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Milestone #1: Become familiar with Unity Linux system, NAMD simulation, and antimicrobial peptides by reading online documents and 1 to 2 related journal papers; launch presentation;

Milestone #2: Set up human beta defensin type 1 to 3 embedding in model Gramnegative bacterial lipid membranes using CHARMM-gui online program; then run NAMD simulations to equilibrate the systems built;

Milestone #3: Run steered MD simulations to pull hBD-1/hBD-2/hBD-3 out of model bacterial lipid membranes;

Milestone #4: Analyze structure, dynamics and free energy profiles from Steered MD simulations; compare results from different defensin simulations, and interpret the results;

Milestone #5: Write a project report, do a presentation based on research findings, wrap up the project, exit interview.

Goals

- Learn background of antimicrobial peptides;
- Learn basic knowledge of Linux system and simulation;
- Learn how to set up simulations on protein in solvents using CHARMM-GUI online program;
- Learn how to run NAMD simulations on UNITY;
- Learn how to run Steered MD simulations;
- Learn how to interpret simulation results;



- Timeframe
 - September 15th, 2023;
 - February 14th, 2024;



- What I hope to learn
 - Basic knowledge of antimicrobial peptides;
 - Basic knowledge of molecular simulations;
 - Basic knowledge of running Steered MD on UNITY;
- What I have learned
 - Read three papers and wrote three reviews on Human Beta Defensin;
 - Learned basic knowledge of Linux systems;



- Goals for Next Month
 - Running MD simulations on UNITY;
 - Analyzing MD simulation results;
 - Continue learning about Human Beta Defensin;



- Help needed (if any)
 - Install VMD on UNITY;
 - Run xmgrace on UNITY;

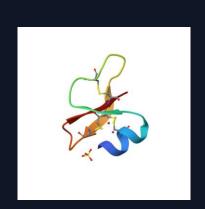


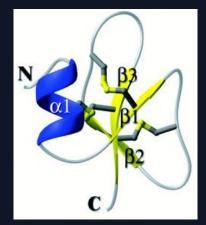
- Human Beta Defensin
 - Peptides that are critical in providing antimicrobial barriers against viruses, fungi, gram-positive/negative bacteria, as well as antibiotic resistant bacteria;
 - Working with 3 defensins: Human Beta Defensin-1,
 Human Beta Defensin-2, and Human Beta Defensin-3.

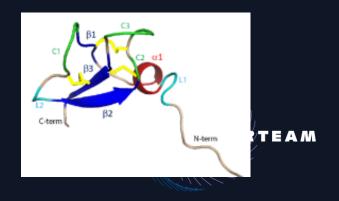
<u>Hoover</u> et al. (2001) J Biol Chem **276**: 39021-39026

Hoover et al. (2000) J. Biol. Chem., 275, 42

Yeasmin et al. (**2018)** *J. Phys. Chem. B,* 122 (50), 11883-11894







- Set up hBD-1 monomer, hBD-2 monomer, and hBD-3 monomer in POPC mixed with POPG membrane using CHARMM-GUI online program;
- Running NAMD simulations on hBD-1 monomer, hBD-2 monomer, and hBD-3 monomer in membrane on UNITY

