Basic theory of MANOVA, LDA and LOO methods

Because of the statistical methods are classical methods that have been widely used in data analyses, their description here is brief. For MANOVA, it is an extension of the univariate analysis of variance. This method is used in the cases where there are two or more dependent variables.

LDA is a classical pattern recognition method and can provide a classification model based on the combination of variables to predict a category or group to which a material belongs (Fisher 1936). The basic theory of LDA is to classify the dependent by dividing a *n*-dimensional parameters space into two regions. That space is separated by a hyperplane defined by a linear discriminant function as follows:

 $Y = a_0 + a_1 X_1 + a_2 X_2 + \dots + a_n X_n$

Where *Y* is a discriminant score, that is the dependant variable; $X_1,...,X_n$ represents the specific descriptor (in this case, metabolite levels); and $a_0...a_n$ are the weights of the corresponding parameters. The two regions formed by the hyperplane correspond to the two classes. Once the model was constructed, it is important to validate the predictabilit of that model. The simplest and a commonly used method of crossvalidation is the LOO method. In this method, the learning algorithm is trained multiple times, using all but one of the training set data points. At last, the mean error over all data points was calculated.

Journal of Neurochemistry

Table 1S: Mass and isotope balance equations for the metabolic calculation:

Mass balance equations:

$$dGlc_{brain}/dt = V_{\max in}Glc_{blood}/(K_M + Glc_{blood}) - V_{\max out}Glc_{brain}/(K_{Mout} + Glc_{brain}) - CMR_{glc}$$

$$d(Glu_N)/dt = d(Glu_A)/dt = d(KG_N)/dt = d(KG_A)/dt = d(Asp_N)/dt = d(Asp_A)/dt$$

$$= d(OAA_N)/dt = d(OAA_A)/dt = 0$$

Isotopic balance equations:

$$\begin{aligned} d(\operatorname{Glc}_{brain,C_{i}^{*}})/dt = V_{\operatorname{maxken}}(\operatorname{Glc}_{brain,C_{i}^{*}})/(\operatorname{Km}_{in} + \operatorname{Glc}_{brain}) - V_{\operatorname{maxken}}(\operatorname{Glc}_{brain,C_{i}^{*}})/(\operatorname{Km}_{out} + \operatorname{Glc}_{brain}) \\ - CMR_{ijk}(\operatorname{Glc}_{brain,C_{i}^{*}}/\operatorname{Glc}_{brain}) + V_{ditLac}(0) - (V_{pdh_{in}} + V_{pdh_{in}} + V_{pc} + V_{dit_{loc}} + V_{pdh_{cisk}})(L_{3}^{*}/L) \\ d(\operatorname{Glu}_{N_{4}^{*}})/dt = CMR_{ijk}(\operatorname{Glc}_{brain,C_{i}^{*}}/\operatorname{Gln}) + V_{x}(\operatorname{KG}_{N_{4}^{*}}/\operatorname{KG}_{N}) - (V_{cycle} + V_{x})(\operatorname{Glu}_{N_{4}^{*}}/\operatorname{Glu}_{N}) \\ d(\operatorname{Glu}_{N_{4}^{*}})/dt = V_{cycle}(\operatorname{Glu}_{N_{4}^{*}}/\operatorname{Glu}_{N}) + (V_{x} + V_{ann})(\operatorname{KG}_{A_{4}^{*}}/\operatorname{KG}_{A}) - (V_{Gin} + V_{x})(\operatorname{Glu}_{A_{4}^{*}}/\operatorname{Glu}_{A}) \\ d(\operatorname{Glu}_{A_{4}^{*}})/dt = V_{cycle}(\operatorname{Glu}_{A_{4}^{*}}/\operatorname{Glu}_{A}) + (V_{x} + V_{ann})(\operatorname{KG}_{A_{4}^{*}}/\operatorname{KG}_{A}) - (V_{Gin} + V_{x})(\operatorname{Glu}_{A_{4}^{*}}/\operatorname{Gln}) \\ d(\operatorname{KG}_{A_{4}^{*}})/dt = V_{goin}(\operatorname{Glu}_{A_{4}^{*}}/\operatorname{Glu}_{A}) + V_{xitGin}(0) - (V_{cycle} + V_{eflux} + V_{cycle}_{cannab}})(\operatorname{Gln}_{4}^{*}/\operatorname{KG}_{A}) \\ d(\operatorname{KG}_{A_{4}^{*}})/dt = V_{goin}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{xitG}(\operatorname{QA}_{A_{2}^{*}}/\operatorname{OAA}_{A}) - (V_{TCA_{A}Net} + V_{x})(\operatorname{KG}_{A_{5}^{*}}/\operatorname{KG}_{A}) \\ d(\operatorname{Glu}_{A_{5}^{*}})/dt = V_{cycle}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{xit}(\operatorname{KG}_{A_{5}^{*}}/\operatorname{KG}_{A}) - (V_{cin} + V_{x})(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) \\ d(\operatorname{Glu}_{A_{5}^{*}})/dt = V_{cycle}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{xit}(\operatorname{KG}_{A_{5}^{*}}/\operatorname{KG}_{A}) - (V_{cin} + V_{x})(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) \\ d(\operatorname{Glu}_{A_{5}^{*}})/dt = V_{cycle}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{dii}}(0) - (V_{cycle} + V_{egitax} + V_{cycle}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) \\ d(\operatorname{Glu}_{A_{5}^{*}})/dt = V_{cycle}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{dii}}(0) - (V_{cycle} + V_{egitax} + V_{cycle}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) \\ d(\operatorname{Glu}_{A_{5}^{*}})/dt = V_{x}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{cod}}(\operatorname{OAA}_{A_{2}^{*}}/\operatorname{OAA}_{A}) - (V_{TCA_{5}} + V_{x})(\operatorname{KG}_{A_{5}^{*}}/\operatorname{KG}_{A}) \\ d(\operatorname{Glu}_{A_{5}^{*}})/dt = V_{x}(\operatorname{Glu}_{A_{5}^{*}}/\operatorname{Glu}_{A}) + V_{cod}}(\operatorname{OAA}_{A_{2}^{*}}/\operatorname{OA$$

 $d(\text{GABA}_{2}^{*})/dt = V_{gad}(\text{Glu}_{GA_{4}}^{*}/\text{Glu}_{GA}) - [V_{shunt} + V_{cycle_{GABAGIn}}](\text{GABA}_{2}^{*}/\text{GABA})$

$$\begin{split} &d(KG_{G_{4}}^{*}/dt) = V_{p\beta\beta_{G_{G_{4}}}}(L_{3}^{*}/L) + V_{x_{G_{G_{4}}}}(Gu_{G_{4}}^{*}/Gu_{G_{4}}) + V_{dig_{G_{4}}}(0) \\ &- [V_{ICA_{G_{M}}NH} + V_{X_{G_{G_{4}}}}](KG_{G_{4}}^{*}/KG_{G_{4}}) \\ &d(Gu_{G_{4}}^{*}/dt) = V_{yeyt_{G_{G_{M}}}}(Gu_{G_{4}}^{*}/Gu_{G_{4}}) + V_{ICA_{G_{M}}}(KG_{G_{4}}) - [V_{g_{g_{g_{4}}}} + V_{X_{G_{G_{M}}}}](Gu_{G_{4}}^{*}/Gu_{G_{4}}) \\ &d(KG_{G_{4}}^{*}/dt) = V_{X_{G_{G_{M}}}}(Gu_{G_{4}}^{*}/Gu_{G_{4}}) + V_{ICA_{G_{M}}}(OAA_{G_{4}}^{*}/OAA_{G_{4}}) \\ &- [V_{ICA_{G_{M}}}N_{H} + V_{X_{G_{G_{G_{4}}}}}(Gu_{G_{4}}) - [V_{f_{G_{M}}} + V_{z_{G_{G_{M}}}}](GABA_{3}^{*}/GABA) \\ &d(GABA_{3}^{*})/dt = V_{g_{g_{G_{G_{M}}}}(Gu_{3}^{*}/Gu_{G_{4}}) - [V_{f_{M}} + V_{z_{G_{G_{M}}}}](GABA_{3}^{*}/ASp_{G_{4}}) \\ &d(OAA_{g_{4}}^{*})/dt = V_{z_{g_{G_{G_{M}}}}(Asp_{g_{3}}^{*}/Asp_{g_{4}}) + 0.5V_{ICA_{a}}(KG_{G_{A}}^{*}/KG_{a}) + 0.5V_{ICA_{a}}(KG_{A_{4}}^{*}/KG_{a}) \\ &- [V_{z_{advac}}}(Asp_{g_{3}}^{*}/Asp_{g_{3}}) + 0.5V_{ICA_{a}}(KG_{g_{4}}^{*}/KG_{g_{4}}) + 0.5V_{ICA_{a}}(KG_{A_{4}}^{*}/KG_{g_{4}}) \\ &- [V_{z_{advac}}(Asp_{g_{3}}^{*}/Asp_{g_{3}}) + 0.5V_{ICA_{a}}(KG_{g_{4}}^{*}/KG_{g_{4}}) + 0.5V_{ICA_{a}}(KG_{A_{4}}^{*}/KG_{g_{4}}) \\ &- [V_{z_{advac}}(Asp_{A_{3}}^{*}/Asp_{g_{3}}) + 0.5V_{ICA_{a}}(KG_{A_{3}}^{*}/Asp_{d_{4}}) + 0.5V_{ICA_{a}}(KG_{A_{4}}^{*}/KG_{g_{4}}) \\ &- [V_{z_{advac}}(Asp_{A_{3}}^{*}/Asp_{g_{3}}) + 0.5V_{ICA_{a}}(KG_{A_{3}}^{*}/KG_{a}) + 0.5V_{ICA_{a}}(KG_{A_{4}}^{*}/KG_{a}) \\ &- [V_{z_{advac}}(Asp_{A_{3}}^{*}/Asp_{d_{4}}) + 0.5V_{ICA_{a}}(KG_{A_{3}}^{*}/Asp_{d_{4}}) \\ &d(Asp_{A_{3}}^{*})/dt = V_{z_{A}OAAAg}(OAA_{A_{3}}^{*}/OAA_{a}) - V_{z_{A}AgOAA}(Asp_{A_{3}}^{*}/Asp_{A}) \\ &d(Asp_{A_{3}}^{*})/dt = V_{z_{A}OAAAg}(OAA_{A_{3}}^{*}/OAA_{a}) - V_{z_{A}AgOAA}(Asp_{A_{3}}^{*}/Asp_{A}) \\ &d(Asp_{A_{3}}^{*})/dt = 0.5V_{ICA_{a}NH}(KG_{A_{3}}^{*}/OAA_{a}) - V_{z_{A}AgOAA}(Asp_{A_{3}}^{*}/Asp_{A}) \\ &d(Asp_{A_{3}}^{*})/dt = 0.5V_{ICA_{a}NH}(KG_{A_{4}}^{*}/KG_{A}) + 0.5V_{ICA_{a}NH}(KG_{A_{4}}^{*}/Asp_{A}) \\ &- [V_{ICA_{a}NH}}(GAA_{A_{3}}^{*}/OAA_{A}) + 0.5V_{z_{a}AMAAA}N(Asp_{A_$$

Combination Pools:

 $Glu_{4_{total}} = Glu_{A4} + Glu_{N4} + Glu_{GA4}$

 $Glu_{3_total} = Glu_{A3} + Glu_{N3} + Glu_{GA3}$

The following are one group of rates and concentrations from the parietal cortex in a saline-treated group using the upper equations and combinational pools.

Rates:

- $CMR_{gl} = (Vpdh_A + Vpdh_N + Vpdh_{GABA} + V_{pc})/2 = 0.54 \ \mu mol/min/g; \ rate \ of \ glucose$ consumption
- $K_{m_{in}} = 13.9 \text{ mM}$; Michaelis-Menten half-saturation constant for blood-bran glucose Transport (Mason et al. 1992b)
- $K_{m_{out}} = K_{m_{in}} * V_d = 10.70 \ \mu mol/g;$ *Michaelis-Menten half-saturation constant for bran*blood glucose transport

 $R_{AspGlu} = 0.29 \mu mol/min/g$; Fraction of Asp that is in glutamatergic neurons

V_{cycle} = 0.49 μmol/min/g; *Rate of Glu-Gln cycling*

V_{cycleGABAGIn} = 0.46*V_{tcaGABA} = 0.053 μmol/min/g; *Rate of GABA-Gln cycling, fraction of* 0.46 from (Patel et al. 2005)

 $V_d = 0.77 \text{ ml/g}$; Brain water space (Buschiazzo et al. 1970)

 $V_{dilGG} = 0.046 \,\mu mol/min/g$; *Rate of dilution in GABAergic neurons*

V_{dilGln} = 0.35 μmol/min/g; Rate of diluting Gln exchange with blood

 $V_{dilN} = 0 \mu mol/min/g$; Rate of dilution in glutamatergic neurons

 $V_{efflux} = V_{pc} + V_{dilGln} = 0.42 \ \mu mol/min/g$; *Rate of loss of carbon from the astrocytic TCA cycle via efflux of Gln from the brain, partly balanced by entry of glutamine from the blood, also at the rate VdilGln.*

 $V_{gad} = V_{shunt} + V_{cycleGABAGIn} = 0.14 \ \mu mol/min/g; Rate of GABA synthesis$

 $V_{gln} = V_{cycle} + V_{cycleGABAGln} + V_{pc} = 0.60 \ \mu mol/min/g; Rate of Gln synthesis$

 $V_{max_{in}} = 5.8*CMR_{gl} = 3.14 \ \mu mol/min/g; V_{max}$ for glucose flow from blood to brain (Mason et al. 1992b).

 $V_{max out} = V_{max in} = 3.14 \mu mol/min/g$; *Vmax for glucose flow from brain to blood*

 $V_{pc} = V_{tcaA} - V_{tcaANet} - V_{cycleGABAGIn} = 0.065 \ \mu mol/min/g; Rate of anaplerosis$

 $V_{pdhA} = 0.176*V_{tcaN} = 0.14 \ \mu mol/min/g; Rate of astrocytic pyruvate dehydrogenase$ (Patel et al. 2005)

V_{pdhGABA} = 0.070 μmol/min/g; *Rate of pyruvate dehydrogenase in the GABAergic* neuron

 $V_{shunt} = V_{tcaGABA} - V_{tcaGABANet} = 0.088 \ \mu mol/min/g;$ Rate of GABA degradation in the GABA ergic neuron

 $V_{tcaGABA} = V_{pdhGABA} + V_{dilGG} = 0.12 \ \mu mol/min/g$; Rate of GABAergic TCA cycle from

citric acid through KG

V_{tcaGABANet} = 0.029 μmol/min/g; *Rate of GABAergic TCA cycle flow from KG to* succinate

 $V_{tcaN} = 0.81 \mu mol/min/g$; Rate of glutamatergic neuronal TCA cycle rate

- $V_{xAspOAAGA} = V_{xOAAAspGA} = 0.5 \ \mu mol/min/g;$ Rate of flow from Glu to KG in GABAergic neurons; value indeterminate
- $V_{XGGKG} = 0.5 \mu mol/min/g$; Rate of flow from Glu to KG in astrocytes; value ndeterminate
- $V_{XKGGG} = V_{XGGKG} + V_{gad} V_{cycleGABAGIn} = 0.59 \ \mu mol/min/g; Rate of flow from KG to Glu in astrocytes$
- $V_{XOAAAspGA} = V_{XGGKG} = 0.5 \ \mu mol/min/g; Rate of flow from OAA to Asp in GABAergic neurons$
- X₁ = V_{max_in}* Glucose_{1blood}/(Km_{_in}+ Glucose_{1Blood}) = 0.89 μmol/min/g; Variable used to calculate steady-state brain glucose concentrations
- X₂ = X₁-CMR_{gl} = 0.35 μmol/min/g; *Variable used to calculate steady-state brain glucose concentrations*

 $V_{dilA} = 0.0058 \ \mu mol/min/g$; Rate of dilution in astrocytes

 $V_{tcaA} = V_{ac} + V_{pdhA} + V_{dilA} = 0.15 \ \mu mol/min/g;$ Rate of TCA cycle in astrocytes

 $V_{tcaANet} = 0.029 \ \mu mol/min/g$; Rate of TCA from citric acid to KG in astrocytes

 $V_{XAAspOAA} = V_{XAOAAAsp} = 1 \mu mol/min/g$; Rate of flow from Asp to OAA in astrocytes

 $V_X = V_{XAGluKG} = 1 \ \mu mol/min/g$; *Rate of flow from OAA to Asp in astrocytes*

 $V_{XAKGGlu} = V_{XAGluKG} + V_{gln} - V_{cycle} = 1.12 \ \mu mol/min/g; Rate of flow from KG to Glu in astrocytes$

 $V_{XAOAAAsp} = V_{XAGluKG} = 1 \mu mol/min/g$; Rate of flow from OAA to Asp in astrocytes

 $V_{pdhN} = V_{tcaN} - V_{dilN} = 0.81 \ \mu mol/min/g$; *Rate of flow through glutamatergic neuronal pyruvate dehydrogenase*

 $V_{XNAspOAA} = V_{XNOAAAsp} = 3.64 \ \mu mol/min/g;$ Rate of flow from Asp to OAA in glutamatergic neurons

 $V_{XNGluKG} = 3.64 \ \mu mol/min/g$; Rate of flow from Glu to KG in glutamatergic neurons

V_{XNKGGlu} = V_{XNGluKG} = 3.64 μmol/min/g; *Rate of flow from KG to Glu in glutamatergic neurons*

 $V_{XNOAAAsp} = V_{XNGluKG} = 3.64 \ \mu mol/min/g$; Rate of flow from OAA to Asp in

1 2 2	
3 4 5	glutamatergic neurons
6 7	Pool Concentrations:
8 9	$Asp_{GA} = Asp_{Total} - Asp_A - Asp_N = 1.43 \ \mu mol/g; GABAergic neuronal Asp$
10 11	Asp _{Total} = 3.16 µmol/g; <i>Total tissue Asp (measured)</i>
12	Glucose _{brain} = Km _{out} * $X_2/(V_{max out} - X_2) = 1.33 \mu mol/g; Brain glucose (Mason et al.$
13 14	1992a)
15 16	GABA = 1.29 umol/s GABA concentration (measured)
17	Gln = 6.52 umol/s: Gln concentration (measured)
18 19	Gin = 0.02 µmol/g, Gin concentration (measured)
20	$Glu_{GA} = 0.02*Glu_{Total} = 0.21 \ \mu mol/g; Glu concentration in GABAergic neurons (small$
21 22	but indeterminate concentration)
23	$Glu_{Total} = 10.48 \ \mu mol/g; Total tissue Glu (measured)$
24 25	$KG_{GA} = 0.01 \ \mu mol/g; KG$ in GABAergic neurons, estimated from (Hawkins & Mans 1983)
26 27	L = 1.5 µmol/g; Tissue lactate concentration (Hawkins & Mans 1983)
28	$OAA_{GA} = 0.05 \mu mol/g$; OAA concentration in GABAergic neurons, estimated from
29 30	(Hawkins & Mans 1983)
31	(14000000000000000000000000000000000000
32 33	OAA _{Total} – 0.2 µmol/g, <i>Total tissue OAA</i> (<i>Hawkins & Mans</i> 1985)
34	$Asp_A = R_AspGlu*Glu_A = 0.30 \ \mu mol/g; Asp in astrocytes$
35 36	$Glu_A = 0.1*Glu_{Total} = 1.05 \ \mu mol/g; Glu in astrocytes (Patel et al. 2010, Lebon et al. 2002)$
37	$KG_A = 0.09 \ \mu mol/g$; KG in astrocytes, estimated from (Hawkins & Mans 1983)
30 39	$OAA_A = OAA_{Total}$ - OAA_N - $OAA_{GA} = 0.05 \mu mol/g$; <i>OAA in astrocytes, estimated from</i>
40 41	(Hawkins & Mans 1983)
42	$Asp_{M} = 0.5*(Asp_{Tatal} - Asp_{A}) = 1.43 \ \mu mol/\sigma$. Asp in alutamatergic neurons
43 44	$Glu_{r} = Glu_{r}$, Glu_{r} , $Glu_{r} = 0.22$ umol/g: Glu_{r} in glutamatorois neurons
45	$Glu_N = Glu_{Total} - Glu_GA = 9.22 \ \mu mol/g, Glu in gluiamatergic neurons$
46 47	(Patel et al. 2010, Lebon et al. 2002)
48	$KG_N = 0.1 \ \mu mol/g$; KG in glutamatergic neurons, estimated from (Hawkins & Mans 1983)
49 50	$OAA_N = 0.1 \ \mu mol/g$; OAA in glutamatergic neurons, estimated from (Hawkins & Mans
51 52	1983)
52 53	
54 55	Note: Glc glucose: L. lactate: Glu glutamate: Gln glutamine: Asp aspartate: OAA oxaloacetate:
56	KG_{α} ketoglutarate: Subscript 'N' 'A' and 'GA' stand for glutamateraic neuron astrogytes and
57 58	No, u-kelogiularate, subscript in, A and OA stand for giulamatergic neuron, astrocytes and
59	
60	

GABAergic neurons, respectively. Numeric subscripts represent the position of ¹³C in the corresponding molecule. Parameters listed in bold-faced type were determined by iterative fitting.

Table 2S.	Mass of	tissue sam	ples (mg) from each	region	across all	the rats.
			-r)			

No.	OB	CE	MEA	THA	HYP	HP	FC	PC	OC	ST	TC	MID
Mean	65.6	217.5	182.8	92.4	65.5	110.1	153.7	152.2	108.4	151.1	195.8	98.2
SD	9.7	21.7	21.4	12.7	15.5	10.7	23.1	22.3	14.0	30.9	23.5	12.5

Table 3S. The descriptors selected by the linear discriminant analysis and their corresponding statistical parameters for the whole group. Their selection was based on the ability to differentiate rats that received nicotine from those that received saline.

Region	Matabolita	Tolerance	F to	Wilks'	CDF
Kegion	Wietabolite		remove	Lambda	CDI
ST	GABA	0.7235	9.7047	0.4500	0.7563
ST	NAA	0.6113	20.4191	0.5890	1.0432
THA	GABA	0.5524	10.6467	0.4622	-0.8944
НҮР	Glu	0.4518	11.3550	0.4714	1.0114
TC	GABA	0.6021	7.8397	0.4258	-0.7659

Note: CDF: Standardized Canonical Discriminant Function Coefficients.

Journal of Neurochemistry

Table 4S: Effect of nicotine on cerebral cortical metabolic fluxes (μ mol·g·min⁻¹), including V_X (exchange rate between KG and glutamate in astrocytes), V_{dilGln} (the diluting inflow rate of glutamine), V_{tcaN} (TCA cycle flux in neuron), V_{dil} (total diluting flow into KG),

	V _X		V _{tcaN}		V _{dilGln}		V _{dil}	
Brain	Saline	Nicotine	Saline	Nicotine	Saline	Nicotine	Saline	Nicotine
CE	0.59±0.01	0.59±0.01	0.64±0.03	0.63±0.02	0.33±0.04	0.32±0.03	0.08±0.05	0.03±0.04
FC	0.57±0.01	0.57 ± 0.00	0.77±0.05	0.66±0.02	0.30±0.04	0.27±0.02	0.02±0.04	0.01 ± 0.02
TC	0.58±0.01	0.57±0.01	0.68±0.03	0.59±0.03	0.38±0.03	0.33±0.03	0.01±0.03	0.01±0.03
HP	0.58±0.01	0.57±0.01	0.64±0.03	0.52±0.02	0.37±0.04	0.30 ± 0.03	0.00 ± 0.04	0.02 ± 0.03
HYP	0.58±0.01	0.57±0.01	0.53±0.02	0.47 ± 0.02	0.36±0.03	0.36±0.03	0.01±0.03	0.01±0.03
MEA	0.59±0.01	0.57 ± 0.00	0.53±0.02	0.47 ± 0.02	0.39±0.03	0.34±0.03	0.00±0.03	0.02 ± 0.03
MID	0.61±0.01	0.60±0.01	0.60±0.03	0.54±0.02	0.35±0.03	0.31±0.03	0.02±0.03	$0.00{\pm}0.02$
OB	0.56±0.01	0.57 ± 0.00	0.58±0.04	0.52±0.03	0.33±0.04	0.34±0.04	0.02±0.05	0.02 ± 0.04
OC	0.59±0.01	0.58 ± 0.01	0.67±0.05	0.62±0.03	0.36±0.03	0.30±0.02	0.07±0.02	0.01 ± 0.02
PC	0.59±0.01	0.57±0.01	0.81±0.03	0.67±0.03	0.35±0.03	0.26±0.02	0.01±0.03	$0.00{\pm}0.02$
ST	0.58±0.01	0.55±0.01	0.67±0.02	0.55±0.02	0.30±0.03	0.28±0.02	0.00±0.03	0.01 ± 0.02
THA	0.58±0.01	0.58 ± 0.00	0.64±0.03	0.65±0.03	0.33±0.03	0.33±0.03	0.01±0.02	0.00±0.02

58±0.01 0.55±0.01 0.67±0.02 0.55±0.02 0.30±0.03 0.28±0.02 0.00±0.03 58±0.01 0.58±0.00 0.64±0.03 0.65±0.03 0.33±0.03 0.33±0.03 0.01±0.02

Table 5S: The p-values of pair-wise comparisons of rates of metabolism after nicotine injection in different regions, with indications of which rates survived adjustments for multiple comparisons.

	CE	FC	FDC	HP	HYP	MEA	MID	OB	OC	PC	ST	THA
CMR _{gl(ox)}	0.449	0.007	0.045	0.010	0.031	0.038	0.022	0.008	0.000*	0.000*	0.004	0.433
V _{cycle}	0.196	0.212	0.238	0.303	0.522	0.742	0.153	0.313	0.001*	0.000*	0.267	0.537
V_{gad}	0.578	0.109	0.427	0.314	0.096	0.035	0.354	0.883	0.037	0.039	0.025*	0.717

Note: *: The rates that remained significant after adjustment for multiple comparisons.



Fig. 1S. Time course of the concentration of GABA in striatum (A) Saline, (B) Nicotine.



Fig. 2S. Correlation between the rate of glucose oxidation (CMR_{gl(ox)}) and glutamate neurotransmitter cycling (V_{cycle}). The plot combined the data from the present study (control group = ●; nicotine = ▲) and the data from former studies (*) (Patel et al. 2004, Sibson et al. 1998). The data from other regions were also added (saline = ○; nicotine = Δ). The line represents the regression analysis of both the wake and

anaesthetized rat cortex, given by the equation y = 0.912x + 0.134, with a

correlation coefficient of r = 0.8672.



Fig. 3S: Relationship between glutamine synthetase activity in each region, taken as a ratio relative to the cortex, and the ratio of V_{cycle} in each region as a ratio relative to the value we measured for V_{cycle} in the parietal cortex. The regions for which glutamine synthetase data were available were the OB, ST, CE, HP, HYP, and MEA.

Bibliography

- Buschiazzo, P. M., Terrell, E. B. and Regen, D. M. (1970) Sugar transport across the blood-brain barrier. *Am J Physiol*, **219**, 1505-1513.
- Fisher, R. A. (1936) The use of multiple measurements in taxonomic problems. *Ann. Eugen.*, **7**, 179-188.
- Hawkins, R. A. and Mans, A. M. (1983) Intermediary metabolism of carboyhdrates and other fuels. I. In: *Biochem J*, (A. Lajtha ed.), Vol. 122, pp. 259-294. Plenum Press, New York.
- Lebon, V., Petersen, K. F., Cline, G. W., Shen, J., Mason, G. F., Dufour, S., Behar, K. L., Shulman, G. I. and Rothman, D. L. (2002) Astroglial contribution to brain energy metabolism in humans revealed by ¹³C nuclear magnetic resonance spectroscopy: Elucidation of the dominant pathway for neurotransmitter glutamate repletion and measurement of astrocytic oxidative metabolism. *J. Neurosci.*, **22**, 1523-1531.
- Mason, G. F., Behar, K. L., Rothman, D. L. and Shulman, R. G. (1992a) NMR Determination of Intracerebral Glucose-Concentration and Transport Kinetics in Rat-Brain. J. Cereb. Blood Flow Metab., **12**, 448-455.
- Mason, G. F., Rothman, D. L., Behar, K. L. and Shulman, R. G. (1992b) NMR determination of the TCA cycle rate and alpha-ketoglutarate/glutamate exchange rate in rat brain. *J Cereb Blood Flow Metab*, **12**, 434-447.
- Patel, A. B., de Graaf, R. A., Mason, G. F., Kanamatsu, T., Rothman, D. L., Shulman, R. G. and Behar, K. L. (2004) Glutamatergic Neurotransmission and Neuronal Glucose Oxidation Are Coupled During Intense Neuronal Activation. *J. Cereb. Blood Flow Metab.*, 24, 972-985.
- Patel, A. B., de Graaf, R. A., Mason, G. F., Rothman, D. L., Shulman, R. G. and Behar, K. L. (2005) The contribution of GABA to glutamate/glutamine cycling and energy metabolism in the rat cortex in vivo. *Proc Natl Acad Sci USA*, **102**, 5588-5593.
- Patel, A. B., de Graaf, R. A., Rothman, D. L., Behar, K. L. and Mason, G. F. (2010) Evaluation of Cerebral Acetate Transport and Metabolic Rates in the Rat Brain In Vivo using ¹H-[¹³C]NMR. J. Cereb. Blood Flow Metab., in press.
- Sibson, N. R., Dhankhar, A., Mason, G. F., Rothman, D. L., Behar, K. L. and Shulman, R. G. (1998) Stoichiometric coupling of brain glucose metabolism and glutamatergic neuronal activity. *P. Nati. Cad. Sci. USA*, 95, 316-321.