

Supplementary Figure 1

MORwd induces postsynaptic depression of AMPAR-mediated neurotransmission in medial LHb.

(a) Left: sample traces, box and scatter plots of sEPSCs amplitudes recorded in ^{Lat}LHb (saline+naloxone (n_{cells/mice}=9/3; gray) versus NP-MORwd (n_{cells/mice}=9/4; orange), two-sided t-test., t₁₆=0.098, P=0.923). Right: same but sEPSCs were recorded in ^{Med}LHb (saline+naloxone (n_{cells/mice}=9/4; black) versus NP- MORwd (n_{cells/mice}=9/5; red), two-sided t-test, t₁₆=3.493, **P=0.003). (b) Left: sample traces, box and scatter plots of sEPSCs frequencies recorded in ^{Lat}LHb (saline+naloxone (n_{cells/mice}=9/4, gray) versus NP-MORwd (n_{cells/mice}=9/4, orange), two-sided t-test,, t₁₆=1.331, P=0.202). Right: same but sEPSCs were recorded in ^{Med}LHb (saline+naloxone $(n_{cells/mice}=9/5, black)$ versus NP-MORwd $(n_{cells/mice}=9/4, red)$, two-sided t-test, t₁₆=0.161, P=0.874). (c) Recording map color-coded for the value of AMPAR:NMDAR ratios recorded throughout the LHb. Lighter colors indicate smaller AMPAR:NMDAR ratios, while darker colors represent high AMPAR:NMDAR ratio. (d) Top: Sample traces and normalized EPSC versus pulse number plots recorded at 5, 10 and 20 Hz in ^{Lat}LHb (saline+naloxone (n_{cells/mice}=10/2, gray) versus NP-MORwd (n_{cells/mice}=10/3, orange), 5Hz interaction factor F_{(4,} ₃₆₎=0.227, P=0.921; 10Hz interaction factor F_(4,36)=0.251, P=0.907; 20Hz interaction factor F_(4,36)=0.573, P=0.683 two-way ANOVA Repeated Measures). Bottom: same but in ^{Med}LHb (n_{cells/mice}=10/3, saline+naloxone (black) versus NP- MORwd (red), 5Hz interaction factor F_(4,36)=0.1.183, P=0.334; 10Hz interaction factor F_(4,36)=1.171, P=0.34; 20Hz interaction factor F_(4,36)=0.88, P=0.485 two-way ANOVA Repeated Measures). (e) Spontaneous withdrawal timeline. AMPAR:NMDAR ratios from MedLHb 1 hour, 10, 20 or 30 days post-saline or MOR (saline 1 hour and 10 days pooled (nmice/cells=6/22; black) versus MORwd 1 hour (nmice/cells=5/11; open red) and MOR 10 (n_{mice/cells}=3/12), 20 (n_{mice/cells}=3/11) and 30 days withdrawal (n_{mice/cells}=3/11; red), F_(4, 62)=3.90 one-way ANOVA, **P=0.007). (f) Example of peak-scaled NSFA of ^{Med}LHb neurons in the saline- and NP-MORwd group. Pooled data for conductance (γ) and number of channels (N) open at the peak together with amplitude versus N of channels and conductance plots (Saline+naloxone, $n_{cells/mice}$ =5/4; MORwd, n_{cells/mice}=8/5; N of channels, two-sided t-test, t₁₁=5.67, ***P=0.0001, r²_(N-Channels) = 0.416; *P=0.017; Conductance, t₁₁=0.006, P=0.99, $r^2_{(Conductance)} = 0.03$, P=0.55). (g) Left: sample traces, box and scatter plots for rectification index calculated from AMPAR EPSCs recorded at -70, 0 and 40 mV in ^{Lat}LH (saline+naloxone ($n_{cells/mice}=9/7$, gray) versus NP- MORwd ($n_{cells/mice}=10/7$, orange), two-sided t-test, $t_{17}=0.210$, P=0.836). Right same but recordings in ^{Med}LHb (saline+naloxone ($n_{cells/mice}=12/8$, black) versus NP- MORwd (n_{cells/mice}=9/5, red), two-sided t-test, t₁₉=1.292, P=0.212). (h) Sample traces, box and scatter plots of AMPAR:NMDAR ratios recorded in ^{/led}LHb via 405 nm laser-assisted uncaging of MNI-glutamate, 500μM (saline+naloxone (n_{cells/mice}=8/2, black) versus NP-MORwd (n_{cells/mice}=10/3, red), two-sided t-test, t₁₆=3.521, **P=0.003). Bottom right: Absolute AMPAR versus absolute NMDAR uncaging-evoked current plots from saline+naloxone (open black circles) or NP-MORwd mice (open red circles). The mean with S.E.M. AMPA and NMDA currents are shown with black and red filled circles for saline versus MORwd respectively (saline+naloxone versus NP-MORwd: AMPA, two-sided t-test, t_{16} =3.536, **P=0.003; NMDA, two-sided t-test, t_{16} =0.195, P=0.848). Data are presented as box plots 10-90 percentiles and scatter.





TNF α signaling is necessary and sufficient for MORwd-induced plasticity

(a) Spontaneous MORwd protocol, sample traces, box and scatter plots for AMPAR:NMDAR ratios recorded in ^{Med}LHb slices incubated with (+) or without (-) exogenous TNF α from spontaneous MORwd mice (10 days). (MORwd -TNF α (red) versus +TNF α (pink), n_{cells/mice}=9/4, two-sided t-test, t₁₆=0.986, P=0.339). (b) ^{Med}LHb AMPAR:NMDAR ratios from saline or MPLA-injected MOR-treated mice (MOR/saline (n_{mice/cells}=3/10; shaded blue) versus MOR/MPLA (n_{mice/cells}=4/11; dark blue), two-sided t-test, t₁₉=3.070, **P=0.006). (c) Sample traces, time versus amplitude plot and bar graphs showing the effect of MPLA (1 g/ml) on AMPAR-EPSCs (MOR (n_{mice/cells}=4/11; 63.93 ± 7.06%; open red), NP-MORwd (n_{mice/cells}=3/10 cells; 91.63 ± 8.86%; filled red), MOR versus NP-MORwd, two-sided t-test, t₁₉=2.419, *P=0.026). Data of this panel are represented as mean and sem. (d) Sample traces, time versus amplitude plot and bar graphs showing the effect of MPLA (1µg/ml) on evoked AMPAR-EPSCs (baseline (1) vs 30 min post-MPLA (2)) recorded in ^{Lat}LHb (open orange) or ^{Med}LHb (open red) in slices obtained from morphine-treated animals (n_{cells/mice}=8/4, morphine ^{Lat}LHb 103.98 ± 10.13; n_{cells/mice}=11/5, morphine ^{Med}LHb 63.31 ± 7.06%; morphine ^{Lat}LHb versus morphine ^{Med}LHb, two-sided t-test, t₁₇=3.406, **P=0.003). Note that the data set for ^{Med}LHb is the same as in c and is used for comparison. Data are presented as mean and SEM. (e) Sample traces, time versus amplitude plot and bar graph showing the effect of MPLA (1µg/ml) on evoked AMPAR-EPSCs (baseline (1) vs 30 min post-MPLA (2)) in the presence of TNF α dominant negative peptide (XENP1595, 6mg/1ml) recorded in ^{Med}LHb in slices obtained from morphine-treated animals (n_{cells/mice}=10/3, black) versus N=40. (f) NP-MORwd protocol with dominant-negative TNF α (XENP1595, 30mg/kg) pretreatment, sample traces, box and scatter plots for AMPAR:NDAR ratios recorded in ^{Med}LHb (saline+naloxone (n_{cells/mice}=10/3, black) versus NP-MORwd (n_{nice/cells}=2/2, 98.84



Behavioral assessment of MORwd

(a) Box and scatter plot showing the percent time spent in the compartment containing the social stimulus for C57BI6 mice (N=22 mice/group, saline+naloxone (black) versus NP-MORwd (red), two-sided t-test, t_{42} =2.401, *P=0.021). (b) Box and scatter plot showing the percent time spent in the compartment containing the object stimulus for C57Bl6 mice (N=22 mice/group, saline+naloxone (black) versus NP-MORwd (red), two-sided t-test, t_{42} =2.465, *P=0.02). (c) Box and scatter plot showing the percent time spent in the central compartment for C57Bl6 mice (N=22 mice/group, saline+naloxone (black) versus NP-MORwd (red), two-sided t-test, t_{42} =1.186, P=0.242). (d) Box and scatter plot showing locomotor activity during social preference test for C57Bl6 mice (N=22 mice/group, saline+naloxone (black) versus NP-MORwd (red), two-sided t-test, t_{42} =2.621, *P=0.012). (e) Box and scatter plot showing the percent time spent in the compartment containing the social stimulus for TNF-R1fl/fl mice (AAV-Control: 58.02 \pm 2.96% saline+naloxone (N_{mice}=20, black) versus 40.45 ± 5.08% NP-MORwd (N_{mice}=23, red); AAV-Cre: 54.41 ± 3.15% saline (N_{mice}=13, open gray) versus 64.52 \pm 5.94 % NP-MORwd (N_{mice}=13, open pink), interaction factor F_(1.65)=8.591 two-way ANOVA, **P=0.005). (f) Box and scatter plot showing the percent time spent in the compartment containing the object stimulus for TNF-R1fl/fl mice (N of mice same as panel e. AAV-Control: 26.42 ± 2.47 % saline+naloxone (black) versus 30.9 ± 4.79% NP-MORwd (red); AAV-Cre: 31.19 ± 2.07% saline+naloxone (open gray) versus 20.7 ± 4.13 % NP-MORwd (open pink), interaction factor F_(1.65)=4.136 two-way ANOVA, *P=0.046). (g) Box and scatter plot showing the percent time spent in the central compartment for TNF-R1fl/fl mice (N of mice same as panel e. AAV-Control: 15.38 ± 3.08% saline+naloxone (black) versus 27.82 ± 4.98% NP-MORwd (red); AAV-Cre: 14.23 ± 2.9 % saline+naloxone (open gray) versus 14.62 ± 3.08% NP-MORwd (open pink), interaction factor F_(1,65)=1.748 two-way ANOVA P=0.191). (h) Box and scatter plot showing number of exploration bouts of TNF-R1fl/fl with the juvenile (N of mice same as panel e. AAV-Control: 56 ± 3.51 saline+naloxone (black) versus 39.30 ± 4.8 NP-MORwd (red); AAV-Cre: 50.38 ± 3.78 saline+naloxone (open gray) versus 57.15 ± 7.01 NP-MORwd (open pink), interaction factor F_(1.65)=5.519 two-way ANOVA, *P=0.022). (i) Box and scatter plot showing number of exploration bouts of TNF-R1fl/fl with the object (N of mice is the same as panel e. AAV-Control: 26.9 ± 1.88 saline+naloxone (black) versus 23.35 ± 2.13 NP-MORwd (red); AAV-Cre: 28.08 ± 2.01 saline+naloxone (open gray) versus 21.69 ± 3.16 NP-MORwd (open pink), interaction factor F(1.65)=0.361, P=0.55). (j) Box and scatter plot showing locomotor activity during social preference test for TNF-R1fl/fl mice (N of mice is the same as panel e. AAV-Control: 28.13 ± 1.35m saline+naloxone (black) versus 33.8 ± 5.59m NP-MORwd (red); AAV-Cre: 23.78 ± 1.47m saline+naloxone (open gray) versus 40.17 ± 7.11m NP-MORwd (open pink) interaction factor $F_{(1,65)}$ =1.224 two-way ANOVA, P=0.273). Data are presented as box plots 10-90 percentiles with median and scatter.