

Danna De Boer¹, Parker Ladd Bremer¹, Walter Alvarado², & Eric J. Sorin^{1,*}

Departments of ¹Chemistry & Biochemistry, ²Physics & Astronomy, California State University, Long Beach

BACKGROUND

- Butyrylcholinesterase (BChE) is a nonspecific enzyme known to hydrolyze acetylcholine, a neurotransmitter associated with memory and learning functions,¹ making BChE associated with Alzheimer's Disease and dementia-like symptoms.
- One of our previous studies,² focused on simulations of thirteen organophosphate inhibitor-projects in complex with BChE, but failed to address and solve heuristic problems with the *k*-means clustering algorithm,³ which clusters BChE-inhibitor complexes into binding modes, or average conformations.
- Our last published study addressed the *k*-means' heuristic shortcomings using an intuitive statistical approach that will overcome the heuristic tendencies of *k*-means clustering and qualitatively validate clustering efficacy using internal metrics based on inter- and intra-cluster similarity.
- **Goal:** The study herein will revisit the thirteen organophosphate inhibitor projects and present reproducible and more accurate tabulations of contacts and interactions for each binding mode.

METHODS

- Models:
 - BChE model PDBID# 1P0I
 - Inhibitors modeled & docked with ICM Pro
- Softwares & O 97-BD25-PM-01051498 0 3456 2stud