

Table 1. Characterization of Hippocampus and HPA-axis.

	Cx (N=5-13)	LBW (N=8-14)	SSRI-LBW (N=8-16)	<i>Post hoc test</i> <i>P-values</i>	
<i>Adrenal Corticosterone Release</i>					
RST AUC 60 to 90 minutes (ng/*minutes)	12470±1270	19440±1240*	11180±1030	*P<0.01 vs. Cx & vs. SSRI-LBW	
24 H U-Corticosterone (nmol/[24 H*kg b.w.])	10.1±0.6	13.9±1.2†	8.1±0.5	†P<0.05 vs. Cx & P<0.01 vs. SSRI-LBW	
<i>Pituitary ACTH Release</i>					
Plasma ACTH concentrations (pg/ml)	9 a.m.	15.96±0.65	18.13±1.40	16.87±1.01	---
	9 p.m.	13.18±0.37‡	15.53±1.42	12.89±0.37‡	‡P<0.05 vs. 9a.m.
<i>Hypothalamic Corticosteroid Feedback</i>					
HSD1 mRNA Expression (% of Cx)	100.0±20.4	82.5±16.1	73.6±13.7	---	
MR mRNA Expression (% of Cx)	100.0±21.4	501.7±160.4§	205.8±60.0	§P<0.05 vs. Cx	
GR mRNA Expression (% of Cx)	100.0±15.8	123.9±15.8	121.6±46.5	---	
<i>Hippocampal Corticosteroid Feedback</i>					
HSD1 mRNA Expression (% of Cx)	100.0±15.5	122.2±25.3	67.5±8.6	---	
MR mRNA Expression (% of Cx)	100.0±10.0	123.2±13.3#	85.5±8.9	# P=0.0725 vs. Cx	
GR mRNA Expression (% of Cx)	100.0±14.1	122.5±11.1	88.1±10.9	---	

Data are means±sem. P-values for comparisons reflect the results for the *post hoc* analysis which has been carried out only when the one way of analysis of variance test (ANOVA) results in a p-value less than 0.05.

RST: Restrain Stress Test (Figure 1). 24 H U-Corticosterone: 24 hour urinary corticosterone excretion.

Table 2. Oral Glucose Tolerance Test (OGTT).

	Cx (N=13)	LBW (N=11)	SSRI-LBW (N=13)	Post hoc test P-values
<i>Oral Glucose Tolerance Test</i>				
Fasting Glucose Concentrations (mmol/l)	4.55±0.13	4.37±0.11	4.72±0.13	---
Fasting Insulin Concentrations (mU/l)	7.83±1.12	26.93±5.24*	11.83±2.17	*P<0.01 vs. Cx & P<0.05 vs. SSRI-LBW
OGTT Total AUC _{glucose} (mmol/l glucose*min)	871±20	898±23	767±29†	†P<0.05 vs. Cx & P<0.01 vs. LBW
OGTT Total AUC _{insulin} (mU/l insulin*min)	1,542±39	3,738±79‡	1,996±37	‡P<0.01 vs. Cx & SSRI-LBW
Insulin-Sensitivity Index (ISI) (expressed as % of Cx)	100.0±10.3	51.1±6.3§	85.0±8.1	§P<0.01 vs. Cx & P<0.05 vs. SSRI-LBW

Data are means±sem. P-values for comparisons reflect the results for the *post hoc* analysis which has been carried out only when the one way of analysis of variance test (ANOVA) results in a p-value less than 0.05.

Table 3. Muscle Glucose Metabolism.

	Cx (N=5-14)	LBW (N=8-9)	SSRI-LBW (N=6-8)	Post hoc test P-values
<i>Glycogen & GLUT4-Protein Content in Red (RG) and White (WG) Gastrocnemius Muscle Tissue</i>				
Basal Glycogen WG (µmol/g muscle w.w.)	38.8±1.3	40.5±1.7	42.9±1.8	---
Basal Glycogen RG (µmol/g muscle w.w.)	31.4±2.2*	33.8±1.7*	32.4±2.3*	†P<0.05 vs. WG
Total GLUT4 Content WG (arbitrary units)	9.89±3.36	10.32±0.90	9.97±0.58	---
Total GLUT4 Content RG (arbitrary units)	13.93±0.71*	16.38±1.92*	13.22±0.98*	†P<0.05 vs. WG
<i>PkB Ser (473) Phosphorylation in Red Gastrocnemius Muscle Tissue</i>				
Basal Phosphorylation (arbitrary units)	29,568±3,239	31,691±4,901	22,459±4,429	---
Clamp Phosphorylation (arbitrary units)	69,596±10,600†	56,729±7,641†	76,127±8,287†‡	†P<0.05 vs. Basal ‡P<0.10 vs. LBW
Insulin-Stimulated Increase from Basal (%)	135±4	79±24	239±43§	§P<0.05 vs. LBW

Data are means±sem. P-values for comparisons reflect the results for the *post hoc* analysis which has been carried out only when the one way of analysis of variance test (ANOVA) results in a p-value less than 0.05.

For online publication.

Primer sequences for PCR experiments (Buhl ES et al, 2009)	
mRNA Expression Studies in Liver Tissue	
G6Pase forward primer	AGG GTA AAA GAA AAG AGC GTT G
G6Pase reverse primer	GTA GAC ATG GCT TGC ATA TGG T
PEPCK forward primer	CAG GAA GTG AGG AAG TTT GTG G
PEPCK reverse primer	ATG ACA CCC TCC TCC TGC AT
GAPDH forward primer	TCACCACCATGGAGAAGGC
GAPDH reverse primer	GCTAAGCAGTTGGTGGTGCA
mRNA Expression Studies in Brain Tissue	
GR forward primer	GGTACTCAAGCCCTGGAATG
GR reverse primer	CAATCGTTTCTTCCAGCACA
MR forward primer	TGAGTTCCTTCCCACCTGTC
MR reverse primer	AAGCCTCATCTCCACACACC
HSD1 forward primer	CTCTCTGTGTCCTCGGCTTC
HSD1 reverse primer	TTCCATGATCCTCCTTCCTG
18S forward primer	ACGGACCAGAGCGAAAGCAT
18S reverse primer	TGTCAATCCTGTCCGTGTCC
Beta-Actin forward primer	TGTCACCAACTGGGACGATA
Beta-Actin reverse primer	GGGGTGTGTAAGGTCTCAA
Cyclophilin A forward primer	AGCACTGGGGAGAAAGATT
Cyclophilin A reverse primer	AGCCACTCAGTCTTGGCAGT
GAPDH forward primer	TCACCACCATGGAGAAGGC
GAPDH reverse primer	GCTAAGCAGTTGGTGGTGCA
Hmbs forward primer	TCCTGGCTTTACCATTGGAG
Hmbs reverse primer	TGAATTCCAGGTGAGGGAAC
Hprt1 forward primer	GCAGACTTTGCTTTCCTTGG
Hprt1 reverse primer	CGAGAGGTCC TTTTACCAG
Rpl13A forward primer	ACAAGAAAAAGCGGATGGTG
Rpl13A reverse primer	TTCCGGTAATGGATCTTTC
Ywhaz forward primer	TTGAGCAGAAGACGGAAGGT
Ywhaz reverse primer	GAAGCATTGGGGATCAAGAA

G6Pase: glucose-6-phosphatase; PEPCK: Phosphor-enol-pyruvate-carboxy-kinase; GAPDH: Glyceraldehyde-3-phosphate dehydrogenase; GR: Glucocorticoid-receptor; MR: Mineralo-corticoid-receptor; 18S: 18s subunit ribosomal RNA; HSD1: 11 β hydroxysteroid dehydrogenase 1; Hmbs : Hydroxy-methylbilane synthase; Hprt1: Hypoxanthine guanine phosphoribosyl transferase 1; Rpl13A: Ribosomal protein L13A; Ywhaz :Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta;