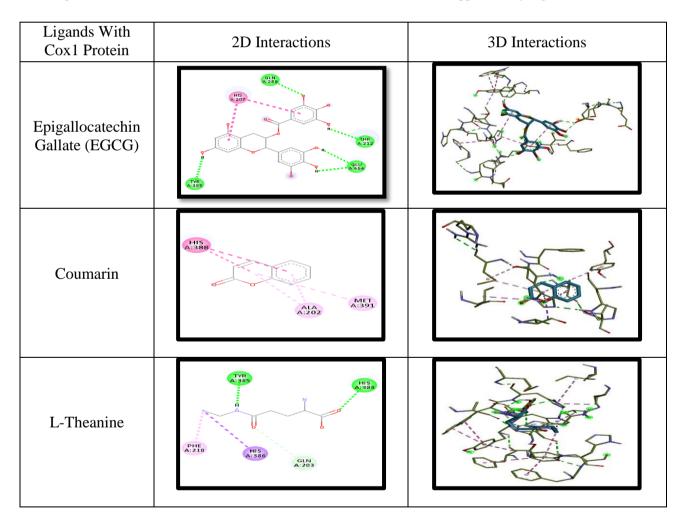
## Green Tea Augments Cognitive Function: An In Silico Model

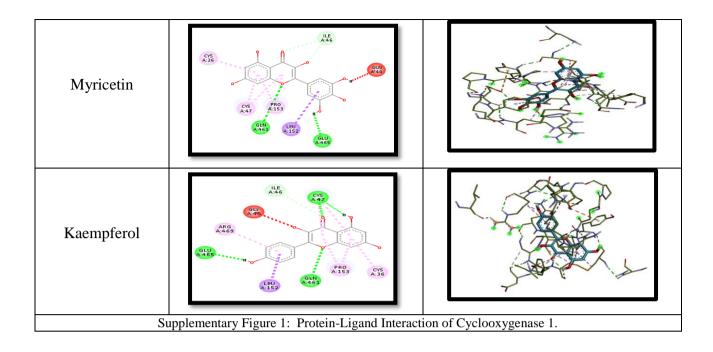
Adittya Sabhayasachi Khan<sup>1,2</sup>, Rezowana Mannan<sup>1,3</sup>, Laila Anjuman Banu<sup>4</sup>, Sohidul Islam<sup>2\*</sup> and Mahmud Hossain<sup>1\*</sup>

## **Supplementary Figures**

Two-dimensional structures simplify the interactions, whereas three-dimensional models amalgamate molecular interactions in ligand binding to the stimulation of the proteins. In the following models, deep green represents conventional hydrogen bonds, light green represents pi-donor hydrogen bond, pink represents a pi-alkyl bond, violet represents pi-sigma, and red represents an unfavorable donor-donor bond. It also simplifies the number of hydrogen bonds and polar interactions favored by ligand-protein interaction.

The ligands' two-dimensional and three-dimensional interaction with Cox 1 (Supplementary Figure 1)





The ligands' two-dimensional and three-dimensional interaction with Cyp4502d6 (Supplementary Figure 2).

Ligands With Cyp4502d6 Protein	2D Interactions	3D Interactions	
Epigallocatechin gallate (EGCG)	A PHE A 366		
Coumarin	ARG LEU A: 444		
L- Theanine	ARG VAL AT19		
Myricetin	A333 A3579		
Kaempferol	ASPA ASSA		
Supplementary Figure 2: Protein-Ligand Interaction of Cytochrome P4502d6.			

The ligands' two-dimensional and three-dimensional interactions with App1 (Supplementary Figure 3).

Ligands With Amyloid-β Protein	2D Interactions	3D Interactions
Epigallocatechin gallate (EGCG)		
Coumarin	LEU A:417	
L-Theanine	VAL A: 460	The state of the s
Myricetin		
Kaempferol	upplementary Figure 3: Protein-Ligand Interacti	on of amulaid & protain

The ligands' two-dimensional and three-dimensional interactions with AChE (Supplementary Figure 4)

Ligands With AChE Protein	2D Interactions	3D Interactions
Epigallocatechin Gallate (EGCG)	ASSO ASSO ASSO	TO STORY
Coumarin	TYR A:124	
L-Theanine	PHE A297  TVR A124  HIS A447	
Myricetin	H A)286  LEU U A)250  LEU A)76  A)341	
Kaempferol	PHE A 293 A 294 PHE A 338 TYR A 124 TYR A 294 PHE A 338 TYR A 124 TYR A 294 PHE A 338 TYR A 124 TYR A 294 PHE A 338 TYR A 124	
Supplementary Figure 4: Protein-Ligand Interaction of acetylcholinesterase.		