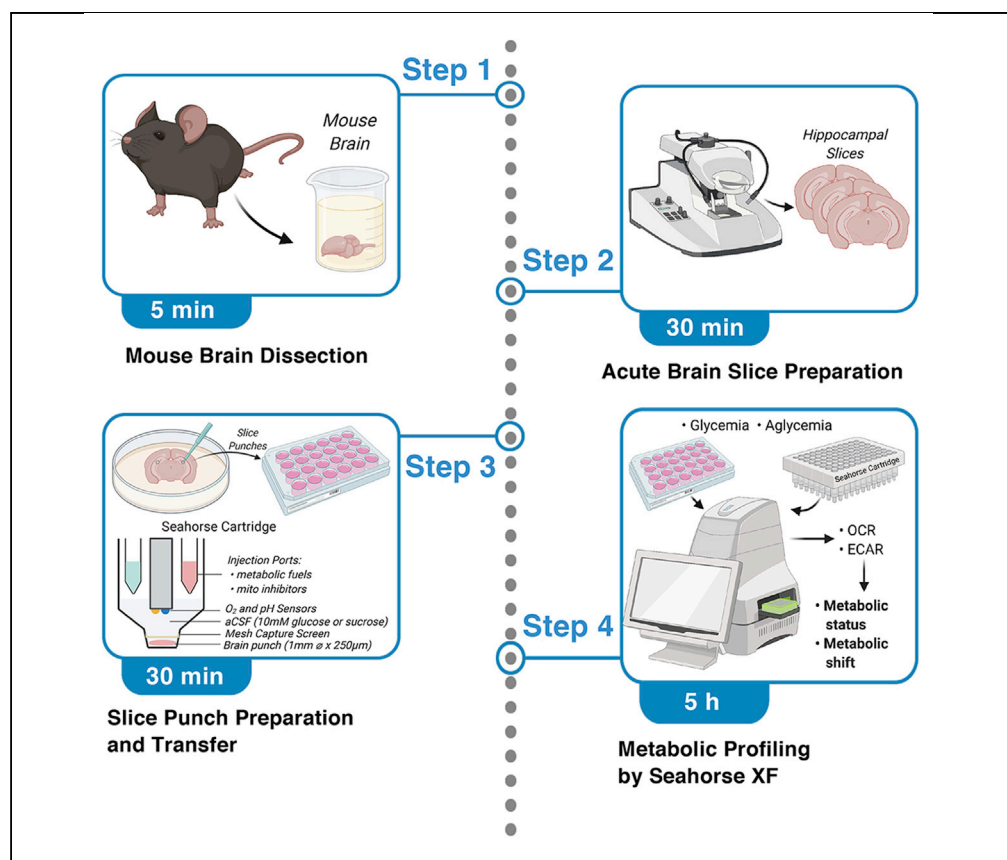


Protocol

Characterizing brain metabolic function ex vivo with acute mouse slice punches



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Highlights

Detailed protocol to determine complex brain metabolic function in mouse

Streamlined method to determine brain metabolic shift or reprogramming

Validated procedure to quantify brain aerobic glycolytic rate ex vivo

Mitochondrial dysfunction and metabolic reprogramming are implicated in a variety of neurological disorders. Here, we present a protocol that enables complex profiling of brain metabolic function using acute mouse brain slices *ex vivo*. Utilizing differential metabolic conditions, substrates, and inhibitors, this protocol can be broadly applied to determine metabolic shift or reprogramming upon genetic manipulations, pathological insults, or therapeutic interventions and could thus further the understanding of the dynamic role of energy metabolism in brain physiological function and diseases.

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Protocol

Characterizing brain metabolic function ex vivo with acute mouse slice punches

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SUMMARY

Mitochondrial dysfunction and metabolic reprogramming are implicated in a variety of neurological disorders. Here, we present a protocol that enables complex profiling of brain metabolic function using acute mouse brain slices *ex vivo*. Utilizing differential metabolic conditions, substrates, and inhibitors, this protocol can be broadly applied to determine metabolic shift or reprogramming upon genetic manipulations, pathological insults, or therapeutic interventions and could thus further the understanding of the dynamic role of energy metabolism in brain physiological function and diseases.

For complete details on the use and execution of this protocol, please refer to Qi et al. (2021).

BEFORE YOU BEGIN

This protocol below describes the specific steps to characterize brain metabolic function *ex vivo* with acute mouse hippocampal slice punches. Depending on the research focus, this protocol can also be applied to other brain regions such as cortex and white matter tracts, or other rodent species (processing of rat brains was described previously (Fried et al., 2014; Underwood et al., 2020)).

Day prior to assay

1. Hydrate Agilent Seahorse XFe24 Sensor Cartridge with water

⌚ Timing: 10 min

- a. Prepare a 24-well Seahorse XFe24 sensor cartridge with utility plate and hydro booster from the Seahorse XFe24 Islet Capture FluxPak.
- b. Add 1 mL sterile water to each well of the utility plate, then put the hydro booster on top of the utility plate and the sensor cartridge through the openings on the hydro booster. Make sure the sensors are submerged in water.
- c. Place the assembled sensor cartridge and utility plate in a non-CO₂ 37°C incubator 12–16 h.
- d. Separately, add 40 mL of Seahorse XF Calibrant (included in the FluxPak) into a 50 mL conical tube and place it a non-CO₂ 37°C incubator 12–16 h.



Table 1. Stock solutions of mitochondrial inhibitors

Reagent	Stock concentration
Oligomycin A	20 mg/mL in 0.2 mL DMSO
FCCP	10 mM in 0.2 mL DMSO
Antimycin A	10 mM in 0.2 mL DMSO

Note: To prevent evaporation of the water, use a properly humidified incubator or place the assembled sensor cartridge and plate back to its packaging to reduce evaporation.

2. Fasting mice 14–16 h (optional)

⌚ Timing: 5 min

Transfer the mice to be assessed to new cages with free access to water but not food ~14–16 h before euthanasia.

Note: This is an optional step aiming to minimize the impact of variable blood glucose and fatty acid levels across individual mice on brain metabolism of exogenous fuels, especially under aglycemic conditions. If glucose availability to brain is not a concern, unfasted mice can also be used. Make sure an approval is obtained from the Institutional Animal Care and Use Committee prior to the fasting of animals.

3. Reagent and buffer preparation

⌚ Timing: 1 h

a. Prepare stock solutions of mitochondrial inhibitors listed in [Table 1](#).

Note: Store at -20°C for up to 6 months and avoid repeated freeze and thaw cycles.

b. Prepare artificial cerebrospinal fluid (aCSF) without glucose or sucrose ([Table 2](#)).

Note: Store at 4°C for up to 4 weeks.

c. Agarose block preparation

Mix 3 g agarose powder with 100 mL aCSF in a conical flask, then microwave the mixture until agarose is completely dissolved. Pour the agarose solution into a 100 mm petri dish at 20°C – 25°C for 0.5 h for solidification. Then cut out an agarose cube with the side length of ~ 1 cm ([Figure 1](#)). Seal the petri dish with parafilm and store it at 4°C . Prepared agarose gel should be used within 2 months.

Table 2. Artificial cerebrospinal fluid (aCSF)

Reagent	Stock concentration	Final concentration	Amount
NaCl	n/a	120 mM	7.00 g
KCl	n/a	3.5 mM	0.26 g
$\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$	n/a	1.3 mM	0.19 g
MgCl_2	1 M	1 mM	1 mL
KH_2PO_4	n/a	0.4 mM	0.054 g
HEPES	1 M	5 mM	5 mL
ddH ₂ O	n/a	n/a	~994 mL
Total	n/a	n/a	1 L

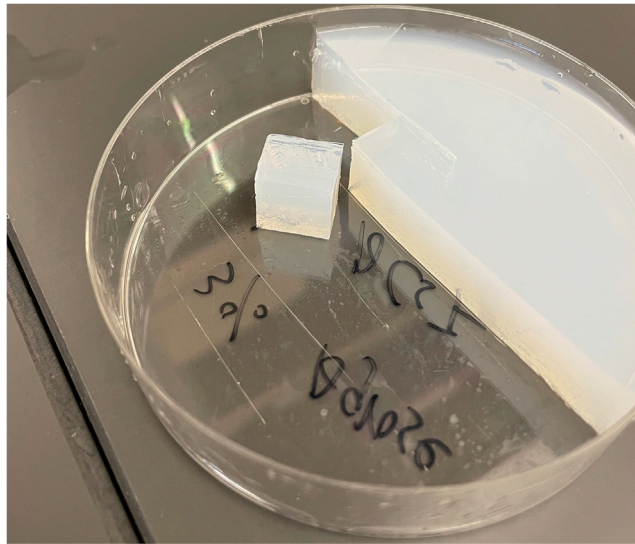


Figure 1. Agarose block preparation
Prepare an agarose cube with side length of ~1 cm.

KEY RESOURCES TABLE

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Chemicals, peptides, and recombinant proteins		
Antimycin A	Sigma-Aldrich	Cat#A8674-50MG
Oligomycin A	Sigma-Aldrich	Cat#75351-5MG
FCCP	Sigma-Aldrich	Cat#C2920-10MG
Fatty acid-free BSA	Fisher Bioreagents	Cat#BP9704-100
NaCl	Sigma-Aldrich	Cat#S7653
KCl	Sigma-Aldrich	Cat#793590
CaCl ₂ ·2H ₂ O	Sigma-Aldrich	Cat#7902
MgCl ₂	Sigma-Aldrich	Cat#M1028
KH ₂ PO ₄	Sigma-Aldrich	Cat#P9791
HEPES	Sigma-Aldrich	Cat#H0887
Sucrose	Sigma-Aldrich	Cat#S9378
PBS (phosphate-buffered saline)	Gibco	Cat#10010023
BSA-Oleate	Cayman Chemical	Cat#29557
Glucose	Sigma-Aldrich	Cat#G7021
Agarose	VWR	Cat#N605-500 g
Critical commercial assays		
Seahorse XFe24 Islet Capture FluxPak	Agilent Technologies	Cat#103518-100
Software and algorithms		
Seahorse Wave	Agilent Technologies	N/A
Other		
Biopsy Punches with Plunger System	Integra LifeSciences	Cat#33-31AA-P/25
Seahorse XFe24 Analyzer	Agilent Technologies	XFe24 Analyzer
Dissecting forceps	Stoelting	Cat#5210206
Vibratome	Leica	VT1200S
Super Glue	Krazy	Cat#KG517
100 mm Petri dish	Thermo Fisher Scientific	Cat#172931

(Continued on next page)

Continued

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Experimental models: Organisms/strains		
3-Month male hApoE3 knockin mouse	The Jackson Laboratory	Stock No: 029018
3-Month male hApoE4 knockin mouse	The Jackson Laboratory	Stock No: 027894

Note: Seahorse XFe24 Analyzer is absolutely required and the XFe24 Islet Capture FluxPak is a critical assay item.

STEP-BY-STEP METHOD DETAILS

Hydrate Agilent Seahorse XFe24 sensor cartridge with XF calibrant

⌚ Timing: 10 min

1. Discard the water from the utility plate prepared previously and fill each well with 1 mL pre-warmed XF calibrant prepared the day before.
2. Place the sensor cartridge utility plate in a non-CO₂ 37°C incubator for 45–60 min.

Preparation of aCSF with different substrates

⌚ Timing: 30 min

3. Freshly add fatty acid-free BSA to previously made aCSF (Table 2) to a final concentration of 4 g/L. Then prepare 200 mL aCSF-glucose and 500 mL aCSF-sucrose by dissolving glucose or sucrose in aCSF (with BSA) to a final concentration of 10 mM glucose or 10 mM sucrose, respectively. Adjust pH to 7.4 using 1 M NaOH.

⚠ **CRITICAL:** When assessing fatty acid-induced oxygen consumption rate (OCR) in brain slices, aCSF-sucrose (aglycemic condition) has to be used to minimize the interference of glucose in the medium. Similarly, for glycolytic activity assessment, Extracellular acidification rate (ECAR) values at aglycemic condition (aCSF-sucrose) will be used as the baseline representing glucose-independent pH changes. aCSF-glucose (glycemic condition) will be used to determine glucose-induced OCR and ECAR. Substrate-free aCSF (no glucose or sucrose) can be prepared and stored at 4°C for up to 4 weeks (Table 2). BSA, glucose or sucrose must be added freshly on the day of experiment.

Vibratome setup

⌚ Timing: 5 min

4. Place a new razor blade in the clamp and fill the vibratome tray with pH 7.4 aCSF-sucrose and set the temperature to 4°C. Adjust the sectioning parameters as follows (Leica VT 1200S): section thickness: 250 μm; cutting speed: 0.20 mm/s; cutting amplitude: 1.50 mm (Figure 2A).

Brain dissection

⌚ Timing: 5 min

5. Euthanize the mouse by cervical dissociation.

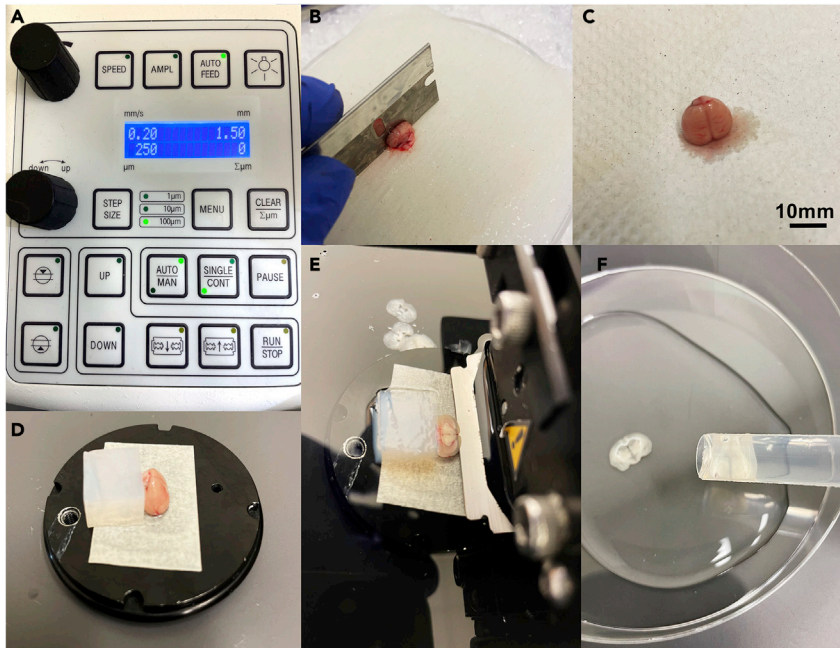


Figure 2. Brain slices preparation

- (A) Experimental settings of the vibratome for sectioning.
 (B) Cerebellum and brain stem are removed and discarded from the brain freshly dissected from the skull.
 (C) Dap the bottom surface of the brain gently with paper towel to remove excess liquid before mount the brain onto the specimen plate.
 (D) A supporting piece of agarose block is added behind the brain to provide structural support during sectioning.
 (E) Sectioning a brain.
 (F) Collect brain slice containing the region of interest from the buffer tray of the vibratome to a petri dish prefilled with aCSF-sucrose using a transfer pipet.

Optional: Anesthesia of mice by isoflurane in an anesthetizing chamber before dissection is optional and should be used with caution, because anesthesia is known to affect brain metabolism.

⚠ **CRITICAL:** If anesthesia is used, isoflurane should be handled in a fume hood and the exposure time of mice to isoflurane should be minimized.

6. Decapitate mouse and dissected the brain rapidly from the skull.
7. Transfer the brain onto phosphate buffered saline (PBS)-soaked paper towel placed in a petri dish on ice. Remove the cerebellum and brain stem (Figure 2B), put the remaining brain into a 50 mL beaker containing ice-cold aCSF-sucrose.

⚠ **CRITICAL:** Brain should be rapidly collected within 30 s–60 s of decapitation.

Vibratome sectioning

⌚ **Timing:** 30 min (per mouse brain)

8. Tape the center of specimen plate and place a small dot of superglue in the center of the tape.

Note: The use of lab tape greatly helps the cleaning of the specimen plate afterwards.

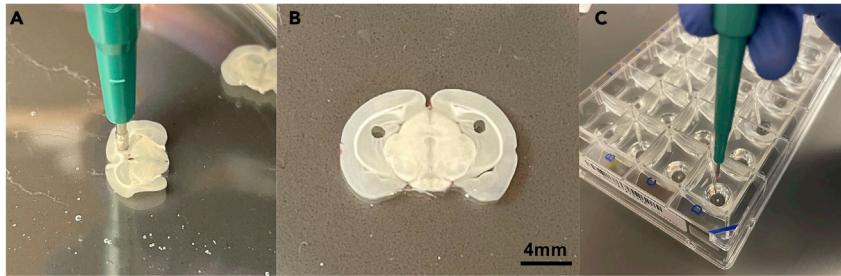


Figure 3. Slice punch preparation and transfer

- (A) Lower 1 mm stainless steel biopsy puncher into the petri dish and press gently overtop the region of interest of the brain slice.
 (B) Brain slice after punch collection of the dentate gyrus of both hemispheres.
 (C) Eject the punched piece into a well of XFe Islet Capture Microplate directly and adjust the brain punches to make sure it's located in the center of each well using the biopsy puncher.

- Take out the brain from aCSF-sucrose from step 7 with forceps and dap the dorsum of the brain gently with paper towel to remove excess buffer (Figure 2C). Carefully place the brain on the superglue (dorsum down), then slide a stabilizing agarose block to the ventral side of the brain (opposite to the blade side) to provide structural support during slicing (Figure 2D).

Note: Remove excess buffer with paper towel will help the brain stick more firmly to the specimen plate.

- Mount and orient the specimen plate to make sure the dorsal side of the brain faces the blade.
- Set the cutting window to define the start and stop positions of the slicing. Make sure the blade cut completely through the tissue so slices will come off freely.
- Adjust the position of the buffer tray until the razor blade is in the cutting place. Start sectioning (Figure 2E).
- Identify and transfer brain slices that contain the region of interest from the buffer tray to a 100 mm petri dish prefilled with 37°C aCSF-sucrose using a transfer pipet (Figure 2F).

△ CRITICAL: Using this protocol, brain slices can maintain their metabolic stability for about 7–8 h after sectioning. The entire procedure needs to be completed within this time frame.

Preparation of brain punches and transfer to seahorse plate

⌚ Timing: 30 min

- Place the mesh inserts of the Seahorse XFe24 Islet Capture Microplate in a 100 mm petri dish containing fresh 37°C aCSF-sucrose, make sure the mesh inserts are completely immersed in the liquid without any bubble under the mesh.

△ CRITICAL: XFe24 Islet Capture Microplate rather than the standard XFe24 cell culture microplate should be used. The XFe Islet Capture Microplate contains depressed chamber and mesh that will help hold the brain punches in place at the center of each well throughout the measurement.

- Add 700 μ L per well 37°C aCSF-sucrose to half of the XFe24 Islet Capture Microplate wells for aglycemic condition and add aCSF-glucose to the other half of wells for glycemic condition.
- Position 1 mm stainless steel biopsy puncher over the area of interest on the brain section from step 13 and press down (Figures 3A and 3B).

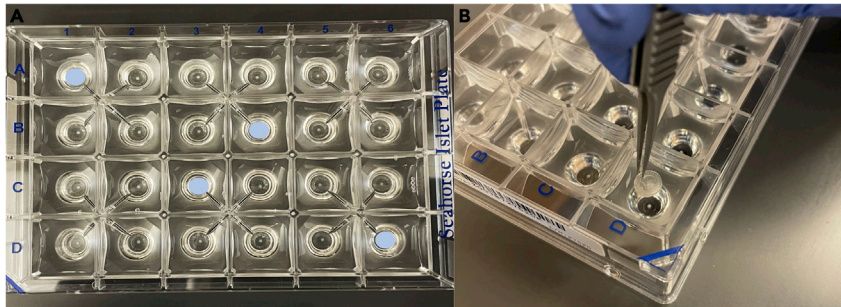


Figure 4. Layout of XFe24 Islet Capture Microplate and insertion of the mesh insert

(A) Four background correction wells (A1, B4, C3, and D6) in the XFe24 Islet Capture Microplate should contain aCSF only (no punches).

(B) Gently place the mesh inserts into each well of the XFe24 Islet Capture Microplate with fine-tipped tweezers above the punch to hold the latter in the center of each well.

⚠ **CRITICAL:** Mechanical damages to the brain tissue during biopsy and transfer can substantially affect tissue respiration. It is recommended to only use brain punches without damage.

17. Expel each punch (1mm diameter) into one well of XFe24 Islet Capture Microplate and adjust the position of brain punch using biopsy puncher to make sure it is located in the center of each well (Figure 3C). Each punch should be submerged in the liquid and centered at the bottom of the wells.

⚠ **CRITICAL:** Do not place brain punches into the four background correction wells (A1, B4, C3, and D6) in the XFe24 Islet Capture Microplate (Figure 4A).

Place the mesh inserts into each well of the XF Islet Capture Microplate with fine-tipped tweezers which will help keep the brain punches in the center of each well throughout the experiment (Figure 4B).

18. Incubate the XF Islet Capture Microplate with punches in a 37°C incubator without CO₂ for 1 h to allow temperature and pH equilibration.

Metabolic profiling by seahorse XFe24 analyzer

⌚ **Timing:** 5 h

19. Prepare injection solutions of BSA-oleate (1.65 mM), oligomycin A (240 µg/mL), FCCP (130 µM) and antimycin A (280 µM) by diluting their stock solutions (prepared as in Table 1) using either aCSF-sucrose or aCSF-glucose (Table 3), corresponding to aglycemic or glycemc conditions assigned in step 15.

20. Load the hydrated sensory cartridge (from step 2) ports with 70 µL of 1.65 mM BSA-oleate in port A, 70 µL of 240 µg/mL oligomycin A in port B, 70 µL of 130 µM FCCP in port C, and 70 µL of 280 µM Antimycin A in port D using a 200 µL pipette and reagent reservoir (Table 4).

⚠ **CRITICAL:** aCSF-sucrose should be used to prepare the injection solutions of substrates and inhibitors for aglycemic conditions. In parallel, for glycemc conditions, aCSF-glucose should be used as diluent of injection solutions.

Table 3. Preparation of injection solutions of BSA-Oleate and mitochondrial inhibitors

Reagent	Volume of reagent (μL)	Volume of aCSF (μL)	Injection concentration
BSA-Oleate	330	670	1.65 mM
Oligomycin A	12	988	240 $\mu\text{g}/\text{mL}$
FCCP	13	987	130 μM
Antimycin A	28	972	280 μM

- Set up the Seahorse XFe24 program as follows: baseline, 4 cycles; inject port A (BSA-oleate), 5 cycles; inject port B (oligomycin), 6 cycles; inject port C (FCCP), 5 cycles; inject port D (Antimycin A), 15 cycles. Each cycle is composed of mix 3 min, wait 3 min, and measure 2 min (Table 5).

Note: Comparing to cellular XF assays, it takes longer time for brain slices to reach stable OCR after oligomycin- and antimycin A injections (Figure 6). Additional cycles are thus needed for measurements after oligomycin (6 cycles) and after antimycin A (15 cycles). However, depending on whether proton leak-related OCR (after oligomycin) and non-mitochondrial OCR (after antimycin A) is of interest, the cycle numbers for these steps can be adjusted.

- Place the utility plate with the loaded sensor cartridge into the XFe24 instrument tray and start the calibration step, which takes approximately 20 min.
- After calibration is completed, replace the utility plate with the XFe24 Islet Capture Microplate with brain punches prepared in step 18 and start the XF assay. Assay duration is around 5 h.

EXPECTED OUTCOMES

Seahorse XFe Analyzer has been widely used in evaluating metabolic function in isolated brain mitochondria and in cultured cells by monitoring OCR and ECAR, indicators of mitochondrial respiration and glycolysis, respectively. As data derived from cultured cells or isolated mitochondria may not recapitulate the complex cellular environment of the brain, and these *in vitro* models cannot reflect the significance of intercellular coordination and metabolic coupling across the heterogenous cell types in the brain (Qi et al., 2019), *ex vivo* methods to measure mitochondrial respiration on glucose-derived substrate have thus been developed with rat brain slices (Fried et al., 2014; Schuh et al., 2011; Underwood et al., 2020). However, over the past decade, the view of brain bioenergetics and its role in diseases has been evolving from the monotonic “respiratory decline” towards a dynamic metabolic reprogramming and adaptation model. Such reprogramming and adaptation involve the differential ability of the mitochondria in utilizing various energy substrates, as well as the allocation of ATP production between glycolysis and OxPhos. To this end, we describe a protocol herein to characterize complex metabolic function in mouse brains beyond glucose-dependent OCR. In our protocol, by incubating punches of acute brain slices under different glucose availability and supplying them with different substrate (e.g., fatty acids), brain metabolic function is quantified and any metabolic shift in the brain’s ability to utilize different fuel substrates can be determined. A typical result of OCR curves for brain slice respiration under glycemic or aglycemic conditions is shown in Figure 5. Further, a metabolic shift towards aerobic glycolysis has been well-established in glial cells upon inflammatory activation in aging and pathological conditions (Qi et al., 2019). This protocol also establishes and validates ECAR measurement as an indicator of brain aerobic glycolysis. By including tissues under both glycemic and aglycemic conditions, our protocol further

Table 4. Port injection volume and final concentrations of BSA-Oleate and inhibitors

Reagent (injection concentration)	Final concentration in well	Volume added to port	Port
BSA-Oleate (1.65 mM)	150 μM	70 μL	A
Oligomycin A (240 $\mu\text{g}/\text{mL}$)	20 $\mu\text{g}/\text{mL}$	70 μL	B
FCCP (130 μM)	10 μM	70 μL	C
Antimycin A (280 μM)	20 μM	70 μL	D

Step	Time	Cycles
Baseline		
Mix	3 min	4
Wait	3 min	
Measure	2 min	
Injection of port A (BSA-Oleate)		
Mix	3 min	5
Wait	3 min	
Measure	2 min	
Injection of port B (Oligomycin A)		
Mix	3 min	6
Wait	3 min	
Measure	2 min	
Injection of port C (FCCP)		
Mix	3 min	5
Wait	3 min	
Measure	2 min	
Injection of port D (Antimycin A)		
Mix	3 min	15
Wait	3 min	
Measure	2 min	

enables quantitative measurements of ECAR (pH) changes specifically due to glycolytic production of lactic acid.

Using this protocol, we revealed a metabolic shift towards enhanced glucose metabolism and diminished fatty acid metabolism induced by Alzheimer’s disease risk gene ApoE4 (Figure 6) (Qi et al., 2021). Hippocampal slices from humanized ApoE4 knockin mice exhibited a higher basal respiration than ApoE3 when glucose, but not FA, was present (Figure 6A). However, upon glucose depletion, ApoE4 slices showed a lower respiration than ApoE3 (Figure 6B). When exogenous FA (oleate-BSA) was supplemented, a lower respiration was seen in ApoE4 slices, but only under aglycemic condition (Figures 6C and 6D). In parallel with increased glucose respiration and decreased FA respiration, ApoE4 brains showed a higher rate of aerobic glycolysis either with or without FA (Figures 6E and 6F).

Collectively, this protocol complements the existing toolbox assessing brain metabolism by measuring multiple aspects of bioenergetics, namely glucose-dependent mitochondrial respiration, alternative-fuel-dependent respiration, and aerobic glycolysis in one streamlined procedure. Utilizing bioenergetic fuels such as ketone body, fatty acids, lactate and amino acids, and pharmacological enhancers or inhibitors of substrate transport, glycolysis, ketolysis, β -oxidation and tricarboxylic

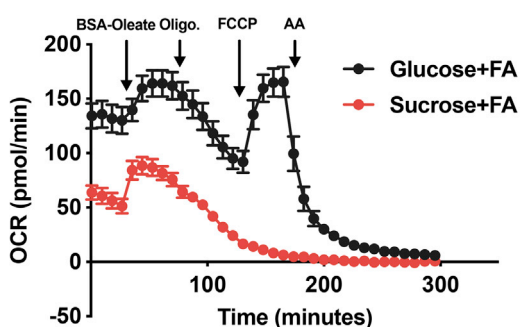


Figure 5. Representative oxygen consumption rate (OCR) with acute hippocampal slice punches under aglycemic (sucrose) and glycemic (glucose) conditions
FA, fatty acid; Oligo., oligomycin; AA, antimycin A.

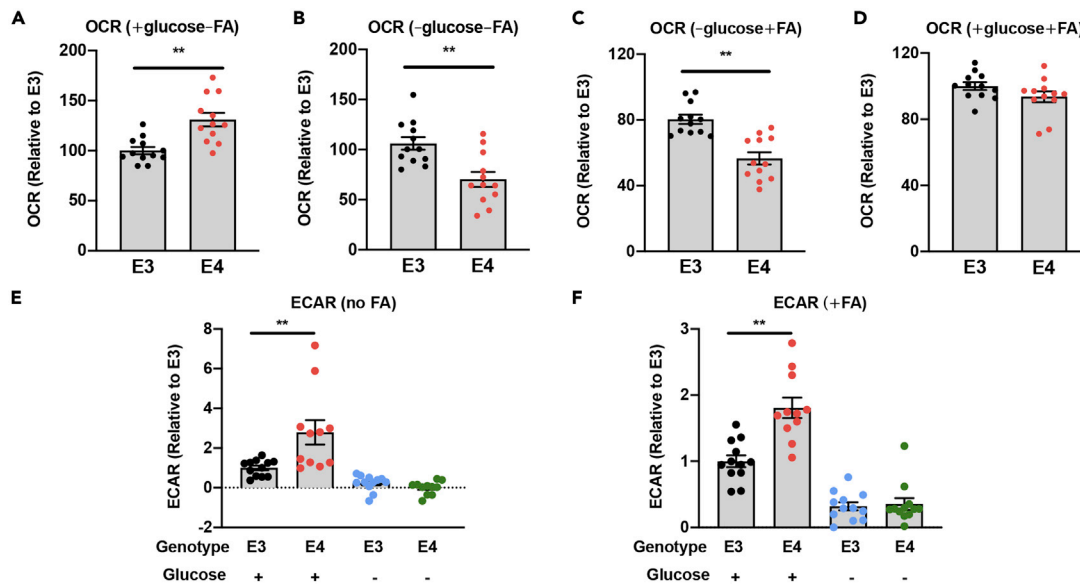


Figure 6. ApoE4 diminishes FA oxidation and promotes glucose metabolism in mouse hippocampus

Figure reprinted with permission from (Qi et al., 2021).

(A) Mitochondrial oxygen consumption rate (OCR) of hippocampal slices with 10 mM glucose but no exogenous FA.

(B) Mitochondrial OCR of hippocampal slices without exogenous glucose or FA.

(C) Mitochondrial OCR of hippocampal slices with 150 mM oleate-BSA but no glucose.

(D) Mitochondrial OCR of hippocampal slices with 10 mM glucose plus 150 mM oleate-BSA.

(E) Extracellular acidification rate (ECAR) in hippocampal slices without exogenous FA.

(F) ECAR in hippocampal slices with exogenous FA.

acid (TCA) cycle, this protocol can be versatilely modified to characterize the multifaceted brain bioenergetic system, which is often reprogrammed and thereby implicated in the pathogenesis of a wide variety of neurological diseases.

QUANTIFICATION AND STATISTICAL ANALYSIS

1. Quantification analysis is performed using Seahorse Wave Desktop. More detailed information on data export with Wave software can be found at Agilent website. https://www.agilent.com/cs/pubimages/misc/User_Guide_Wave_Desktop_2-3.pdf
2. Assay results can be exported to Microsoft Excel or GraphPad Prism for further statistical analyses.

LIMITATIONS

Theoretically, the present method measures the overall metabolic function of the mixed population of cells including neurons, glia, and other cell types present in the tissue punches. Therefore, it cannot stratify the contribution from each cell type to the observed metabolic phenotype. Also, the *ex vivo* setting of this protocol, while advantageous to cell cultures, may not fully recapitulate brain metabolic function *in vivo*. A combination of this protocol with *in vitro* studies of different cell types, and other *in vivo* assessment of brain metabolism, such as metabolomics, should be used to address these limitations.

Technically, the entire procedure, including brain dissection, slicing, punch preparation, and eventually the Seahorse assay, takes approximately 7 h. Helpful practices were discussed previously to keep brain punches metabolically active and obtain reliable and reproducible results

(Underwood et al., 2020). Lastly, the availability of the Seahorse XFe24 Analyzer may limit the usage of this protocol.

TROUBLESHOOTING

Problem 1

High variance in results between punch replicates.

Potential solution

High variance between experimental replicates could arise from the quality control of brain punches. Each biopsy puncher should be used for no more than 4 punches. Brain punches of poor quality or irregular shape should not be used (step 16).

Another step that could cause damage to slice punches is step 17. Make sure the mesh insert is gently placed into each well of the microplate with fine-tipped tweezers and the mesh should only lightly touch the punches. Punches could be damaged due to excessive pressure on the mesh insert.

Problem 2

Lack of response in OCR to substrate or inhibitor injections.

Potential solution

If punches in some wells show normal baseline readouts but fail to respond to the injections of inhibitors or substrates, this may suggest incomplete injections of inhibitors into those wells. It is thus important to always check all injection ports upon completion of the XFe24 assay to make sure all inhibitors or substrates have been injected (steps 20 and 23).

Moreover, the responsiveness of punches to substrates or inhibitors may vary by brain region, as well as the sex, age, and/or genotype of the animal. If a lack of response is found in most wells across multiple experimental groups, a titration should be conducted for the substrate or inhibitor in question to determine the optimal dose (step 19).

Problem 3

Poor metabolic activity due to extended punch preparation

Potential solution

Poor metabolic activity (OCR and ECAR) may be observed if the procedure takes more than 7 h due to the extended time to process additional brains for punches (steps 8–18). In this case, brain sections can be placed in oxygenated aCSF prior to slice punching and Seahorse assays to help preserve their metabolic activity (step 13). In addition to antimycin A, complex I inhibitor rotenone (20 μ M final concentration in well) can be included in port D injection (steps 19 and 20) which may help to reduce the time reaching complete electron transport chain (ETC) inhibition and thus reduce the measurement cycles.

Problem 4

High OCR and ECAR values under aglycemic condition

Potential solution

OCR values under aglycemic condition should be lower than glyceemic groups (Figure 5) and the ECAR values should be minimal (Figure 6). If higher than expected values are observed for aglycemic wells, there might be interference from endogenous glucose remaining in the brain tissue. In this case, a 14–16 h starvation to mice can help control the endogenous glucose available to brain tissues and reduce variance across animals (Day prior to assay step 2 in [before you begin](#)). Also make sure

aCSF-sucrose is used for punches assigned for aglycemic condition throughout the entire protocol including the fatty acid and inhibitor solutions for these wells (steps 3, 4, 7, 13, 14–18, and 19).

Problem 5

Difficulty in obtaining good-quality brain slices

Potential solution

It is important to ensure the integrity and quality of each brain slice. If it is difficult to obtain brain slice with good quality using the method described (step 9), an option is to solidify the brain with 4% low-melting-point agarose before sectioning as previously described (Fried et al., 2014).

RESOURCE AVAILABILITY

Lead contact

Further information and requests for resources and reagents should be directed to and will be fulfilled by the lead contact, Fei Yin (feiyin@arizona.edu)

Materials availability

This study did not generate new unique reagents.

Data and code availability

This study did not generate or analyze any datasets or code.

ACKNOWLEDGMENTS

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AUTHOR CONTRIBUTIONS

Conceptualization, F.Y.; methodology, G.Q., Y.M., and F.Y.; investigation, G.Q. and Y.M.; writing, G.Q., Y.M., and F.Y.; funding acquisition, F.Y.

DECLARATION OF INTERESTS

The authors declare no competing interests.

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