

**Table S6**, parameter settings for the simulations of post exercise recovery dynamics (Figure 6)

Control dataset

Model configuration	Parameter name					
	ATPase demand flux (mmol (L cell water) <sup>-1</sup> )	Mitochondrial volume (mL mito / mL cell)	End exercise pH (unitless)	Cr <sub>tot</sub> (mmol (L cell water) <sup>-1</sup> )	Pi initial (mmol (L cell water) <sup>-1</sup> )	λ (mM s <sup>-1</sup> pH unit <sup>-1</sup> )
Original model	0.90	0.056	6.65	41.2	4.0	0.33
Model configuration 1 (substrate feedback)	0.90	0.056	6.65	41.2	4.0	0.33
Model configuration 2 (parallel activation)	0.90	0.056	6.65	41.2	4.0	0.12

Athlete dataset

Model configuration	Parameter name					
	ATPase demand flux (mmol (L cell water) <sup>-1</sup> )	Mitochondrial volume (mL mito / mL cell)	End exercise pH (unitless)	Cr <sub>tot</sub> (mmol (L cell water) <sup>-1</sup> )	Pi initial (mmol (L cell water) <sup>-1</sup> )	λ (mM s <sup>-1</sup> pH unit <sup>-1</sup> )
Original model	1.1	0.085	6.95	38.2	4.0	1.33
Model configuration 1 (substrate feedback)	1.1	0.085	6.95	38.2	4.0	1.33
Model configuration 2 (parallel activation)	1.1	0.080	6.95	38.2	4.0	1.33

Subject with sedentary lifestyle dataset

Model configuration	Parameter name					

	ATPase demand flux (mmol (L cell water) <sup>-1</sup> )	Mitochondrial volume (mL mito / mL cell)	End exercise pH (unitless)	Cr <sub>tot</sub> (mmol (L cell water) <sup>-1</sup> )	Pi initial (mmol (L cell water) <sup>-1</sup> )	$\lambda$ (mM s <sup>-1</sup> pH unit <sup>-1</sup> )
Original model	0.35	0.0235	6.70	47.0	4.0	0.1
Model configuration 1 (substrate feedback)	0.35	0.0235	6.70	47.0	4.0	0.1
Model configuration 2 (parallel activation)	0.42	0.030	6.70	47.0	4.0	0.017