



SUIT reference assay for OXPHOS analysis by high-resolution respirometry

Doerrier C, Sumbalova Z, Lamberti G, Krumschnabel G, Gnaiger E

OROBOROS INSTRUMENTS
high-resolution respirometry
Schöpfstr 18, A-6020 Innsbruck, Austria
Email: instruments@orooboros.at
www.orooboros.at

Contents

1. SUIT_RP1: CI-coupling/substrate control	2
2. Concept SUIT_RP2	5
3. Test experiments	7
3.1. SUIT RP1	7
3.2. SUIT RP2	7
4. Technical details	8
Supplement	1

Summary: A substrate-uncoupler-inhibitor titration (SUIT) reference assay (SUIT-RA) is developed to provide a common baseline for comparison of mitochondrial respiratory control in a large variety of species, tissues and cell types, mt-preparations and laboratories, for establishing a database on comparative mitochondrial physiology. The SUIT-RA is applied in the MitoFit proficiency test with HEK 293T cells. It includes a large number of chemicals used in various specific SUIT protocols, subjecting these chemicals under quality control in the MitoFit proficiency test.

The SUIT-RA consists of two coordinated SUIT reference protocols (SUIT-RP). These are harmonized such that they can be statistically evaluated as replicate measurement for carefully selected respiratory states, while additional information is obtained when the two protocols are conducted in parallel. Therefore, the two SUIT-RP are complementary with their focus on specific respiratory coupling and substrate control aspects, extending previous strategies for respirometric OXPHOS analysis.

SUIT reference protocols

RP1: 1PM 2D 2c (2NADH) 3U 4Oct 5G 6S 7Rot 8Gp 9Ama 10Tm 11Azd

RP2: 1D 2Oct 3MtIt 3c (3NADH) 4P 5G 6S 7U 8Gp 9Rot 10Ama 11Tm 12Azd

1. SUIT_RP1: CI(LPE) substrate control

RP1: 1PM 2D 2c (2NADH) 3U 4Oct 5G 6S 7Rot 8Gp 9Ama 10Tm 11Azd

<i>E</i>	3U	4Oct	5G	6S	7Rot	8Gp	9Ama	10Tm	11Azd
<i>P</i>	2D+c								
<i>L</i>	1PM								
	CI	CI&FAO	CI&FAO	CI&II&FAO	CII	CII&Gp	ROX	CIV	ROX

RP1 spotlights

- CI-linked linear coupling control: $L - P - E$, thus separating coupling control (CI-linked) and substrate control (in the ETS state).
- If Oct is without effect on (3U) PM_E , then we can conclude that it is also without effect on PGM_E and $PGMS_E$. On the other hand, if Oct added after $PGMS_E$ is without effect, then it is not clear if it would be without effect on PM_E and PGM_E .
- (3U) CI_E , (6S) $CI&II&FAO_E$ and (7Rot) CII_E are measured. If $CI&II&FAO_E \approx CI&II_E$, then this sequence allows calculation of the additivity index of CI- and CII- linked ETS capacity (related to supercomplex-channeling). This criterium is tested in step 4Oct, evaluating the effect of Oct on (3U) PM_E .
- Oct is added before G and S to compare RP1 with RP2: harmonization between protocols in states (RP1:6S, RP2:7U) $PGMSOct_E$, (RP1:8Gp, RP2:9Rot) SGp_E , and (10Tm or 11Tm) CIV_E .
- If Oct is without effect on PM_E (expected in many types of mt), then additional harmonization between protocols is obtained in states (RP1:2D) $PM_P = (RP2:4P) PMOct_P$.
- Harmonization with many previous protocols up to step 7Rot.

Limitations:

- Depletion of endogenous substrates with D is not possible, to obtain (1PM) CI_L without an inhibitor of ATP synthase (Omy).
- $PGMSOctGpDH_E$ is not obtained (substrate combination for maximum ETS capacity), in favour of measuring (7Rot) CII_E . This reference state has to be calculated using the $PGMSOct_E/PGMSOctGpDH_E$ ratio from RP2.

Substrate type: CI-CII-FAO-CGpDH-CIV

Substrate state category:

CI + CI&FAO + CI&II&FAO + CII + CII&GpDH + CIV

SUIT protocol name:

PM(LP)+c(PE)+Oct+G+S+Rot+Gp+Ama+AsTm+Azd

RP1-Pfi

Step	State	Comment
PM		CI-linked substrates are added to the medium before the mt-preparation (mt). The state without added substrates is not well defined, slightly higher than ROX due to the presence of some endogenous substrates (shown by a slight decline of respiration and mt-membrane potential upon inhibition by Rot; Krumschnabel et al 2014).
+mt		Incubation for about 20 min to allow stabilization of flux at high oxygen and during slow exhaustion of endogenous substrates, to obtain CI_L .
1PM	PM_L	CI-linked LEAK state.
2D	PM_P	OXPPOS coupling efficiency ($P-L$ or $\approx P$ control factor), $j_{\approx P} = \approx P/P = (P-L)/P = 1-L/P$, is measured in the CI-linked substrate state, with defined coupling sites (CI, CIII, CIV) and at high flux compared to FAO (human skeletal muscle: compare with FAO(LP) and CII(PL); Gnaiger et al 2015).
2c	PM_{C_P}	Quality control test early in the SUIT protocol. In pathological states with c release, early addition of c provides information on the $FCF_c = 1-CI/CI_c$, and separates the FCF_c from other injuries in the subsequent respiratory states. All subsequent states contain c, which is not explicitly written in the following substrate states.
2NADH		NADH is titrated only in case of a high cytochrome c control factor, $FCF_c > 0.1$, to check for the presence of inverted mitochondrial particles and permeability of the inner mt-membrane. If $FCF_c < 0.1$, then the high intactness of the outer mt-membrane provides an indication for intactness of the inner mt-membrane, and thus NADH does not have to be added.
3U	PM_E	CCCP is titrated stepwise to maximum flux, to evaluate limitation of OXPPOS by the phosphorylation system, expressed as the apparent excess $E-P$ capacity factor ($E-P$ coupling control factor), $j_{EXP} = (E-P)/E = 1-P/E$. If $j_{EXP} > 0$, then the ETS coupling efficiency rather than the OXPPOS coupling efficiency is the proper expression of coupling, $j_{\approx E} = \approx E/E = (E-L)/E = 1-L/E$.
4Oct	PM_{Oct_E}	$FCF_{Oct} = 1-CI/CI\&FAO$ low or zero in many mt-

		types. Inhibition is observed at higher FA concentrations. Then also state PM_p is identical to PM_{Oct_p} in RP2, and may thus further link the two protocols in the SUIT reference assay for statistical analysis (protocol harmonization).
5G	$PGMOct_E$	$FCR_G = 1 - PM_{Oct}/PGMOct$, reveals an additive effect of convergent electron flux through NADH (CI-linked), with a possible contribution by partially activating CII-linked respiration.
6S	$PGMSOct_E$	$FCF_{CII} = 1 - CI\&FAO/CI\&II\&FAO$. Additive effect of CI&II. It may be important if the uncoupler concentration titrated in the PM substrate state is also sufficient for this substrate state.
7Rot	S_E	$FCF_{CI} = 1 - CII/CI\&II\&FAO$. Rot inhibits CI and FAO simultaneously. Additive effect of $CI\&II\&FAO$. In some cases it takes very long, until a steady state is reached after inhibition by Rot. Addition of Gp before Rot would not allow a valid estimation of CII-linked capacity (compare RP2).
8Gp	SGp_E	CGpDH capacity is not measured in the SUIT reference assay. $FCF_{CGpDH} = 1 - CII/CII\&GpDH$. This late addition of Gp is a compromise for evaluation of CGpDH capacity. Malonic acid does not effectively inhibit CII at S_{50} (competitive inhibition). Little is known about the diagnostic value of this CGpDH-flux control factor. Gp is expensive.
9Ama	ROX	Inhibition may take very long, particularly in human muscle fibres (Pesta et al 2011; Lemieux et al 2011). This may make ROX correction questionable, particularly if ROX is high in comparison with the initial LEAK state.
10Tm	CIV_E	Ascorbate (As) is added before TMPD (Tm). $Tm_{0.5}$ is not saturating CIV, and thus represents a compromise, to prevent a too high chemical O_2 background.
11Azd	$cAsTm_{ROX}$	Cyanide is avoided due to the presence of P, but very high Azd concentrations are required. The oxygen dependence of the chemical O_2 is evaluated by a reoxygenation soon after titration of Azd, and is automatically performed by using the DatLab background calibration function (Slope).

2. SUIT_RP2: FAO-CI substrate control

RP2: 1D 2Oct 3Mtit 3c (3NADH) 4P 5G 6S 7U 8Gp 9Rot 10Ama 11Tm 12Azd

<i>E</i>					7U	8Gp	9Rot	10Ama	11Tm	12Azd	
<i>P</i>	1D	2Oct	3M+c	4P	5G	6S					
<i>L</i>											
	ROX	FAO	FAO	CI &FAO	CI &FAO	CI&II &FAO	CI&II &FAO&Gp	CII &Gp	ROX	CIV	ROX

RP2 spotlights:

- Depletion of endogenous substrates with D (1D; State 2).
- (3M) FAO_P compared to (4P) CI&FAO_P.
- Measurement of maximum ETS capacity, obtained in state (8Gp) PGMSOctGp_E.
- Harmonization between protocols RP1 and RP2 in states (RP1:6S, RP2:7U) PGMSOct_E, (RP1:8Gp, RP2:9Rot) SGp_E, and (10Tm or 11Tm) CIV_E. Harmonization with many previous protocols up to (+S).
- P/E (6S/7U) at high ETS capacity compared to RP1.

Limitations:

- CII_E is not obtained (but see RP1).
- The full substrate combination, PGMSOctGp_P and PGMSOctGp_E, is not covered, and thus the maximum apparent excess E-P capacity factor, $j_{Exp} = 1 - P/E$, may be missed.

Substrate type: CI-CII-FAO-CGpDH-CIV

Substrate state category:

FAO + CI&FAO + CI&II&FAO + CI&II&FAO&GpDH + CII&GpDH + CIV

SUIT protocol name:

D(ROX)+Oct(P)+M+P+c+G+S(PE)+Gp+Rot+Ama+AsTm+Azd

RP2-Pfi

Step	State	Comment
+D		In experiments with mt-preparations (mt), ADP is added to the medium before the mt.
+mt		D accelerates the depletion of endogenous substrates.
1D	D _{ROX}	Substrate depleted ROX state (State 2; Chance, Williams 1955).
2Oct	Oct _{ROX}	Oct alone does not establish an ETS (and OXPHOS) competent substrate state in many mt-types, since M is required to form oxaloacetate and prevent accumulation of acetyl-Co A by the citrate synthase reaction.

	Oct _p	Stimulation of OXPHOS by Oct alone in the presence of D indicates an obscure mechanism of anaplerosis.
3M	OctM _p	M is titrated stepwise: M.05; M.1; M2. Note that M alone can support OXPHOS if mt-malic enzyme is active, and thus FAO may be overestimated.
3c		See RP1.
3NADH		See RP1.
4P	PMOct _p	M ₂ is required to reduce flux through CII (minimize inhibition by malonate), such that CI-linked OXPHOS capacity can be estimated without high scope of compensation by CII-linked respiration. GM _p includes a higher share of CII-linked respiration in comparison with PM _p . $FCF_{CI} = 1 - FAO/CI$, important information on training status or cardiac failure (Pesta et al 2011; Lemieux et al, 2011).
5G	PGMOct _p	See RP1.
6S	PGMSOct _p	See RP1. The state CI&II&FAO _p is identical in RP1 and RP2, and may thus link the two protocols in the SUIT reference assay for statistical analysis (protocol harmonization).
7U	PGMSOct _E	CCCP is titrated in the CI&II state with high ETS capacity, to evaluate limitation of OXPHOS by the phosphorylation system. The apparent excess <i>E-P</i> capacity factor (<i>E-P</i> coupling control factor), $j_{EXP} = (E-P)/E = 1 - P/E$, is not measured in the state of maximum ETS capacity, if Gp exerts an additional stimulation (RP1). If Gp stimulates ETS further, the <i>E-P</i> coupling control factor is underestimated in the absence of Gp.
8Gp	PGMSOctGp _E	RP2 focuses on maximum <i>E</i> . $FCF_{CGpDH} = 1 - CI&II&FAO/CI&II&FAO\&GpDH$. This CGpDH-flux control factor evaluates additivity at high ETS capacity, which can be compared with additivity on the basis of CII _E (RP1).
9Rot	SGp _E	This state is not a generally valid estimate of CII _E (compare RP1).
Further steps		See RP1.

RP2-Pc

Step	State	Comment
0Ce	R	In experiments with intact cells (Ce), ROUTINE respiration (R) is measured initially, based on endogenous substrates.

1Dig	ROX	Digitonin permeabilizes the plasma membrane. Endogenous substrates are depleted and diluted in the incubation medium.
1D	D _{ROX}	D accelerates the depletion of endogenous substrates (State 2; Chance, Williams 1955; ROX; Gnaiger 2014).

3. Test experiments

Test experiments: Test experiments are required to finalize the RP2 for specific applications.

G 10 mM may not be saturating, and higher concentrations should be checked in a test experiment.

Gp Different sources of GP are tested. Gp (type) is expensive.

3.1. SUIT RP1

D D₅ (D_{7.5} in Pfi) is tested to be saturating CI&II&FAO&CGpDH_p. 7.5 mM may not be saturating in all cases, and higher concentrations of ADP should be checked.

Depletion of endogenous substrates with D is not possible, to obtain CI_L without inhibitors. Check with Omy after (+c).

Oct Oct_{0.5} (0.5 mM) might be generally applicable, but in preliminary experiments a higher concentration (1 mM) should be evaluated to check for saturation of flux.

+S Step titration from S₁₀ to S₅₀ to test if S₁₀ is saturating CI&II- and CII-linked respiratory capacity. If fluxes with S₅₀ > S₁₀ in CI&II_E, then S₅₀ is added immediately in the OXPHOS state. If S₅₀ < S₁₀, then it is tested if S₅₀ > S₁₀ in CII_E, in which case S₅₀ is only added in CII_E. If S₁₀ is saturating in all states, S₅₀ may be tested only occasionally, to exclude a shift in the succinate kinetics (in pathologies, ageing, etc).

3.2. SUIT RP2

Oct Oct_{0.5} is tested to be saturating in OXPHOS and not inhibiting or uncoupling (titration of high Oct after M.05 or M.1).

M M_{0.1} is tested to be saturating FAO in OXPHOS without activating CI-linked respiration beyond FAO capacity (HEK: mtME). M should be titrated stepwise (M.05;

M.1; M2) in the presence of D, to compare the malate kinetics of FAO_p and CI_p.

4. Technical details

Temperature: 37 °C.

Data recording interval: 2 s.

Effective chamber volume: 2 ml

Stirrer speed: 750 rpm.

DatLab file: The default name of the DatLab file contains the date, Power-O2k number and serial experimental number for each day.

2016-01-17 P1-02.DLD

O2k: Enter Power-O2k number: P1, P2, P3, P4, ..

Experiment: Experimental code, as in DatLab [F3].

Event: Set an 'Event' in DatLab at the time of titration. Use the abbreviated event name, and add information in the comment.

MiR05+CtlCr:

Ctl is present in all cells, hence addition of Ctl is considered physiological, even if reoxygenations are not required with H₂O₂.

Cr is present in many vertebrate cells, and thus should be added generally. With Cr, lower ADP concentrations are saturating for OXPHOS. It may be argued that it should be replaced in invertebrates (*Drosophila*, *C. rabditis*).

MiR / O2: Mitochondrial respiration medium, 2 ml in the O2k-chamber, plus 100 µl in the capillary of the stopper (more accurately: 88 µl without meniscus). Increase the oxygen concentration to ~450 µM. Close the chamber.

mt mt-preparation: Imt, Pfi, Pc.

D If there is time available (20 min), this period may yield a single point for the instrumental high-O2k background. D may be added just before titrating mt or before opening the chamber for addition of Pfi.

Pfi / O2: During addition of Pfi, the O₂ concentration drops and should be increased immediately to ~450 µM before closing the O2k-chamber.

U: 'Slope smoothing' may be reduced, e.g. to 25 (=25 data points used for calculation of the slope), to evaluate very quickly the stimulation of respiration and the need for additional titration steps of CCCP. If only FCCP (more expensive) is available, this can be used and be fully

compared with CCCP titrations (a minimally high CCCP than FCCP concentration may be required for maximum flux).

Cleaning

After the experiment clean the O2k-chambers: 3x water, 1x liver homogenate (20 min), 3x water, 3x EtOH 70% (5 min), 1x EtOH 100% (15 min).



O2k-cleaning SOP

» http://bioblast.at/index.php/MiPNet19.03_O2k-cleaning_and_ISS

5. Author contributions, publication versions, references

This communication is a pre-publication prepared by CD and EG. CD, ZS, GL and GK performed test experiments, contributed to the concept and co-wrote the manuscript.



Contribution to the project MitoFit, funded by the Tyrolian Government within the program K-Regio of Standortagentur Tirol.

<http://www.mitofit.org/index.php/O2k-MitoFit>



Full version with references

» http://wiki.oroboros.at/index.php/MiPNet21.06_SUIT_reference_assay

RP1: CI(LPE) substrate control

2016-01-23

E	3U	4Oct	5G	6S	7Rot	8Gp	9Ama	10Tm	11Azd
P	2D+c								
L	1PM								
	CI	CI&FAO	CI&FAO	CI&II&FAO	CII	CII&Gp	ROX	CIV	ROX

 PM + mt: **1PM** **2D** **2c** (2NADH) **3U** **4Oct** **5G** **6S** **7Rot** **8Gp** **9Ama** **10Tm** **11Azd**
Sample mt=Permeabilized fibres, RP1-Pfi:

O2k and DatLab file: P___(A / B) 2016-								
Experimental code:								
Operator:								
MiR: MiR05+CtlCr								
Event	Mark name	LPE	Final conc. 2 ml O2k	Stock [mM]	Comment	Tit. [µl]	A	B
MiR								
O2			~450 µM					
P			5 mM	2000		5		
M			2 mM	400		10		
mt								
O2	1PM	L	~450 µM					
D	2D	P	7.5 mM	500		30		
c	2c	P	10 µM	4		5		
NADH	2NADH	P	2.8 mM	280	NADH only if $FCF_c > .1$	20		
U	3U	E	Δ0.5 µM	1	CCCP	Δ1 µl		
Oct	4Oct	E	0.5 mM	100		10		
G	5G	E	10 mM	2000		10		
S	6S	E	50 mM	1000		100		
Rot	7Rot	E	0.5 µM	1		1		
Gp	8Gp	E	10 mM	1000		20		
Ama	9Ama	ROX	2.5 µM	5		1		
O2			~450 µM					
As			2 mM	800		5		
Tm	10Tm	E	0.5 mM	200	~20 min	5		
Azd	11Azd	ROX	≥100 mM	4000	~10 min	100		
O2	12Azd	ROX	~450 µM		400 -> 250 µM			

RP2: FAO-CI substrate control

2016-01-23

<i>E</i>					7U	8Gp	9Rot	10Ama	11Tm	12Azd	
<i>P</i>	1D	2Oct	3M+c	4P	5G	6S					
<i>L</i>											
	ROX	FAO	FAO	CI &FAO	CI &FAO	CI&II &FAO	CI&II &FAO&Gp	CII &Gp	ROX	CIV	ROX

D + mt: 1D 2Oct 3Mtit 3c (3NADH) 4P 5G 6S 7U 8Gp 9Rot 10Ama 11Tm 12Azd

Sample mt=Permeabilized fibres, RP2-Pfi:

O2k and DatLab file: P__(A / B) 2016-								
Experimental code:								
Operator:								
MiR: MiR05+CtlCr								
Event	Mark name	LPE	Final conc. 2 ml O2k	Stock [mM]	Comment	Tit. [µl]	A	B
MiR								
O2			~450 µM					
D			7.5 mM	500		30		
mt								
O2	1D	ROX	~450 µM					
Oct	2Oct	<i>P</i>	0.5 mM	100		10		
M.05	3M.05	<i>P</i>	0.05 mM	50		2		
M.1	3M.1	<i>P</i>	0.1 mM	50		2		
M2	3M2	<i>P</i>	2 mM	400		9.5		
c	3c	<i>P</i>	10 µM	4		5		
NADH	3NADH	<i>P</i>	2.8 mM	280	NADH only if $FCF_c > .1$	20		
P	4P	<i>P</i>	5 mM	2000		5		
G	5G	<i>P</i>	10 mM	2000		10		
S	6S	<i>P</i>	50 mM	1000		100		
U	7U	<i>E</i>	Δ0.5 µM	1	CCCP	Δ1 µl		
Gp	8Gp	<i>E</i>	10 mM	1000		20		
Rot	9Rot	<i>E</i>	0.5 µM	1		1		
Ama	10Ama	ROX	2.5 µM	5		1		
O2			~450 µM					
As			2 mM	800		5		
Tm	11Tm	<i>E</i>	0.5 mM	200	~20 min	5		
Azd	12Azd	ROX	≥100 mM	4000	~10 min	100		
O2	13Azd	ROX	~450 µM		400 -> 250 µM			

Supplement

A. General links

Introduction

» http://wiki.orooboros.at/index.php/Gnaiger_2014_MitoPathways

Respiratory substrate-coupling states

» [http://www.bioblast.at/index.php/MitoPedia: Respiratory substrate-coupling states](http://www.bioblast.at/index.php/MitoPedia:Respiratory_substrate-coupling_states)

Table of titrations

» http://wiki.orooboros.at/index.php/MiPNet09.12_O2k-Titrations

Definition

» [http://www.bioblast.at/index.php/Substrate-uncoupler-inhibitor titration](http://www.bioblast.at/index.php/Substrate-uncoupler-inhibitor_titration)

Context

» http://www.mitofit.org/index.php/SUIT_protocol_library

Abbreviations

» <http://www.bioblast.at/index.php/MitoPedia>

