

Challenges Predicting Ligand-Receptor Interactions of Promiscuous Proteins:

The Nuclear Receptor PXR

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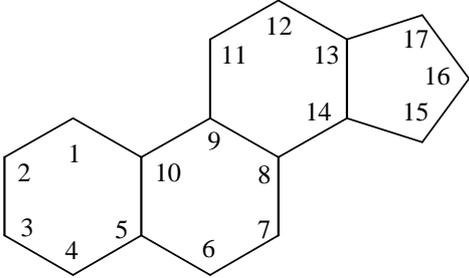
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Table S9. Three-ordered atom alignments (based on the steroidal core) used in the 4D- QSAR analysis.

				
Alignment	Atom 1	Atom 2	Atom 3	
1	1	3	5	
2	5	9	7	
3	1	6	12	
4	9	12	14	
5	2	7	17	
6	3	10	7	
7	14	13	17	
8	5	8	13	
9	8	13	15	
10	5	12	15	