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Supplemental Information

Hypothalamic Control of Conspecific Self-Defense

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Figure S1. Additional histological images from the c-Fos experiment. Related to Figure 1. Leftmost column shows the Esr1 expression along the anterior-posterior axis of the VMHvI. The remaining columns, from left to right, show Images of c-Fos expression in the VMHvI of a control animal with no intruder exposure and four test animals that interacted with a non-aggressive Balb/C male intruder, a non-aggressive C57BL/6 male intruder, an aggressive C57BL/6 male intruder and an aggressive SW male intruder. Scale bar: 100 µm.



Figure S2. Behavioral characterization of the defending mice. Related to Figure 2.

- (A) Images showing the characteristic behaviors during aggressor-defender interaction.
- (B) The distribution of behaviors of the defending mouse when being attacked.
- (C) The behaviors that were employed by the defending mice to terminate an episode of attack.
- (D) The behaviors shown by the defending mice when they were approached by an aggressor.
- (E) The probability of upright posture (top) and dashing (bottom) aligned to the aggressor approach offset. Black traces are constructed using shuttled time points. Shades represent ± SEM.



Figure S3. Control data for the fiber photometry recording. Related to Figure 2.

- (A) Viral construct and experimental schematics
- (B) Expression of GFP (green) is overlapped with Esr1 staining (red). Blue: NissI. Scale bar: 50 µm
- (C) Average GFP signals during various behaviors are not significantly different. One way ANOVA. p > 0.05.
- (D) Heat maps showing the Z scored GFP signals in the VMHV of individual animals aligned to the onsets of various behaviors during aggressor interaction.
- (E) PETHs showing the average Z scored GFP signals aligned to various behavioral onset across all animals. Shade: ± SEM.



Figure S4: *In vivo* population recording of the VMHvI Esr1+ cells during encounters with an aggressor or a predator. Related to Figure 2.

- (A) Schematics of viral injection and implantation location and a representative image showing the optic fiber track (yellow arrows) above the VMHV and the virally expressed GCaMP6f. Scale bars: 500 µm.
- (B) The distribution of behaviors of 3 repeatedly defeated Esr1-2A-Cre C57 mice when they were being attacked.
- (C) A representative GCaMP6f trace during encounters with a C57 aggressor. Color shades indicate behavioral events.
- (D) PSTHs of Z scored GCaMP6f aligned to various behaviors of the mouse shown in C. Shades represents ± SEM.
- (E) The GCaMP6f trace from the same animal as in C and D during close interactions with a hand-held rat. Color shades indicate behavioral events.
- (F) PSTHs of Z scored GCaMP6f signal aligned to behavioral onsets of the mouse in the presence of a rat. Shades represent ± SEM.
- (G) The average Z scored GCaMP6 responses during various behaviors observed in the presence of a conspecific aggressor or a rat (n = 3). One way ANOVA, Post-hoc pairwise comparison with Tukey-Kramer correction. **p< 0.01, ***p<0.001. One sample *t-test*. #p< 0.05. Error bars: ± SEM.



Figure S5. Control data for optogenetic activation experiment. Related to Figure 4.

- (A) Viral construct and experimental deign. Image showing the track of a cannula. Scale bar: 100 μm.
- (B) Behavior raster during light-on and light-off period from a representative animal. Scale bar: 10 s.
- (C) Accumulated probability of attack from 60 s before VMHvI stimulation to the stimulation offset.
- (D) Comparison of percentage of upright posture (Left) and latency to upright postures (right) between light-on and light-off periods. Paired t-test.
- (E) Comparison of the percentage of trials that animals showed dashing behaviors (left) and latency to dash between light-on and light-off periods. Paired t-test.
- Error bars: ± SEM.



Figure S6. Projection pattern of the VMHvI Esr1⁺ cells. Related to Figure 5.

- (A) Viral construct and experimental design.
- (B) Example images showing GFP expressing cells in the VMHvI. Right shows the enlarged view of the boxed area. Scale bars: 500 μm (left) and 100 μm (right).
- (C) The axons from the VMHvI Esr1+ cells in the LS, MPOA, AHN, PVN, aPAG and cPAG. Dashed red lines indicate the regions that were targeted in the retrograde experiment. Scale bars: 100 µm.

LS: lateral septum; MPOA: medial preoptic area; AHN: anterior hypothalamic nucleus; PVN: periventricular nucleus; rPAG and cPAG: rostral and caudal periaqueductal gray.