Immune activation state modulates infant engram expression across development

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Abstract

Infantile amnesia is possibly the most ubiquitous form of memory loss in mammals. We investigated how memories are stored in the brain throughout development by integrating engram labeling technology with mouse models of infantile amnesia. Here, we discovered a phenomenon in which male offspring in maternal immune activation models of autism spectrum disorder do not experience infantile amnesia. Maternal immune activation altered engram ensemble size and dendritic spine plasticity. We rescued the same apparently forgotten infantile memories in neurotypical mice by optogenetically reactivating dentate gyrus engram cells labeled during complex experiences in infancy. Further, we were able to permanently reinstate lost infantile memories by artificially updating the memory engram, demonstrating that infantile amnesia is a reversible process. Our findings suggest that not only is infantile amnesia due to a reversible retrieval deficit in engram expression, but that immune activation during development modulates innate, and reversible, forgetting switches that determine whether infantile amnesia will occur.

Introduction

Infantile amnesia is the rapid forgetting of memories formed during early development; and is a largely neglected form of memory loss that seemingly affects the entirety of the human population (*1*, *2*). Not uniquely a human phenomenon, this form of amnesia has been documented in rodents, which show forgetting of contextual and fear memories formed during the infant period (*3–6*). Little is known about the basic neurobiology of infantile amnesia or its effect on the engrams cell ensembles that encode specific memories. Thanks

to the integration of activity-dependent ensemble labeling with optogenetics it is now possible to investigate if a memory engram is still present and/or functional in the brain even in the case of amnesia (7). Using this methodology, it has been shown that memory recall can be induced after amnesia by optogenetic activation of engram cells in the hippocampus and other brain regions, demonstrating that these memories are not only still present in the brain, but are recoverable (8–11). This framework provides an opportunity to investigate how development affects the storage and retrieval of early childhood memories. Environmental conditions strongly influence both learning and forgetting rates, but less is known about how forgetting occurs under altered developmental trajectories (12). During embryonic and postnatal development, there are periods in which the developing brain has heightened sensitivity to environmental influences (6, 13–18). Infantile amnesia has been shown to be preventable through postnatal pharmacological interventions using GABA agonists (19–21) and corticosteroids (6) or ectopic administration of neurotrophins (22), but the relevance of these plasticity switches in connecting environmental experiences with memory function across natural development is unknown. Events during embryonic development such as the activation of the immune system, and subsequent cytokine release, is known to induce altered developmental trajectories associated with autism spectrum disorder (ASD) and schizophrenia, but the impact on engram function has not been explored (13, 23–26). Here, we sought to identify naturally occurring variation in infantile amnesia caused by the developmental experience of the animal, and then investigate the effects on engram cell function.

Results

Maternal immune activation during the embryonic period alleviates infantile amnesia in male offspring

Using a contextual fear conditioning (CFC) paradigm (Fig. 1A, B), we trained and tested infant (P17) (27–30) and adult (P63) mice for fear memory recall 1 or 8 days post-training (Fig. 1C). At both time points, the experimental adult shock group showed significantly more freezing than the no shock control group (Fig. 1C). Infant mice tested for memory recall 1 day following training showed significantly higher levels of freezing compared to the control group (Fig. 1C); while the group tested 8 days after training showed robust infantile amnesia, with similar levels of freezing to the no shock controls (Fig. 1C). Consistent with the literature, infant mice demonstrate forgetting as early as one week after training, while adult mice show continuous memory retention (4, 31).

We tested whether altering external environmental influences, such as environmental enrichment during infancy and the time of weaning from maternal care, affected the occurrence of infantile amnesia (SFig.1). We next looked at the effect of altered developmental trajectories resulting from embryonic challenges. During pregnancy, exposure to pathogens or inflammation during critical periods of brain development (maternal immune activation (MIA)) has been shown to alter postnatal brain growth and cognitive development through the release of pro-inflammatory cytokines such as the interleukin 6 (IL-6) and interleukin 17 (IL-17) families (32). Delivery of the viral-mimetic compound, Polynosinic-polycytidylic acid (Poly I:C), at E12.5 is a well-established method for the experimental induction of MIA in rodents (33–35). More recently the maternally derived cytokine, IL-17a, has been shown to mediate Poly I:C's effect on maternal immune activation and is sufficient to induce an altered brain state in offspring (13, 17, 23, 24, 36). Here we employed the administration of either Poly I:C, or recombinant IL-17a, as two

distinct methods to stimulate MIA in pregnant dams at E12.5 in C57BL/6J mice (Fig. 1D). To first validate our method, we measured changes in IL-17a blood serum after injection of Poly I:C or PBS in female mice. Injection of Poly I:C resulted in an increase in IL-17a in the blood serum compared to PBS controls. In line with published studies, male, but not female, offspring from pregnant dams treated with Poly I:C or IL-17a demonstrated repetitive behavior and deficits in social behavior, behavioral phenotypes indicative of ASD (Fig. 1E, F, SFig. 2A-F) (13, 23, 24). To test if MIA causes any effect on memory retention, we next trained infant offspring from MIA dams using contextual fear conditioning at P17 (Fig. 1G, SFig. 2G-I). When tested 1 day after training, all cohorts showed similar levels of freezing, irrespective of sex (SFig. 2G, H). However, male, but not female offspring from pregnant dams treated with Poly I:C or IL-17a showed retention of infant memory, like that of adult mice, when tested 8 days after training (Fig. 1G, SFig. 2I). This effect was also true when MIA male offspring were tested for recall of a CFC memory 15 days after training (SFig. 2J). Regardless of whether male MIA offspring were tested for recall 8 or 15 days after training, the fear memory was retained. Further, the effect was sex-specific (interaction $p = 0.0119$) (SFig. 2K). This represents an unknown phenomenon in which male, but not female, offspring from Poly I:C injected dams do not demonstrate infantile amnesia for a contextual fear memory. This phenomenon was also witnessed in offspring from dams injected with IL-17a, indicating the mediating role of IL-17a signaling.

MIA male offspring show an increase in engram labeling and spine density in the DG

We characterized and employed a Cre-based engram labeling strategy for whole-brain tagging of engrams cells in infant mice, by crossing a FosTRAP transgenic line that expresses iCre from an inducible *c-fos* promoter with an Ai32 transgenic mice express channelrhodopsin-2/EYFP (ChR2-EYFP) (Fig. 2A-C) (37, 38). Intraperitoneal (IP) injection of the tamoxifen metabolite, 4-hydroxytamoxifen (4-OHT), allows iCre to recombine the ChR2-EYFP transgene (39, 40). Using this engram labeling method, ChR2-EYFP expression can be detected in engram cells 3 days after 4-OHT injection. Firstly, to ensure infant mice still show memory retention at P20, we trained infant mice using CFC at P17 and tested them for memory recall 3 days later (P20) (SFig. 3A). Infant mice trained at P17 show similar memory retention for a fear memory when tested 1 or 3 days after training (SFig. 3B, C). We evaluated the efficiency of the Ai32-FosTRAP engram labeling system in P17 (infant) and P42 mice by quantifying the number of ChR2-EYFP$^+$ cells in brain regions associated with context and fear memory, including the dentate gyrus (DG), amygdala (AMG), retrosplenial cortex (RSC) and the periaqueductal gray (PAG) at P63 for a home cage, contextual or contextual fear memory (Fig. 2D-F, SFig. 4). Exposure to context A resulted in an activity-dependent increase in engram labeling (EYFP$^+$ cells) in the DG at both P17 (Fig. 2E) and P42 (SFig. 4B). Similarly, there was an activity-dependent increase in EYFP$^+$ labeling in the AMG after contextual fear conditioning at both P17 (Fig. 2F) and P42 (SFig. 4C). These results demonstrate that the Ai32-FosTRAP line is an efficient engram labeling system for activity-dependent engram labeling in both infant and adult mice.

We next investigated whether engram cells formed at the time of infant memory encoding are reactivated by exposure to training cues (Fig. 2H). Cellular activation of engram cells was examined by quantifying the number of EYFP$^+$ c-Fos$^+$ cells before (P20) and after (P63) the infantile amnesia period across multiple brain regions (Fig. 2G, SFig. 5, 6). Natural recall cues result in above chance EYFP$^+$ and c-Fos$^+$ overlap in the DG and AMG at P20 (Fig. 2I, J, SFig. 5D, G), an effect that was not seen at P63 (Fig. 2I, J, SFig. 6D, G). We next compared engram reactivation after natural recall for a contextual fear memory in male offspring from MIA dams (Fig. 2K). Interestingly, male, but not female, offspring from

MIA dams have an increased number of EYFP$^+$ engram cells in the DG compared to controls (Fig. 2L, SFig. 7A, B), indicating an altered engram ensemble also reported for a genetic model of ASD (*41*). Further, there was a positive correlation between freezing behavior (%) and the total number of engram cells (EYFP$^+$) in the DG of male MIA offspring (Fig. 2M); r(30) = 0.36, p = 0.043. A larger engram size was related to an increase in freezing behavior during recall of a contextual fear memory in P25 male MIA offspring. Although no difference was seen between groups in the total level of cell activation (c-Fos$^+$) (Fig. 2N), MIA male offspring demonstrated a decrease specifically in non-engram cell activation in the DG during recall (Fig. 2O). No difference was seen in the relative level of engram reactivation between groups as both demonstrated an above chance level of EYFP$^+$ and c-Fos$^+$ overlap in the DG (Fig. 2P, SFig. 7C). While prior research has examined alterations in dendritic spines within the cortex of infant MIA offspring (*42*), we sought to investigate the specific modifications in dendritic spines related to engram cells within the DG of MIA male offspring after recall (Fig. 2Q). Dendritic spine analysis indicated that engram cells in MIA offspring had significantly increased dendritic spine density (Fig. 2R) as well as larger spine head diameter and volume relative to control animals at P25 (Fig. 2S, T). Thus, a combination of larger engram size, lower non-engram cell activation, and increased dendritic plasticity may contribute to enhanced functional engram expression of infant engrams in MIA offspring.

Optogenetic stimulation of an infant engram allows for acute and permanent reinstatement of the forgotten memory

To investigate the functionality of memories after naturally occurring infantile amnesia, we tested the behavioral effect of optogenetically stimulating infant-labeled DG engram cells in adult mice. Engram cells for context A were labeled in Ai32-FosTRAP mice at P17 using CFC (Fig. 3A-C). Although both cohorts were placed in context A, only the experimental group received foot shocks (S). At P63, both cohorts were placed in context C for a 12 min test session where they received four 3 min epochs of blue light on or off. During the light-off epochs, both groups showed background levels of freezing (Fig. 3D). The experimental group showed significantly higher levels of freezing during the light-on epochs (Fig. 3D, E). Despite the same number of EYFP$^+$ cells being labeled (Fig. 3F), this light-induced freezing was not observed for animals that underwent the same behavioral paradigm but did not receive any footshocks (NS) at P17 (Fig. 3D, E). While stimulation of a neutral memory, labeled in infancy, does not result in freezing behavior (SFig. 8A-C). We extended this paradigm by optogenetically stimulating infant-labeled CA1 engram cells in adult mice (SFig. 8D) (*8*, *43*, *44*). A previous study demonstrated that the application of the standard 20 Hz protocol for activating CA1 engram cells did not effectively induce memory recall (*43*). However, implementing a 4 Hz protocol to stimulate CA1 engram cells elicited comparable improvements in recovering memory deficits from anisomycin-induced amnesia (*8*). Optogenetic stimulation at 4 Hz (SFig. 8H, I), but not 20 Hz (SFig. 8F), of infant-labeled engram cells in the CA1 of adult mice resulted in light-induced freezing. Based on these data, 'lost' infantile memories can be acutely recalled by optogenetic stimulation in either the DG or CA1.

Although optogenetic stimulation of a forgotten infant memory in adulthood resulted in a specific behavioral response, it is unclear how the information survives. We investigated the connectivity between engram cells in downstream regions, after infantile amnesia, by histologically assessing engram cell reactivation following optogenetic stimulation of upstream engram cells (Fig. 3F-K) (*8*, *45*). The resulting c-Fos$^+$ counts were equivalent in the hippocampus after light activation of a CFC (shock, S) or a contextual (no shock, NS) memory (Fig. 3G). In contrast, c-Fos$^+$ cell counts were significantly higher in the AMG

after light activation of an infant CFC memory (Fig. 3J), demonstrating an increase in both amygdala activity and the behavioral expression of freezing. Light activation of DG engram cells resulted in an above chance c-Fos$^+$ and EYFP$^+$ overlap in the DG (Fig. 3H, SFig. 9B) and crucially in other downstream brain regions including the AMG (Fig. 3K, SFig. 9C), RSC (SFig. 9F, G) and PAG (SFig. 9J, K). These results were consistent for light activation of both a CFC (shock, S) and a contextual (no shock, NS) memory. These results demonstrate that the connectivity pattern of the engram survives infantile amnesia and persists into adulthood, and optogenetic stimulation of a lost infantile memory is sufficient to reactivate the functional connectivity patterns. As a positive control, we optogenetically stimulated an engram encoded after the infantile amnesia period (P29) (SFig. 10A). By selecting this specific stage of development for labeling, we were able to target a memory engram that was not only formed post-infantile amnesia, and therefore retained over a long delay (SFig. 3D, E), but also underwent subsequent developmental processes. Optogenetic stimulation of an engram encoded after (P29) the infantile amnesia period also increased engram reactivation to above chance levels in the DG, AMG, RSC, and PAG (SFig. 10B-Q).

Since optogenetic stimulation results in memory recall under artificial conditions, we sought to permanently reinstate an infant memory by adopting an optogenetic induction procedure to induce plasticity in engram cells (SFig. 11A, B) (*8, 43*). Infant mice can form purely contextual memories that can be updated with shock information (SFig. 11C, D), provided updating occurs before the onset of infantile amnesia (SFig. 11E, F). At P63 and in a novel context, we optogenetically stimulated DG engram cells for context A, originally encoded at P17, while simultaneously delivering footshocks to create a false association of an infant context engram with adult shock experience (Fig. 3L). Animals subsequently froze significantly more when exposed to context A even though they were shocked in context C (Fig. 3M). This increased freezing was not due to generalization, since the control group did not freeze in context B. There was also an increased level of overlap of c-Fos$^+$ and EYFP$^+$ cells in the DG (Fig. 3N, P, SFig. 12D) and RSC (Fig. 3O, Q) after exposure to context A, demonstrating that permanent engram reinstatement was also reflected at a cellular level (SFig. 12B-O). This histological result demonstrates that engram cells active at the time of encoding were reactivated during adult exposure to context A. Together, these findings show that the plasticity that accompanies the targeted updating of an infant engram restores the natural accessibility of that engram to the appropriate perceptual cues and that context specificity is maintained into adulthood.

MIA offspring do not show infantile amnesia for spatial or object memories and these memories can be optogenetically activated in control offspring

Although fear conditioning is a robust assay, we wanted to further expand our knowledge by testing the relevance of infantile amnesia to other types of memory. To do this, we next looked at this form of forgetting in a novel object recognition behavioral paradigm (Fig. 4A). Although studies have looked at novel object location in infant rats, there has been no clear demonstration of infantile amnesia for novel object recognition in mice at this stage of development (*46, 47*). Infant mice demonstrated rapid forgetting for novel object recognition tasks, following the developmental trend like that of other forms of infant encoded memories. Both adult and infant cohorts spent more time exploring a novel object relative to a familiar object when tested 1 day after acquisition, demonstrating that a long-term memory was formed (Fig. 4B, SFig. 13A-C). When tested again 8 days later, only the adult cohort spent significantly more time exploring the novel object (Fig. 4B, SFig. 13C). We then investigated whether male offspring from MIA dams show infantile amnesia for object memory (Fig. 4C). When tested for object memory retention, male offspring from

MIA dams spent significantly more time with the novel object 8 days after acquisition (Fig. 4D, SFig. 13E). Like contextual fear memory, male offspring from Poly I:C or IL-17a injected dams do not show infantile amnesia for an object memory.

We then investigated whether it was possible to rescue a 'lost' infant object memory by optogenetic stimulation of infant-labeled engram cells (Fig. 4E, SFig. 13F). On the test day, animals that received light stimulation 3 min before the test demonstrated a preference for the novel object (Fig. 4F). While animals that did not receive light stimulation before the test spent the same amount of time exploring both the novel and familiar object. These object memory engrams survive infantile amnesia, can be retrieved artificially, and their forgetting can be prevented by MIA.

Spatial and object tasks in rodents are useful for the development of parallel, translational object/context and maze learning assays in human behavior experiments (*48*). To expand our analysis of infantile amnesia to an active, maze-based task we probed memory for a spatial location in infant and adult C57BL/6J mice using a cued version of the Barnes maze (Fig. 4G, SFig. 14 G). All cohorts sufficiently learned the escape hole's location during training over five sessions, evidenced by a gradual decrease in the latency, and distance traveled, to reach the escape hole (SFig. 14A, B, H, I). Unlike adults, infant mice demonstrated memory retention for the location of the escape hole when tested 1, but not 8, days later (Fig. 4H, I, SFig. 14 C-F, J-O). To the best of our knowledge, this is the first demonstration of infantile amnesia using the Barnes maze paradigm. We assessed memory content profiling changes in navigational strategies taken by the animal during the task (Fig. 4J). Navigational strategies can be categorized as random, procedural, or contextual (*49–51*). Infant mice predominantly used contextual search strategies when tested for spatial memory on the Barnes maze 1 day after training (Fig. 4K). However, during the 8-day probe trial infant mice show a switch in navigational strategy, moving to procedural and random strategies (Fig. 4K). This adaptation in navigational search strategies reflects the forgetting of the location of the escape platform. In contrast, adult mice show similar search patterns both 1 and 8 days after training (Fig. 4L). Like other forms of memory, we also found that male offspring from MIA models do not show infantile amnesia for a spatial navigation task (Fig. 4M-S).

By labeling the engram ensembles for the location of the escape platform in the Barnes Maze in infants, we next investigated whether DG optogenetic stimulation of the engram in adulthood results in the location of the escape hole (Fig. 4T). The experimental light group received 3 min light stimulation before being placed on the Barnes maze for a probe test. The experimental light group spent significantly more time in the zone where the escape hole should be when compared to zone 2 and 3 (Fig. 4W). Although there was no significant difference in the total time spent between zone 4 and zone 1, heat map analysis of the mean time spent in each location showed a concentration around the location where the escape hole previously was located during training (Fig. 4X). No difference was seen in the total time spent between zones for the no-light control group (Fig. 4U, V). In a separate experiment (SFig. 14P), a different cohort of mice were trained where the location of the escape hole during memory encoding differed from experiment Fig. 4T. The location of the escape hole during encoding did not affect successful recall of the location of the escape hole after light stimulation (SFig. 14Q-T). Optogenetic stimulation of EYFP-expressing neurons in the DG (neurons labeled during infant encoding of the Barnes maze memory) allows for recall of the location of the escape hole in the Barnes maze.

Investigating the effect of IL-17a on infantile amnesia

Having established that immune activation, and direct administration of the downstream cytokine Il-17a, are sufficient to induce altered infantile amnesia, we next sought to test whether Il-17a is necessary for this effect to occur. We injected pregnant *Il17a* knockout (KO) dams with Poly I:C at E12.5 (Fig. 5A). Importantly adult male *Il17a* KO mice showed normal memory retention for a CFC memory when trained at P63 (SFig. 15). This suggests that the *Il17a* genotype does not affect long-term memory retention in adult *Il17a* KO mice. Adult male and female offspring from MIA *Il17a* KO dams did not demonstrate repetitive behavior (SFig. 16E) or deficits in social behavior (SFig. 16A-D). Crucially, infant offspring showed normal infantile amnesia for a CFC memory at P25 (Fig. 5B, C). Preventing maternal IL-17a signaling, using a genetic model, occluded the MIA effect on infantile amnesia in the offspring. Therefore, the behavioral effect seen following MIA is specifically dependent on IL-17a signaling. We investigated whether altering the timing of immune activation, from maternal to postnatal, would have the same effect on memory retention (Fig. 5D) (*52*). Postnatal immune activation in C57BL/6J male mice did not affect infantile amnesia, indicating that there is a critical window during prenatal development where IL-17a and immune activation can affect a mammal's propensity for infantile amnesia (Fig. 5E-F). Induction of an inflammatory response in MIA offspring has been shown to temporarily rescue deficits in sociability in ASD (*23*). Therefore, we investigated whether we could reverse the effect of MIA on infantile amnesia in the offspring through the administration of IL-17a at the time of memory encoding (P17) or memory recall (P25) (Fig. 5G). MIA male offspring were injected with IL-17a three hours before CFC (Fig. 5H) or the recall test (Fig. 5I). Regardless of whether injected with IL-17a or PBS, MIA male infant offspring showed normal memory retention for a CFC memory 8 days after training (Fig. 5H, I), indicating that the effect of MIA on infantile amnesia cannot be reversed through acute IL-17a administration.

Discussion

We provide an integrative set of analyses that triangulates infantile amnesia, engram expression, and immune activation state *in utero*. Together our findings indicate that infantile amnesia, for a wide range of memory types, and engram cells as the substrate of memory, can be modulated by the immunological experiences of the subject during embryonic development. The effects of MIA preserve the natural retrievability of infant engrams across postnatal development. Furthermore, we find that MIA directly modulates engram plasticity at a structural level; altering engram ensemble size and engram dendritic spine density. In addition, engram reactivation shows greater potential specificity under MIA as non-engram cell activity is significantly reduced during recall. This in turn may be reflective of a reduction in interference of competing ensembles (*53, 54*). Furthermore, our data shows that many types of infant memories are naturally suppressed during development, but that their engrams persist and can be acutely activated in adulthood by direct optogenetic stimulation (*31*). Finally, we show that the downstream cytokine, IL-17a, is both sufficient and necessary for the effect of MIA on infantile amnesia.

These findings demonstrate that optogenetic reactivation of an infant memory is not specific to fear conditioning, as memories for more complex navigational and recognition tasks can also be rescued. Furthermore, artificially updating an engram restored natural access to the target memory at both a behavioral and cellular level, reversing infantile amnesia. The specific connectivity patterns between engram cells, formed during infancy across distributed brain regions, also remain intact into adulthood. This kind of distributed engram

connectivity pattern has been shown to survive other kinds of amnesia and may account for specific stored information to be retained in the brain (the engram itself) (*8*, *9*, *45*, *55–58*).

Based on our analysis and others, infantile amnesia seems to be an innately driven form of natural forgetting and is not modulated by specific environmental experiences during postnatal development. Immune activation *in utero* results in the development of an altered brain state where the capacity to form engrams is altered, and infantile amnesia does not occur Postnatal immune activation does not seem to affect infantile amnesia, making it plausible that an innate predetermined forgetting switch is fixed from birth. However, interactions between MIA and genetic risk factors for ASD might allow for postnatal inflammation to alter forgetting rates (*52*, *59*).

Our findings suggest that the brain state that permits infantile amnesia is absent in immunological models of ASD, and these findings are reminiscent of transgenic fly experiments and retrospective human studies that have provided evidence for improved long-term memory retention in cases of ASD (*60*, *61*). It is conceivable that the MIA offspring's brain state may mirror that of precocial mammals, who do not seem to show infantile amnesia (*4*, *62*). Infantile amnesia may represent a genetically canalized form of natural forgetting that can be modulated by developmental plasticity during critical periods. Future studies should determine the nature of the switching mechanism that determines when infantile amnesia occurs, its interaction with the immune system, and its reversible effect on engram cell function and engram ensemble expression. In the future, maternal immune activation models may offer new opportunities for translational studies of memory and forgetting across the lifespan in rodents and humans (*63*).

Materials and Methods

Experimental Design
The objective of this study was to investigate how memories are stored in the brain throughout development, specifically focusing on infantile amnesia and its modulation by immune activation. Mice were randomly assigned to experimental groups with littermate controls or otherwise relevant controls. During behavioral testing, investigators were blinded to the treatment conditions in the experimental groups. Histological analysis was conducted individually for each mouse using nested analysis, and the same approach was applied to maternal immune activation experiments where analysis was performed per litter. Throughout all experiments, data collection and analysis were conducted blindly.

Subjects
For wild-type experiments, the C57BL/6J substrain C57BL/6JOlaHsd was used. The Ai32-FosTRAP mice were generated by crossing Fos-iCre (*37*, *38*) mice with Ai32(RCL-ChR2(H134R)/EYFP) (*39*, *40*) and selecting offspring carrying both the CreETR and EYFP transgenes. The Fostrap transgenic line Fos-iCre expresses iCre from an inducible *c-fos* promoter. Intraperitoneal (IP) injection of the tamoxifen metabolite, 4-hydroxytamoxifen (4-OHT), allows iCre to translocate into the nucleus. Ai32 transgenic mice express channelrhodopsin-2/EYFP (ChR2-EYFP) following exposure to Cre recombinase. Crossing TRAP mice with Ai32 mice makes it possible to produce a stable transgenic line containing both Fos-iCre and ChR2-EYFP. All mice were socially housed in numbers 2-5 littermates, on a 12 h light/dark cycle with access to food and water ad libitum. The day of birth was designated postnatal day 0 (P0). Infant (P17) mice were housed with parents until the time

of weaning when they were group-housed by the same sex. Litters size was controlled, and no litter consisted of more than 6 pups. For infant experiments, each litter was counterbalanced across groups to limit litter effects. Each experimental group consisted of 3-5 liters on average. For maternal immune activation experiments refer to Table S1 for the exact number of litters per experimental group. All mice were 7 weeks old before undergoing surgery. Post-operation mice were allowed to recover for a minimum of 2 weeks in their homecage before experimentation. All procedures relating to mouse care and treatment conformed with the health produces regulatory authority (HPRA) Ireland guidelines.

ChR2-EYFP expression in Ai32-FosTRAP mice
To label memory engram cells with this system, mice were injected intraperitoneally with 4-OHT (50mg/kg) 2 h post the learning event. The Fos-iCre line expresses the inducible *c-fos* promoter. Injection of 4-OHT activates iCre recombinase. Activated iCre recombinase translocates to the nucleus and acts on two lox sites removing the stop codon that otherwise prevents expression of the ChR2/EYFP transgene. ChR2/EYFP transgene expression is driven by the pCAG promoter in iCre-expressing tissue 72 h after 4-OHT injection. For activity-dependent expression of ChR2-EYFP in Ai32-FosTRAP mice, Kainic acid (5mg/kg) was injected intraperitoneally followed by 4-OHT injection 2 h post kainic acid injection.

Stereotactic optical fiber implant
Mice were anesthetized using 500 mg kg^{-1} avertin. Each animal underwent bilateral craniotomies using a 0.5 mm diameter drill bit at -2.0 mm anteroposterior (AP), $\pm$ 1.3 mm mediolateral (ML) for DG injections. A bilateral patch cord optical fiber implant (200 μm core diameter) was lowered above the injection site (-1.9 mm DV). A layer of adhesive cement was applied to secure the optical fiber implant to the skull. Dental cement was applied to secure a protective cap to the implant and close the surgical site. Each animal received meloxicam analgesic (0.075 mL/5g) via subcutaneous injection and remained on a heating pad until fully recovered from anesthesia. All mice were allowed to recover for two weeks before any subsequent experiments. All fiber sites were verified histologically.

Immunohistochemistry
Mice were dispatched by overdose with 50 μl sodium-pentobarbital and perfused transcardially with phosphate-buffered saline (PBS), followed by 4 % paraformaldehyde (PFA) in PBS. Extracted brains were kept in 4 % PFA at 4 ^{o}C overnight and stored in PBS. 50 μm coronal slices were cut using a vibratome and collected in PBS. Slices were washed in PBS-Triton X-100 (PBS-T) 0.2% followed by a 1 h blocking in PBS-T with 10% normal goat serum at room temperature before being incubated with the primary antibody at 4 oC overnight. Slices were washed in PBS-T 0.1 % followed by a 1 h incubation with the secondary antibody before undergoing another round of washing using PBS-T 0.1%. Vectashield DAPI was used to mount the slices onto super frost slides before visualization using an Olympus BX51 upright microscope. For cell counting experiments and dendritic spine analysis, an additional incubation with DAPI in PBS (1:1000) was carried out to label nuclei, and images were acquired using Leica SP8 gated stimulated emission depletion (STED) nanoscope. All images were taken at 40 X magnification.

Cell counting
To measure the extent to which populations of active cells overlap between the exposure to the same or different contexts, the number of EYFP$^+$ and c-Fos$^+$ immunoreactive neurons

in the DG, AMG, RSC, and PAG were counted. All animals were sacrificed 45 min post-assay or optical stimulation for immunohistochemical analyses. Four coronal slices were taken from the dorsal hippocampus of each animal. For optogenetic stimulation experiments, only slices with accurate implant location sites were used for counting. Sections were imaged on a Leica SP8 gated stimulated emission depletion (STED) nanoscope at a magnification of 40X. The area of interest was manually identified and the area of each region of interest was calculated using Fiji. To calculate the total number of DAPI cells in the DG, the average diameter of a sample of EYFP-positive cells in the DG was taken from each animal and the area of the cell was calculated. The total number of cells was estimated as the area of the DG divided by the area of the cell. For the AMG, PAG, and RSC, the number of DAPI cells in 3 randomly selected regions of interest were counted and used along with the total area of the region to determine the total number of DAPI cells. The total number of c-Fos positive and EYFP positive cells were manually identified and counted for each region using Adobe Photoshop CC 2018. To calculate the percentage of cells expressing EYFP in each region, the total number of EYFP-positive cells was divided by the total number of DAPI-positive cells for each area. To calculate the percentage of cells expressing c-Fos in each region, the total number of c-Fos positive cells was divided by the total number of DAPI-positive cells for each area. Non-engram cell activation was quantified by subtracting the total number of cells positive for both EYFP and c-Fos from the total number of c-Fos positive cells. To quantify engram reactivation, the number of EYFP and c-Fos positive cells were quantified as a percentage of total EYFP or DAPI positive cells. The level of chance was calculated by multiplying the percentage of EYFP$^+$ cells with the total percentage of c-Fos$^+$ cells and dividing by 100. This allowed for the determination of the chance of engram reactivation relative to the total number of EYFP$^+$ and c-Fos$^+$ labeled cells.

Dendritic spine analysis
The dendrites of DG engram cells in Ai32-FosTRAP mice were analyzed after recall at P25. Dendritic spines were imaged using the Leica SP8 gated STED scope. Z-stack images (10 μm) of 50 μm coronal brain sections were taken using Leica Application Suite X (LasX) software (line average 2, zoom factor 1.7-1.9) at 40X. Imaris software (Oxford Instruments, Imaris v9.5) was used to carry out dendritic spine analysis. Dendrites were traced (10 μm) using a semi-automated neurofilament tracer tool and dendritic spines were individually highlighted and manually traced with the software. Each fragment represents a different ChR2-EYFP$^+$ cell. Parameters were generated automatically using Imaris software. For analysis of dendritic spine head diameter and volume and average for each fragment was taken and plotted.

Behavioral assays
All behavioral experiments were conducted during the light cycle of the day (07:00 am to 7:00 pm). All behavioral subjects were individually habituated to handling by the investigator for 3 min on 3 consecutive days. Handling took place in the breeding room. Immediately before each handling session mice were transported in separate cages to and from the vicinity of the experimental room to habituate them to the journey.

Contextual fear conditioning
Three distinct contexts were used for contextual fear conditioning (CFC). Context A was a 31 x 24 x 21 cm Med Associates chamber with a removable grid floor (bars 3.2 mm diameter, spaced 7.9 mm apart), opaque triangular ceiling, and scented with 0.25 % benzaldehyde. Context B chambers were 29 x 25 x 22 cm Coulbourne Instruments chambers

with Perspex white floors, bright white lighting, and scented with 1 % acetic acid. All conditioning sessions were conducted in Context A. Each session was 330 s in duration with three 0.75 mA shocks of 2 s duration delivered at 150 s, 210 s, and 270 s. Mice were removed from the conditioning chamber 60 s following the final footshock and returned to their homecage. All recall or context generalization tests were 180 s in duration. Testing conditions were identical to training conditioning except that no shocks were presented. Up to 4 mice were run simultaneously in the 4 identical chambers. Floors of chambers were cleaned with Trigene before and between runs. Mice were transported to and from the experimental room in separate Perspex cages. All experimental groups were counter-balanced for chambers within contexts.

Contextual pre-exposure
For context pre-exposure mice were exposed to Context A (31 x 24 x 21 cm Med Associates chamber with a removable grid floor) for 10 min. Mice were removed from the chamber and placed back in their homecage. The following day, mice were placed back in Context A where an immediate foot shock (1mA, 2s) was delivered. Mice were removed from the chamber 1 min after the shock and placed back in their homecage. Up to 4 mice were run simultaneously in the 4 identical chambers except for during the update when one mouse was run at a time. All recall or context specificity tests were 180 s in duration. Floors of chambers were cleaned with Trigene before and between runs.

Novel object recognition
Each subject was allowed to habituate to the rectangular testing arena for 10 min on the day before object acquisition. During object acquisition, two identical objects were positioned on adjacent walls of the arena. Mice were allowed to explore both objects freely for a 10 min period before being placed back in their homecage. On testing, one familiar object was replaced with a novel object and each subject was introduced for a 5 min exploration period. Object exploration was defined as the time when the subject's nose came within a 2 cm radius of the object. In between each trial, the testing arena and objects were cleaned with Trigene. For engram labeling experiments, P17 mice received a 4-OHT injection 2 h after object acquisition. For optogenetic reactivation experiments, animals received 3 min of blue light stimulation directly before being placed into the testing arena.

Barnes maze
Training consisted of five trials with a 10 min limit per trial. At the start of each trial, the mouse was placed in a black PVC start chamber located in the center of the maze apparatus. After 15 secs, the start chamber was removed, and the latency and distance traveled to enter the escape tunnel were recorded. Subjects were placed back in their homecage and allowed 40 min in between each trial. The position of the escape tunnel remained in a fixed location relative to spatial cues in the room. For each probe trial, the escape tunnel was removed from the apparatus. Each mouse was allowed 5 min to freely explore the four quadrants of the maze before being removed and placed back in its homecage. The latency and distance traveled before reaching the location of the removed escape tunnel, along with the time spent in each quadrant were recorded. The apparatus and escape tunnel were cleaned with Trigene after each trial. For engram labeling experiments, P17 mice received a 4-OHT injection 2 h after the 5th trial. For optogenetic reactivation experiments, animals received 3 min of blue light stimulation directly before being placed into the testing arena.

Three-chamber social interaction

Mice were habituated to the chamber with two empty holders for 10 min. The next day, mice were placed in the middle of the chamber and allowed to explore the three chambers for a period of 5 min. During this testing period, a social object (novel mouse) was contained in one holder in one chamber and an inanimate object (lego blocks) was contained in a holder in the other chamber. Object exploration and distance traveled were tracked using ANY-maze video tracking software.

Repetition assay
A large homecage was filled 5 cm deep with wood chipping bedding and lightly tamped down to make an even flat surface. A consistent pattern of 20 identical glass marbles (15mm diameter) was evenly placed (4 cm apart) on the surface of the wood chip bedding. Mice were left alone in the testing arena for 30 min. A picture was taken before and after the test for analysis. A marble was considered buried if 2/3rds of the depth of the marble was buried.

Optogenetic engram reactivation
Optogenetic engram reactivation was carried out in Context C, a 31 x 24 x 21 cm Med Associates chamber with a white Perspex floor and curved insert, a dim light level and scented with 0.25 % acetophenone. The optical fiber implant was connected to a 45 nm laser diode fiber light source (Doric LDFLS 450/080). Habituation sessions lasted for 12 min. During testing, the 12 min session was divided into four 3 min epochs split into two light-off and two light-on epochs. During the light-on epochs, blue light stimulation (20 Hz) with a pulse width of 15 ms was delivered through the optical fiber implant for the entire 3 min duration. At the end of the 12 min session, the mouse was immediately detached from the laser and returned to its homecage.

Artificial updating
Context A was a 31 x 24 x 21 cm Med Associates chamber with a removable Perspex black floor, black triangular ceiling, and scented with 0.25 % benzaldehyde. Context B exposure was carried out in a rectangular testing arena. Artificial updating of a contextual memory was carried out in Context C, a 29 x 25 x 22 cm Coulbourne Instruments chamber with removable grid floors, bright white lighting, and scented with 1 % acetic acid. Contexts A, B, and C were all located in different behavioral rooms. The optical fiber implant was connected to a 45 nm laser diode fiber light source (Doric LDFLS 450/080). The 7 min session consisted of a 2 min acclimatization period followed by 5 min of blue light stimulation (20Hz) with a pulse width of 15 ms delivered through the optical fiber implant for the entire 5 min duration. Three shocks, 0.75 mA of 2 s duration, were delivered at minutes 4, 5, and 6 of the session. At the end of the 7 min session, the mouse was immediately detached from the laser and returned to its homecage.

Maternal immune activation
Mice were mated overnight for one night. Females were weighed and checked for seminal plugs, noted as E0.5. Pregnant dams were injected SC with a single dose of Poly I:C HMW (InvivoGen) at 20mg/kg (*13*, *64*, *65*), rmlIL-17A (Immunotools) at 50 ug/kg, or control PBS on E12.5. For more detailed information refer to Table S2 (*66*). For rmlIL-17A injections at P17 or P25, mice were injected SC with a single dose of rmlIL-17A (Immunotools) at 50 ug/kg or control PBS 3 h before the recall test.

Postnatal immune activation

Mice were injected SC with a single dose of Poly I:C HMW (InvivoGen) at 20 mg/kg or control PBS on P3, P7, and P14. For serum analysis by ELISA, blood serum was collected from female mice 24 h after subcutaneous administration of either PBS or Poly (I:C) (20 mg/kg), and frozen at -20 oC until further use. Blood serum was first diluted 1:5 and IL-17a levels were measured using the ELISA MAX Deluxe Set (Biolegend) as per the manufacturers' instructions.

Genotyping
Genomic DNA was extracted from the ear punches of each mouse and sent to Transnetyx for genotyping.

Statistical analysis
All experiments were analyzed blind to the experimental group. All videos were randomized before manual scoring. Behavioral performance was recorded by a digital video camera. Videos were manually scored individually, and investigators were blind to the experimental condition and test day during all manual scoring. Data analysis and statistics were conducted using GraphPad Prism 6.00 (GraphPad software) or RStudio v2023.03.0+386. Unpaired Student's t-tests were used for independent group comparisons. Paired Student t-tests were used to assess differences within groups. ANOVA followed by a Bonferroni post hoc test was used to determine conditions that were significant from each other where appropriate. Outliers were detected using Grubbs' test where $p < 0.05$. For cell counting analysis results were analyzed per mouse using nested t-test or nested ANOVA followed by Bonferroni post hoc test, where each symbol represents N=1. For maternal immune activation behavioral analysis, results were analyzed per litter (Table S1) using nested t-test or nested ANOVA followed by Bonferroni post hoc test. Pearson correlation coefficient was used to measure the relationship between two variables. Results are displayed as mean with SEM and deemed significant when $p < 0.05$.

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Competing interests:

The authors declare that they have no competing interests.

Data and materials availability:

All data needed to evaluate the conclusions in the paper are present in the paper and/or the Supplementary Materials.

Figures and Tables

Fig. 1. MIA male offspring do not demonstrate infantile amnesia in a contextual fear paradigm.

(**A**) Developmental trajectory of infant mice. (**B**) Behavioral schedule. The black lightning symbol represents foot shocks. Mice are trained using contextual fear conditioning (CFC) in context A and tested for recall 1 or 8 days later. Shock (S), no shock (NS). (**C**) Memory recall in context A. Adult (P63) C57BL/6J male mice (n=8) froze significantly more in context A than no shock controls (n=8) both 1 and 8 days after training. Infant (P17) C57BL/6J male mice (n=9) froze significantly more than no shock controls during recall 1 day after training. No significant difference in freezing between groups (n=9) when tested for recall 8 days after training. (**D**) Representative diagram of MIA in pregnant dams. The syringe indicates Poly I:C (PIC) injection at E12.5. (**E**) Social preference index (% time spent exploring social stimulus out of total object investigation time) of MIA adult C57BL/6J male offspring (n=10). (**F**) Marble-burying behavior of MIA adult C57BL/6J male offspring (n=10). Percentage of the number of buried marbles is plotted on the y-axis. (**G**) Memory recall in male C57BL/6J MIA offspring in context A after CFC at P17 (n=10). $P < 0.05, P < 0.01, P < 0.001$ calculated by (**C**) two-way ANOVA (**E**, **F**, **G**) or nested ANOVA with Bonferroni post hoc tests. Data presented as mean ±SEM.

Fig. 2. Natural reactivation of engram cells for an infant encoded memory.
(**A**) Engram labeling method in Ai32-FosTRAP mice. (**B**) Whole-brain engram labeling of an infant engram. Scale bar 100 μm. (**C**) DG engram cells (ChR2-EYFP$^+$ green) in Ai32-FosTRAP mice. Scale bar 100 μm. (**D**) Behavioral schedule for engram labeling of a homecage (HC), contextual (Cxt), or contextual fear memory (CFC) in P17 Ai32-FosTRAP mice. The black lightning symbol represents foot shocks. Syringe symbol represents 4-OHT injection 2 h after training. (**E, F**) ChR2-EYFP$^+$ cell counts in the DG (N=4-6, n=4) and AMG (N=4, n=4). (**G**) Example images of DG ChR2-EYFP$^+$ (green) and c-Fos$^+$ (red) cell counts after recall at P20 and P63. Infantile Amnesia (IA). Scale bar 20 μm. (**H, K**) Behavioral schedule. (**I, J**) Engram reactivation (c-Fos$^+$ChR2-EYFP$^+$/ChR2-EYFP$^+$) in the DG (N=4-6, n=4) and AMG (N=4, n=4) after recall or no recall (homecage) at P20 and P63. (**L**) ChR2-EYFP$^+$ cell counts in the DG in MIA male offspring (N=5-6, n=4). (**M**) Scatter plot of relationship between the number of EYFP$^+$/DAPI cells and freezing behavior (%) (N=8, n=4). (**N**) c-Fos$^+$ cell counts in the DG (N=4-5, n=4). (**O**) c-Fos$^+$ cell counts in non-engram cells the DG (N=10, n=4). (**P**) c-Fos$^+$ChR2-EYFP$^+$/ChR2-EYFP$^+$ cell counts in the DG (N=4-5, n=4). (**Q**) Engram cell dendritic spines in the DG in MIA male offspring after recall at P25 (N=4, n=10). Scale bar 20 μm (**R**) Spine density per 10μm. (**S**) Average dendritic spine head diameter (μm). (**T**) Average dendritic spine head volume (μm^2). $P < 0.05, P < 0.01, P < 0.001$ calculated by (**E, L, N-P**) nested Student t-test, (**R-T**) Student t-test, (**M**) Pearson's correlation or (**F, I, J**) nested ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. 3. Artificial updating of infant encoded engram cells permanently reinstates a 'lost' infantile memory.

(**A**) Behavioral schedule for optogenetic reactivation of infant DG engram cells in adulthood. (**B**) Representative image of optogenetic stimulation of DG engram cells in Ai32-FosTRAP labeled mice. Scale bar 100 μm. (**C**) Freezing levels of infant mice (n=10) during natural memory recall. No shock (NS), shock (S). (**D**) Memory recall in context C (engram reactivation) with light-off and light-on epochs. (**E**) Freezing for the two light-off and light-on epochs averaged. (**F, I**) ChR2-EYFP$^+$ cell counts in the DG (N=4, n=4) and AMG (N=5, n=4) after optogenetic stimulation at P63. (**G, J**) c-Fos$^+$ cell counts in the DG and AMG. (**H, K**) c-Fos$^+$ChR2-EYFP$^+$/ChR2-EYFP$^+$ cell counts in the DG and AMG. (**L**) Behavioral schedule for artificial updating of an infant engram. (**M**) Freezing levels during recall test in context B and context A (n=10). Experimental light group froze significantly more in context A. (**N, O**) Histological representation of engram reactivation in the (**N**) DG and (**O**) RSC during recall in context A. DAPI$^+$ (blue), ChR2-EYFP$^+$ (green), c-Fos$^+$ (red). Scale bar 20 μm. (**P, Q**) DG (N=6, n=4) and RSC (N=4-5, n=4) c-Fos$^+$ChR2-EYFP$^+$/ChR2-EYFP$^+$ cell counts after recall in context A. No light (NL), light (L). *P < 0.05, P < 0.01, P < 0.001* calculated by (**F-K, P, Q**) nested Student t-test or (**C, E, M**) two-way ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. 4. Infantile amnesia for context, object, and maze memory.

(**A**, **C**) Behavioral schedule for novel object recognition. (**B**) Interaction index (%) of infant C57BL/6J mice during novel object recognition test 1 or 8 d after acquisition (n=9). (**D**) Interaction index of MIA male infant offspring (n=9-10). (**E**) Behavioral schedule for DG optogenetic reactivation of an infant encoded engram for an object. (**F**) Interaction index of adult mice after optogenetic stimulation (n=9-10). No light (NL), light (L). (**G**) Behavioral schedule for the Barnes maze in infant C57BL/6J mice. (**H**, **I**) Time spent in each zone during the test (**H**) 1 d or (**I**) 8 d after training. (**J**) Example representations of navigational search strategies. (**K**, **L**) Infant and adult search strategies during testing 1 and 8 d after training. (**M**) Behavioral schedule for the Barnes maze in infant C57BL/6J male MIA offspring (**N**, **P**, **R**) Time spent in each zone by MIA male offspring during test 8 d after training (n=9-11). (**O**, **Q**, **S**) Heat map analysis of probe test in (**O**) Ctrl, (**Q**) PIC (**S**) IL-17a male MIA offspring. (**T**) Behavioral schedule for DG optogenetic reactivation of an infant encoded engram for the Barnes maze. (**U**, **W**) Time spent in each zone during the test (n=8-9). (**V**, **W**) Heat map analysis of probe test in (**V**) no light and (**X**) light groups. *P < 0.05, P < 0.01,* P < 0.001 calculated by (**H**, **I**, **U**, **W**) ANOVA, (**D**, **N**, **P**, **R**) nested ANOVA, (**B**) two-way repeated measures ANOVA or (**F**) two-way ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. 5. IL-17a is required for MIA effects on infantile amnesia.

(**A**) Representative diagram of MIA in *Il17a* KO mice. (**B**) Behavioral schedule for CFC in *Il17a* KO male mice. (**C**) Freezing levels of male *Il17a* KO infant mice during recall 1 or 8 d after training. (**D**) Behavioral schedule for postnatal immune activation in C57BL/6J infant mice. Syringe symbol represents Poly I:C (PIC) injection at P3, P7, and P14. (**E**) Behavioral schema for CFC. (**F**) Freezing levels of infant male mice when tested 8 d after training. (**G**) Behavioral schedule for IL-17a injection 3 h before memory encoding (P17) or recall (P25) in MIA C57BL/6J offspring. (**H**, **I**) Freezing levels of infant male mice when tested 8 d after training. *P < 0.05, P < 0.01, P < 0.001* calculated by (**F**, **H**, **I**) nested Student t-test or (**C**) nested ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Supplementary Materials for

Immune activation state modulates the retrieval of infant engrams

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Figs. S1 to S16
Tables S1 to S2

Fig. S1. Delayed weaning or environmental enrichment does not affect infantile amnesia.
(**A**) Behavioral schedule for CFC in infant (P17) C57BL/6J male mice subjected to normal (P21) or delayed (P24) weaning. The lightning symbol represents footshocks. (**B-E**) Freezing levels during recall tests (n=10) at Test 1 (P18), CxtB (P19), Test 3 (P25), and Test 4 (P32). (**F**) Behavioral schedule for CFC in infant C57BL/6J male mice. (**G-J**) Freezing levels during recall tests (n=10). *P < 0.05, P < 0.01, P < 0.001* calculated by (**B-E**) Student t-test or (**G-J**) ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S2. MIA female offspring do not demonstrate social interaction deficits or repetitive behavior but do show infantile amnesia for a fear memory.

(**A**) IL-17a cytokine levels in blood serum of female C57 mice injected with PBS (Ctrl n=5) or Poly I:C (PIC n=5). (**B, E**) Total distance (m) traveled during social interaction test in adult (P63) C57BL/6J (**B**) male (n=10) and (**E**) female (n=10) MIA offspring. (**C, F**) Heat map analysis of time spent in each chamber during social interaction task. (**D**) Social interaction index (%) of adult (P63) C57BL/6J female MIA offspring (n=10). (**G**) Number of marbles buried by adult female MIA offspring during marble burying task (n= 8-10). (**H**) Memory recall for CFC in context A for male MIA offspring testing 1 d after training (n=10). (**I, J**) Recall for CFC in context A for female MIA offspring tested (**I**) 1 d (n=9-10) or (**J**) 8 d (n= 9) after training. (**K**) Memory recall for CFC in context A for male MIA offspring tested 8 d (P25) (n=10) or 15 d (P32) (n=13) after training. (**L**) Memory recall comparison in male and female MIA offspring 8 d after training. *P < 0.05, P < 0.01, P < 0.001* calculated by (**A**) Student t-test, (**K**) nested Student t-test, (**B, D, E, G-J**) nested ANOVA, or (**L**) two-way ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S3. Infant mice show similar memory retention when tested 1 or 3 d after training at P17.
(**A**) Behavioral schedule for CFC in C57BL/6J P17 mice. The syringe symbol represents 4-OHT injection 2 h after exposure. The black lightning symbol represents foot shocks. (**B**, **C**) Freezing levels (%) during recall in context A. No shock (NS), shock (S). (**D**) Behavioral schedule for CFC in C57BL/6J P29 mice. (**E**) Freezing levels (%) during recall in context A at P63. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**, **C**, **E**) Student t-test. Data presented as ±SEM.

Fig. S4. Adult Ai32-FosTRAP mice show activity-dependent increases in ChR2-EYFP labeling.
(**A**) Behavioral schedule. The syringe symbol represents 4-OHT injection 2 h after exposure. The black lightning symbol represents foot shocks. (**B**-**E**) Quantification of ChR2-EYFP+ cells in the (**B**) DG (N=4-7, n=4), (**C**) AMG (N=4-5, n=4), (**D**) RSC (N=4-6, n=4) and (**E**) PAG (N=3-4, n=4) for an adult (P42) encoded homecage (HC), context (Cxt) and contextual fear (CFC) memory. Exposure to a context resulted in significantly more ChR2-EYFP+ cells in the (**B**) DG (p < 0.001) and (**D**) RSC (p < 0.001). Quantification of ChR2-EYFP+ cells after CFC was significantly higher in the (**B**) DG (p < 0.05), (**C**) AMG (p < 0.01), and the (**D**) RSC (p < 0.05) compared to HC control. (**F**, **G**) ChR2-EYFP+ cell counts (N=4, n=4) in the RSC and PAG for an infant (P17) encoded HC, Cxt, and CFC memory. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**-**G**) nested ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S5. Engram reactivation is increased in the DG and AMG after natural recall at P20.

(**A**) Behavioral paradigm. (**B**-**O**) Histological analysis of cell counts after recall at P20. Percentage of (**B**) ChR2-EYFP⁺ and (**C**) c-Fos⁺ cells in the DG (N=5-6, n=4). There was a significant increase in c-Fos⁺ cells in the DG after recall at P20 (p < 0.05). (**D**) Percentage of DAPI cells positive for both ChR2-EYFP and c-Fos. (**D**) Engram reactivation as a percentage of DAPI⁺ was significantly higher in the DG after recall at P20 (p < 0.05). Percentage of (**E**) ChR2-EYFP⁺ and (**F**) c-Fos⁺ cells in the AMG (N=4, n=4). There was a significant increase in c-Fos⁺ cells in the AMG after recall at P20 (p < 0.05). (**G**) Percentage of DAPI cells positive for both ChR2-EYFP and c-Fos. (**G**) Engram reactivation as a percentage of DAPI⁺ in the AMG was significantly higher after recall at P20 (p < 0.05). Percentage of (**H**) ChR2-EYFP⁺ and (**I**) c-Fos⁺ cells in the RSC (N=4, n=4). Engram reactivation in the RSC as a percentage of (**J**) ChR2-EYFP⁺ and (**K**) DAPI⁺ cells. Percentage of (**L**) ChR2-EYFP⁺ and (**M**) c-Fos⁺ cells in the PAG (N=4, n=4). Engram reactivation in the PAG as a percentage of (**N**) ChR2-EYFP⁺ and (**O**) DAPI⁺ cells. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**-**O**) nested t-test. Data presented as ±SEM.

Fig. S6. Presentation of recall cues for an infant encoded context does not result in engram reactivation at P63.

(**A**) Behavioral paradigm. (**B**-**O**) Histological analysis of cell counts after recall at P63. Percentage of (**B**, **E**) ChR2-EYFP+ and (**C**, **F**) c-Fos+ cells in the DG (N=4-5, n=4) and AMG (N=4, n=4). (**C**) There was a significant increase in c-Fos+ cells in the DG after recall at P63 (p < 0.05). (**D**, **G**) Engram reactivation as a percentage of DAPI+ in the DG and AMG. Percentage of (**H**, **L**) ChR2-EYFP+ and (**I**, **M**) c-Fos+ cells in the RSC (N=4-5, n=4) and PAG (N=3-4, n=4). (**I**) There was a significant increase in c-Fos+ cells in the RSC after recall at P63 (p < 0.05). Engram reactivation in the RSC and PAG as a percentage of (**J**, **N**) ChR2-EYFP+ and (**K**, **O**) DAPI+ cells. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**-**O**) nested Student t-test. Data presented as ±SEM.

Fig. S7. MIA female offspring do not show an increased number of EYFP⁺ cells in the DG.

(**A**) Behavioral paradigm for labeling engram cells in MIA female Ai32-FosTRAP offspring. (**B**) ChR2-EYFP⁺ cell counts in the DG (N= 4, n=4) of female MIA offspring after recall at P25. (**C**) Engram reactivation in the DG of male MIA offspring as a percentage of DAPI⁺ cells. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**, **C**) nested Student t-test. Data presented as ±SEM.

Fig. S8 Optogenetic reactivation of an infant engram in the CA1 elicits freezing behavior in adult mice.

(**A**) Behavioral schedule for engram reactivation of a neutral context B in the DG of Ai32-FosTRAP male mice. The black lightning symbol represents foot shocks. The syringe symbol represents 4-OHT injection 2 h after context exposure. (**B**) Memory recall in context C (engram reactivation) with light-off and light-on epochs (n=9). (**C**) Freezing for the two light-off and light-on epochs averaged. (**D**) Behavioural schedule for infant engram reactivation in the CA1 of adult Ai32-FosTRAP male mice (n=7) (NS), (n=9) (S). (**E, G**) Freezing levels during natural memory recall. (**F, H**) Memory recall in context C during engram reactivation at (**F**) 20 Hz or (**H**) 4 Hz with light-off and light-on epochs. (**I**) Freezing for the two light-off and light-on epochs during 4 Hz stimulation averaged. *P < 0.05, P < 0.01, P < 0.001* calculated by (**C, G**) Student t-test or (**E, I**) two-way ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S9. Optogenetic reactivation of an infant encoded engram in adult mice.
(**A**) Behavioral paradigm. (**B**-**K**) Cell counts after optogenetic reactivation of an infant encoded engram in adult Ai32-FosTRAP mice. No shock (NS), shock (S). (**B**, **C**) Percentage of DAPI cells positive for both ChR2-EYFP and c-Fos in the DG and AMG. (**C**) Engram reactivation as a percentage of DAPI$^+$ cells was significantly higher (p < 0.05) in the AMG (N=5, n=4) after optogenetic stimulation of an infant-encoded fear memory. Percentage of (**D**) ChR2-EYFP$^+$ and (**E**) c-Fos$^+$ cells in the RSC (N=4, n=4). Engram reactivation in the RSC as a percentage of (**F**) ChR2-EYFP$^+$ and (**G**) DAPI$^+$ cells. Percentage of (**H**) ChR2-EYFP$^+$ and (**I**) c-Fos$^+$ cells in the PAG (N=4, n=3-4). Engram reactivation in the PAG as a percentage of (**J**) ChR2-EYFP$^+$ and (**K**) DAPI$^+$ cells. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**-**K**) nested Student t-test. Data presented as ±SEM.

Fig. S10 Optogenetic stimulation of an CFC engram encoded after the infantile period causes an increase in activity in the amygdala in adult mice.

(**A**) Behavioral schedule for optogenetic reactivation of DG engram cells, labeled at P29, in adult (P63) Ai32-FosTRAP mice. The black lightning symbol represents foot shocks. The syringe symbol represents 4-OHT injection 2 h after training. Percentage of ChR2-EYFP+ cells labeled during training in the (**B**) DG (N=4, n=4), (**F**) AMG (N=5-6, n=4) ($p < 0.05$), (**J**) RSC (N=4, n=4), and (**N**) PAG (N=4, n=4). Percentage of c-Fos+ cells in the (**C**) DG, (**G**) AMG, (**K**) RSC, and (**O**) PAG after optogenetic stimulation at P63. Engram reactivation as a percentage of (**D, H, L, P**) EYFP+ cells and (**E, I, M, Q**) DAPI+ cells. *$P < 0.05$, $P < 0.01$, $P < 0.001$* calculated by (**B-Q**) nested Student t-test. Data presented as ±SEM.

Fig. S11. Infant mice can form context-specific memories that can be behaviorally updated.
(**A**) Behavioral schedule for artificial updating of an adult engram in Ai32-FosTRAP mice. (**B**) Freezing levels during recall in context B and context A (n=10). Experimental light group froze significantly more in context A (*P < 0.001*). (**C**) Behavioral schedule for updating paradigm in C57BL/6J infant (P17) mice. (**D**) Freezing levels (n=10) during recall tests at P19 (Recall 1), P20 (CxtB), and P63 (Recall 2). Infant mice that were pre-exposed (PE) to context A froze significantly (*P < 0.01*) more during recall 1. (**E**) Behavioral schedule for updating paradigm when update occurs after the infantile amnesia period. (**F**) Freezing levels (n=9) during recall tests. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**, **D**, **F**) two-way ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S12. Artificial updating of infant engram cells in adult mice results in an above-chance level of engram reactivation in the DG, AMG, RSC, and PAG.

(**A**) Behavioral paradigm. (**B**-**O**) Histological analysis of cell counts after context A recall in adult Ai32-FosTRAP mice. Percentage of (**B**) ChR2-EYFP+ and (**C**) c-Fos+ cells in the DG (N=6, n=4). No light (NL), Light (L). (**D**) Percentage of DAPI cells positive for both ChR2-EYFP and c-Fos. The experimental light group showed a significant increase (p < 0.05) in engram reactivation in the DG. Percentage of (**E**) ChR2-EYFP+ and (**F**) c-Fos+ cells in the AMG (N=4, n=4). Engram reactivation in the AMG as a percentage of (**G**) ChR2-EYFP+ cells and (**H**) DAPI+cells. Percentage of (**I**) ChR2-EYFP+ and (**J**) c-Fos+ cells in the RSC (N4-5, n=4). (**K**) Percentage of DAPI cells positive for both ChR2-EYFP and c-Fos. Percentage of (**L**) ChR2-EYFP+ and (**M**) c-Fos+ cells in the PAG (N=4, n=4). Engram reactivation in the PAG as a percentage of (**N**) ChR2-EYFP+ cells and (**O**) DAPI+ cells. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**-**O**) nested Student t-test. Data presented as ±SEM.

Fig. S13. Adult mice retain an object memory 8 days after training.

(**A**) Behavioral schedule for novel object recognition task in adult C57BL/6J male mice. (**B**) Object interaction during acquisition in infant (P17) (n=10) and adult (P63) (n=9) mice. (**C**) Object interaction (%) during novel object recognition test in adult mice (n=9). Mice spend significantly more time exploring the novel object when tested 1 d (p < 0.001) or 8 d (p < 0.001) after acquisition. (**D**) Behavioral schedule for novel object recognition task in infant (P17) MIA offspring. (**E**) Object interaction (%) during acquisition in infant (P17) C57BL/6J MIA male offspring (n=9-10). (**F**) Object interaction (%) during acquisition in infant (P17) male Ai32-FosTRAP mice (n=8-9). No light (NL), light (L). *P < 0.05, P < 0.01, P < 0.001* calculated by (**E**) nested ANOVA or (**B, C, F**) two-way ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S14. Adult mice retain both a spatial memory for the location of the escape hole on the Barnes maze task 8 days after training.

(**A**) Time (s) and (**B**) distance (m) taken by infant C57BL/6J mice to reach the escape hole during each trial (n=18-20). (**C**) Time (s) and (**D**) distance (m) taken to first reach where the escape hole should be located during probe tests (n=9-10). (**E, F**) Heat map analysis of probe test (**E**) 1 d or (**F**) 8 d after training. (**G**) Behavioral schedule for the Barnes maze in adult (P63) C57BL/6J male mice. (**H**) Time (s) and (**I**) distance (m) taken to reach the escape hole during each trial (n=20). (**J-M**) Time spent in each zone during probe test (**J, K**) 1 d (*P < 0.001*) or (**L, M**) 8 d (*P < 0.05, P < 0.01*) after training. (**N**) Time (s) and (**O**) distance (m) taken to first reach where the escape hole should be located during probe tests. (**P**) Behavioral schedule for DG optogenetic reactivation of an infant encoded engram for the Barnes maze. (**Q, S**) Time spent in each zone during the test (n=8-9). (**R, T**) Heat map analysis of probe test. *P < 0.05, P < 0.01, P < 0.001* calculated by (**C, D, N, O**) Student t-test or (**J, L, Q, S**) ANOVA with Bonferroni post hoc tests. Data presented as ±SEM.

Fig. S15 *Il17a* **KO adult mice show natural memory retention for contextual fear memory when tested 8 d after training.**

(**A**) Behavioral schedule for CFC in *Il17a* KO adult (P63) mice. The black lightning symbol represents foot shocks.

(**B**) Freezing levels (%) during recall in context A. *P < 0.05, P < 0.01, P < 0.001* calculated by (**B**) Student t-test. Data presented as ±SEM.

Fig. S16. *Il17a* KO MIA offspring do not show social interaction deficits or repetitive behavior.
(**A**) Total distance (m) traveled during social interaction task in adult (P63) male and female *Il17a* KO MIA offspring
(n=9-10). (**B**) Social interaction index (%) of adult male and female *Il17* KO MIA offspring. (**C, D**) Heat map analysis
of time spent in each chamber for (**C**) male and (**D**) female *Il17a* KO adult MIA offspring. (**E**) Number of marbles
buried during marble burying task (n=10). *P < 0.05, P < 0.01, P < 0.001* calculated by (**A, B, E**) nested ANOVA with
Bonferroni post hoc tests. Data presented as ±SEM.

Figure	Sex	Strain	Injection	Compound	Cohort	Litters	n
1E	M	C57BL/6J	E12.5	PBS	Ctrl	2	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC	5	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
1F	M	C57BL/6J	E12.5	PBS	Ctrl	2	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC	5	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
IG	M	C57BL/6J	E12.5	PBS	Ctrl NS	3	10
	M	C57BL/6J	E12.5	PBS	Ctrl S	3	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC NS	3	9
	M	C57BL/6J	E12.5	poly (I:C)	PIC S	4	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a NS	4	9
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a S	4	10
2L-N	M	Ai32FosTRAP	E12.5	PBS	Ctrl	4	5
	M	Ai32FosTRAP	E12.5	poly (I:C)	PIC	3	5
2P-R	M	Ai32FosTRAP	E12.5	PBS	Ctrl	4	4
	M	Ai32FosTRAP	E12.5	poly (I:C)	PIC	2	4
4D	M	C57BL/6J	E12.5	PBS	Ctrl	4	9
	M	C57BL/6J	E12.5	poly (I:C)	PIC	3	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	9
4L	M	C57BL/6J	E12.5	PBS	Ctrl	5	10
4M	M	C57BL/6J	E12.5	poly (I:C)	PIC	5	9
4N	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	3	11
5C	M	Il17a-/-	E12.5	PBS	Ctrl 1d	4	9
	M	Il17a-/-	E12.5	PBS	Ctrl 8d	3	9
	M	Il17a-/-	E12.5	poly (I:C)	PIC 1d	4	8
	M	Il17a-/-	E12.5	poly (I:C)	PIC 8d	4	8
5F	M	C57BL/6J	P3, P7, P14	PBS	Ctrl	5	11
	M	C57BL/6J	P3, P7, P14	poly (I:C)	PIC	6	10
5H	M	C57BL/6J	E12.5, P17	poly (I:C), PBS	PIC+PBS	5	9
	M	C57BL/6J	E12.5, P17	poly (I:C), rmIIL-17A	PIC+IL-17a	5	8
5I	M	C57BL/6J	E12.5, P25	poly (I:C), PBS	PIC+PBS	7	11
	M	C57BL/6J	E12.5, P25	poly (I:C), rmIIL-17A	PIC+IL-17a	7	12
S2A, B	M	C57BL/6J	E12.5	PBS	Ctrl	2	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC	5	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
S2C-E	F	C57BL/6J	E12.5	PBS	Ctrl	5	10
	F	C57BL/6J	E12.5	poly (I:C)	PIC	4	10
	F	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
S2F	F	C57BL/6J	E12.5	PBS	Ctrl	4	8
	F	C57BL/6J	E12.5	poly (I:C)	PIC	4	9
	F	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
S2G	M	C57BL/6J	E12.5	PBS	Ctrl	4	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC	5	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
S2H	F	C57BL/6J	E12.5	PBS	Ctrl	5	10
	F	C57BL/6J	E12.5	poly (I:C)	PIC	3	9
	F	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	10
S2I	F	C57BL/6J	E12.5	PBS	Ctrl	5	9
	F	C57BL/6J	E12.5	poly (I:C)	PIC	3	9
	F	C57BL/6J	E12.5	rmIIL-17A	IL-17a	5	9
S2J	M	C57BL/6J	E12.5	poly (I:C)	PIC P25	4	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC P32	4	13
S2K	M	C57BL/6J	E12.5	PBS	Ctrl	3	10
	M	C57BL/6J	E12.5	poly (I:C)	PIC	4	10
	F	C57BL/6J	E12.5	PBS	Ctrl	5	9
	F	C57BL/6J	E12.5	poly (I:C)	PIC	3	9
S7	F	Ai32FosTRAP	E12.5	PBS	Ctrl	3	4
	F	Ai32FosTRAP	E12.5	poly (I:C)	PIC	3	4
S13E	M	C57BL/6J	E12.5	PBS	Ctrl	4	9
	M	C57BL/6J	E12.5	poly (I:C)	PIC	3	10
	M	C57BL/6J	E12.5	rmIIL-17A	IL-17a	4	9
S15A, B	M	Il17a-/-	E12.5	PBS	Ctrl	4	9
	M	Il17a-/-	E12.5	poly (I:C)	PIC	4	10
	F	Il17a-/-	E12.5	PBS	Ctrl	2	9
	F	Il17a-/-	E12.5	poly (I:C)	PIC	4	10
S15E	M	Il17a-/-	E12.5	PBS	Ctrl	5	10
	M	Il17a-/-	E12.5	poly (I:C)	PIC	4	10
	F	Il17a-/-	E12.5	PBS	Ctrl	2	10
	F	Il17a-/-	E12.5	poly (I:C)	PIC	4	10

Table S1. Record of litters used for each immune activation experiment.

Maternal Immune Activation Model Reporting Guidelines Checklist

ARRIVE Reporting Guideline & Recommendation	Arrive Item	MIA Model Specific Reporting Recommendation *Please complete this chart for each point outlined below. If not applicable, write N/A*
Study design ➢ Overview of immune activation issues For each experiment, give brief details of the study design including: a. The number of experimental and control groups. b. Any steps taken to minimize the effects of subjective bias when allocating animals to treatment (e.g. randomization procedure) and when assessing results (e.g. if done, describe who was blinded and when). c. The experimental unit (e.g. a single animal, group or cage of animals). A time-line diagram or flow chart can be useful to illustrate how complex study designs were carried out.	6	MIA Specific Reporting: a. General need for improved reporting in MIA model methods + reporting pilot data ○ Details on pilot data: Mice were randomly assigned to experimental groups with littermate controls or otherwise relevant controls. During behavioral analysis, investigators were blinded to the treatment conditions for all experimental groups. Histological analysis was conducted individually for each mouse using nested analysis, where > 4 slices per mouse were analyzed. The same approach was applied to maternal immune activation experiments where nested analysis was performed per litter. Throughout all experiments analysis was conducted blindly. For number of experimental and control groups please refer to Table 1 in supplementary materials.

Experimental procedures
➢ Compounds
➢ Validation measures
 	
For each experiment and each experimental group, including controls, provide precise details of all procedures carried out. For example:
a. How (e.g. drug formulation and dose, site and route of administration, anaesthesia and analgesia used [including monitoring], surgical procedure, method of euthanasia). Provide details of any specialist equipment used, including supplier(s).
b. When (e.g. time of day).
c. Where (e.g. home cage, laboratory, water maze).
d. Why (e.g. rationale for choice of specific anaesthetic, route of administration, drug dose used).
	7	Provide details of:
		a. Compounds – source, vehicle, preparation/storage, administration route, volume administered, whether anesthetics were used at time of immune challenge.
		    ○ Name of compound: Polyinosinic:polycytidylic acid or rmIL-17A
		    ○ Catalogue number: InvivoGen 31852-29-6 or ImmunoTools 12340174
		    ○ Lot number: 5934-42-01/2, 5934-43-01/2/3/4/5, 5934-44-01
		    ○ Vehicle control used: Phosphate buffer solution
		    ○ Route of administration: subcutaneous
		    ○ Volume administered: 20mg/kg (200ul) or 50ug/kg (200ul)
		    ○ Storage conditions: -20°C
		    ○ Anesthetic (type, dose, duration) used: n/a
		b. Housing variables at injection - temperature of room at injection time, cage change at time of injection or not
		    ○ Light cycle of animal housing room: 07:00-19:00 (12 h)
		    ○ Time of day of injection: 2-3pm
		    ○ Room temperature at injection time: 20-21°C
		    ○ Did a cage change occur at time of injection: No

		c. Validation of immune activation – behavior, physiological indices and/or cytokine data, including pilot dosing data
		    ○ Method used to verify immune activation:
		
		Blood serum IL-17a cytokine levels were measured by ELISA from female mice 24 h after subcutaneous administration of either Poly IC or PBS control.
		
		d. Validation of gestational timing – vaginal plug, estrous cycle, weight gain
		    ○ Method of validating gestational timing:
		Weight gain: Dams were time mated for one night and weighed at E0 and then again at E11.
		
		Additional comments:
		N/A

Experimental animals
➢ Species/strain/vendor
 	
a. Provide details of the animals used, including species, strain, sex, developmental stage (e.g. mean or median age plus age range) and weight (e.g. mean or median weight plus weight range).
b. Provide further relevant information such as the source of animals, international strain nomenclature, genetic modification status (e.g. knock-out or transgenic), genotype, health/immune status, drug or test naïve, previous procedures, etc.
	8	Provide details of:
		a. Species – considerations for appropriate species (mouse, rat, non human primate, other)
		    ○ Species: mouse
		
		b. Strain – variability in strain can influence model
		    ○ Strain: C57BL/6JOlaHsd, Ai32(RCL-ChR2(H134R)/EYFP or *Il17a* KO
		
		c. Maternal/Offspring Physiological Variables at time of immune challenge – age, body weight
		    ○ Maternal Age at challenge: 8-12 weeks
		    ○ Maternal Body weight: 24-28g (dependent on pregnant dams litter size)
		    ○ Offspring Age at challenge: E12.5
		    ○ Offspring Sex: Both males and females tested
		    ○ Offspring Body weight: n/a
		
		d. Vendor – even within the same strain, vendor can influence endpoints
		    ○ Vendor: Jackson Labratory
		    ○ Location of Vendor: USA
		    ○ Room/area where animals originated from: Refer to comments

		Additional Comments: Mouse strains were obtained from JAX and rederived in to the facility. Each line was then bred within the facility and never underwent previous procedures.

Housing and husbandry
➤ Cage, ventilation, bedding, enrichment
 	
Provide details of:
a. Housing (type of facility e.g. specific pathogen free [SPF]; type of cage or housing; bedding material; number of cage companions; tank shape and material etc. for fish).
b. Husbandry conditions (e.g. breeding program, light/dark cycle, temperature, quality of water etc for fish, type of food, access to food and water, environmental enrichment).
c. Welfare-related assessments and interventions that were carried out prior to, during, or after the experiment.
	9	Provide details of:
		a. Caging systems
		    ○ *At breeding*
		        Material of cage: GM500 cages
		        Cage dimensions: Floor area: 501 cm2
		    ○ *After parturition*
		        Material of cage: GM500 cages
		        Cage dimensions: Floor area: 501 cm2
		    ○ *At weaning*
		        Material of cage: GM500 cages
		        Cage dimensions: Floor area: 501 cm2
		
		b. Animal Holding room
		    ○ Temperature in room: 20-21°C
		    ○ Humidity in room: 50-55%
		    ○ Ventilation system: IVC ventilation
		    ○ Specific pathogen free [SPF]: YES
		    ○ Are males & females housed in the same or separate rooms:
		        Housed in same room after weaning
		c. Bedding exchanges/bedding type
		    ○ Type of cage bedding used: SAFE® Select Bedding Rettenmaier
		    ○ Frequency of cage changes per week
		        during gestation: 1
		        during neonatal period: 0 in 1st week, then every second week
		        following weaning: every second week
		
		d. Breeding - bred on site or timed pregnant, how many different sires (are the same fathers breeding with both experimental and control dams)
		    Breeding location: on site

		○ Gestational age at shipping: N/A
		    ○ Biological age of dams (if not listed in Section 8c): 8-12 weeks
		    ○ Number of Dams bred: 160
		    ○ How many times have dams been mated previously: 0 (virgins used)
		    ○ How many times did the dams mate and not become pregnant: 1-2
		    ○ Are the dams primiparous or multiparous? All dams are multiparous
		    ○ What was the frequency of maternal handling during the gestational/neonatal period (e.g. cage cleanings, weighing, blood collection manipulations): weighing at E11, change cage at E13
		    ○ Biological age of sires: 7-12
		    ○ Number of sires bred: 55
		    ○ How many times have sires been mated previously: 2-3
		    ○ How many times did the sires mate successfully (e.g. mating resulted in pregnancy, full term birth): 2-3
		    ○ If bred previously, what was the interval between mating times:
		    ○ Are sires matched to experimental and control dams: No
		    ○ Describe the mating design (1:1, 1:2 etc): paired 1:1 for one night
		
		e. Social enrichment – number of cage companions
		    ○ Number of cage companions prior to breeding: 1-5
		    ○ Gestational age when dam separated for parturition: E13
		    ○ Number of cage companions at weaning: 1-5
		
		f. Physical enrichment – describe enrichment devices, and when enrichment is in the cage (removed when pups born? Or present throughout study), does the enrichment type change? How frequently?
		    ○ Describe what type of enrichment devices (and how many) are included in cage/housing room:
		
		Enrichment stays consistent throughout experiment. Each cage contains one tunnel tube and bedding.

		○ Does enrichment type/access change across study? No ○ If so, when does enrichment type/access change (e.g. enrichment removed prior to parturition and replaced in late neonatal period): N/A Additional Comments: Only virgin dams were used for mating purposes and were not mated until 8 weeks of age to ensure survival of the litter. Dams were time mated 1:1 for one night and then placed back in their homecage. Dams were housed with cage mates until E13 where they were transfered to a new cage with one other dam (cage mate). It was important to ensure that the cage mate had not been time mated to prevent discrepancies with P0 of the litter.

Sample size
➢ Litter versus offspring
 	
a. Specify the total number of animals used in each experiment, and the number of animals in each experimental group.
b. Explain how the number of animals was arrived at. Provide details of any sample size calculation used.
c. Indicate the number of independent replications of each experiment, if relevant.
	10	Provide details of:
		a. Maternal N vs offspring N
		    ○ What is the total number of dams/litters included in the study: 160
		    ○ What is the total number of offspring per litter included the study: 2-6
		b. Litter size and sex distribution
		    ○ What size was each litter maintained at: 6
		    ○ What age did culling take place at: P10-P12
		    ○ How many males and females were maintained in each litter: N/A
		c. Cross fostering
		    ○ Did cross fostering occur: No
		    ○ If so, at what age did cross fostering occur: N/A
		
		Additional Comments:

Allocating animals to experimental groups
 	
a. Give full details of how animals were allocated to experimental groups, including randomization or matching if done.
b. Describe the order in which the animals in the different experimental groups were treated and assessed.
	11	a. How many offspring per litter were used in each measure: refer to S.Table 1
		
		b. Randomization/Matching procedures
		    ○ What procedures were used to assign animals to groups:
		Litters from damns that were injected on the same day were counter balanced across groups.
		
		c. Sex as a biological variable (behavioral and physiological outcomes)
		    ○ Were both males and females evaluated in each behavioral and physiological outcome: Yes
		
		Additional Comments:

Experimental outcomes
➢ Behavioral testing
➢ Physiological endpoints
 	
Clearly define the primary and secondary experimental outcomes assessed (e.g. cell death, molecular markers, behavioral changes).
	12	a. Maternal behavior and pup interactions
		    ○ If maternal care was evaluated, were there differences following immunogen challenge (if so, please briefly describe):
		N/A
		
		b. Age(s) of offspring at behavioral testing/physiological evaluation endpoints:
		CFC: P17 and recall at P18, P25 or P32 ; Social interaction and marble burying: P63-P65 Engram labeling: P17 ; Perfusions P25
		
		c. Order of testing (e.g. behavioral test order)
		    ○ Were animals evaluated in a counter-balanced order in terms of:
		      presentation of tests to each animal: Yes
		      order of experimental/control groups run through each test: Yes
		    ○ What was the inter-test interval if a single animal underwent a battery of tests:
		38 days between CFC and sociability test; 1 day in between sociability and marble burying tests.

		Additional Comments:
Statistical methods a. Provide details of the statistical methods used for each analysis. b. Specify the unit of analysis for each dataset (e.g. single animal, group of animals, single neuron). c. Describe any methods used to assess whether the data met the assumptions of the statistical approach.	13	a. Unit of analysis for each data set ○ Is the unit (n) of each analysis based on number of litters, or number of animals used per group: For behavioral analysis, results were analyzed per litter using nested t-test or nested ANOVA followed by Bonferroni post hoc test. For cell counting analysis results were analyzed per mouse using nested t-test or nested ANOVA followed by Bonferroni post hoc test. Results were considered significant if $P<0.05$.
Other Disclosures		Please make note of any other extraneous variables that you would like to report (e.g. fire alarms, construction, temporary relocations, other variables that you think we should be considering in our studies etc.): N/A

Kentner, A. C., Bilbo, S. D., Brown, A. S., Hsiao, E. Y., McAllister, A. K., Meyer, U., Pearce, B. D., Pletnikov, M. V., Yolken, R. H., & Bauman, M. D. (2019). Maternal immune activation: reporting guidelines to improve the rigor, reproducibility, and transparency of the model. *Neuropsychopharmacology : official publication of the American College of Neuropsychopharmacology*, *44*(2), 245–258. https://doi.org/10.1038/s41386-018-0185-7

Table S2. Maternal immune activation model reporting guidelines checklist.