

Supporting Information

Structure based prediction of subtype-selectivity for Adenosine receptor antagonists.

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Supporting Information includes:

- Figure SM1: full-length alignment for AR subfamily
- Figure SM2: set of 88 subtype specific AR antagonists used in the study
- A zip file with 88 ligand complexes with corresponding AR subtype models (PDB format).

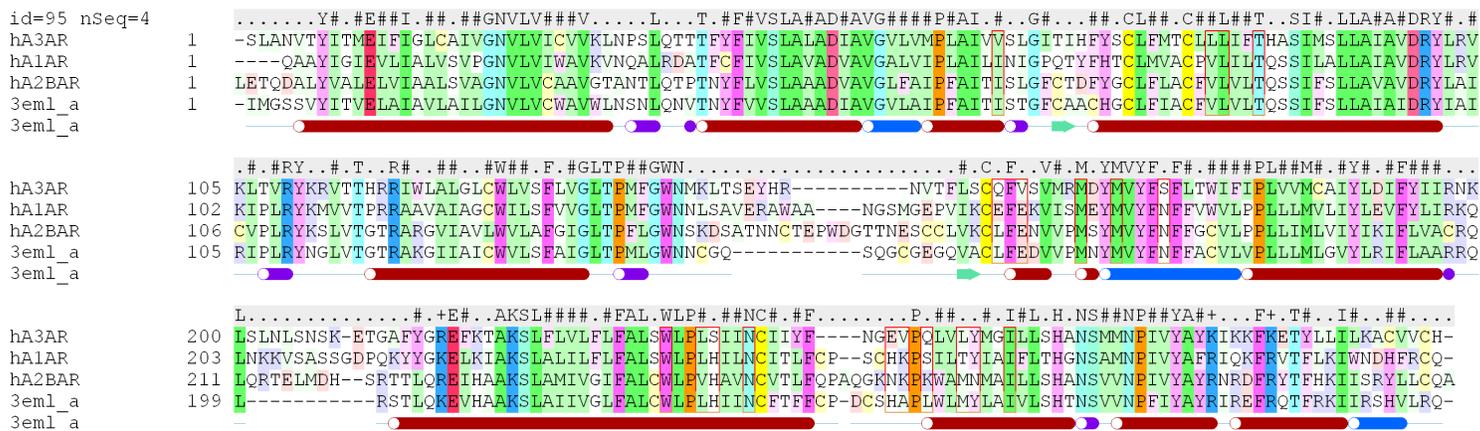
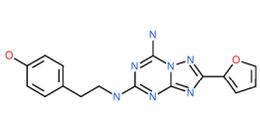
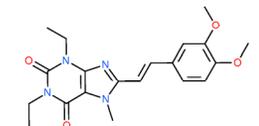
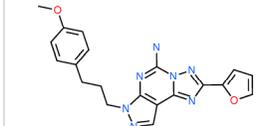
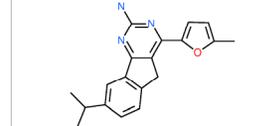
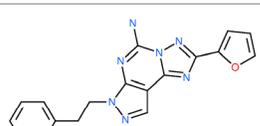
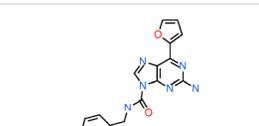
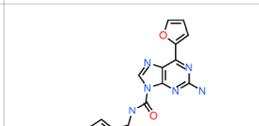
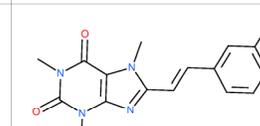
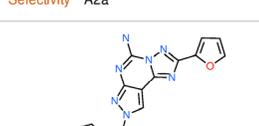
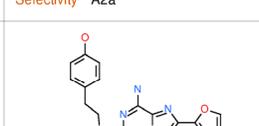
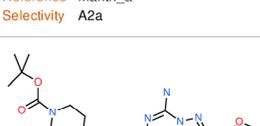
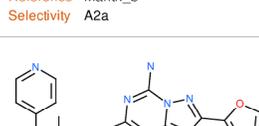
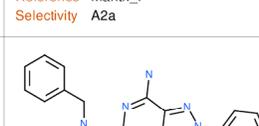
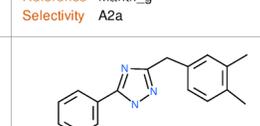
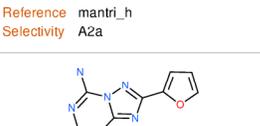
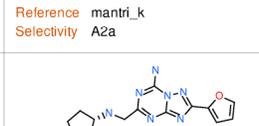
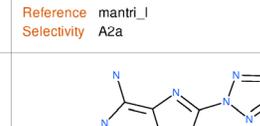
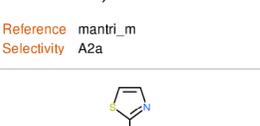
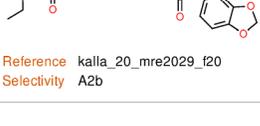
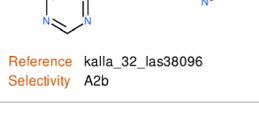
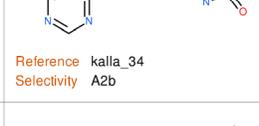
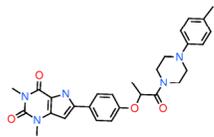
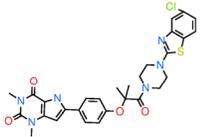
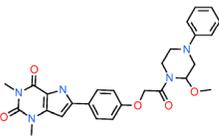
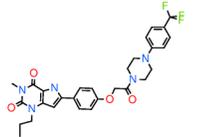
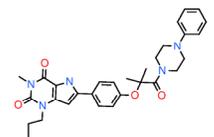
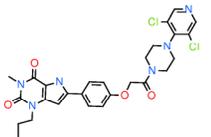
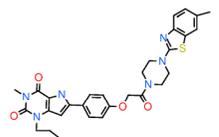
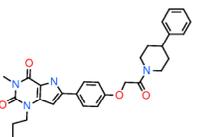
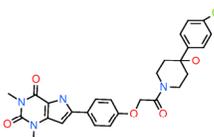
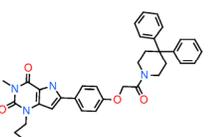
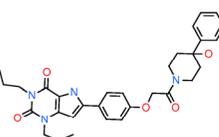
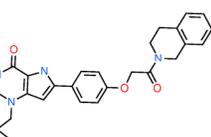
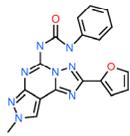
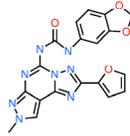
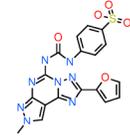
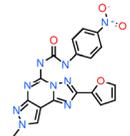
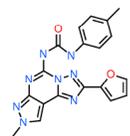
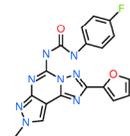
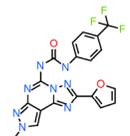
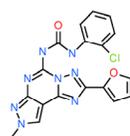
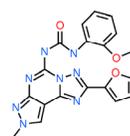
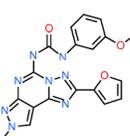
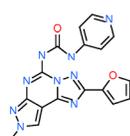
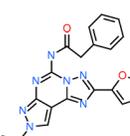
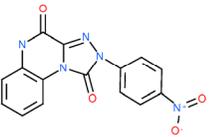
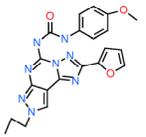
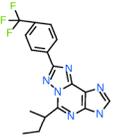


Figure SM1. Protein sequence alignment of all four human adenosine receptors, used to generate homology models. Red boxes highlight residues of the binding pocket within 6Å distance from ZM241385 compound in A2AR crystal structure (PDB code 3eml).

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 <p>Reference jac0_sch58261 Selectivity A2a</p>	 <p>Reference jac0_ver6947 Selectivity A2a</p>	 <p>Reference jac0_ver7835 Selectivity A2a</p>	 <p>Reference jac0_csc Selectivity A2a</p>
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 <p>Reference stef_11 Selectivity A2b</p>	 <p>Reference stef_23 Selectivity A2b</p>	 <p>Reference stef_29 Selectivity A2b</p>	 <p>Reference stef_52 Selectivity A2b</p>
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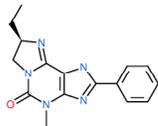
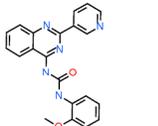
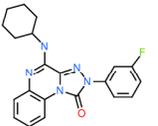
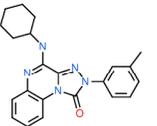
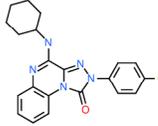
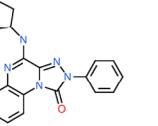
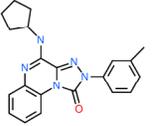
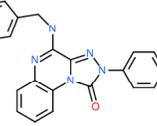
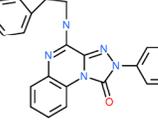
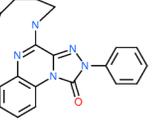
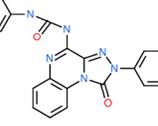
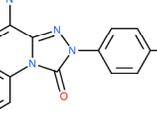
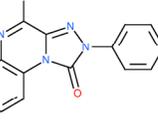
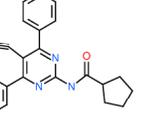
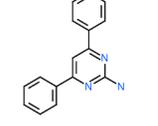
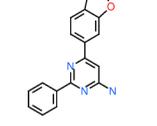
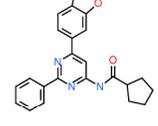
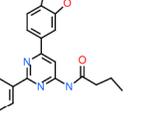
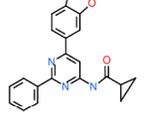
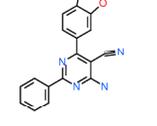
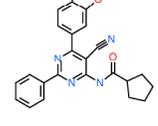
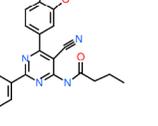
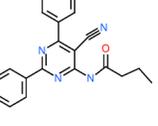
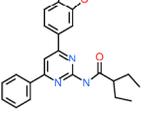
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 Reference veld_44 Selectivity A1	 Reference veld_45 Selectivity A1	 Reference veld_47 Selectivity A1	 Reference veld_52 Selectivity A1
 Reference veld_53 Selectivity A1	 Reference veld_54 Selectivity A1	 Reference veld_57 Selectivity A1	 Reference veld_7 Selectivity A1

Figure SM2. The set of 88 known subtype specific AR antagonists used in the study, comprised of 22 compounds for each AR subtype. The names of the compounds include the source reference and the compound number in the reference study, as well as the common name of the compound, if known. The reference codes are as follows: jaco¹, mantri², veld³, coll⁴, bara⁵, kalla⁶ and stef⁷.

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