

# Anti-HIV Drug Design

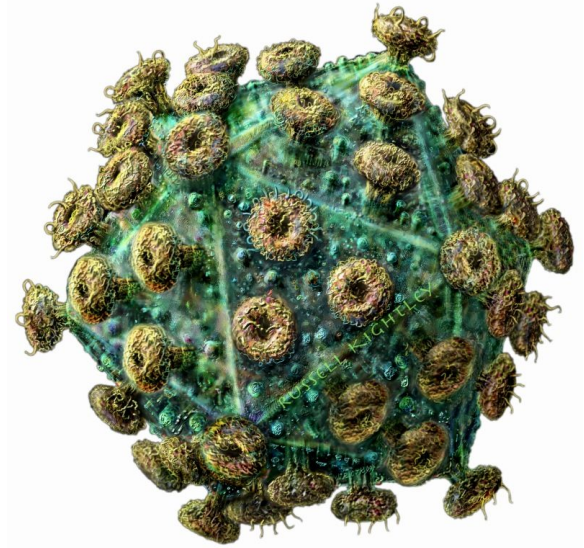
**Group Members:**

- 1. ED15B001 Adarsh**
- 2. CY16C016 Hruday**



# The Human Immunodeficiency Virus

- First clinically observed in USA in 1981
- Causes AIDS, a condition in which progressive failure of the immune system allows life-threatening opportunistic infections and cancers to thrive
- Most commonly spread via sexual contact
- Also transferred from infected mother to baby during pregnancy, by using infected needles & through blood transfusions



**Fig:** The HIV Virus



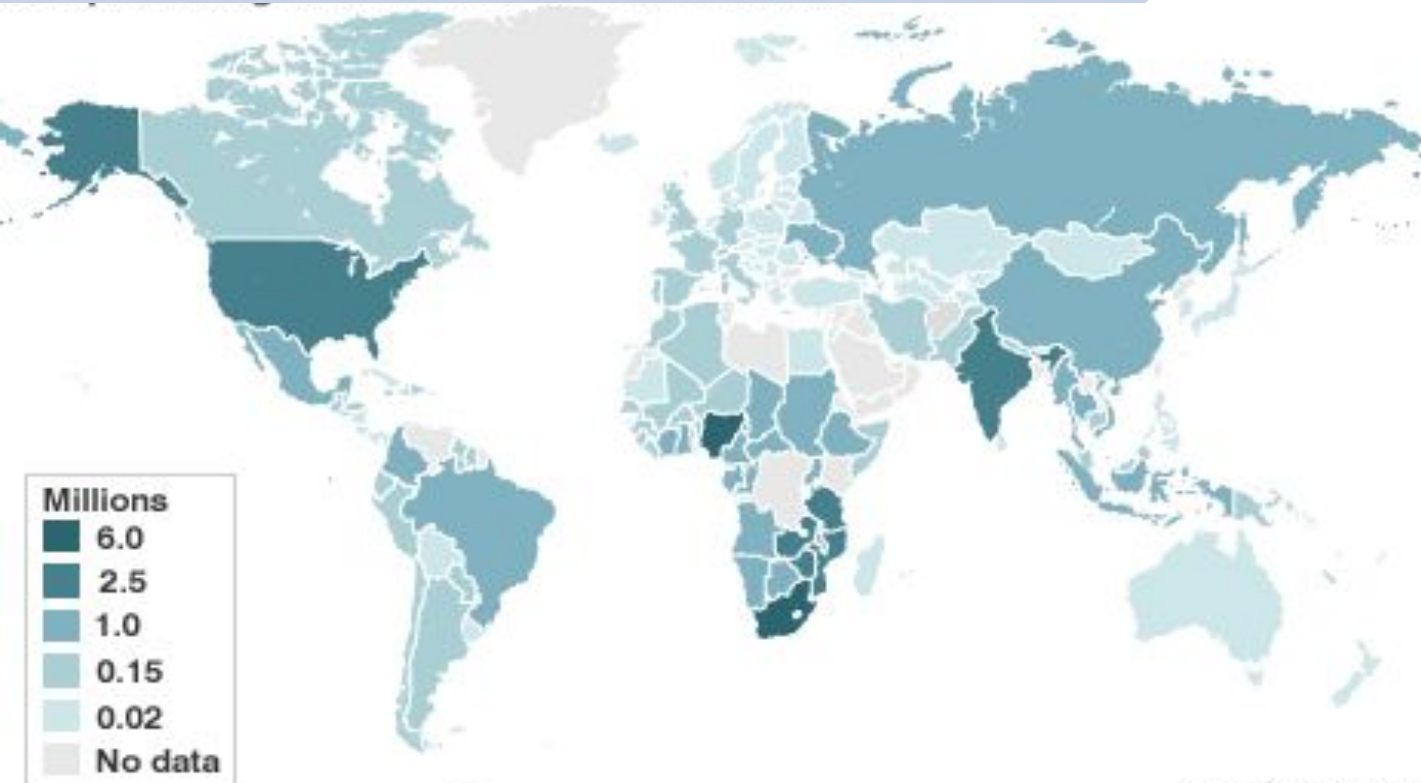
# HIV/AIDS Statistics

- Approximately 36.7 million people living with HIV/AIDS at the end of 2016. Of these, around 2.1 million were children(<15 years)
- Around 5000 new cases reported everyday
- Only 60% of people infected know their statuses; remaining 40% don't even have access to HIV testing services
- 1 million people died from AIDS-related illnesses in 2016

**Source:** UNAIDS Foundation



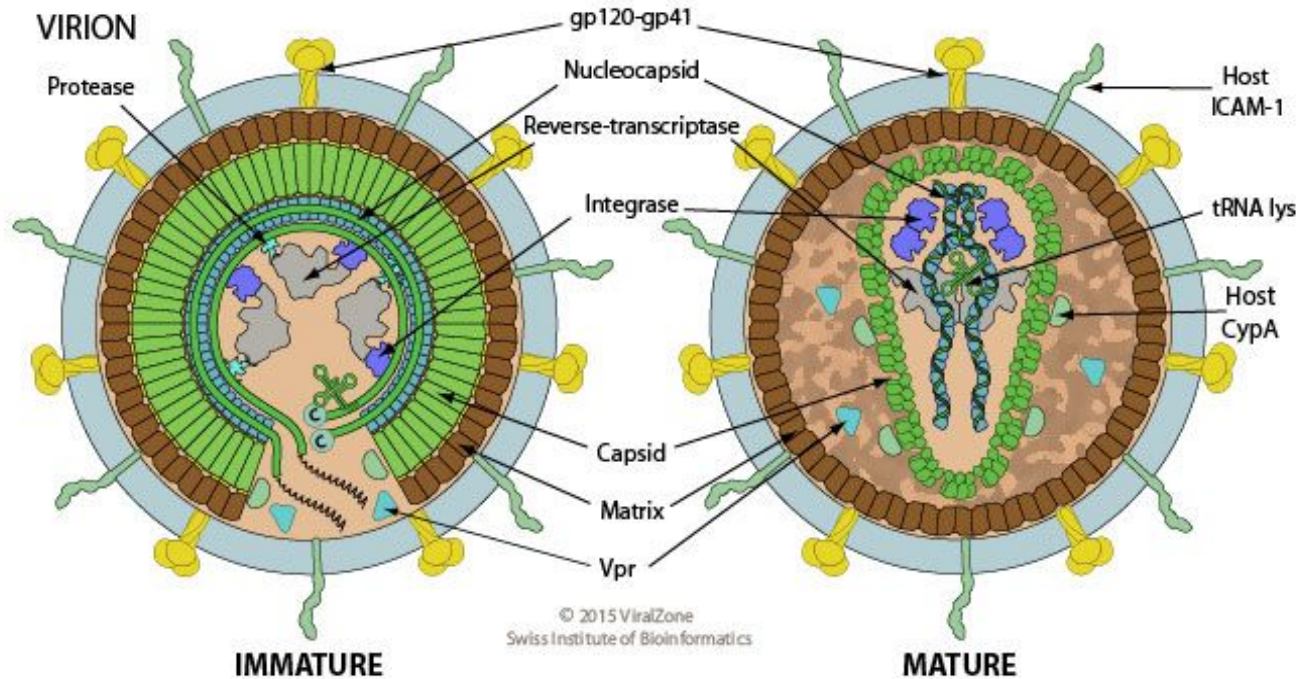
# HIV Spread Map



*Source: UNAIDS*



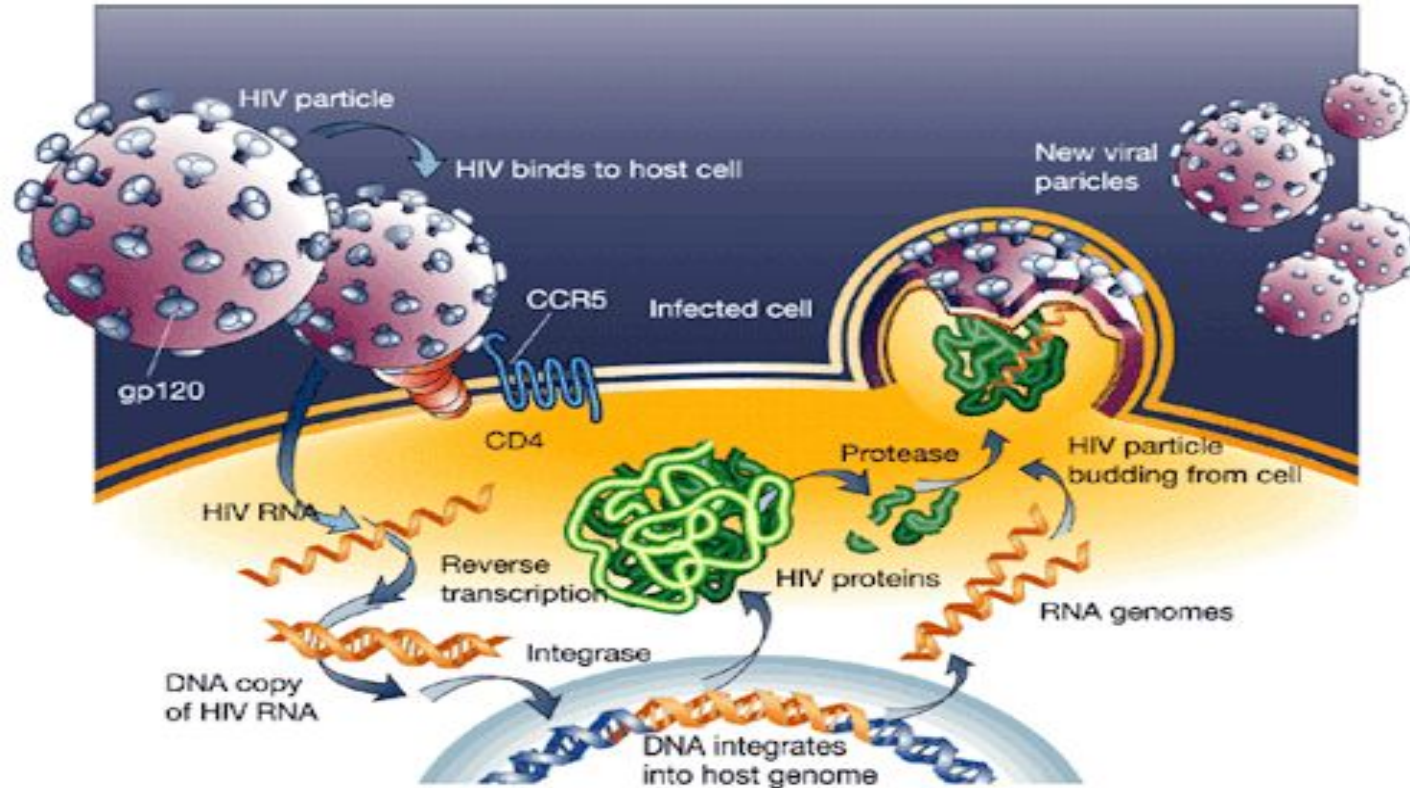
# HIV Cell Structure



**Ref:** [https://viralzone.expasy.org/7?outline=all\\_by\\_species](https://viralzone.expasy.org/7?outline=all_by_species)



# HIV Replication Life Cycle





# Existing Treatment Methods

## Antiretroviral Therapy

- **Entry Inhibitors:** Block entry of HIV into immune cells. Ex: Enfuvirtide
- **Nucleoside RT Inhibitors:** Block the reverse transcriptase proteins that HIV needs to multiply. Ex: Abacavir, Didanosine
- **Non-Nucleoside RT Inhibitors:** Bind to and disable the reverse transcriptase proteins that HIV needs to multiply. Ex: Delavirdine, Efavirenz
- **Integrase Inhibitors:** Block the enzyme that HIV needs to infect immune cells with its genetic material. Ex: Dolutegravir, Raltegravir
- **Protease Inhibitors:** Inhibit HIV Protease, that HIV needs to make copies of itself. Ex: Saquinavir, Darunavir

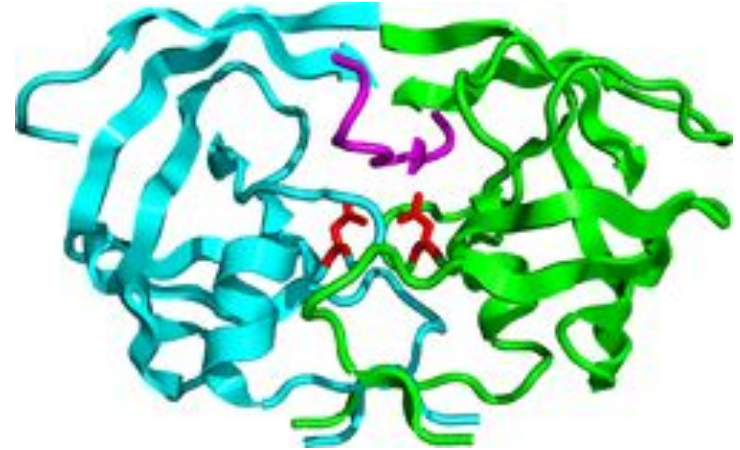




# Target Molecule

## HIV Protease

- Essential for life cycle replication of HIV
- Cleaves newly formed polyproteins at the appropriate locations to create mature components of an infectious HIV Virion
- Without effective HIV Protease, HIV Virions remain non-infectious

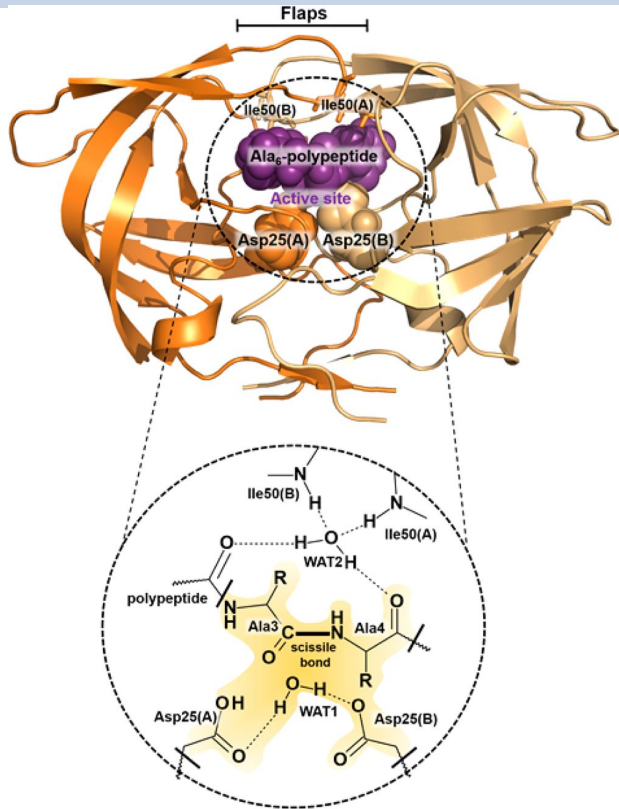


**Fig:** HIV protease dimer(green & blue)  
with active site marked in red





# Target Molecule - Active Site

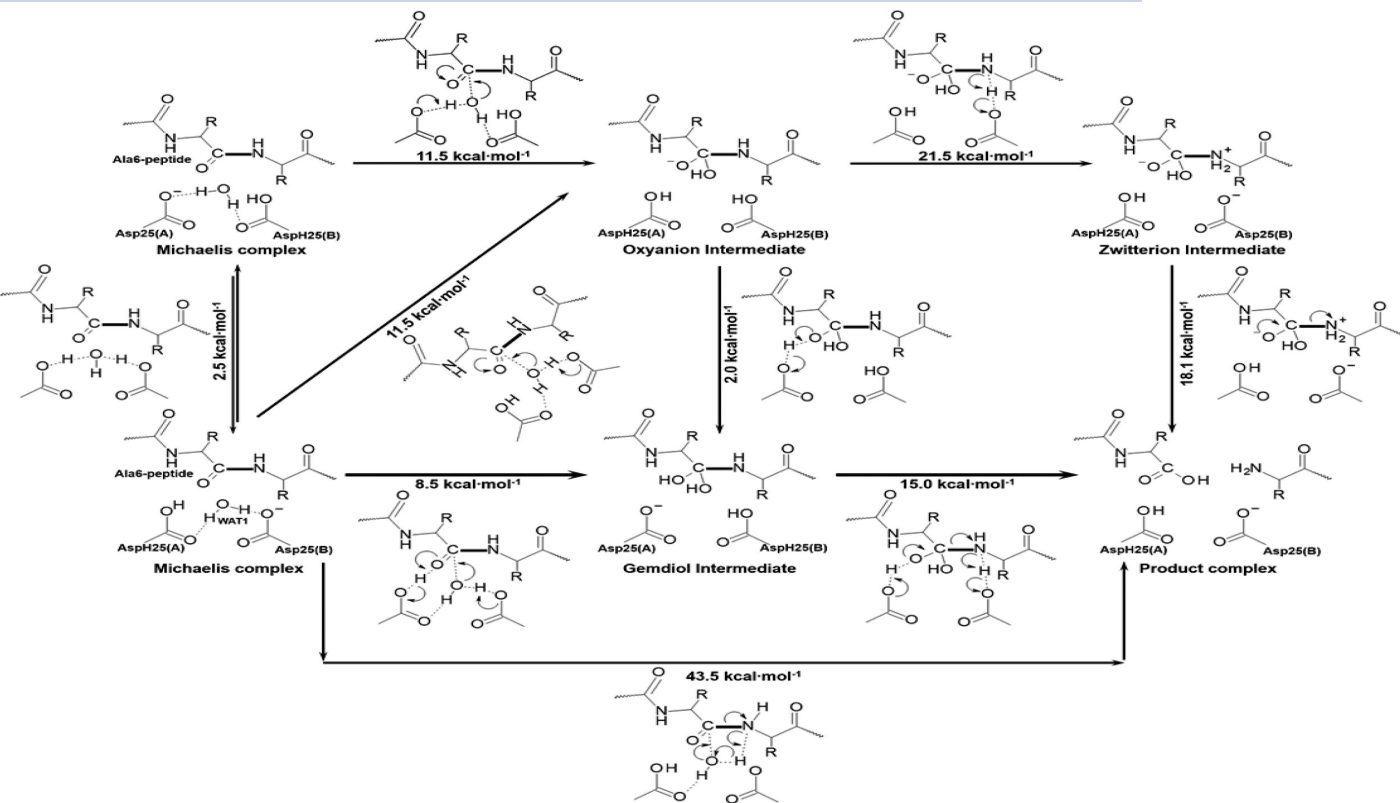


**Fig:**

*Schematic representation of HIV Protease and detail of the active site, with protonated Asp-25(A), and Ala6 peptide as substrate*



# Action of HIV Protease



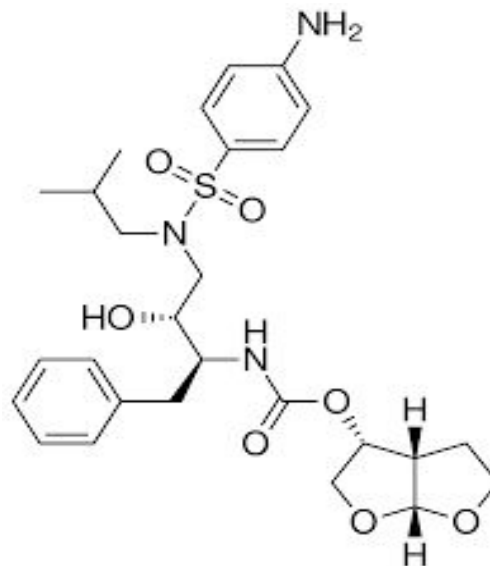
**Ref:** Agnieszka Krzeminska, Vicent Moliner, Katarzyna Swiderek, "Dynamic and Electrostatic Effects on the Reaction Catalyzed by HIV-1 Protease", *Journal of the American Chemical Society*-2016



# Methods of Drug Development

## Lead Compound: Darunavir

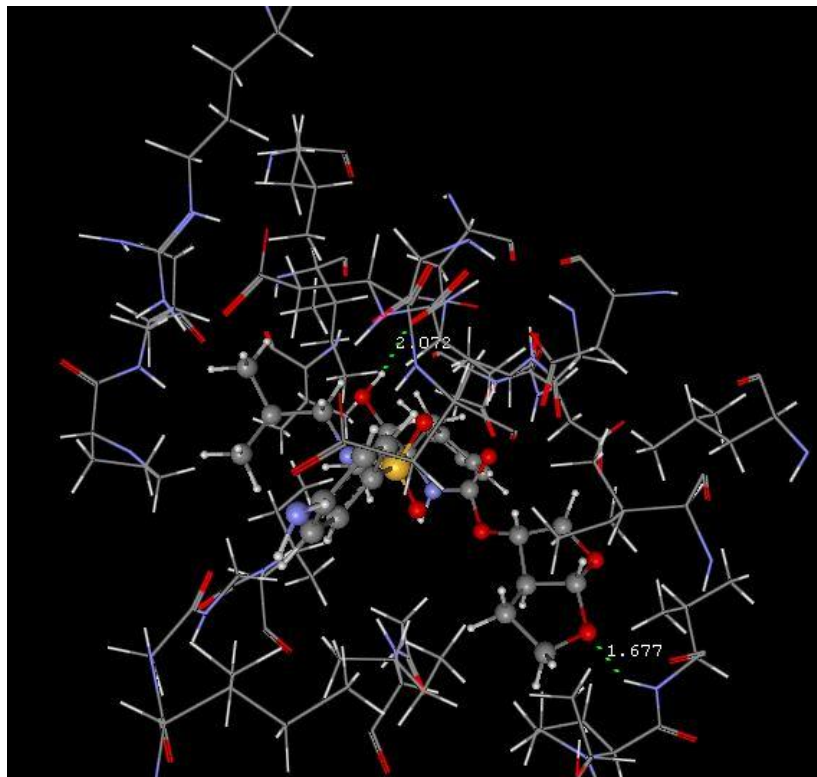
- Darunavir was chosen as a base molecule from which analogues were constructed
- The docking of Darunavir was studied and possible analogues were designed considering gaps between the drug & the target as well as potential new interactions



**Fig:** Structure of Darunavir



# Docking of Darunavir



## Interactions

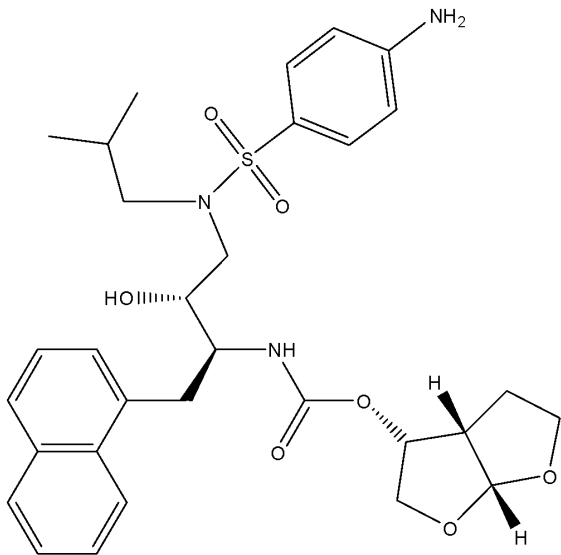
- 2 Hydrogen Bonds
- No Pi-Pi Stacking

**Docking Score:**  
-65.69

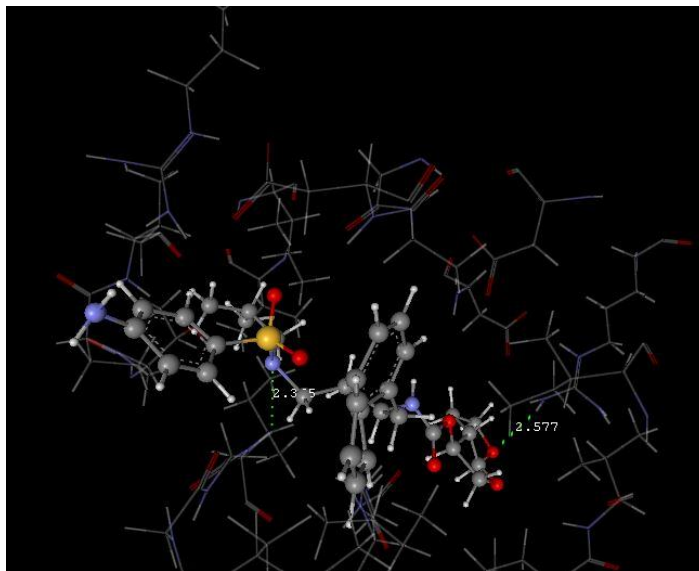


# New Drug Design

## Analogue 1



## Docking Study



## Interactions

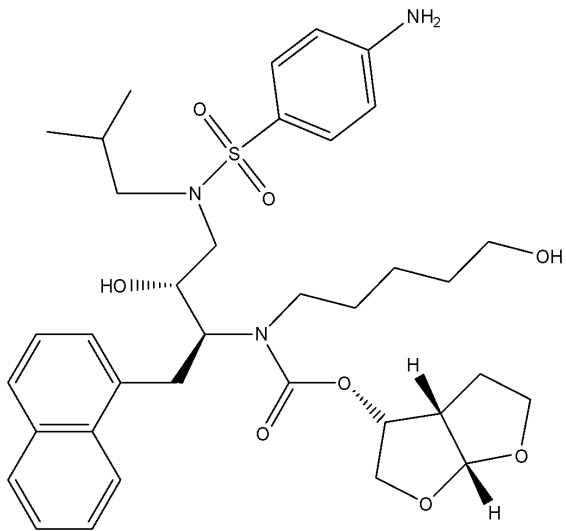
- 2 Hydrogen Bonds
- No Pi-Pi Stacking

**Docking Score:**  
-41.24

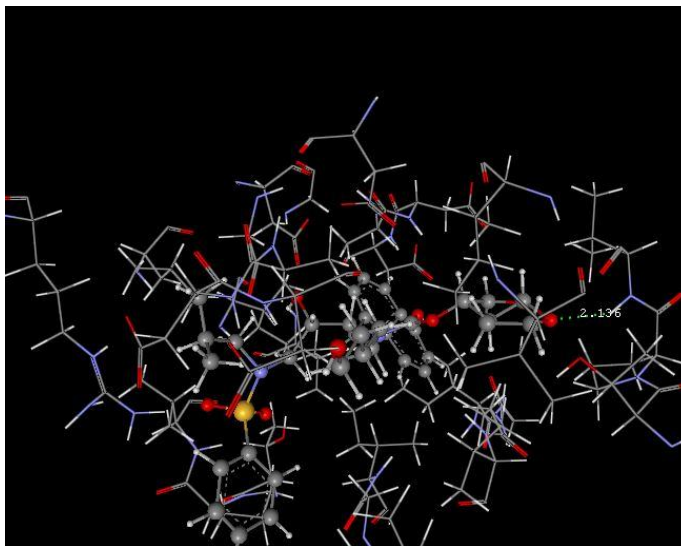


# New Drug Design

## Analogue 2



## Docking Study



## Interactions

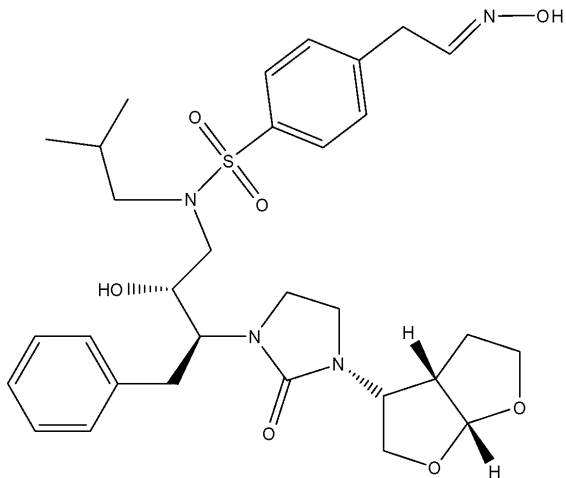
- 1 Hydrogen Bond
- No Pi-Pi Stacking

**Docking Score:**  
-4.44

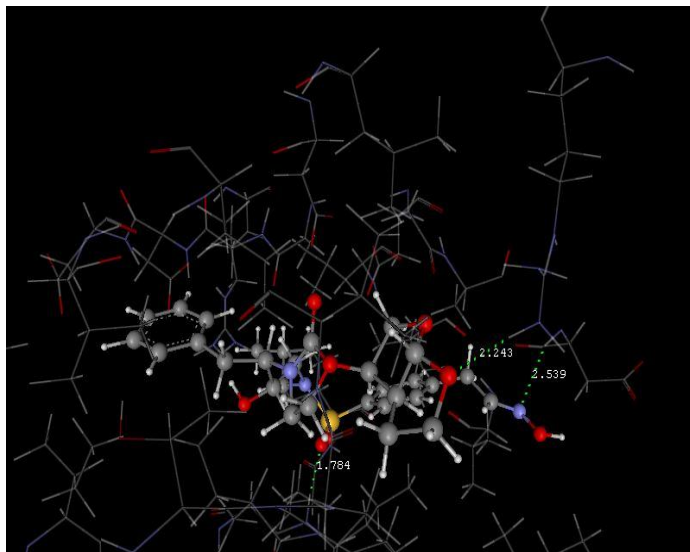


# New Drug Design

## Analogue 3



## Docking Study



## Interactions

- 3 Hydrogen Bonds
- No Pi-Pi Stacking

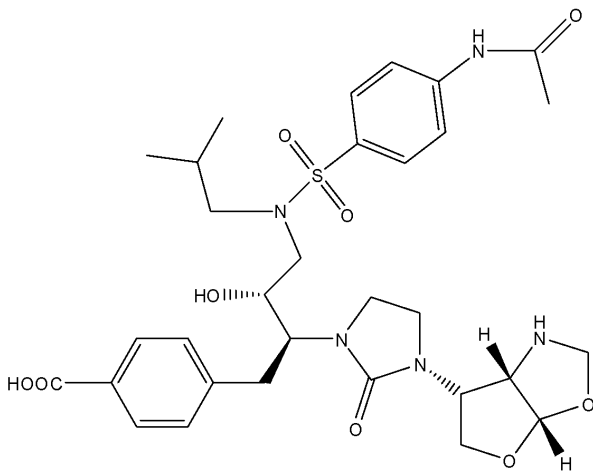
**Docking Score:**  
-56.84



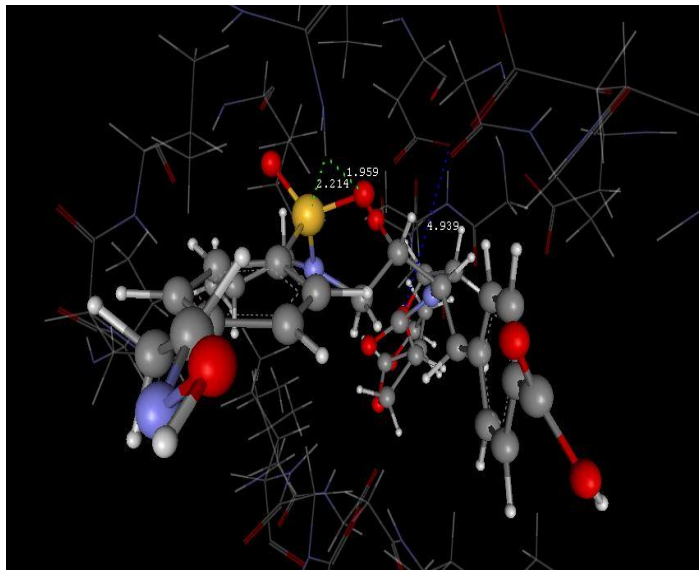


# New Drug Design

## Analogue 4



## Docking Study



## Interactions

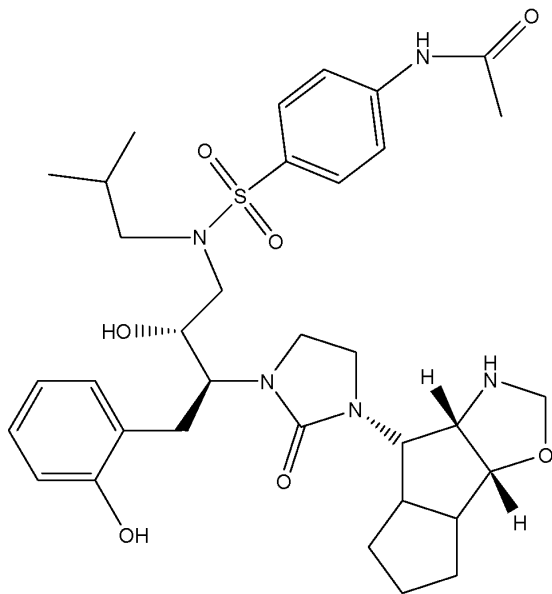
- 2 Hydrogen Bonds
- No Pi-Pi Stacking

**Docking Score:**  
-53.23

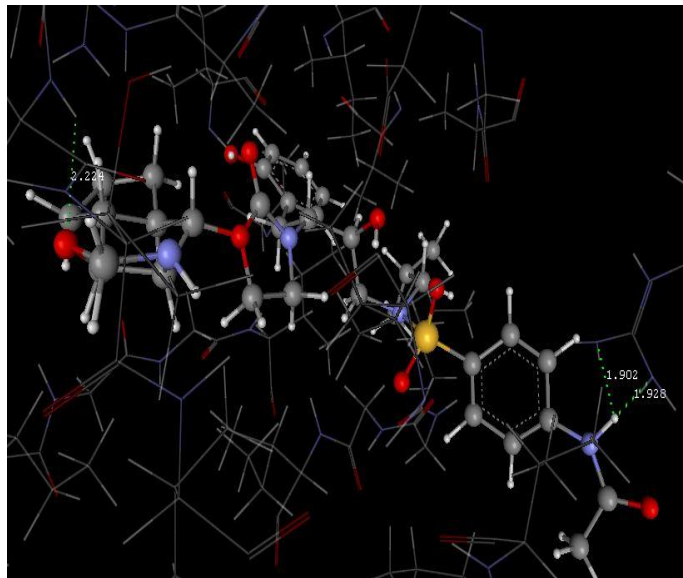


# New Drug Design

## Analogue 5



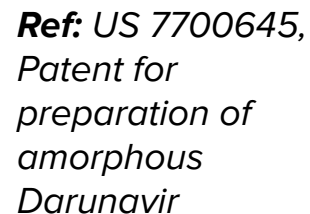
## Docking Study



## Interactions

- 3 Hydrogen Bonds
- No Pi-Pi Stacking

**Docking Score:**  
-59.14





# Drug Metabolism

- Darunavir is extensively metabolized by CYP enzymes, primarily CYP3A
- The terminal elimination half-life of darunavir is approximately 15 hours
- Around 48% Darunavir excreted unchanged from the human body

***Ref:*** Marc Vermeir, Sophie Lachau-Durand et al, "Absorption, Metabolism, and Excretion of Darunavir, a New Protease Inhibitor", *Drug Metabolism & Disposition* April 2009



# Conclusions

- HIV infection mechanism was studied & target HIV Protease was identified
- HIV Protease action was reviewed
- Mechanism of inhibition of Darunavir was examined
- Analogues of Darunavir were designed & their interactions were investigated



**THANK YOU**