Introduction to HIV Integrase Inhibitors for SAMPL

Background:

HIV integrase is one of the viral targets for which small molecule therapeutics have now been approved to treat AIDS. Integrase is a critical enzyme in the HIV life cycle as it is required for the integration of viral DNA into the host chromatin and performs both the 3' viral processing step as well as the strand transfer step to insert viral genome into the host DNA. The structure of HIV integrase has been thoroughly investigated [Jenkins, Godgur, Chen, Cherepanov 2005, Wang], and consists of three domains (N-terminal DNA binding, catalytic core (CCD) and C-terminal DNA binding). The CCD has several pockets to which small molecules have been shown to bind and inhibit the enzymatic activity [Christ, Wielens, Rhodes, Schrijvers, Kessl]. Integrase forms a complex with viral DNA and several host cellular factors that has been termed the pre-integration complex (PIC) [Cherepanov 2003]. One component of this complex is lens epithelium derived growth factor (LEDGF/p75), which has a conserved integrase binding domain (IBD, residues 347-429) that mediates binding [Cherepanov 2005]. Within the IBD is a loop (residues 362 to 369) that binds in a pocket formed by a dimer of the HIV integrase CCD. Small peptides composed of the residues in this IBD loop can bind to the integrase CCD with micromolar affinity and can compete with the IBD for this binding site [Tsiang 2009]. Several crystal structures with various peptide sequences have been solved showing the interactions between these peptides and integrase [Rhodes 2011]. Several groups have also found small molecule inhibitors of HIV integrase that bind in the LEDGF pocket [Schrijvers, Christ, De Luca, Fan, De Luca, Tsantrizos, Peat].

The challenge in this case is based on the intermediate level compounds that were developed to bind to HIV integrase during the med chem program conducted by CSIRO and Avexa Ltd. Various aspects of this program have been published previously (Peat PONE 2012, Wielens J.Bio.Screen. 2013, Rhodes ChemBiochem 2011, Rhodes AVCC 2011) but the compounds selected for this challenge are not incorporated in these previous publications. The compounds were developed from a fragment screen run against the Maybridge fragment library and compounds were found in multiple sites on HIV integrase noted in the references above, including the LEDGF pocket, an allosteric site, and a so-called fragment site.

References:

Jenkins TM, Hickman AB, Dyda F, Ghirlando R, Davies DR, et al. (1995) Catalytic domain of human immunodeficiency virus type 1 integrase: identification of a soluble mutant by systematic replacement of hydrophobic residues. Proc Natl Acad Sci USA 92: 6057–6061.

Goldgur Y, Dyda F, Hickman AB, Jenkins TM, Craigie R, et al. (1998) Three new structures of the core domain of HIV-1 integrase: an active site that binds magnesium. Proc Natl Acad Sci USA 95: 9150–9154.

Chen JC-H, Krucinski J, Miercke LJW, Finer-Moore JS, Tang AH, et al. (2000) Crystal structure of the HIV-1 integrase catalytic core and C-terminal domains: a model for viral DNA binding. Proc Natl Acad Sci USA 97: 8233–8238.

Cherepanov P, Ambrosio AL, Rahman S, Ellenberger T, Engelman A (2005) Structural basis for the recognition between HIV-1 integrase and transcriptional coactivator p75. Proc Natl Acad Sci USA 102: 17308-17313.

Wang J-Y, Ling H, Yang W, Craigie R (2001) Structure of a two-domain fragment of HIV-1 integrase: implications for domain organization in the intact protein. EMBO J. 20:7333-7343.

Christ F, Voet A, Marchand A, Nicolet S, Desimmie BA, et al. (2010) Rational design of small-molecule inhibitors of the LEDGF/p75-integrase interaction and HIV replication. Nature Chemical Biology 6: 442–448.

Wielens J, Headey SJ, Jeevarajah D, Rhodes DI, Deadman J, et al. (2010) Crystal structure of the HIV-1 integrase core domain in complex with sucrose reveals details of an allosteric inhibitory binding site. FEBS Lett 584: 1455-1462.

Rhodes DI, Peat TS, Vandegraaff N, Jeevarajah D, Le G, et al. (2011) Structural basis for a new mechanism of inhibition of HIV-1 integrase identified by fragment screening and structure-based design. Antiviral Chemistry and Chemotherapy 21: 155-168.

Schrijvers R, De Rijck J, Demeulemeester J, Adachi N, Vets S, et al (2012) LEDGF/p75-Independent HIV-1 Replication Demonstrates a Role for HRP-2 and Remains Sensitive to Inhibition by LEDGINs. PLoS Path. 8: e1002558.

Kessl JJ, Jena N, Koh Y, Taskent-Sezgin H, Slaughter A, et al. (2012) A multimode, cooperative mechanism of action of allosteric HIV-1 integrase inhibitors. J. Biol. Chem 287: 16801-16811.

Cherepanov P, Maertens G, Proost P, Devreese B, Van Beeumen J, et al. (2003) HIV-1 integrase forms stable tetramers and associates with LEDGF/p75 protein in human cells. J Biol Chem 278: 372–381.

Cherepanov P, Sun Z-YJ, Rahman S, Maertens G, Wagner G, et al. (2005) Solution structure of the HIV-1 integrase-binding domain in LEDGF/p75. Nature Struct Mol Bio 12: 526-532.

Tsiang M, Jones GS, Hung M, Mukund S, Han B, et al. (2009) Affinities between the binding partners of the HIV-1 integrase dimer-lens epithelium-derived growth factor (IN dimer-LEDGF) complex. J Biol Chem 284: 33580-33599.

Rhodes DI, Peat TS, Vandegraaff N, Jeevarajah D, Newman J, et al. (2011) Crystal structures of novel allosteric peptide inhibitors of HIV integrase identify new interactions at the LEDGF binding site. ChemBiochem 12: 2311-2315.

De Luca L, Ferro S, Morreale F, De Grazia S, Chimirri A (2011) Inhibitors of the interactions between HIV-1 IN and the cofactor LEDGF/p75. ChemMedChem 6: 1184-91.

Fan X, Zhang FH, Al-Safi RI, Zeng LF, Shabaik Y, et al. (2011) Design of HIV-1 integrase inhibitors targeting the catalytic domain as well as its interaction with LEDGF/p75: a scaffold hopping approach using salicylate and catechol groups. Bioorg Med Chem 19: 4935-52.

De Luca L, Gitto R, Christ F, Ferro S, De Grazia S, et al. (2011) 4-[1-(4-Fluorobenzyl)-4-hydroxy-1H-indol-3-yl]-2-hydroxy-4-oxobut-2-enoic acid as a prototype to develop dual inhibitors of HIV-1 integration process. Antiviral Res 92: 102-7.

Tsantrizos YS, Boes M, Brochu C, Fenwick C, Malenfant E, et al. (2007), pp. 1-116, International Patent Application PCT/CA2007/000845.

Peat TS, Rhodes DI, Vandegraaff N, Le G, Jones ED, Smith J, Coates JAV, Thienthong N, Newman J, Dolezal O, Ryan JH, Savage GP, Francis CL, Deadman JJ (2012) Small molecule inhibitors of the LEDGF site of Human Immunodeficiency Virus Type 1 Integrase identified by fragment screening and structure based design. PLoS ONE 7: e40147; doi:10.1371/journal.pone.0040147

J Wielens, SJ Headey, DI Rhodes, RJ Mulder, O Dolezal, JJ Deadman, J Newman, DK Chalmers, MW Parker, TS Peat, MJ Scanlon (2013) Parallel screening of a low molecular weight compound library. Do differences in methodology affect hit identification? Journal of Biomolecular Screening, 18:147-159. doi:10.1177/1087057112465979