

Kinetic description of the dynamic model of oxidative phosphorylation in isolated skeletal muscle mitochondria respiring on pyruvate.

Subscripts: e, external (suspension); i, internal (mitochondrial); t, total; f, free; m, magnesium complex; j, monovalent; a.u., arbitrary units.

Respiration rate (v_{C4}) scaled to 100 a.u. in state 3. All metabolite concentrations in μM .

DH, NADH supply; C1, complex I; C3, complex III; C4, complex IV; SN, ATP synthase; EX, ATP/ADP carrier; PI, P_i carrier; UT, ATP usage; LK, proton leak; AK, adenylate kinase.

Constants

$$k_{DH} = 997.5 \text{ a.u.}$$

$$K_{mN} = 100$$

$$p_D = 0.8$$

$$k_{C1} = 8.287 \text{ a.u.}$$

$$k_{C3} = 4.731 \text{ a.u.}$$

$$k_{C4} = 0.1215 \text{ a.u.}$$

$K_{mO} = 120 \mu\text{M}$ (mechanistic K_m for O_2 , much higher than apparent K_m)

$$k_{SN} = 1070 \text{ a.u.}$$

$n_A = 2.5$ (phenomenological H^+ /ATP stoichiometry of ATP synthase)

$$k_{EX} = 2112 \text{ a.u.}$$

$$K_{mADP} = 3.5 \mu\text{M}$$

$$k_{PI} = 1.502 \text{ a.u.}$$

$$pK_a = 6.8$$

$$k_{UT} = 0 \text{ a.u. (state 4), } 275 \text{ a.u. (state 3.5, 50 \% of state 3 respiration), } 900 \text{ a.u. (state 3)}$$

$$K_{mA} = 150 \mu\text{M}$$

$$k_{LK1} = 0.1456 \text{ a.u.}$$

$$k_{LK2} = 0.038 \text{ mV}^{-1}$$

$$k_{fAK} = 139.2 \text{ a.u.}$$

$$k_{bAK} = 3.673 \text{ a.u.}$$

$$K_{DTe} = 24 \mu\text{M} \quad (\text{magnesium dissociation constant for external ATP})$$

$$K_{DDe} = 347 \mu\text{M} \quad (\text{magnesium dissociation constant for external ADP})$$

$$K_{DTi} = 17 \mu\text{M} \quad (\text{magnesium dissociation constant for internal ATP})$$

$$K_{DDi} = 282 \mu\text{M} \quad (\text{magnesium dissociation constant for internal ADP})$$

$$R_{sm} = 340 \text{ (suspension volume/mitochondria volume ratio)}$$

$$B_N = 5 \text{ (buffering capacity coefficient for NAD)}$$

$$T = 298$$

$$R = 0.0083 \text{ kJ}\cdot\text{mol}^{-1}\cdot\text{K}^{-1}$$

$$F = 0.0965 \text{ kJ}\cdot\text{mol}^{-1}\cdot\text{mV}^{-1}$$

$$S = 2.303 \cdot R \cdot T$$

$$Z = 2.303 \cdot R \cdot T / F$$

$$u = 0.861 \quad (= \Delta\Psi / \Delta p)$$

$$c_{buffi} = 0.022 \text{ M H}^+/\text{pH unit} \quad (\text{buffering capacity for H}^+ \text{ in matrix})$$

$$\Delta G_{P0} = 31.9 \text{ kJ}\cdot\text{mol}^{-1}$$

$$\begin{aligned}E_{mN0} &= -320 \text{ mV} \\E_{mU0} &= 85 \text{ mV} \\E_{mc0} &= 250 \text{ mV} \\E_{ma0} &= 540 \text{ mV}\end{aligned}$$

Constant metabolite concentrations

$$O_2 = 240 \mu\text{M}$$

$$pH_e = 6.9$$

$$\begin{aligned}c_t &= 270 \mu\text{M} \quad (= c^{2+} + c^{3+}, \text{ total concentration of cytochrome c}) \\U_t &= 1350 \mu\text{M} \quad (= UQH_2 + UQ, \text{ total concentration of ubiquinone}) \\N_t &= 2970 \mu\text{M} \quad (= NADH + NAD^+, \text{ total concentration of NAD}) \\a_t &= 135 \mu\text{M}\end{aligned}$$

$$\begin{aligned}Mg_{fe} &= 4000 \mu\text{M} \quad (\text{free external magnesium concentration}) \\Mg_{fi} &= 380 \mu\text{M} \quad (\text{free internal magnesium concentration})\end{aligned}$$

$$\begin{aligned}A_{isum} &= 16260 \mu\text{M} \quad (= ATP_{ti} + ADP_{ti}, \text{ total internal adenine nucleotide concentration}) \\A_{esum} &= 2000 \mu\text{M} \quad (= ATP_{te} + ADP_{te} + AMP_e, \text{ total external adenine nucleotide concentration})\end{aligned}$$

Initial values of independent variables (μM), respiration rate (v_{C4} , a.u.) and AMP_e (μM)

State:	State 4 ($k_{UT} = 0$)	State 3.5 ($k_{UT} = 275$)	State 3 ($k_{UT} = 900$)
v_{C4} (= vO_2)	10.48	50.02	99.86
NADH	1643.7	425.4	180.4
UQH_2	1150.1	1160.3	1194.4
c^{2+}	54.913	63.137	58.152
H^+_i	0.0446	0.0492	0.0529
ATP_{ti}	15410	8662	1890
Pi_{ti}	56501	33460	27901
ATP_{te}	1999.868	1977.605	238.605
Pi_{te}	9939	10050	13239
ADP_{te}	0.132	21.726	347.130
AMP_e	0.0000244	0.668	1414.3

Calculations

$$\begin{aligned}c^{3+} &= c_t - c^{2+} \\UQ &= U_t - UQH_2 \\NAD^+ &= N_t - NADH\end{aligned}$$

$$\begin{aligned}AMP_e &= A_{esum} - ATP_{te} - ADP_{te} \\ADP_{ti} &= A_{isum} - ATP_{ti}\end{aligned}$$

$$\begin{aligned}ATP_{fe} &= ATP_{te}/(1+Mg_{fe}/k_{DTe}) \\ATP_{me} &= ATP_{te} - ATP_{fe} \\ADP_{fe} &= ADP_{te}/(1+Mg_{fe}/k_{DDe}) \\ADP_{me} &= ADP_{te} - ADP_{fe}\end{aligned}$$

$$\begin{aligned}ATP_{fi} &= ATP_{ti}/(1+Mg_{fi}/k_{DTi}) \\ATP_{mi} &= ATP_{ti} - ATP_{fi}\end{aligned}$$

$$\text{ADP}_{\text{fi}} = \text{ADP}_{\text{ti}}/(1+\text{Mg}_{\text{fi}}/\text{k}_{\text{DDi}})$$

$$\text{ADP}_{\text{mi}} = \text{ADP}_{\text{ti}} - \text{ADP}_{\text{fi}}$$

$$\text{pH}_i = -\log(H_i/10^6) \quad (H_i \text{ expressed in } \mu\text{M})$$

$$\Delta \text{pH} (\text{mV}) = Z \cdot (\text{pH}_i - \text{pH}_e)$$

$$\Delta \text{p} (\text{mV}) = 1/(1-u) \cdot \Delta \text{pH}$$

$$\Delta \Psi (\text{mV}) = -(\Delta \text{p} - \Delta \text{pH})$$

$$\Psi_i (\text{mV}) = 0.65 \cdot \Delta \Psi$$

$$\Psi_e (\text{mV}) = -0.35 \cdot \Delta \Psi$$

$$c_{0i} = (10^{-\text{pH}_i} - 10^{-\text{pH}_i - \Delta \text{pH}})/\Delta \text{pH} \quad (\text{'natural' buffering capacity for H}^+ \text{ in matrix})$$

$$\Delta \text{pH} = 0.001$$

$$r_{\text{buffi}} = c_{\text{buffi}}/c_{0i} \quad (\text{buffering capacity coefficient for H}^+ \text{ in matrix})$$

$$P_{i,e} = P_{i,e}/(1 + 10^{\text{pH}_i - \text{pK}_a})$$

$$P_{i,i} = P_{i,i}/(1 + 10^{\text{pH}_i - \text{pK}_a})$$

$$\Delta G_{SN} = n_A \cdot \Delta p - \Delta G_P \quad (\text{thermodynamic span of ATP synthase})$$

$$\Delta G_P = \Delta G_{P0}/F + Z \cdot \log(10^6 \cdot \text{ATP}_{ti}/(\text{ADP}_{ti} \cdot P_{i,i})) \quad (\text{concentrations expressed in } \mu\text{M})$$

$$E_{mN} = E_{mN0} + Z/2 \cdot \log(NAD^+/NADH) \quad (\text{NAD redox potential})$$

$$E_{mU} = E_{mU0} + Z/2 \cdot \log(UQ/UQH_2) \quad (\text{ubiquinone redox potential})$$

$$E_{mc} = E_{mc0} + Z \cdot \log(c^{3+}/c^{2+}) \quad (\text{cytochrome c redox potential})$$

$$E_{ma} = E_{mc} + \Delta p \cdot (2 + 2u)/2 \quad (\text{cytochrome a}_3 \text{ redox potential})$$

$$A_{3/2} = 10^{(E_{ma} - E_{ma0})/Z}$$

$$a^{2+} = a_t / (1 + A_{3/2})$$

(a^{3+}/a^{2+} ratio)

(concentration of reduced cytochrome a₃)

$$\Delta G_{C1} = E_{mU} - E_{mN} - \Delta p \cdot 4/2 \quad (\text{thermodynamic span of complex I})$$

$$\Delta G_{C3} = E_{mc} - E_{mU} - \Delta p \cdot (4 - 2u)/2 \quad (\text{thermodynamic span of complex III})$$

Kinetic equations

(All reaction rates are expressed in a.u.).

Substrate dehydrogenation:

$$v_{DH} = k_{DH} \frac{1}{\left(1 + \frac{K_{mN}}{NAD^+/NADH} \right)^{p_D}}$$

Complex I:

$$v_{C1} = k_{C1} \cdot \Delta G_{C1}$$

Complex III:

$$v_{C3} = k_{C3} \cdot \Delta G_{C3}$$

Complex IV:

$$v_{C4} = k_{C4} \cdot a^{2+} \cdot c^{2+} \frac{1}{1 + \frac{K_{mO}}{O_2}}$$

ATP synthase:

$$v_{SN} = k_{SN} \frac{\gamma - 1}{\gamma + 1},$$

$$\gamma = 10^{\Delta G_{SN}/Z}$$

ATP/ADP carrier:

$$v_{EX} = k_{EX} \cdot \left(\frac{ADP_{fe}}{ADP_{fe} + ATP_{fe} \cdot 10^{-\Psi_e/Z}} - \frac{ADP_{fi}}{ADP_{fi} + ATP_{fi} \cdot 10^{-\Psi_i/Z}} \right) \cdot \left(\frac{1}{1 + K_{mADP}/ADP_{fe}} \right)$$

Phosphate carrier:

$$v_{PI} = k_{PI} \cdot (Pi_{je} \cdot H_e - Pi_{ji} \cdot H_i)$$

ATP usage:

$$v_{UT} = k_{UT} \frac{1}{1 + \frac{K_{mA}}{ATP_{te}}}$$

Proton leak:

$$v_{LK} = k_{LK1} \cdot (e^{k_{LK2} \cdot \Delta p} - 1)$$

Adenylate kinase:

$$v_{AK} = k_{fAK} \cdot ADP_{fe} \cdot ADP_{me} - k_{bAK} \cdot ATP_{me} \cdot AMP_e$$

Set of differential equations

$$\dot{NADH} = (v_{DH} - v_{C1}) \cdot R_{sm} / B_N$$

$$\dot{UQH_2} = (v_{C1} - v_{C3}) \cdot R_{sm}$$

$$\dot{c^{2+}} = (v_{C3} - 2 \cdot v_{C4}) \cdot 2 \cdot R_{sm}$$

$$\dot{O_2} = 0 \quad (\text{constant saturated oxygen concentration} = 240 \mu\text{M}) \text{ or } \dot{O_2} = -v_{C4}$$

$$\dot{H_i^+} =$$

$$- (2 \cdot (2 + 2 \cdot u) \cdot v_{C4} + (4 - 2 \cdot u) \cdot v_{C3} + 4 \cdot v_{C1} - n_A \cdot v_{SN} - u \cdot v_{EX} - (1 - u) \cdot v_{PI} - v_{LK}) \cdot R_{sm} /$$

$$r_{buffi}$$

$$\dot{ATP}_t = (v_{SN} - v_{EX}) \cdot R_{sm}$$

$$\dot{Pi}_{ti} = (v_{PI} - v_{SN}) \cdot R_{sm}$$

$$\dot{ATP}_t = v_{EX} - v_{UT} + v_{AK}$$

$$\dot{ADP}_t = v_{UT} - v_{EX} - 2 \cdot v_{AK}$$

$$\dot{Pi}_{te} = v_{UT} - v_{PI}$$