

Supplementary Material

Mollusc Derived Brominated Indoles for the Selective Inhibition of Cyclooxygenase: A Computational Expedition

Md. Mominur Rahman ¹, Md. Junaid ², S. M. Zahid Hosen ^{2,3}, Mohammad Mostafa ², Lei Liu ⁴ and Kirsten Benkendorff ^{1,5,*}

¹ Marine Ecology Research Centre, Faculty of Science and Engineering, Southern Cross University, Lismore, NSW 2480, Australia; m.rahman.23@student.scu.edu.au

² Molecular Modeling Drug-design and Discovery Laboratory, Pharmacology Research Division, BCSIR Laboratories Chattogram, Bangladesh Council of Scientific and Industrial Research, Chattogram, Bangladesh; md.junaid@northsouth.edu (M.J.); drmostafabcsir@yahoo.com (M.M.)

³ Pancreatic Research Group, South Western Sydney Clinical School, and Ingham Institute for Applied Medical Research, Faculty of Medicine, University of New South Wales, Australia; s.hosen@student.unsw.edu.au

⁴ Southern Cross Plant Science, Southern Cross University, Lismore, NSW 2480, Australia; ben.liu@scu.edu.au

⁵ National Marine Science Centre, Southern Cross University, Coffs Harbour, New South Wales, Australia; kirsten.benkendorff@scu.edu.au

* Correspondence: kirsten.benkendorff@scu.edu.au

Table S1. Summary of nonbonding interactions analysis for Cyclooxygenase-1 (PDB ID: 3N8X), for the reference molecule aspirin and four *D. orbita* compounds.

Compounds Name	Hydrogen Bond				Bonding type	Hydrophobic Bond			Bonding type	Electrostatic Bond (EB) / Halogen Bond (HB) / Other Bond (OB)		
	Bonding type	Protein	Ligand	Distance		Protein	Ligand	Distance		Protein	Ligand	Distance
		Interacting amino acids	Interact- ing atoms or ring			Interacting amino ac- ids	Interact- ing atoms or ring			Interacting amino acids	Interact- ing atoms or ring	
Aspirin	Conven- tional	Arg-120	H...O	2.04	Pi-Alkyl	Leu-352	X	5.07				
		Arg-120	H...O	2.44		Ile-523	X	4.70				
	Carbon Hydro- gen	Ser-530	H...O	2.97		Ala-527	X	4.49				
Tyrindoxyl Sul- fate	Conven- tional	Arg-120	H...O	2.41	Amide Pistacked	Gly-526	X1	4.47	Attrac- tive Charge	Arg-120	H...O (EB)	2.13
	Carbon Hydro- gen	Arg-120	H...O	2.13		Ala-527	X1	4.47				
	Salt Bridge	Ser-353	H...O	2.76	Alkyl	Ile-523	X3	3.83	Pi-Sulfur	His-90	X3 (OB)	5.87
					Pi-Alkyl	Leu-352	X2	5.04				
						Ile-523	X2	4.56				
						Ala-527	X2	4.27				
Tyrindoleninone	Carbon Hydro- gen	Ile-523	H...O	2.73	Amide Pistacked	Gly-526	X1	4.87	Pi-Sulfur	Met-522	X1 (OB)	5.98
						Ala-527	X4	4.87				
						Gly-526	X4	3.96				
						Ala-527	X1	3.96				
					Alkyl	Ile-523	X3	4.77				
						Tyr-355	X3	4.19				
						Val-349	X4	5.21				
						Leu-352	X4	4.82				
					Pi-Alkyl	Ala-527	X4	3.93				
						Leu-352	X1	5.18				
						Ala-527	X1	4.71				
6-Bromoisatin					Amide Pi- stacked	Gly-526	X5	3.88				
						Ala-527	X1	3.88				
						Gly-526	X1	4.97				
						Ala-527	X5	4.97				
					Pi-Alkyl	Leu-352	X5	5.17				
						Ala-527	X5	4.38				
						Val-349	X1	5.05				
						Leu-352	X1	5.04				
						Ala-527	X1	3.71				
						Gly-526	X6	4.83	Attrac- tive charge	Arg-120	N-H...O (EB)	5.41
6,6' dibromoindi- rubin	Carbon Hydro- gen	Ser-353	H...O	2.59	Amide Pi- stacked	Ala-527	X1	4.83				
						Gly-526	X1	3.96				
						Ala-527	X6	3.96				
						Ala-527	X6	3.96				
	Pi-donor Hydro- gen	Tyr-355	H...O	2.59	Alkyl	Leu-384	Br	4.86	Pi-An- ion	Tyr-355	O (EB)	4.20
						Phe-381	Br					
						Tyr-385	Br					
						Trp-387	Br					
6,6' dibromoindi- rubin (Continue)					Pi-Alkyl	Val-349	X6					
						Ala-527	X6					
						Val-349	X6					
						Leu-352	X6					
						Ile-523	X6					
						Ala-527	X6					
						Val-349	X1					
						Leu-359	X1					
						Ala-527	X1					
						Leu-531	X1					
						Leu-352	X1					

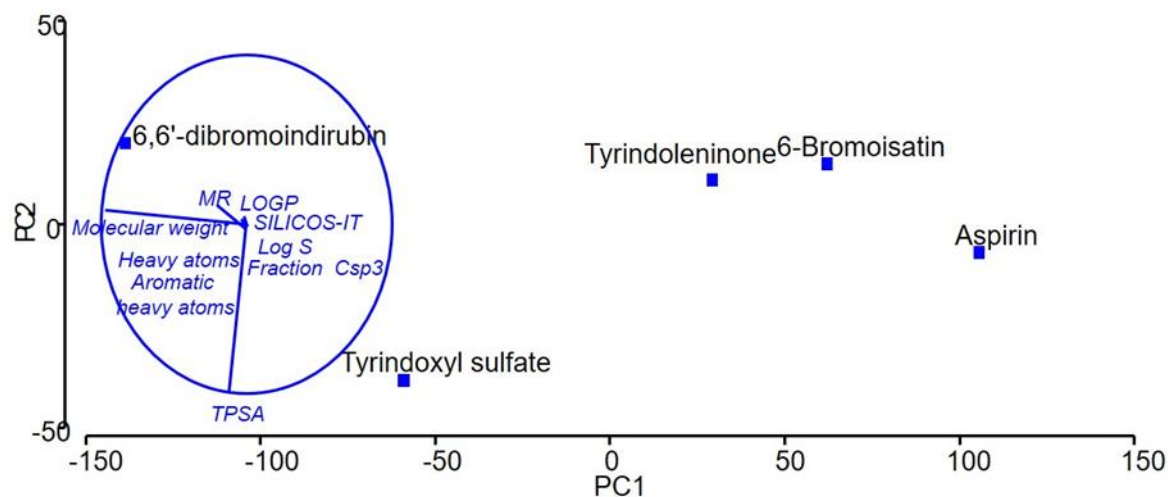
Here, X, X1, X2, X3 indicates that, X = Benzene, X1 = 6-Bromo benzene, X2 = Indole-3-ol, X3 = (Methylthio)methane, X4 = Indole-3-one, X5 = Indoline-2,3-dione, X6 = Indolin-2-one.

Table S2. Summary of nonbonding interactions analysis for Cyclooxygenase-2 (PDB ID: 5IKR), for the reference molecule aspirin and four *D. orbita* compounds.

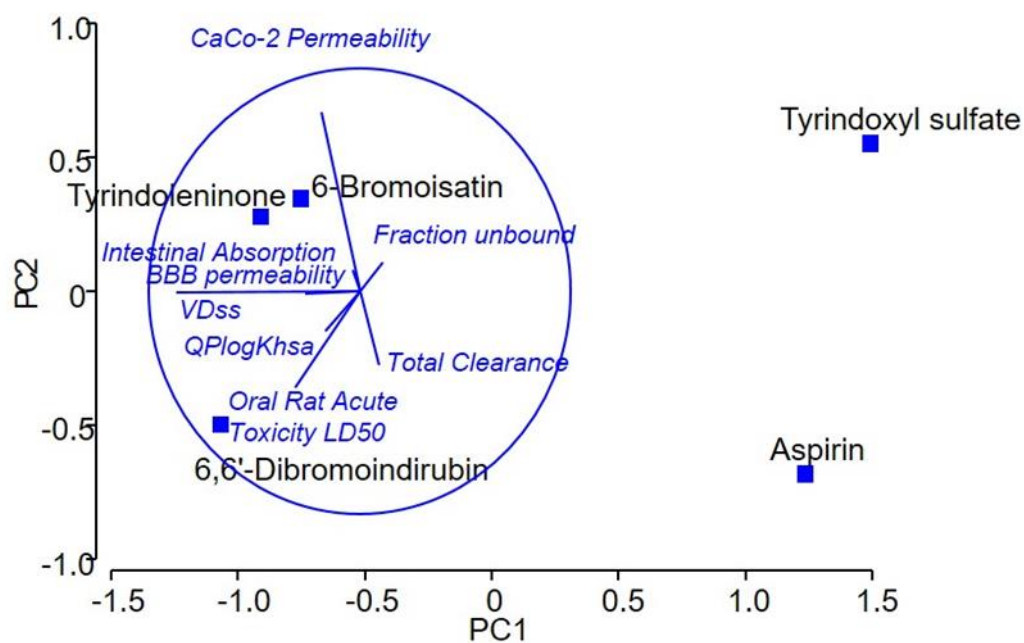
Compounds Name	Hydrogen Bond				Hydrophobic Bond				Electrostatic Bond (EB) / Halogen Bond (HB) / Other Bond (OB)											
	Bonding Type	Protein	Ligand	Dis - tance	Bonding Type	Protein	Ligand	Distance	Bonding Type	Protein	Ligand	Distance								
		Inter- acting Amino Acids	Interact- ing Atoms or Rings	Interact- ing Amino Acids		Interacting Atoms or Rings	Inter- acting Amino Acids	Interact- ing Atoms or Rings												
Aspirin	Conventional Hydrogen	Tyr-385	H...O	1.79	Amide Pi-stacked	Gly-526	X	3.76												
		Ser-530	H...O	1.83		Ala-527	X	3.76												
		Pi-Alkyl	Leu-352	X	5.13															
			Ala-527	X	4.82															
Tyrindoxyl Sulfate	Conventional Hydrogen	Tyr-385	H...O	2.11	Pi-Alkyl	Val-349	X2	5.03	Pi-Sulfur	Trp-387	X3 (OB)	5.72								
		Ser-530	H...O	2.22		Leu-352	X2	4.85		Trp-387	O (OB)	5.83								
		Carbon Hydrogen	Ser-530	H...O		2.82	Val-523	X1		5.38										
	Ser-530		H...O	2.50		Ala-527	X1	3.57												
	Ser-530		H...O	2.93		Val-349	X	4.25												
		Ala-527	X	3.47																
		Leu-531	X	4.12																
		Conventional Hydrogen	Ser-530	H...O	2.10	Amide Pi-stacked Alkyl	Gly-526	X1	4.12	Halogen	Met-522	Br	3.12							
							Ala-527	X4	4.12											
							Gly-526	X4	3.80											
							Ala-527	X1	3.80											
	Hydrogen		Ser-530	H...O	2.71	Alkyl	Ala-527	X3	3.63											
				Val-349	X3		4.39													
				Leu-531	X3		3.72													
					Pi-Alkyl	Val-349	X4	5.15												
						Leu-352	X4	4.83												
						Ala-527	X4	3.92												
						Leu-352	X1	5.17												
						Ala-527	X1	5.14												
6-Bromoisatin	Conventional Hydrogen	Ser-530	H...O	2.12	Amide Pi-stacked	Gly-526	X1	4.12	Halogen	Met-522	Br	3.13								
						Ala-527	X4	4.12												
						Gly-526	X4	3.80												
						Ala-527	X1	3.80												
	Carbon Hydrogen	Ser-530	H...O	2.70	Pi-Alkyl	Val-349	X5	5.10												
						Leu-352	X5	4.78												
						Ala-527	X5	3.92												
						Leu-352	X1	5.14												
6,6′dibromoindi- rubin	Carbon Hydrogen	Ser-530	H...O	2.92	Pi-pi T shaped	Tyr-355	X1	4.74												
						Gly-526	X6	4.95												
	Pi-donor Hydrogen	Arg-120	H...O	3.00	Amide pi- stacked	Ala-527	X6	4.95												
						Gly-526	X1	4.06												
						Ala-527	X1	4.06												
						Leu-352	Br	5.09												
					Alkyl	Met-522	Br	4.95												
						Trp-387	Br	5.22												
					Pi-Alkyl	Phe-518	Br	4.46												
						Val-116	X6	4.86												

	Val-349	X6	5.35
	Leu-359	X6	4.98
	Ala-527	X6	4.93
	Leu-531	X6	4.71
	Val-349	X6	4.32
	Leu-352	X1	4.81
	Ala-527	X1	3.97

Here, X, X1, X2, X3 indicates that, X = Benzene, X1 = 6-Bromo benzene, X2 =Indole-3-ol, X3 = (Methylthio)methane, X4 = Indole-3-one, X5 = Indoline-2,3-dione, X6 = Indolin-2-one.



a)



b)

Figure S1. Principle coordinate ordination showing the difference in a) physicochemical properties and b) pharmacokinetic properties of the brominated indoles from *Dicathais orbita*.