

TABLE S1. Equations Used in CWave Software to Calculate Metabolic Rates

Mass Balance Equations

$$\begin{aligned} d\text{Glu}_N/dt &= V_{\text{cycle}} + V_{\text{pdhN}} + V_{\text{dilN}} - (V_{\text{tcaN}} + V_{\text{cycle}}) = 0 \\ d\text{Glu}_A/dt &= V_{\text{cycle}} + V_{\text{pdhA}} + V_{\text{dilA}} - (V_{\text{gln}} + V_{\text{tcaANet}}) = 0 \\ d\text{Gln}/dt &= V_{\text{gln}} - (V_{\text{cycle_GABA}} + V_{\text{cycle}} + V_{\text{efflux}}) = 0 \\ d\text{GluGABA}/dt &= V_{\text{cycle_GABA}} + V_{\text{pdhGABA}} + V_{\text{dilGluGABA}} - (V_{\text{gad}} + V_{\text{tcaGABA}}) = 0 \\ d\text{GABA}/dt &= V_{\text{gad}} - (V_{\text{shunt}} + V_{\text{cycle_GABA}}) = 0 \\ d\text{TCANIntmd}_N/dt &= 0 \\ d\text{TCAIntmd}_{\text{gaba}}/dt &= 0 \\ d\text{TCAIntmd}_A/dt &= 0 \\ d\text{Lactate}/dt &= 2 \times \text{CMR}_{\text{gl}} - (V_{\text{ana}} + V_{\text{pdhA}} + V_{\text{pdhN}} + V_{\text{pdhGABA}}) = 0 \\ d\text{Glucose}/dt &= V_{\text{max_Glc_in}} (\text{Plasma_Glucose}) / (K_m_{\text{Glc_in}} + \text{Plasma_Glucose}) - [\text{CMR}_{\text{gl}} + (V_{\text{max_Glc_out}}) (\text{Glucose}) / (K_m_{\text{Glc_out}} + \text{Glucose})] = 0 \end{aligned}$$

Isotope Balance Equations

$$\begin{aligned} d\text{GluN}_4/dt &= V_{\text{cycle}} (\text{Gln}_4/\text{Gln}) + V_{\text{pdhN}} (\text{Lactate}_3/\text{Lactate}) + V_{\text{dilN}} (0) - [V_{\text{tcaN}} + V_{\text{cycle}}] (\text{GluN}_4/\text{GluN}) \\ d\text{GluA}_4/dt &= V_{\text{cycle}} (\text{GluN}_4/\text{GluN}) + V_{\text{pdhA}} (\text{Lactate}_3/\text{Lactate}) + V_{\text{dilA}} (\text{NA}_0/\text{NA}) - [V_{\text{gln}} + V_{\text{tcaANet}}] (\text{GluA}_4/\text{GluA}) \\ d\text{Gln}_4/dt &= V_{\text{gln}} (\text{GluA}_4/\text{GluA}) - [V_{\text{cycle_GABA}} + V_{\text{cycle}} + V_{\text{efflux}}] (\text{Gln}_4/\text{Gln}) \\ d\text{GluN}_3/dt &= V_{\text{tcaN}} (\text{TCANIntmd}_{N2}/\text{TCANIntmd}_N) + V_{\text{cycle}} (\text{Gln}_3/\text{Gln}) - [V_{\text{tcaN}} + V_{\text{cycle}}] (\text{GluN}_3/\text{GluN}) \\ d\text{Gln}_3/dt &= V_{\text{gln}} (\text{GluA}_3/\text{GluA}) - [V_{\text{cycle_GABA}} + V_{\text{cycle}} + V_{\text{efflux}}] (\text{Gln}_3/\text{Gln}) \\ d\text{GluA}_3/dt &= V_{\text{cycle}} (\text{GluN}_3/\text{GluN}) + V_{\text{tcaANet2}} (\text{TCAIntmd}_{A2}/\text{TCAIntmd}_A) - [V_{\text{gln}} + V_{\text{tcaANet}}] (\text{GluA}_3/\text{GluA}) \\ d\text{GluGABA}_4/dt &= V_{\text{cycle_GABA}} (\text{Gln}_4/\text{Gln}) + V_{\text{pdhGABA}} (\text{Lactate}_3/\text{Lactate}) + V_{\text{dilGluGABA}} (\text{NA}_0/\text{NA}) - [V_{\text{gad}} + V_{\text{tcaGABA}}] (\text{GluGABA}_4/\text{GluGABA}) \\ d\text{GluGABA}_3/dt &= V_{\text{cycle_GABA}} (\text{Gln}_3/\text{Gln}) + V_{\text{tcaGABA}} (\text{TCAIntmd}_{\text{gaba}2}/\text{TCAIntmd}_{\text{gaba}}) - [V_{\text{gad}} + V_{\text{tcaGABA}}] (\text{GluGABA}_3/\text{GluGABA}) \\ d\text{GABA}_2/dt &= V_{\text{gad}} (\text{GluGABA}_4/\text{GluGABA}) - [V_{\text{shunt}} + V_{\text{cycle_GABA}}] (\text{GABA}_2/\text{GABA}) \\ d\text{GABA}_3/dt &= V_{\text{gad}} (\text{GluGABA}_3/\text{GluGABA}) - [V_{\text{shunt}} + V_{\text{cycle_GABA}}] (\text{GABA}_3/\text{GABA}) \\ d\text{TCANIntmd}_{N2}/dt &= 1/2 V_{\text{tcaN}} (\text{GluN}_4/\text{GluN}) + 1/2 V_{\text{tcaN}} (\text{GluN}_3/\text{GluN}) + V_{\text{dilC3}} (\text{NA}_0/\text{NA}) - [V_{\text{tcaN}} + V_{\text{dilC3}}] (\text{TCANIntmd}_{N2}/\text{TCANIntmd}_N) \\ d\text{TCAIntmd}_{\text{gaba}2}/dt &= 1/2 V_{\text{tcaGABA}} (\text{GluGABA}_4/\text{GluGABA}) + 1/2 V_{\text{tcaGABA}} (\text{GluGABA}_3/\text{GluGABA}) + 1/2 V_{\text{shunt}} (\text{GABA}_2/\text{GABA}) + 1/2 V_{\text{shunt}} (\text{GABA}_3/\text{GABA}) - V_{\text{tcaGABA}} (\text{TCAIntmd}_{\text{gaba}2}/\text{TCAIntmd}_{\text{gaba}}) \end{aligned}$$

$$\begin{aligned}
dTCAIntmd_{A2}/dt &= 1/2 V_{cycle_GABA} (GABA_2/GABA) + 1/2 V_{cycle_GABA} (GABA_3/GABA) + 1/2 \\
&\quad V_{tcaANet} (GluA_4/GluA) + 1/2 V_{tcaANet} (GluA_3/GluA) + V_{ana} (\text{Lactate}_3/\text{Lactate}) - V_{tcaANet2} \\
&\quad (TCAIntmd_{A2}/TCAIntmd_A) \\
d\text{Lactate}_3/dt &= CMR_{gl} (\text{Glucose}_1/\text{Glucose}) - [V_{ana} + V_{pdhA} + V_{pdhN} + V_{pdhGABA}] (\text{Lactate}_3/\text{Lactate}) \\
d\text{Glucose}_1/dt &= V_{max_Glc_in} (\text{Plasma_Glucose}_1) / (K_{m_Glc_in} + \text{Plasma_Glucose}) - [CMR_{gl} + \\
&\quad V_{max_Glc_out} (\text{Glucose}) / (K_{m_Glc_out} + \text{Glucose})] (\text{Glucose}_1/\text{Glucose})
\end{aligned}$$

Variables

$\text{Glu}_{\text{Total}4} = \text{GluA}_4 + \text{GluN}_4 + \text{GluGABA}_4$; $\text{Glu}_{\text{Total}3} = \text{GluA}_3 + \text{GluN}_3 + \text{GluGABA}_3$; $\text{Glx}_3 = \text{Gln}_3 + \text{GluA}_3 + \text{GluGABA}_3 + \text{GluN}_3$; $CMR_{gl} = (V_{pdhA} + V_{pdhN} + V_{pdhGABA} + V_{ana}) / 2$ =cerebral metabolic rate of glucose; V_{cycle} =glutamate/glutamine neuron transmitter cycle, iterated; V_{pdhN} =neuronal pyruvate dehydrogenase rate, iterated; $V_{tcaN} = V_{pdhN} + V_{dilN}$ =neuronal TCA cycle rate; V_{shunt} =GABA shunt rate, iterated; V_{cycle_GABA} =rate of GABA/glutamine cycling; $V_{gad} = V_{shunt} + V_{cycle_GABA}$ =GABA synthesis rate; V_{ana} =anaplerotic flux; $V_{efflux} = V_{ana}$ =glutamine efflux, balanced by V_{ana} ; $V_{gln} = V_{cycle} + V_{cycle_GABA} + V_{efflux}$ =glutamine synthesis rate; V_{pdhA} =astrocytic pyruvate dehydrogenase rate; $V_{tcaA} = V_{pdhA} + V_{dilA}$ =astrocytic TCA cycle rate; $V_{pdhGABA} = V_{tcaGABA} - V_{dilGluGABA}$ =GABAergic pyruvate dehydrogenase rate; $V_{tcaANet} = V_{tcaA} - V_{gln} + V_{cycle}$ =astrocytic TCA cycle rate from alpha-KG to succinate; $V_{tcaANet2} = V_{tcaANet} + V_{cycle_GABA} + V_{ana}$ =astrocytic TCA cycle rate from succinate to oxaloacetate; $V_{tcaGABA_{net}} = GABA_{\text{ergic}} \text{ neuronal TCA cycle rate from a-KG to succinate}$; $V_{tcaGABA} = V_{tcaGABA_{net}} + V_{shunt}$ =GABAergic TCA cycle rate; $FracGlu_A$ =fraction of total glutamate that resides in astroglial; $FracGlu_{GABA}$ =fraction of total glutamate that resides in GABAergic neuron; $K_{m_Glc_in}$ =Michaelis-Menten half-saturation constant for blood-brain glucose transport; $K_{m_Glc_out} = V_d \times K_{m_Glc_in}$ =Michaelis-Menten half saturation constant for brain-blood glucose transport; V_d =distribution space of brain water; V_{dilA} =astrocytic dilution of acetyl CoA, iterated; V_{dilC3} =neuronal dilution of glutamate C3 relative to C4, iterated; $V_{dilGluGABA}$ =GABAergic neuronal dilution of acetyl CoA, iterated; V_{dilN} =glutamatergic neuronal siliation of acetyl CoA, iterated; $V_{max_Glc_in}$ =maximum rate of blood-brain glucose transport; $V_{max_Glc_out} = V_{max_Glc_in}$ =maximum rate if brain-blood glucose transport; $GABA$ =total brain GABA concentration; Gln =total brain glutamine concentration; Glu =total brain glutamate concentration; $Glu_A = FracGlu_A \times Glu_{\text{Total}}$ =astrocytic glutamate concentration; $Glu_N = Glu_{\text{Total}} - Glu_A - Glu_{GABA}$ =neuronal glutamate concentration; $Glu_{GABA} = FracGlu_{GABA} \times Glu_{\text{Total}}$ =GABAergic glutamate concentration; $TCAIntmd_A$ =astrocytic TCA cycle intermediates; $TCAIntmd_{gaba}$ =GABAergic neuronal TCA cycle intermediates; $TCANIntmd_N$ =glutamatergic neuronal TCA cycle intermediates

TABLE S2. Comparison of Occipital Metabolites

Metabolite	Depression Group (N=23)	Healthy Group (N=17)	<i>t</i>	<i>df</i>	<i>p</i>
	Mean ± SD	Mean ± SD			
GABA (mmol/kg)	1.12 ± 0.10	1.12 ± 0.12	0.04	38	0.97
Glutamine (mmol/kg)	2.32 ± 0.72	2.51 ± 0.70	0.54	38	0.41
Glutamate (mmol/kg)	7.42 ± 1.44	7.47 ± 1.57	0.36	38	0.92

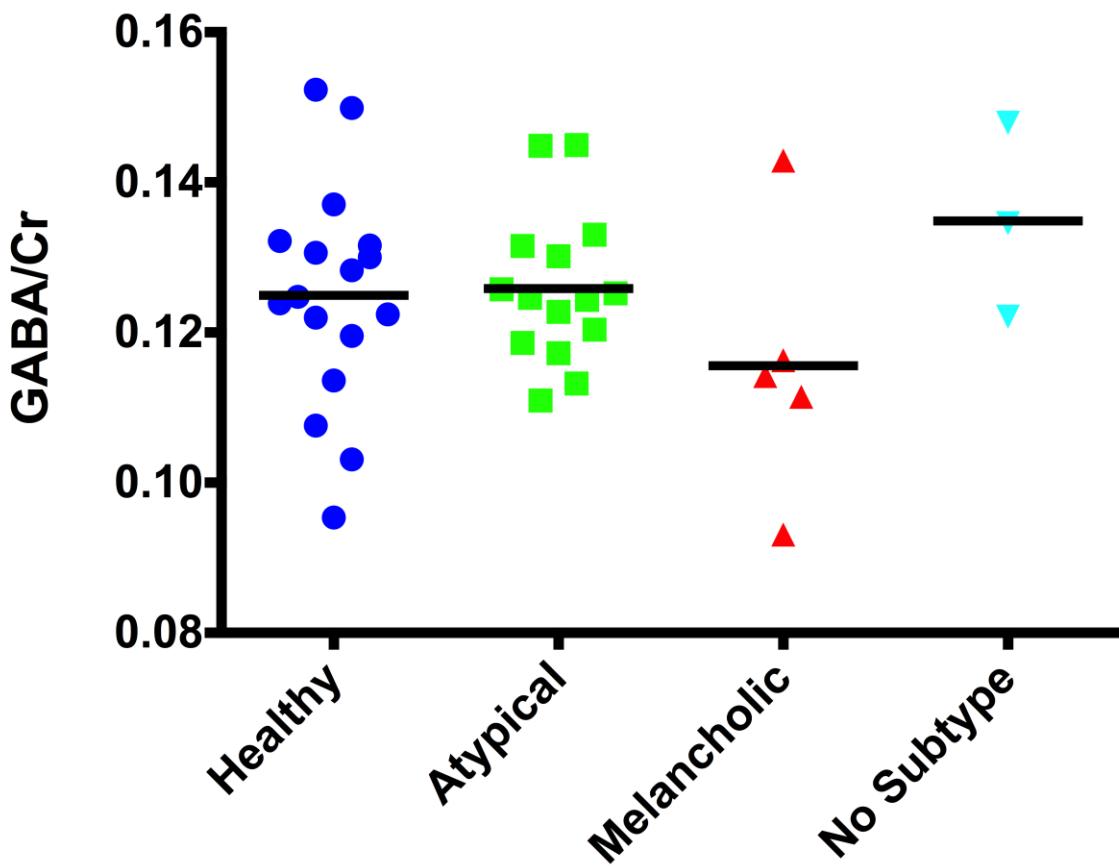
FIGURE S1. Scatter of Occipital GABA Levels

TABLE S3. Correlations (Spearman's Correlation Coefficient) Between Clinical And Spectroscopy Measures in the Depression Group

	HAM-D	HAM-A	BDI	Number of Episodes (Lifetime)
V _{TCAN}	0.19	-0.37 [‡]	0.09	0.19
V _{cycle}	0.15	0.10	0.29	-0.26
V _{GAD^a}	0.14	-0.17	0.07	-0.21
GABA	0.03	-0.14	-0.09	-0.10
Glutamine	-0.03	-0.47 *	-0.23	0.23
Glutamate ^b	0.39 [‡]	0.19	0.36	-0.59 *

^a V_{GAD} was excluded for three participants with noise level higher than 0.05 μmol/g/min.

^b One subject had poor spectral fitting for Glu and Glx.

Abbreviations: HAM-D: Hamilton Depression Rating Scale; HAM-A: Hamilton Anxiety Rating Scale; BDI: Beck Depression Inventory.

*p<0.05. ‡p<0.1

TABLE S4. Correlations Between V_{TCAN} and V_{cycle}

		Healthy	MDD
		V _{TCAN}	V _{TCAN}
Healthy	V _{cycle}	r _s =0.44, N=17, p=0.08	
Depression	V _{cycle}		r _s =0.25, N=23, p=0.25