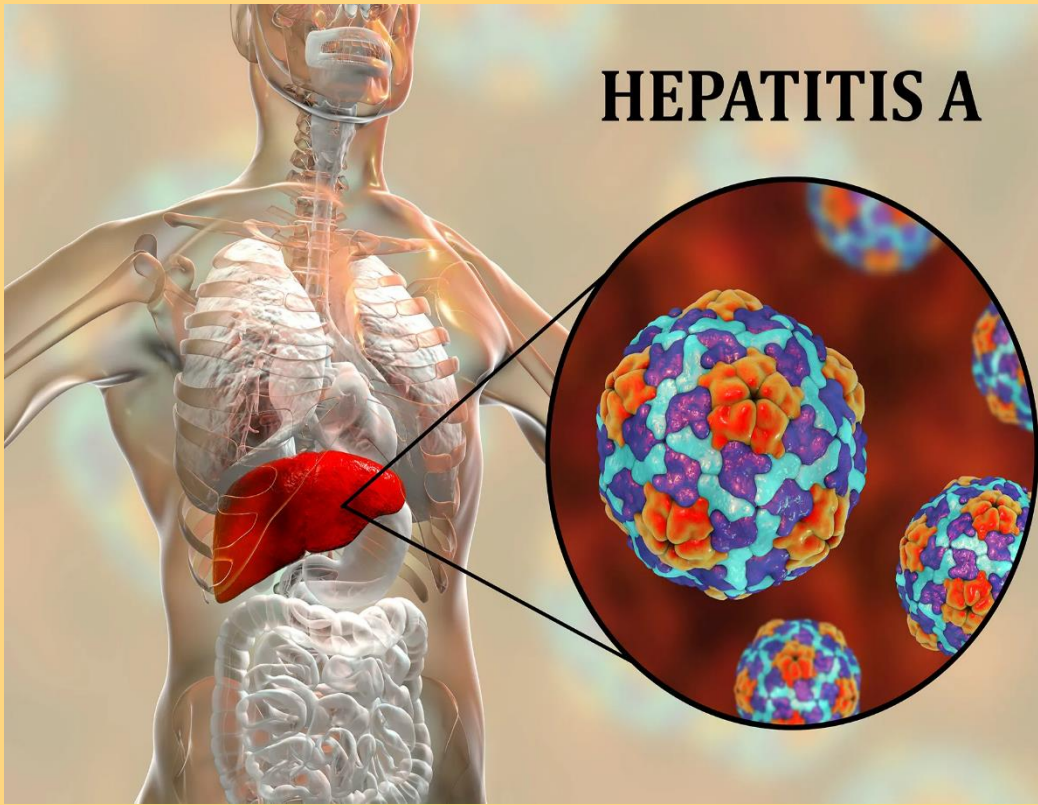


# Assessing the Potential of Anti Hepatitis-A activity of *Catharanthus roseus* Phytochemicals using Immunoinformatics

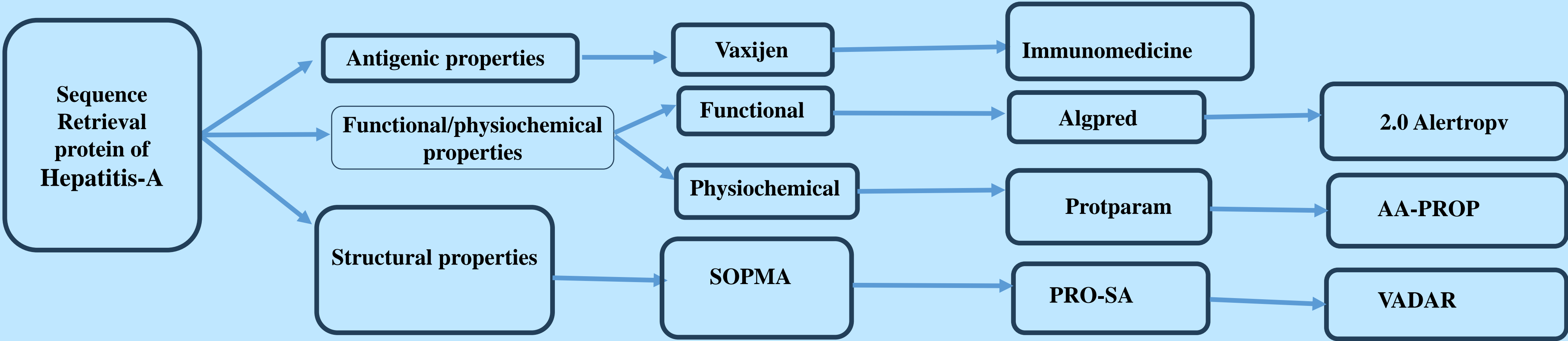
## INTRODUCTION

- Hepatitis A is a highly contagious liver infection caused by the hepatitis A virus (HAV). It is typically transmitted through the ingestion of contaminated food or water or through close contact with an infected person.
- Existing commercially available antiviral drug for Hepatitis-A include Havrix that contains inactivated hepatitis A virus. The main drawback of using the Havrix vaccine is that it provides protection specifically against hepatitis
- A virus infection but not against other forms of hepatitis, such as hepatitis B or C.
- The medicinal properties of *Catharanthus roseus* and its inherent alkaloids, such as vincristine and vinblastine have been used in the treatment of certain cancers
- *Catharanthus roseus* also plays a major role in traditional medicine in various culture for treating a range of ailment.
- The aim of this present research introduces an *insilico* approach to identify a novel potential ligand to target the disease causing protein thereby giving a futuristic perspective to develop a novel sustainable medication against effective diseases.



Characteristics of Hepatitis-A Virus

## MATERIALS AND METHODS



## RESULTS

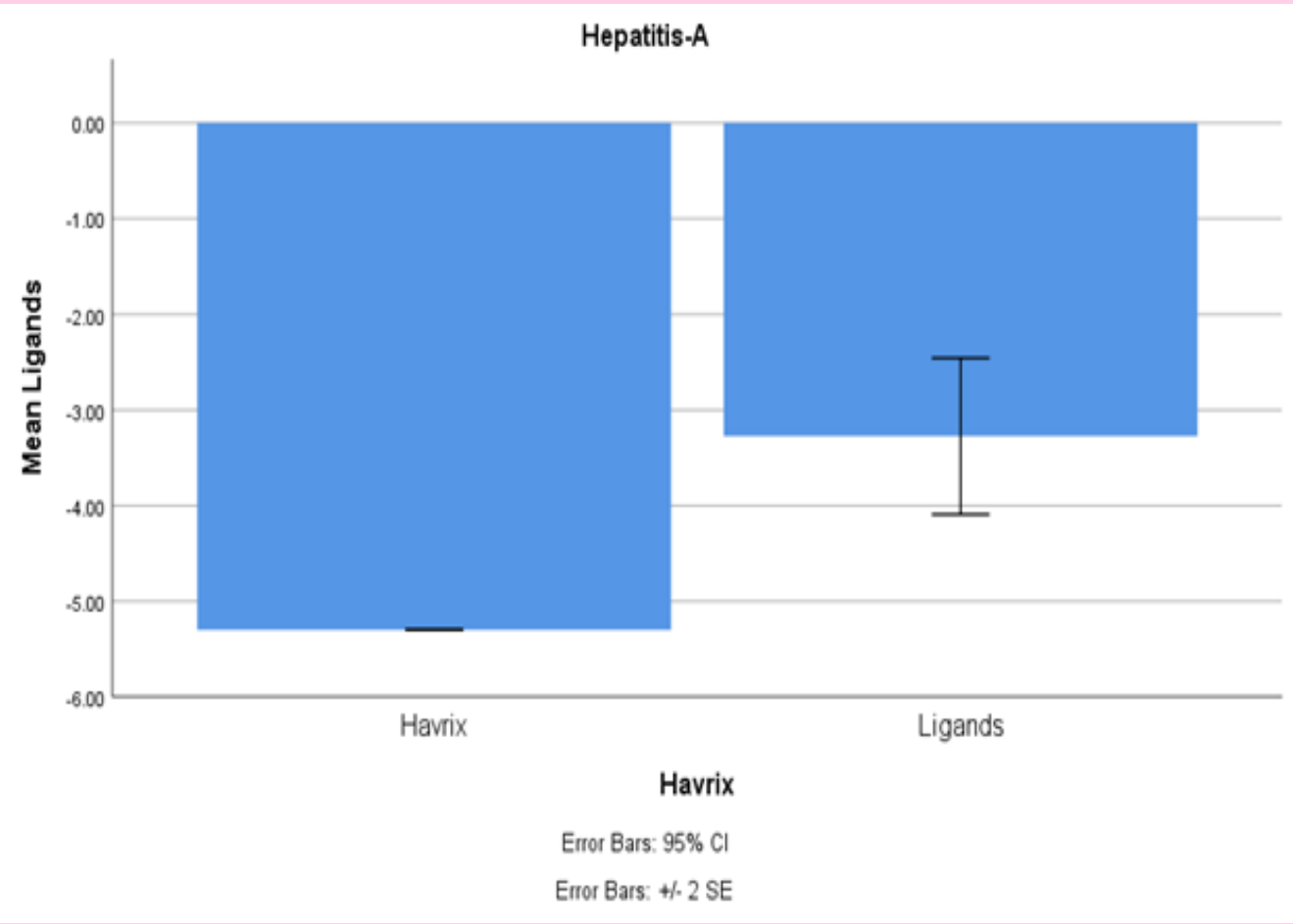


Fig. 1. Bar chart representing the mean binding affinity of Havrix and Ligand

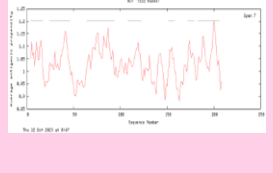
Protein sequence	Antigenic properties				
	VaxiJen		Immunomedicine		
	Antigenic score Threshold = 0.4	Antigenic nature	Average antigenic propensity	Antigenic determinants	Antigenic plot sequence
3C protease	0.4981	Probable antigen	1.0296	7	

Table 1. Antigenic Properties of the selected epitopes of Hepatitis-A virus

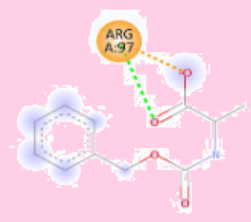
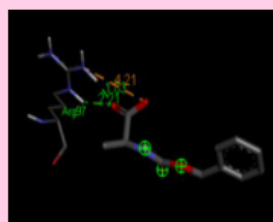
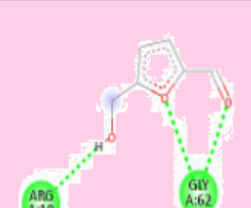

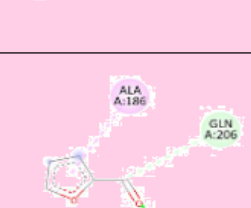
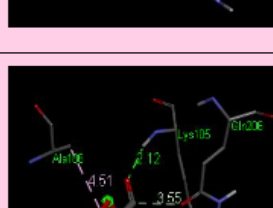

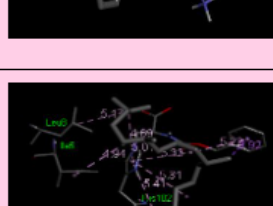
		1H-pyrrole, 2,5-dihydro-1-nitroso.
		2-Furancarboxaldehyde
		3,7,11,15-Tetramethyl-2-hexadecen-1-ol
		5-(hydroxymethyl)

Table 2. 2D and 3D representations of docking results of HAV

## DISCUSSION AND CONCLUSION

- Four compounds were identified through GC-MS analysis and were subjected as a ligand for docking studies, the highest binding affinity of -5.1 kcal/mol was observed.
- The secondary protein structures with access to the number of alpha helices, beta turns, and coils of the antigenic chains of HAV 3C protease were obtained from tabulated SOPMA findings.
- The comparison between the antidrug against Hepatitis-A as control group and ligands of *Catharanthus roseus* as study group indicates a statistical significance between the two groups (p=0.018, p < 0.05)
- Table 2. shows the 2D and 3D representations of docking findings of phytochemical substances docked to HAV 3C protease of Hepatitis.
- According to structural and epitope modelling, the epitope with PDB ID 2CXV is the most suitable receptor for antiviral medication targeting in the treatment of hepatitis-A.
- The current study merely demonstrates an immunoinformatic method for assessing a potential drug as an alternative to Havrix and their interactions with the phytochemicals found in *C.roseus*, which will spur further *insilico* research to develop an effective Hepatitis-A antiviral drug.

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