



## INTRODUCTION AND OBJECTIVES

STARD11 is a protein domain that transfers ceramide from the ER membrane to the Golgi membrane. It belongs to the superfamily of StAR-related lipid-transfer domains (STARD) which contains 13 other members. Other STARD proteins transfer different types of lipids.<sup>1</sup>

The main aim of this study was to investigate the START-domain of the human STARD11. Questions to be answered:

1. **Is the membrane-binding region conserved?** Investigated STARD11 in other organisms as well as human STARD proteins.
2. **Is the ceramide-binding region conserved?** Investigated STARD11 in other organisms as well as human STARD proteins.
3. **Is the membrane binding region flexible?** Investigated STARD11 in WebNma and presenting a fluctuation diagram.

## MATERIALS AND METHODS

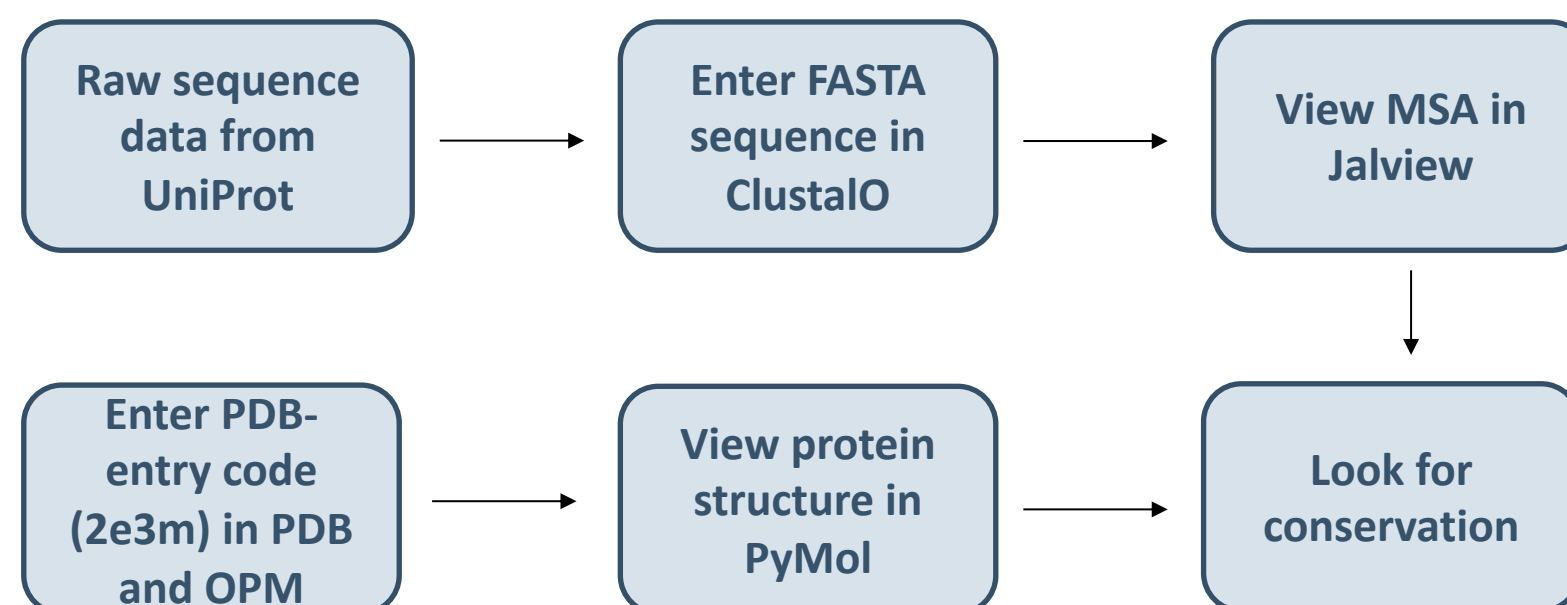


Figure 1: Flow chart of the bioinformatic strategy used in this research. Tools: UniProt, ClustalO<sup>2</sup>, Jalview<sup>3</sup>, Protein Data Bank, PyMol, OPM<sup>4</sup>, WebNma<sup>5</sup> and WebLogo<sup>5</sup>.

## REFERENCES

1: Wong LH et al., 2019 (2):85-101.  
2: Madeira et al., (2019). Nucleic Acids Research. 47(W1):W636-W641

## RESULTS OVERVIEW

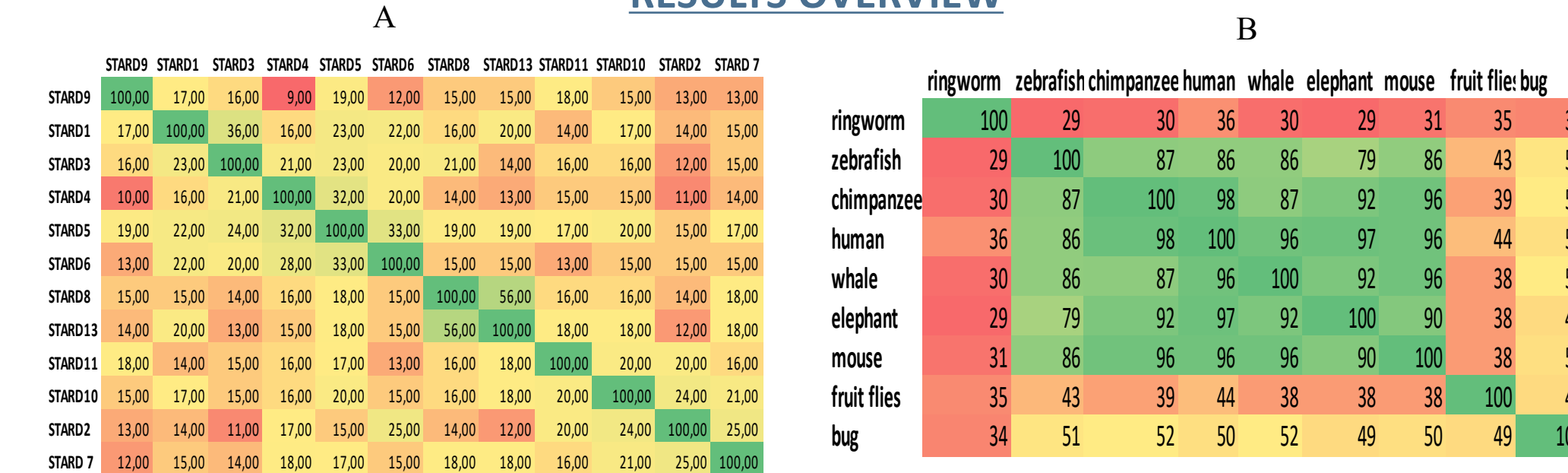


Figure 2: Heat map based on a percentage identity matrix obtained from ClustalO of different human STARD proteins (A) and different species of STARD11 (B). This heatmap illustrates which proteins have low sequence similarity (red) and which proteins have high sequence similarity (green).

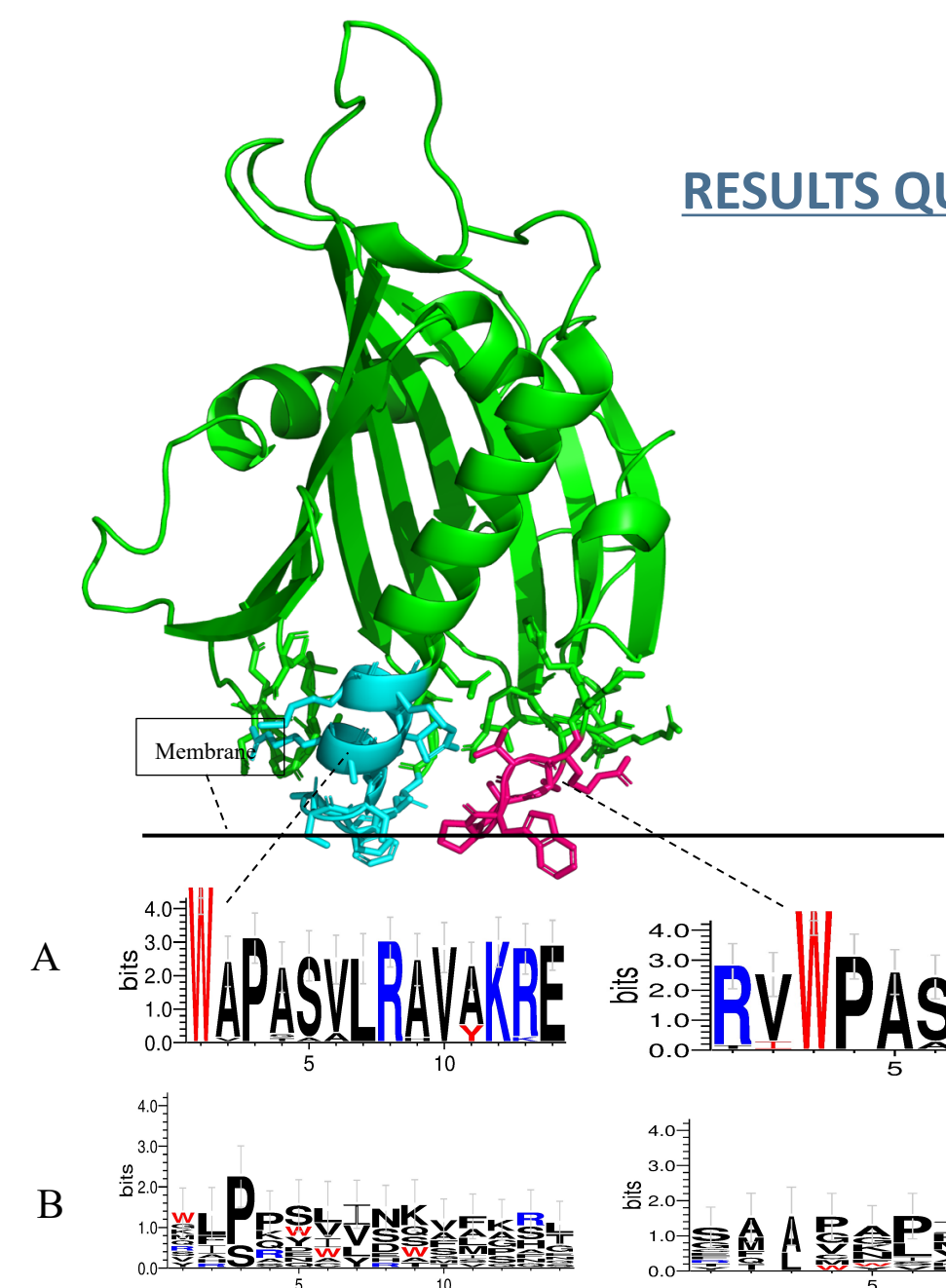


Figure 3: Comparison of the membrane binding site in various STARD11 species (A) and STARD proteins (B). The membrane binding site is more conserved in the different species compared to different human STARD proteins.

## RESULTS QUESTION 1 AND 2

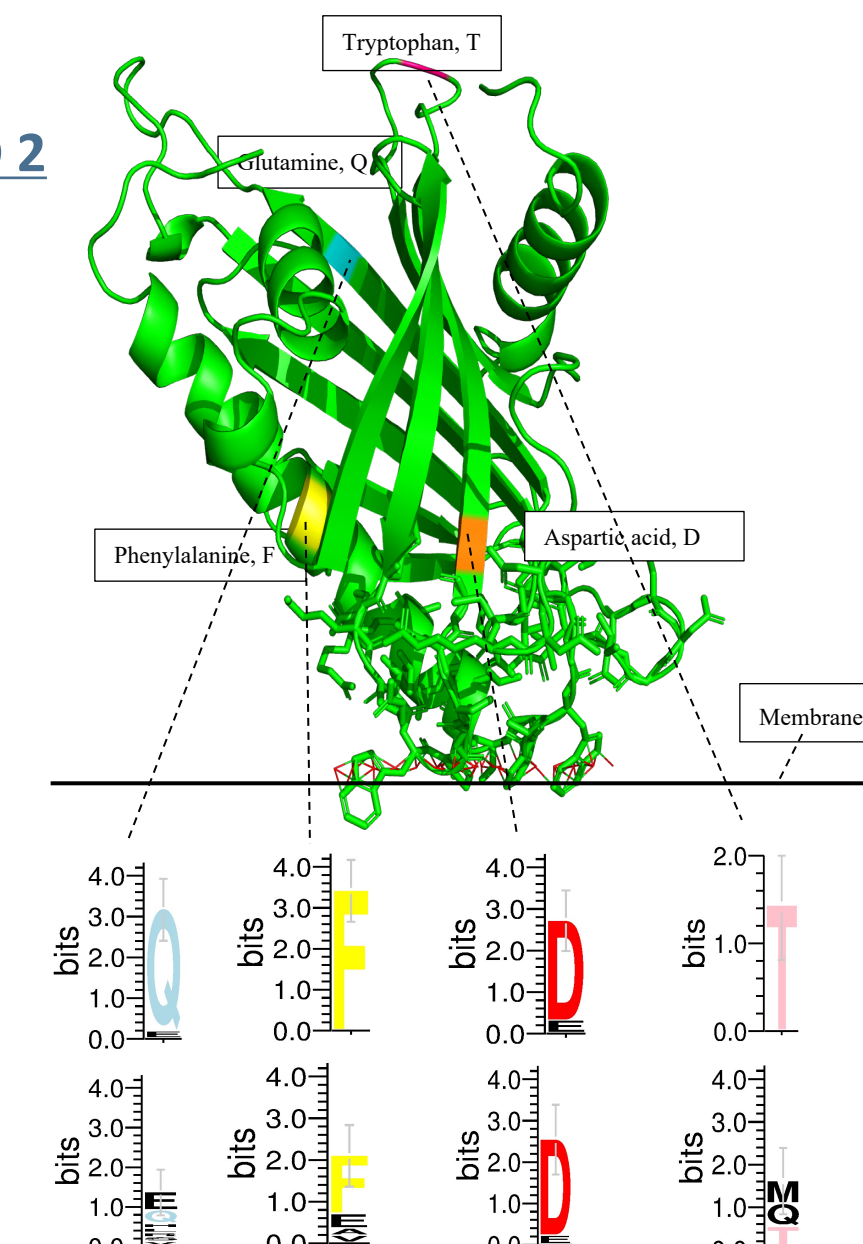


Figure 4: Comparison of the Ceramide binding site in various STARD11 species (A) and STARD proteins (B). The ligand binding site is more conserved in the different species compared to different human STARD proteins.

## RESULTS QUESTION 3

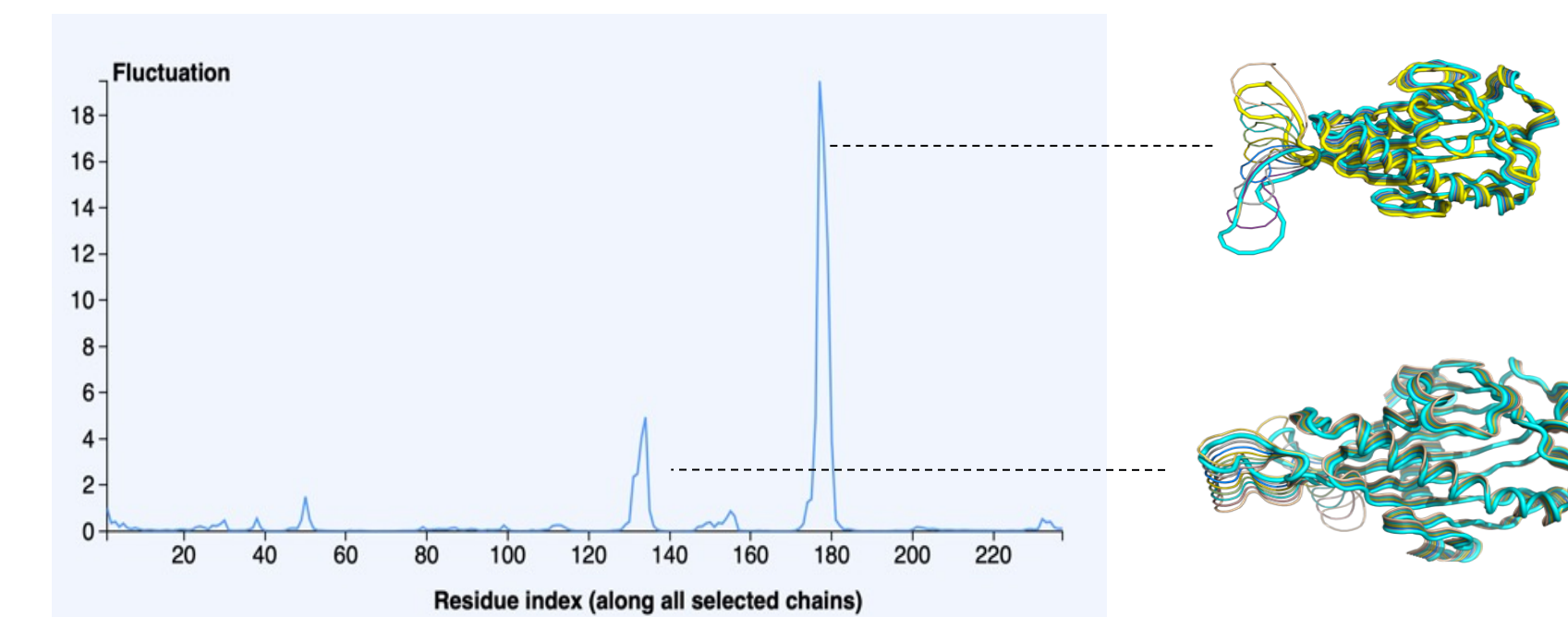


Figure 5: Flexibility profile of the STARD11 domain and PyMol representation of the two most flexible nodes. Ligand binding site 479 (D) and 579 (F) is present in node 1 and 2, respectively. These two nodes are located at the opposite site of the membrane binding region.

## CONCLUSION

- **Overview:** Other organisms that contain STARD11 have conserved both the membrane and the ceramide binding site, but not other human STARD proteins.
- **Q1:** The amino acid W is highly conserved in the membrane binding site between the different species<sup>1</sup>.
- **Q2:** The ligand binding site 479 (D) and 579 (F) are conserved in both other organisms with STARD11 and other human STARD proteins.
- **Q3:** Two ligand binding sites are found in the two flexible regions, node 1 and 2. This could explain how STARD11 attracts ceramides and shelters them.

## FURTHER STUDIES

- Investigate why the most flexible regions are located at the opposite site of the membrane binding region.