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Electron Transport Chain Protocol

DOI

<https://dx.doi.org/10.17504/protocols.io.6qpvrw39olmk/v1>

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Protocol status: Working

We use this protocol and it's working

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Abstract

This protocol describes the procedure for analyzing Electron Transport Chain activity of Complexes 1, 2, and 4 in mitochondria isolated from fibroblasts used by the Seifert Lab. It follows the "Electron Transport Chain Protocol" (ETCP) described in:

Vekaria, H.J., Kalimon, O.J., Prajapati, P., Velmurugan, G.V., Sullivan, P.G., 2024. An efficient and high-throughput method for the evaluation of mitochondrial dysfunction in frozen brain samples after traumatic brain injury. *Front Mol Biosci* 11, 1378536.

which is a modified version of the "Respirometry in Frozen Samples" (RIFS) assay from:

Acin-Perez, R., Benador, I.Y., Petcherski, A., Veliova, M., Benavides, G.A., Lagarrigue, S., Caudal, A., Vergnes, L., Murphy, A.N., Karamanlidis, G., Tian, R., Reue, K., Wanagat, J., Sacks, H., Amati, F., Darley-USmar, V.M., Liesa, M., Divakaruni, A.S., Stiles, L., Shiriha, O.S., 2020. A novel approach to measure mitochondrial respiration in frozen biological samples. *Embo j* 39, e104073.



- 1 Thaw frozen mitochondrial samples
- 2 Add  25 μL Mitochondria Isolation Buffer (MIB;  215 millimolar (mM) mannitol,  75 millimolar (mM) sucrose, 0.1% BSA,  20 millimolar (mM) HEPES, and  1 millimolar (mM) EGTA, adjusted to pH 7.2 with KOH
- 3 Gently resuspend pellet
- 4 Measure concentration of  4 μL of each mitochondrial mix by BCA
- 5 Prepare

Equipment

Seahorse XFe96 Analyzer

NAME

Agilent

BRAND

BLE2100298

SKU

<https://www.agilent.com/en/product/cell-analysis/real-time-cell-metabolic-analysis/xf-analyzers/seahorse-xfe96-analyzer-740879>

LINK

96-well cartridge ports as follows:

Port A (ETC 1):  3220 μL Mitochondrial Respiration buffer w/o BSA,  2.8 μL

Rotenone,  280 μL succinate;

Port B:  3500 μL Mitochondrial Respiration buffer w/o BSA,  3.5 μL Antimycin A;

Port C:  3325 μL Mitochondrial Respiration buffer w/o BSA,  88 μL Ascorbate,

 88 μL TMPD;

Port D:  3033 μL Mitochondrial Respiration buffer w/o BSA,  480 μL Azide.



- 6 Load cartridge well ports with $25\ \mu\text{L}$ of port mixes
- 7 0°C On ice Load $75\ \mu\text{L}$ of mitochondrial sample ($2\ \mu\text{g}$ protein) per well
- 8 Balance the plate and spin at $3214\ \text{rcf}$, 4°C , 00:10:00
- 9 Prepare $11\ \text{mL}$ Alamethicin Assay solution ($10.6\ \text{mL}$ Mitochondrial Respiration buffer w/o BSA, $40\ \text{mg/mL}$ Alamethicin $9.63\ \mu\text{L}$, $1.5\ \text{mM}$ NADH $385\ \mu\text{L}$, $10\ \mu\text{M}$ cytochrome c $48\ \mu\text{L}$)
- 10 Add $100\ \mu\text{L}$ assay solution to each well
- 11 Run plate on

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LINK

sequentially injecting the contents of Ports A-D over $01:00:00$



Protocol references

Vekaria, H.J., Kalimon, O.J., Prajapati, P., Velmurugan, G.V., Sullivan, P.G., 2024. An efficient and high-throughput method for the evaluation of mitochondrial dysfunction in frozen brain samples after traumatic brain injury. *Front Mol Biosci* 11, 1378536.