

Supplementary Table S1. Overall linear mixed model analysis for *Tph2* mRNA expression using a first-order ante-dependence covariance structure by REML, selected by best -2 log likelihood criterion.

Source	Test statistic	p-value
Cohort	$F(5, 211.906) = 31.090$	0.001***
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 258.786) = 3.665$	0.057
Chronic disruption of rhythms (CDR)	$F(1, 309.394) = 23.620$	0.001***
Acute social defeat (SD)	$F(1, 289.818) = 4.263$	0.040*
Subregion	$F(5, 169.079) = 26.768$	0.001***
Rostrocaudal level	$F(13, 200.949) = 25.673$	0.001***
<i>Mv</i> × Subregion	$F(5, 169.077) = 0.359$	0.876
CDR × Subregion	$F(5, 169.077) = 1.058$	0.385
SD × Subregion	$F(5, 169.076) = 0.274$	0.927
Rostrocaudal level × Subregion	$F(23, 221.234) = 22.749$	0.001***
<i>Mv</i> × CDR	$F(1, 260.102) = 0.019$	0.889
<i>Mv</i> × SD	$F(1, 263.026) = 0.050$	0.823
<i>Mv</i> × Rostrocaudal level	$F(13, 200.452) = 2.014$	0.021*
CDR × SD	$F(1, 280.639) = 0.194$	0.660
CDR × Rostrocaudal level	$F(13, 200.561) = 3.557$	0.001***
SD × Rostrocaudal level	$F(13, 200.636) = 1.452$	0.138
<i>Mv</i> × CDR × Subregion	$F(5, 169.074) = 0.085$	0.995
<i>Mv</i> × SD × Subregion	$F(5, 169.076) = 0.326$	0.897
<i>Mv</i> × CDR × SD	$F(1, 271.492) = 0.093$	0.760
<i>Mv</i> × CDR × Rostrocaudal level	$F(13, 200.581) = 3.474$	0.001***
CDR × SD × Rostrocaudal level	$F(13, 200.437) = 1.462$	0.134
<i>Mv</i> × Subregion × Rostrocaudal level	$F(23, 221.225) = 0.616$	0.916
<i>Mv</i> × CDR × SD × Subregion	$F(10, 169.287) = 0.353$	0.964
<i>Mv</i> × CDR × Subregion × Rostrocaudal level	$F(46, 228.603) = 1.269$	0.132
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(26, 203.560) = 1.467$	0.075
<i>Mv</i> × CDR × SD × Subregion × Rostrocaudal level	$F(92, 226.588) = 0.386$	1.000

* $p < 0.05$, ** $p < 0.01$, * $p < 0.001$

Supplementary Table S2. Subregional linear mixed model analyses for *Tph2* mRNA expression using unstructured covariance models by REML and best -2 log likelihood criteria.

Dorsal raphe nucleus, dorsal part	Test statistic	p-value
Cohort	$F(5, 19.784) = 2.121$	0.105
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 42.367) = 0.54$	0.466
Chronic disruption of rhythms (CDR)	$F(1, 43.394) = 6.191$	0.017*
Acute social defeat (SD)	$F(1, 42.635) = 1.048$	0.312
Rostrocaudal level	$F(7, 38.479) = 12.869$	0.001***
<i>Mv</i> × CDR	$F(1, 42.426) = 0.036$	0.851
<i>Mv</i> × SD	$F(1, 42.412) = 0.001$	0.982
<i>Mv</i> × Rostrocaudal level	$F(7, 38.475) = 1.068$	0.402
CDR × SD	$F(1, 42.609) = 0.025$	0.876

CDR × Rostrocaudal level	$F(7, 38.445) = 1.883$	0.099
SD × Rostrocaudal level	$F(7, 38.466) = 1.090$	0.389
<i>Mv</i> × CDR × SD	$F(1, 42.730) = 0.643$	0.427
CDR × SD × Rostrocaudal level	$F(7, 38.460) = 0.387$	0.904
<i>Mv</i> × CDR × Rostrocaudal level	$F(7, 38.492) = 1.297$	0.278
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(14, 38.396) = 1.469$	0.170
Dorsal raphe nucleus, ventral part		
Cohort	$F(5, 23.731) = 1.684$	0.177
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 42.634) = 0.594$	0.445
Chronic disruption of rhythms (CDR)	$F(1, 43.397) = 5.7$	0.021*
Acute social defeat (SD)	$F(1, 42.844) = 1.413$	0.241
Rostrocaudal level	$F(6, 37.567) = 18.784$	0.001***
<i>Mv</i> × CDR	$F(1, 42.629) = 0.060$	0.808
<i>Mv</i> × SD	$F(1, 42.634) = 0.013$	0.910
<i>Mv</i> × Rostrocaudal level	$F(6, 37.571) = 1.262$	0.298
CDR × SD	$F(1, 42.863) = 0.017$	0.896
CDR × Rostrocaudal level	$F(6, 37.553) = 1.294$	0.284
SD × Rostrocaudal level	$F(6, 37.553) = 1.431$	0.229
<i>Mv</i> × CDR × SD	$F(1, 42.781) = 0.392$	0.535
CDR × SD × Rostrocaudal level	$F(6, 37.570) = 0.748$	0.615
<i>Mv</i> × CDR × Rostrocaudal level	$F(6, 37.557) = 3.022$	0.016*
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(12, 37.541) = 1.573$	0.142
Dorsal raphe nucleus, ventrolateral part/ Ventrolateral periaqueductal grey		
Cohort	$F(5, 43.017) = 5.736$	0.001***
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 39.746) = 1.813$	0.186
Chronic disruption of rhythms (CDR)	$F(1, 44.11) = 5.425$	0.024*
Acute social defeat (SD)	$F(1, 41.083) = 0.435$	0.513
Rostrocaudal level	$F(3, 44.673) = 3.650$	0.019*
<i>Mv</i> × CDR	$F(1, 39.98) = 0.131$	0.719
<i>Mv</i> × SD	$F(1, 40.431) = 0.529$	0.471
<i>Mv</i> × Rostrocaudal level	$F(3, 44.663) = 1.855$	0.151
CDR × SD	$F(1, 41.118) = 0.020$	0.889
CDR × Rostrocaudal level	$F(3, 44.713) = 0.340$	0.797
SD × Rostrocaudal level	$F(3, 44.662) = 1.198$	0.321
<i>Mv</i> × CDR × SD	$F(1, 41.181) = 0.188$	0.667
CDR × SD × Rostrocaudal level	$F(3, 44.665) = 0.178$	0.911
<i>Mv</i> × CDR × Rostrocaudal level	$F(3, 44.704) = 0.391$	0.760
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(6, 44.362) = 1.463$	0.213
Dorsal raphe nucleus, caudal part		
Cohort	$F(5, 30.659) = 3.893$	0.008**
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 34.889) = 2.115$	0.155
Chronic disruption of rhythms (CDR)	$F(1, 38.005) = 6.502$	0.015*
Acute social defeat (SD)	$F(1, 35.983) = 0.224$	0.639

Rstrocaudal level	$F(5, 27.88) = 52.541$	0.001***
<i>Mv</i> × CDR	$F(1, 34.554) = 0.328$	0.571
<i>Mv</i> × SD	$F(1, 35.305) = 0.008$	0.931
<i>Mv</i> × Rstrocaudal level	$F(5, 27.868) = 2.671$	0.043*
CDR × SD	$F(1, 36.132) = 1.473$	0.233
CDR × Rstrocaudal level	$F(5, 27.887) = 3.816$	0.009**
SD × Rstrocaudal level	$F(5, 27.895) = 0.506$	0.769
<i>Mv</i> × CDR × SD	$F(1, 36.077) = 0.730$	0.399
CDR × SD × Rstrocaudal level	$F(5, 27.903) = 2.069$	0.100
<i>Mv</i> × CDR × Rstrocaudal level	$F(5, 27.878) = 3.279$	0.019*
<i>Mv</i> × CDR × SD × Rstrocaudal level	$F(10, 27.900) = 0.820$	0.613
Dorsal raphe nucleus, interfascicular part		
Cohort	$F(5, 37.701) = 11.198$	0.001***
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 37.738) = 1.038$	0.315
Chronic disruption of rhythms (CDR)	$F(1, 44.437) = 9.975$	0.003**
Acute social defeat (SD)	$F(1, 38.795) = 1.323$	0.257
Rstrocaudal level	$F(8, 32.831) = 12.876$	0.001***
<i>Mv</i> × CDR	$F(1, 37.880) = 0.167$	0.685
<i>Mv</i> × SD	$F(1, 38.778) = 0.628$	0.433
<i>Mv</i> × Rstrocaudal level	$F(8, 32.822) = 0.533$	0.823
CDR × SD	$F(1, 38.836) = 0.068$	0.795
CDR × Rstrocaudal level	$F(8, 32.751) = 3.110$	0.010*
SD × Rstrocaudal level	$F(8, 32.678) = 1.048$	0.422
<i>Mv</i> × CDR × SD	$F(1, 39.377) = 0.115$	0.736
CDR × SD × Rstrocaudal level	$F(8, 32.742) = 1.535$	0.183
<i>Mv</i> × CDR × Rstrocaudal level	$F(8, 32.837) = 2.782$	0.018*
<i>Mv</i> × CDR × SD × Rstrocaudal level	$F(16, 32.310) = 0.944$	0.533
Median raphe nucleus		
Cohort	$F(5, 39.975) = 3.024$	0.021*
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 24.542) = 0.114$	0.739
Chronic disruption of rhythms (CDR)	$F(1, 25.475) = 1.736$	0.199
Acute social defeat (SD)	$F(1, 24.829) = 0.836$	0.369
Rstrocaudal level	$F(7, 31.870) = 11.334$	0.001***
<i>Mv</i> × CDR	$F(1, 24.677) = 0.058$	0.811
<i>Mv</i> × SD	$F(1, 24.705) = 0.798$	0.380
<i>Mv</i> × Rstrocaudal level	$F(7, 31.974) = 1.613$	0.168
CDR × SD	$F(1, 24.848) = 1.731$	0.200
CDR × Rstrocaudal level	$F(7, 31.946) = 0.947$	0.485
SD × Rstrocaudal level	$F(7, 31.991) = 1.204$	0.329
<i>Mv</i> × CDR × SD	$F(1, 24.900) = 1.035$	0.319
CDR × SD × Rstrocaudal level	$F(7, 31.969) = 1.672$	0.152
<i>Mv</i> × CDR × Rstrocaudal level	$F(7, 32.003) = 0.973$	0.467
<i>Mv</i> × CDR × SD × Rstrocaudal level	$F(14, 31.232) = 0.458$	0.939

* $p < 0.05$, ** $p < 0.01$, * $p < 0.001$

Supplementary Table S3. Overall linear mixed model analysis for *Slc6a4* mRNA expression using a first-order ante-dependence covariance structure by REML, selected by best -2 log likelihood criterion.

Source	Test statistic	p-value
Cohort	$F(5, 119.812) = 20.467$.000***
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 286.887) = 2.513$.114
Chronic disruption of rhythms (CDR)	$F(1, 356.592) = 45.539$.000***
Acute social defeat (SD)	$F(1, 327.888) = 1.984$.160
Subregion	$F(5, 156.763) = 32.663$.001***
Rostrocaudal level	$F(13, 213.074) = 46.419$.001***
<i>Mv</i> × Subregion	$F(5, 156.766) = 0.322$.899
CDR × Subregion	$F(5, 156.764) = 1.461$.206
SD × Subregion	$F(5, 156.763) = 0.198$.963
Rostrocaudal level × Subregion	$F(23, 268.966) = 30.625$.001***
<i>Mv</i> × CDR	$F(1, 289.011) = 0.290$.591
<i>Mv</i> × SD	$F(1, 286.801) = 5.369$.021*
<i>Mv</i> × Rostrocaudal level	$F(13, 212.975) = 2.256$.009**
CDR × SD	$F(1, 313.071) = 4.867$.028*
CDR × Rostrocaudal level	$F(13, 212.922) = 4.854$.001***
SD × Rostrocaudal level	$F(13, 212.603) = 1.398$.162
<i>Mv</i> × CDR × Subregion	$F(5, 156.766) = 0.104$.991
<i>Mv</i> × SD × Subregion	$F(5, 156.765) = 0.291$.917
<i>Mv</i> × CDR × SD	$F(1, 299.027) = 7.710$.006**
<i>Mv</i> × CDR × Rostrocaudal level	$F(13, 212.832) = 1.490$.123
CDR × SD × Rostrocaudal level	$F(13, 212.683) = 1.902$.031*
<i>Mv</i> × Subregion × Rostrocaudal level	$F(23, 268.963) = 0.590$.934
<i>Mv</i> × CDR × SD × Subregion	$F(10, 157.013) = 0.195$.996
<i>Mv</i> × CDR × Subregion × Rostrocaudal level	$F(46, 268.202) = 1.345$.079
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(26, 212.473) = 2.061$.003**
<i>Mv</i> × CDR × SD × Subregion × Rostrocaudal level	$F(92, 265.55) = 0.721$.966

* $p < 0.05$, ** $p < 0.01$, * $p < 0.001$

Supplementary Table S4. Subregional linear mixed model analyses for *Slc6a4* mRNA expression using unstructured covariance models by REML and best -2 log likelihood criteria.

Dorsal raphe nucleus, dorsal part	Test statistic	p-value
Cohort	$F(5, 27.194) = 2.477$	0.057
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 43.808) = 0.526$	0.472
Chronic disruption of rhythms (CDR)	$F(1, 45.978) = 13.306$	0.001***
Acute social defeat (SD)	$F(1, 44.112) = 0.421$	0.520
Rostrocaudal level	$F(7, 40.513) = 33.702$	0.001***
<i>Mv</i> × CDR	$F(1, 43.878) = 0.001$	0.981
<i>Mv</i> × SD	$F(1, 43.862) = 0.923$	0.342

<i>Mv</i> × Rostrocaudal level	$F(7, 40.363) = 1.992$	0.080
CDR × SD	$F(1, 44.036) = 0.917$	0.344
CDR × Rostrocaudal level	$F(7, 40.459) = 5.477$	0.001***
SD × Rostrocaudal level	$F(7, 40.340) = 1.219$	0.315
<i>Mv</i> × CDR × SD	$F(1, 44.489) = 0.128$	0.722
CDR × SD × Rostrocaudal level	$F(7, 40.332) = 0.525$	0.810
<i>Mv</i> × CDR × Rostrocaudal level	$F(7, 40.408) = 0.321$	0.940
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(14, 40.529) = 0.898$	0.567
Dorsal raphe nucleus, ventral part		
Cohort	$F(5, 28.039) = 2.711$	0.040*
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 45.882) = 0.51$	0.479
Chronic disruption of rhythms (CDR)	$F(1, 46.823) = 9.29$	0.004**
Acute social defeat (SD)	$F(1, 46.054) = 0.533$	0.469
Rostrocaudal level	$F(6, 40.270) = 34.237$	0.001***
<i>Mv</i> × CDR	$F(1, 45.924) = 0.004$	0.951
<i>Mv</i> × SD	$F(1, 45.897) = 0.851$	0.361
<i>Mv</i> × Rostrocaudal level	$F(6, 40.253) = 0.884$	0.515
CDR × SD	$F(1, 45.988) = 0.376$	0.543
CDR × Rostrocaudal level	$F(6, 40.253) = 3.378$	0.009**
SD × Rostrocaudal level	$F(6, 40.227) = 1.552$	0.186
<i>Mv</i> × CDR × SD	$F(1, 46.171) = 0.294$	0.590
CDR × SD × Rostrocaudal level	$F(6, 40.252) = 0.856$	0.535
<i>Mv</i> × CDR × Rostrocaudal level	$F(6, 40.230) = 0.573$	0.749
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(12, 40.256) = 1.322$	0.244
Dorsal raphe nucleus, ventrolateral part/ Ventrolateral periaqueductal grey		
Cohort	$F(5, 42.157) = 13.173$	0.001***
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 46.626) = 0.396$	0.532
Chronic disruption of rhythms (CDR)	$F(1, 51.324) = 3.998$	0.051
Acute social defeat (SD)	$F(1, 47.330) = 0.182$	0.672
Rostrocaudal level	$F(3, 47.492) = 5.661$	0.002**
<i>Mv</i> × CDR	$F(1, 46.775) = 1.440$	0.236
<i>Mv</i> × SD	$F(1, 47.047) = 0.184$	0.670
<i>Mv</i> × Rostrocaudal level	$F(3, 47.475) = 0.808$	0.496
CDR × SD	$F(1, 47.286) = 0.137$	0.713
CDR × Rostrocaudal level	$F(3, 47.357) = 0.160$	0.923
SD × Rostrocaudal level	$F(3, 47.399) = 0.368$	0.776
<i>Mv</i> × CDR × SD	$F(1, 48.036) = 0.052$	0.821
CDR × SD × Rostrocaudal level	$F(3, 47.403) = 0.790$	0.505
<i>Mv</i> × CDR × Rostrocaudal level	$F(3, 47.429) = 0.799$	0.501
<i>Mv</i> × CDR × SD × Rostrocaudal level	$F(6, 47.529) = 0.708$	0.645
Dorsal raphe nucleus, caudal part		
Cohort	$F(5, 24.962) = 10.162$	0.001***
<i>Mycobacterium vaccae</i> NCTC 11659 (<i>Mv</i>)	$F(1, 43.422) = 0.919$	0.343

Chronic disruption of rhythms (CDR)	$F(1, 45.568) = 7.038$	0.011*
Acute social defeat (SD)	$F(1, 43.419) = 0.168$	0.684
Rostrocaudal level	$F(5, 39.587) = 20.033$	0.001***
$Mv \times CDR$	$F(1, 43.402) = 1.896$	0.176
$Mv \times SD$	$F(1, 43.495) = 2.432$	0.126
$Mv \times Rostrocaudal level$	$F(5, 39.756) = 1.693$	0.159
$CDR \times SD$	$F(1, 43.437) = 0.590$	0.447
$CDR \times Rostrocaudal level$	$F(5, 39.705) = 2.223$	0.071
$SD \times Rostrocaudal level$	$F(5, 39.468) = 0.602$	0.699
$Mv \times CDR \times SD$	$F(1, 43.669) = 5.705$	0.021*
$CDR \times SD \times Rostrocaudal level$	$F(5, 39.520) = 1.300$	0.283
$Mv \times CDR \times Rostrocaudal level$	$F(5, 39.722) = 2.117$	0.083
$Mv \times CDR \times SD \times Rostrocaudal level$	$F(10, 39.399) = 1.881$	0.078
Dorsal raphe nucleus, interfascicular part		
Cohort	$F(5, 39.994) = 4.533$	0.002**
<i>Mycobacterium vaccae</i> NCTC 11659 (Mv)	$F(1, 34.006) = 0.309$	0.582
Chronic disruption of rhythms (CDR)	$F(1, 37.093) = 7.705$	0.009**
Acute social defeat (SD)	$F(1, 34.167) = 0.447$	0.508
Rostrocaudal level	$F(8, 37.499) = 15.442$	0.001***
$Mv \times CDR$	$F(1, 34.038) = 0.236$	0.630
$Mv \times SD$	$F(1, 34.092) = 1.75$	0.195
$Mv \times Rostrocaudal level$	$F(8, 37.563) = 1.794$	0.109
$CDR \times SD$	$F(1, 34.250) = 0.241$	0.627
$CDR \times Rostrocaudal level$	$F(8, 37.475) = 3.038$	0.010*
$SD \times Rostrocaudal level$	$F(8, 37.430) = 2.126$	0.058
$Mv \times CDR \times SD$	$F(1, 34.551) = 1.663$	0.206
$CDR \times SD \times Rostrocaudal level$	$F(8, 37.461) = 3.26$	0.006**
$Mv \times CDR \times Rostrocaudal level$	$F(8, 37.608) = 1.237$	0.305
$Mv \times CDR \times SD \times Rostrocaudal level$	$F(16, 37.432) = 2.954$	0.003**
Median raphe nucleus		
Cohort	$F(5, 30.960) = 1.625$	0.183
<i>Mycobacterium vaccae</i> NCTC 11659 (Mv)	$F(1, 38.743) = 0.223$	0.640
Chronic disruption of rhythms (CDR)	$F(1, 41.042) = 9.004$	0.005**
Acute social defeat (SD)	$F(1, 39.010) = 0.617$	0.437
Rostrocaudal level	$F(7, 35.622) = 22.651$	0.001***
$Mv \times CDR$	$F(1, 38.833) = 0.027$	0.871
$Mv \times SD$	$F(1, 38.683) = 2.059$	0.159
$Mv \times Rostrocaudal level$	$F(7, 35.561) = 3.03$	0.013*
$CDR \times SD$	$F(1, 39.016) = 2.468$	0.124
$CDR \times Rostrocaudal level$	$F(7, 35.546) = 1.628$	0.159
$SD \times Rostrocaudal level$	$F(7, 35.460) = 1.898$	0.099
$Mv \times CDR \times SD$	$F(1, 39.320) = 0.872$	0.356
$CDR \times SD \times Rostrocaudal level$	$F(7, 35.507) = 0.751$	0.631

$Mv \times CDR \times$ Rostrocaudal level	$F(7, 35.531) = 1.372$	0.247
$Mv \times CDR \times SD \times$ Rostrocaudal level	$F(14, 35.378) = 1.888$	0.063
$*p < 0.05, **p < 0.01, *p < 0.001$		

Supplementary File S1. Data inclusion criteria table listing data used for analysis throughout 2015–2017 and calculated n values for each analysis.