

Discovery of NAD(P)H:quinone oxidoreductase 1 (NQO1) inhibitors with novel chemical scaffolds by shape-based virtual screening combined with cascade docking

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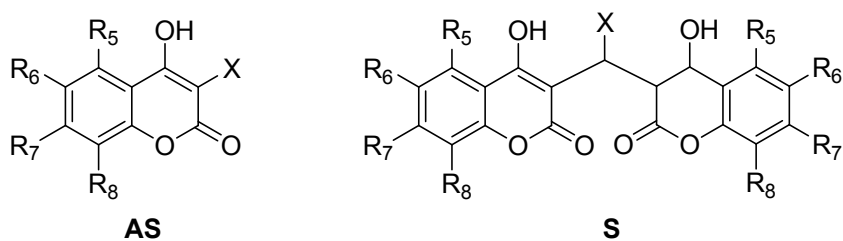
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

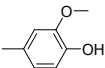
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Table S1

Structures of NQO1 inhibitors used to develop the active set for ROCS model and docking model. (Biological values used in this study were from references 1 and 2).



Cmpd	R5	R6	R7	R8	X	pIC ₅₀ (M)
S1	H	H	H	H	H	6.39
S2	H	CH ₃	H	H	H	5.25
S3	OCH ₃	H	H	H	H	7.42
S4	H	OCH ₃	H	H	H	5.48
S5	H	F	H	H	H	5.80
S6	H	CH ₃	CH ₃	H	H	6.63
S7	H	H	7,8-C ₄ H ₄		H	6.43
S8	H	H	CH ₃	CH ₃	H	7.02
S9	H	H	OH	H	H	6.46
S10	H	H	H	H		5.30
S11	H	H	H	H		5.23
S12	H	H	H	H		6.19

AS1	H	CH ₃	CH ₃	H	1-naphthyl	5.96
AS2	H	CH ₃	CH ₃	H	2-naphthyl	6.78
AS3	H	H	7,8-C ₄ H ₄		2-naphthyl	6.59
AS4		5,6-C ₄ H ₄	H	H	1-naphthyl	6.21
AS5		5,6-C ₄ H ₄	H	H	Phenyl	6.33
AS6	H	CH ₃	CH ₃	H	3,4-dimethyl phenyl	6.72

Reference

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