

## BINDING THERMODYNAMICS AT THE HUMAN A<sub>2B</sub> ADENOSINE RECEPTOR

<u>Eleonora Fogli<sup>1</sup></u>, Stefania Gessi<sup>1</sup>, Katia Varani<sup>1</sup>, Stefania Merighi<sup>1</sup>, Valeria Sacchetto<sup>1</sup>, Carolina Simioni<sup>1</sup>, Edward Leung<sup>2</sup>, Stephen Mac Lennan<sup>2</sup>, Pier Andrea Borea<sup>1</sup>

<sup>1</sup>Department of Clinical and Experimental Medicine, Pharmacology Unit and Interdisciplinary Center for the Study of Inflammation, University of Ferrara, Italy; <sup>2</sup>King Pharm. Research and Development, Inc., Cary, North Carolina

The thermodynamic parameters  $\Delta G^{\circ}$ ,  $\Delta H^{\circ}$  and  $\Delta S^{\circ}$  of the binding equilibrium of four adenosine receptor agonists and five antagonists at adenosine A<sub>2B</sub> receptors were determined by means of affinity measurements at six different temperatures (4, 10, 15, 20, 25 and 30) and van't Hoff plots were constructed. Affinity constants were measured on human embryonic kidney (HEK 293) cells transfected with the human A<sub>2B</sub> receptors by inhibition assays of the binding of the selective A<sub>2B</sub> antagonist [<sup>3</sup>H]MRE 2029F20. Van't Hoff plots were linear for agonists and antagonists in the temperature range 4-30 degree. The thermodynamic parameters of radioligand were  $\Delta H^{\circ}$  –16 kJmol<sup>-1</sup> and  $\Delta S^{\circ}$  106 J(K/mol)<sup>-1</sup>, showing that antagonist binding is enthalpy- and entropy-driven. This binding behaviour has previously been found to be typical of adenosine A<sub>1</sub>, A<sub>2A</sub> and A<sub>3</sub> receptor antagonists. The results are discussed with the aim of obtaining new details on the nature of the forces driving the A<sub>2B</sub> binding at a molecular level.