



Covalent Docking for Novel Series of Spiro-butenolides on Trx R1 for QSAR Modelling

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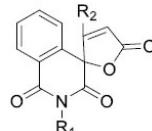
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Table 1. Structure for modeling.

#	R ₁	R ₂	R ₃	R ₄
1	4-F-Ph-	-H	-H	-H
2	Ph-	-H	-H	-H
3	CH ₃ -	-H	-H	-H
4	4-Me-Ph-	-H	-H	-H
5	4-OMe-Ph	-H	-H	-H
6	3-CF ₃ -Ph	-H	-H	-H
7	2-OMe-Ph	-H	-H	-H
8	(CH ₃) ₂ -CH-	-H	-H	-H
9	4-OMe-Ph-CH ₂	-H	-H	-H
10	(CH ₃) ₂ N-	-H	-H	-H
11	3,4-(OMe) ₂ -Ph-CH ₂ -CH ₂	-H	-H	-H
12	4-Cl-Ph-	-H	-F	-H
13	Ph-	-H	-Cl	-H
14	Ph	Me	-H	-H
15	Ph-	Ph-	-H	-H
16	Ph-	(CH ₂) ₂ CH ₃	-H	-H



#	R ₁	R ₂	R ₃	R ₄
17	Ph	Me	-H	-F
18	4-F-Ph	4-F-Ph	-H	-H
19	4-Me-Ph	4-OMe-Ph	-H	-H
20	(CH ₃) ₂ -CH-	2-Cl-Ph	-H	-H

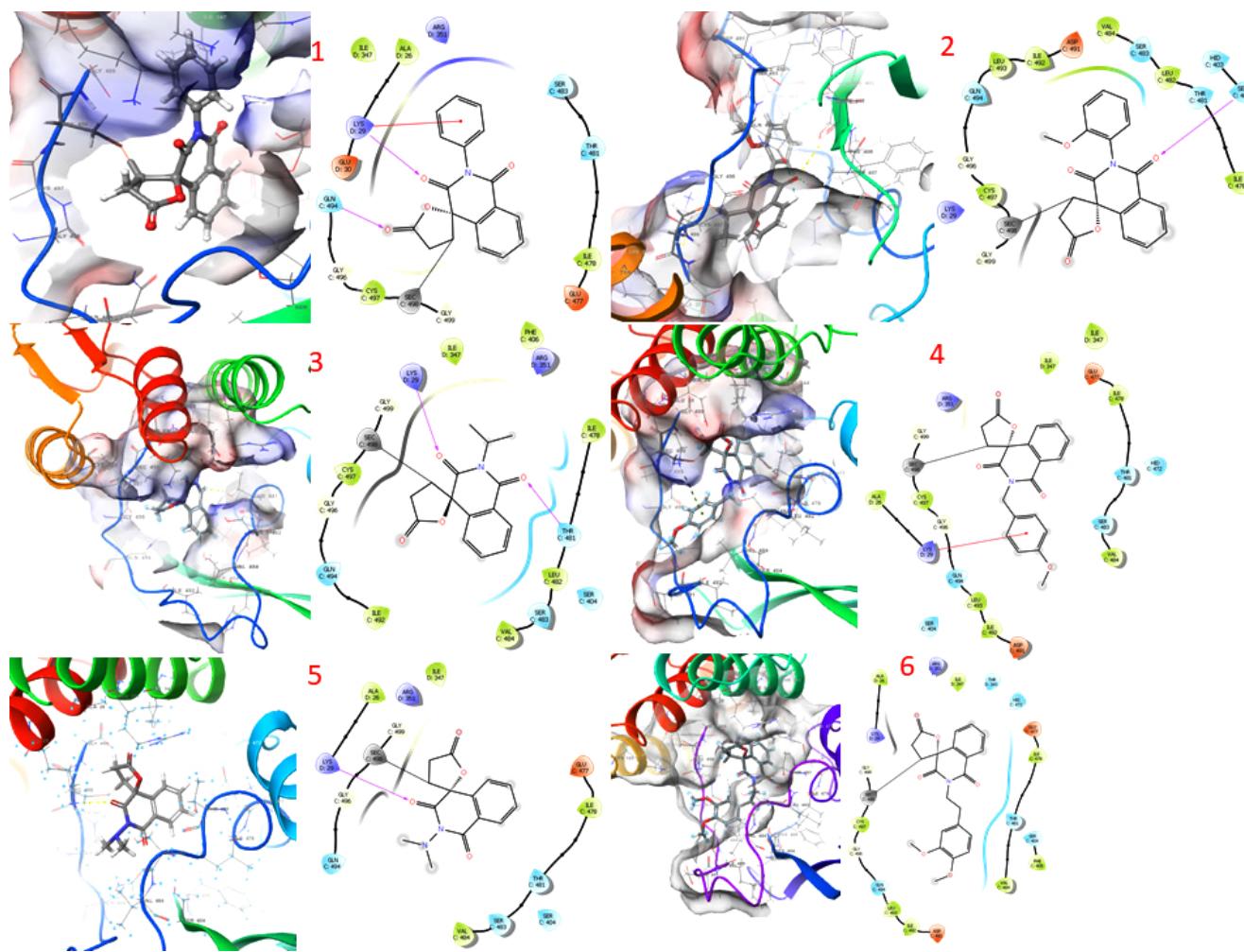


Fig. (1). Visualization of docking results.