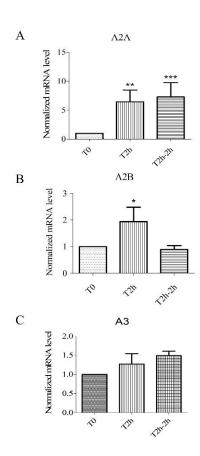
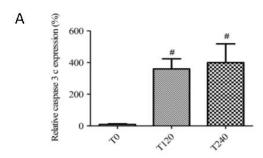
Supplemetary Table 1: List of the primers used in the real time PCR experiements

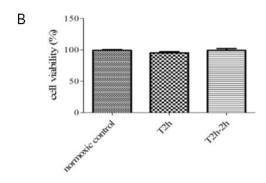
Gene	Primer sequence
EEF2	Forward: GGCCCTCTTATGATGTATATTTCC
	Reverse: CTGACCTTCAGGCCAGT
424	Forward: TCGCCATTGACCGCTAC
AZA -	Reverse: CCAGCAGATGGCAATGAT
A2B -	Forward: CATCTGTGTCCCGCTCA
AZD -	Reverse: CCACCCAGGAATGGA
A3 -	Forward: AGAGCTGATTGTAACTGACG
A3 -	Reverse: GACAACTTTGGGAGCTT

Supplementary data



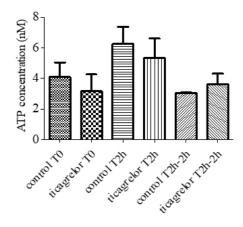
Complement of Figure 1 : mRNA expression after 2 hours of hypoxia and 2 hours of hypoxia followed by 2 hours of reoxygenation for A2A, A2B and A3 receptors.



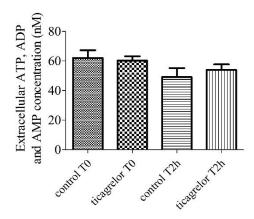


Complement of Figure 2 : Apoptosis and cell viability after 2 hours of hypoxia (T2h) followed by 2 hours of reoxygenation (T2h-2h), determined by the relative expression of cleaved caspase 3 by immunoblotting (A) and PrestoBlue assays (B). Results are expressed as means \pm sem (n = 6/group). #: p < 0.05, compared to normoxic control.

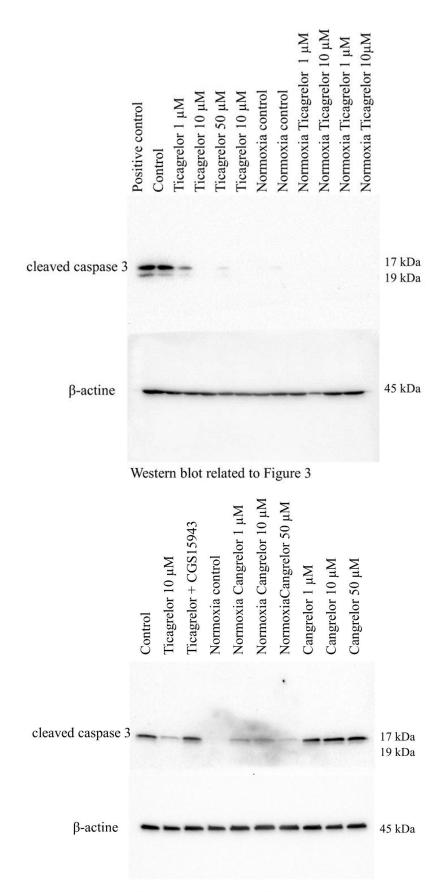
Extracellular ATP concentrations estimated using LC-HRMS assay.



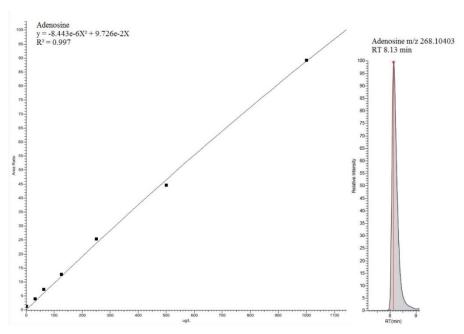
Anova p >0.1 for difference between groups



Anova p >0.1 for difference between groups



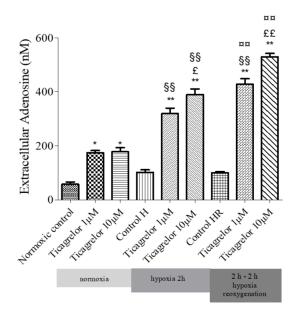
Western blot related to Figure 3



Quantification using liquid chromatography coupled with high resolution mass spectrometry:

Calibration curve of adenosine, coefficient of correlation, retention time, high-resolution masses.

Related to Figure 4



Complement of Figure 4: Extracellular adenosine concentration: effect of 1 μ M and 10 μ M ticagrelor during a 2 hours of hypoxia followed by 2 hours of reoxygenation stress in HUVECs. Extracellular adenosine concentrations are expressed in nM. Results are expressed as means \pm sem (n = 6/group). *: p < 0.05, **: p < 0.01, compared to the control groups of the corresponding times; £: p < 0.05, ££: p < 0.01 compared to ticagrelor normoxia for each corresponding concentration. **\maxxix*: p < 0.01 compared to ticagrelor hypoxia for each corresponding concentration.

