| | Base model (37 ℃) | MAP model (4℃) | PAGGGM model (4℃) |
|-----------------------------------|--|--|----------------------------|
| Enzymatic activities | 100% | 24% | 25% |
| Purine metabolism activities | 100% | 34% | 30% |
| Sodium/pottasium pump activity | 100% | 6.0% | 1.0% |
| Extracellular glucose | 5 mM (fixed) | 40 mM | 47.5 mN |
| Extracellular adenine | 0.015 mM (fixed) | 1.04 mM | 1.44 mN |
| Extracellular inorganic phosphate | 1.1 mM (fixed) | 6 mM | 16 mM |
| Extracellular sodium ion | 135 mM (fixed) | 106 mM | 64 mN |
| Extracellular guanosine | - | - | 1.44 mM |
| Guanosine phosphorylation | - | - | First-order kinetic |
| pH setting | pH = 7.2 (fixed) | Initial pH = 6.95, | Initial $pH = 7.62$ |
| | | | $pH(t) = at^2 + bt + 7.6$ |
| | | $pH(t) = at + 6.95$ $a = -8.66 \times 10^{-8}$ | $a = 8.79 \times 10^{-1}$ |
| | | | $b = -6.01 \times 10^{-3}$ |
| State of hemoglobin | Transition between T- and R-state | R-state | R-state |
| | in response to various parameters; | | |
| | pO ₂ , pCO ₂ , pH, temperature and | | |
| | 2,3-BPG | | |