

Table S1. Primer references for TaqMan® Gene Expression Assays (Applied Biosystems).

Gene description	Assay ID	No accession GenBank	Amplicon Length
<i>Actb</i>	Rn00667869_m1	NM_031144.3	91
<i>Cnr1</i>	Rn02758689_s1	NM_012784.4	92
<i>Cnr2</i>	Rn01637601_m1	NM_020543.4	68
<i>Dagla</i>	Rn01454304_m1	NM_001005886.1	67
<i>Daglb</i>	Rn01453771_m1	NM_001107120.1	98
<i>Mgll</i>	Rn00593297_m1	NM_138502.2	78
<i>Napepld</i>	Rn01786262_m1	NM_199381.1	71
<i>Faah</i>	Rn00577086_m1	NM_024132.3	63
<i>Ppara</i>	Rn00566193_m1	NM_013196.1	98
<i>Crh</i>	Rn01462137_m1	NM_031019.1	112
<i>Crhr1</i>	Rn00578611_m1	XM_006247542.2	58
<i>Npy</i>	Rn00561681_m1	NM_012614.2	63
<i>NpyIr</i>	Rn02769337_s1	NM_001113357.1	98
<i>Grm5</i>	Rn00566628_m1	NM_017012.1	112
<i>Grin1</i>	Rn01436034_m1	NM_001270602.1	73
<i>Grin2b</i>	Rn00680474_m1	NM_012574.1	79
<i>Gria2</i>	Rn00568514_m1	NM_001083811.1	122

Table S2. Summary of changes in the gene expression of components of the endocannabinoid, glutamatergic, CRH and NPY systems in the amygdala and the mPFC of male rats exposed to restraint stress and/or intermittent alcohol during adolescence.

Signaling systems (genes)	AMYGDALA*			mPFC*		
	Stress	Alcohol	Stress+Alcohol	Stress	Alcohol	Stress+Alcohol
ECS						
<i>Cnr1</i>	▼	▼	▼	—	▲	▼
<i>Cnr2</i>	▲	▲	▲	—	↓	▼
<i>Ppara</i>	—	▼	▼	▲	▲	▼
<i>Napepld</i>	↓	▼	▼	—	▲	▼
<i>Dagla</i>	—	—	↓	—	▲	—
<i>Daglb</i>	▼	▼	▼	—	▲	—
<i>Faah</i>	▼	▼	▼	—	▼	▼
<i>Mgll</i>	▼	▼	▼	▲	▲	▼
Glutamate						
<i>Grm5</i>	—	▲	—	—	▲	↑
<i>Grin1</i>	—	—	▼	↓	▼	▼
<i>Grin2B</i>	—	—	▲	↓	—	↓
<i>Gria2</i>	↑	↑	↑	↑	▲	▲
CRH						
<i>Crh</i>	↓	▼	▼	X	X	X
<i>Crhr1</i>	↓	▲	▲			
NPY						
<i>Npy</i>	↓	▼	▼	X	X	X
<i>Npyr1</i>	—	▼	—			

(*) The direction of the changes was determined in relation to the non-stress saline subgroup in the amygdala and the mPFC. Symbols: (▼▲) = significant difference, (↓↑) = tendency, (—) = no change, (X) = not determined.