnitude of this improvement is partly a consequence of the highly active antiretroviral therapy (HAART), which targets the activities of the viral reverse transcriptase, integrase and protease enzymes that are critical for the viral replication². However there is a need for new treatments as drug resistant strains are becoming more common². Retroviral integrase (IN) is an enzyme responsible for the insertion of the viral DNA into the host chromosomal DNA, an essential step for HIV replication^{2,3}. Inhibitors currently exist which target the active site of the viral-encoded integrase in the early phase ³. This project however focuses on the new approach which targets the interaction between integrase and the cellular host factor LEDGF/p75.

Methods

The crystal structure of IN was used with the target protein LEDGF/P75 which was extracted from Protein Data Base

Autodock was used to generate a computer model of IN with LEDGF/P75 and ran this model against a pool of zinc based compounds using Rocket HPC

Hits were generated and filtered based on binding energy to produce lead compounds

The lead compounds were refined by screening through ADMET to choose the most appropriate physiochemical parameters

The most suitable compound (N-(4-chlorophenyl)4phenoxybenzenesulfonamide was chosen and synthesized in the laboratory

Overview of in silico screening

Results

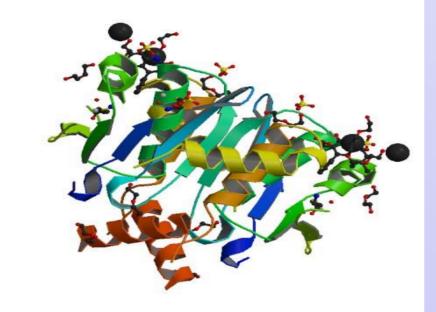


Figure 1. Interaction between LEDGF/p75 and IN⁴

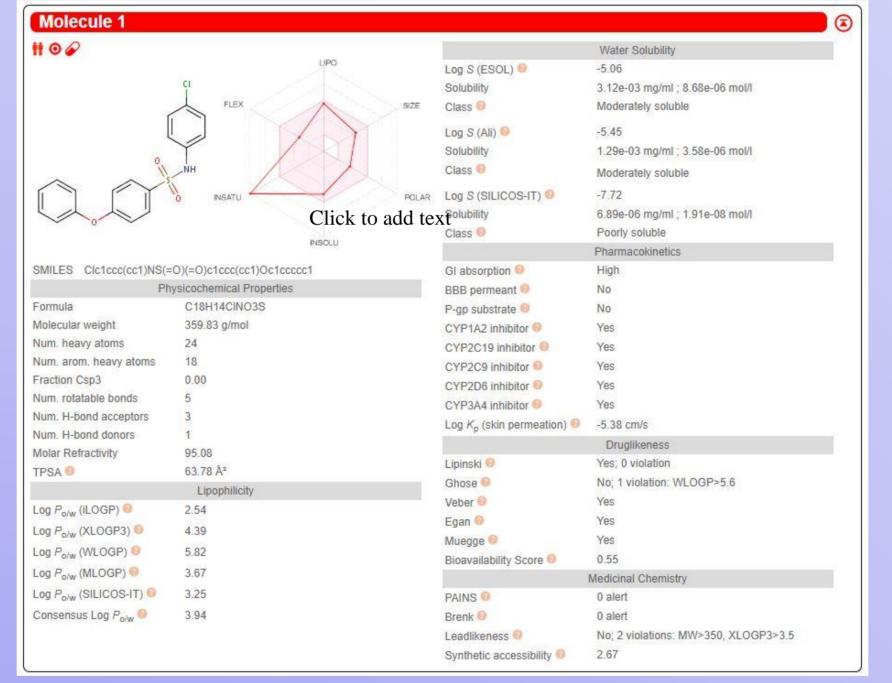


Figure 2. Predicted properties of the chosen drug⁵

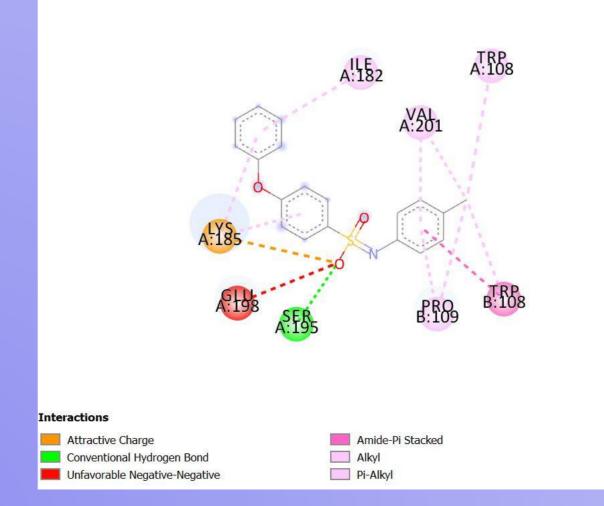


Figure 3. Binding properties of the drug

The chosen compound was found to be a suitable inhibitor using ADMET (figure 2, 3), however the compound inhibits CYP3A4 which is a key enzyme involved in the metabolism of various drugs (figure 2). Therefore it could lead to drug-drug interactions in patients. The compound is also predicted to have a good GI absorption hence would be a suitable candidate for oral administration. In addition, the compound is predicted not to penetrate the blood brain barrier therefore eliminating the associated side effects.

The chosen compound has proven to be an effective inhibitor of LEDGF/P75 and could potentially be developed. Due to a time constraint, we were unable to complete the synthesis and in vivo testing of the proposed compound. The drug will also need to be tested with live tissue to ensure efficacy and safety before it can be further developed.

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This research made use of the Rocket High Performance Computing service at Newcastle University



Discussion

Conclusion

References

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3. Gupta K, Brady T, Dyer B, Malani N, Hwang Y, Male F et al. Allosteric Inhibition of Human Immunodeficiency Virus Integrase. Journal of Biological Chemistry. 2014;289(30):20477-20488.

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