In Silico ADME Prediction and Molecular Docking of 1,2,3-Triazole-based Compounds Against Human Aromatase Cytochrome P450

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Table S1. Violations to medicinal chemistry rules and bioavailability score.

Molecule	Number of	Bioavailability				
	Lipinski	Ghose	Veber	Egan	Muegge	Score
Letrozole	0	0	0	0	0	0.55
Anastrozole	0	0	0	0	0	0.55
Compound 1	0	0	0	0	0	0.55
Compound 2	0	0	0	0	0	0.55
Compound 3	0	0	0	0	0	0.55
Compound 4	0	0	0	0	0	0.55
Compound 5	0	0	0	0	0	0.55
Compound 6	0	0	0	0	0	0.55
Compound 7	0	0	0	0	0	0.55
Compound 8	2	2	1	1	2	0.17
Compound 9	0	0	0	0	0	0.55
Compound 10	0	0	0	0	0	0.55
Compound 11	0	0	0	0	0	0.55

Table S2. Select pharmacokinetic properties.

Molecule	GI Absorption	BBB Permeant	P-gp Substrate	CYP1A2 Inhibitor	CYP2C19 Inhibitor	CYP2C9 Inhibitor	CYP2D6 Inhibitor	CYP3A4 Inhibitor
Letrozole	High	Yes	No	Yes	Yes	Yes	Yes	No
Anastrozole	High	Yes	No	No	No	Yes	No	No
Compound 1	High	Yes	No	Yes	Yes	Yes	No	No
Compound 2	High	Yes	No	Yes	Yes	Yes	No	No
Compound 3	High	Yes	No	Yes	Yes	Yes	No	No
Compound 4	High	No	No	No	No	Yes	No	Yes
Compound 5	High	No	No	No	No	Yes	No	Yes
Compound 6	High	No	No	No	No	Yes	No	Yes
Compound 7	High	No	No	No	No	Yes	No	Yes
Compound 8	Low	No	No	No	Yes	No	No	No
Compound 9	High	Yes	No	Yes	Yes	No	No	No
Compound 10	High	No	No	Yes	No	No	No	No
Compound 11	High	Yes	No	Yes	Yes	Yes	No	No

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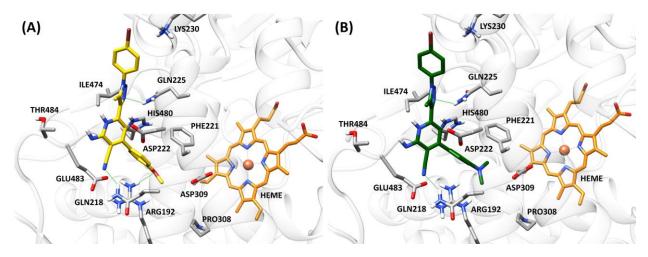


Figure S1. Docked poses of **(A)** compound **6** and **(B)** compound **7** in the potential allosteric site 1 of the enzyme.

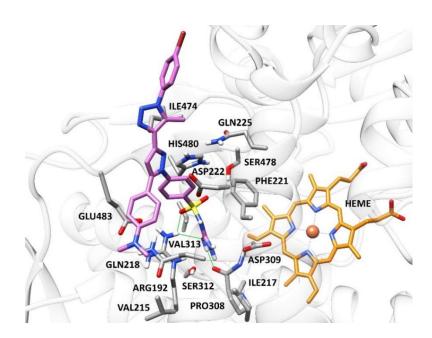


Figure S2. Compound 8 in the allosteric site 1 of the enzyme.

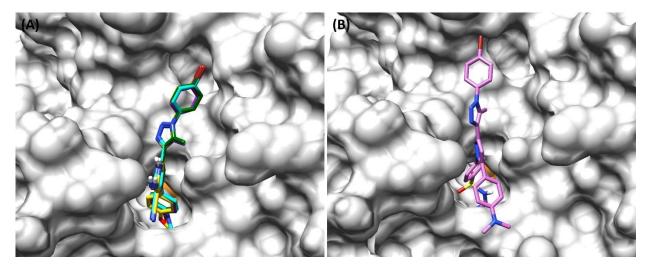


Figure S3. Potential allosteric site 1 of the enzyme with bound ligands: (A) Overlaid compounds 4 (purple), 5 (cyan), 6 (gold), and 7 (forest green). (B) Compound 8 (orchid).

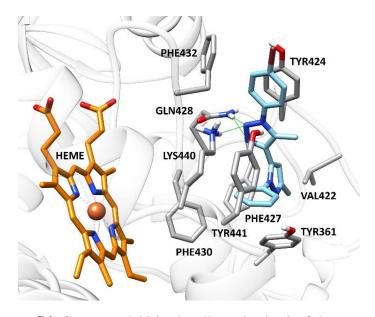


Figure S4. Compound **11** in the allosteric site 2 of the enzyme.

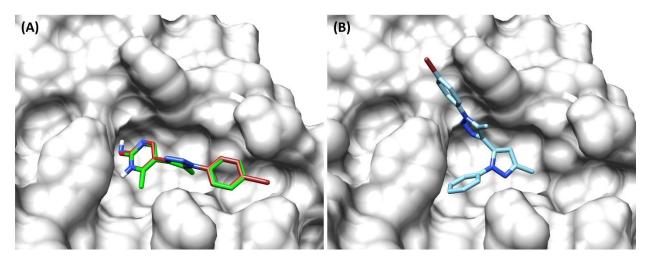


Figure S5. Potential allosteric site 2 of the enzyme with bound ligands: **(A)** Overlaid compounds **9** (green) and **10** (brown). **(B)** Compound **11** (sky blue).