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Engrams and Circuits Crucial for Systems Consolidation of a Memory

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Abstract

Episodic memories initially require rapid synaptic plasticity within the hippocampus for their formation and are gradually consolidated in neocortical networks for permanent storage. However, the engrams and circuits that support neocortical memory consolidation remain unknown. We found that neocortical prefrontal memory engram cells, critical for remote contextual fear memory, were rapidly generated during initial learning via inputs from both hippocampal-entorhinal cortex and basolateral amygdala. After their generation, the prefrontal engram cells, with support from hippocampal memory engram cells, became functionally mature with time. Whereas hippocampal engram cells gradually became silent with time, engram cells in the basolateral amygdala, which were necessary for fear memory, are maintained. Our data provide new insights into the functional reorganization of engrams and circuits underlying systems consolidation of memory.

> Memories are thought to be initially stored within the hippocampal-entorhinal cortex (HPC-EC) (recent memory) and over-time are slowly consolidated within the neocortex for permanent storage (remote memory) (1-7). Systems memory consolidation models suggest that the interaction between the HPC-EC and the neocortex during and after an experience is crucial (8–12). Experimentally, prolonged inhibition of hippocampal or neocortical networks during the consolidation period produces deficits in remote memory formation (13–15). However, little is known regarding specific neural circuit mechanisms underlying the formation and maturation of neocortical memories through interactions with the HPC-EC network. By employing activity-dependent cell labeling technology (16-18) combined with viral vector-based transgenic, anatomical (19, 20), and optogenetic strategies (19, 21) for

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SUPPLEMENTARY MATERIALS Materials and Methods

Figs. S1 to S12 Statistics and Sample Sizes

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circuit-specific manipulations and *in vivo* calcium imaging (22), we investigated the nature and dynamics of neocortical and subcortical memory engram cells (a population of neurons that are activated by learning, have enduring cellular changes, and are reactivated by a part of the original stimuli for recall (18)) and their circuits for systems consolidation of memory.

We first traced entorhinal projections to frontal cortical structures (the medial prefrontal cortex (PFC), caudal anterior cingulate cortex (cACC), retrosplenial cortex (RSC)) involved in contextual fear memory, and the basolateral amygdala (BLA), with injections of the retrograde tracer cholera toxin subunit B (CTB)-Alexa555 into these regions (fig. S1). CTB injections resulted in labeling in the medial entorhinal cortex (MEC) specifically in cells in layer Va (Fig. 1A–D, H and fig. S1A–D), indicating that MEC-Va cells have extensive projections to the neocortex and BLA (23). We then sought to inhibit these specific projections by bilaterally injecting adeno-associated virus 8 (AAV₈)-calcium/calmodulindependent protein kinase II (CaMKII):eArchT-enhanced yellow fluorescent protein (eYFP) in the deep layers of the MEC in wild-type (WT) mice with bilaterally implanted optic fibers above the PFC, cACC, or RSC (Fig. 1E, J, and fig. S2G). Expression of eArchT-eYFP was abundant in MEC-Va terminals located in the PFC, cACC and RSC (Fig. 1B, I, and fig. S2D). These mice were then subjected to contextual fear conditioning (CFC) while we delivered green light bilaterally to the different cortical areas receiving MEC-Va projections during either the conditioning period (Day-1) (fig. S2E) or recall test period (Day-2, Day-8, Day-15 and Day-22) (fig. S2F). Axon terminal inhibition with optogenetics of MEC-Va cells within the PFC during Day-1 of CFC disrupted memory at Day-15 and Day-22, but not at Day-2 or Day-8 (Fig. 1F). Terminal inhibition during memory recall tests did not affect memory retrieval (Fig. 1G). Finally, terminal inhibition in the cACC or RSC during CFC or recall had no effect on memory throughout these periods (Fig. 1J-L and fig. S2G-I).

The above results suggest that MEC-Va input into the PFC during CFC is crucial for the eventual formation of remote memory. This hypothesis was supported by several findings. First, CFC increased the number of c-Fos⁺ cells in the PFC compared to that of homecage mice (Fig. 1M–O), whereas context only exposure did not increase c-Fos activity in the PFC (Fig. 10). Second, optogenetic terminal inhibition of MEC-Va projections within the PFC during CFC inhibited the observed increase of c-Fos⁺ cells in the PFC (Fig. 10). Finally, we identified CFC engram cells in the PFC. We targeted injections of AAV₉-c-fos:tTA and AAV₉-TRE:channelrhodopsin-2 (ChR2)-mCherry (Fig. 1P, Q) and optic fibres to the PFC of WT mice and labeled the PFC cells activated by CFC with ChR2 while the mouse was off doxycycline (OFF-Dox) (Fig. 1R). Blue light stimulation at 4 Hz, but not the conventional 20 Hz, of ChR2-mCherry-expressing cells in the PFC induced freezing behavior on Day-2 and Day-12 in an unconditioned context (Fig. 1S, fig. S3) compared with freezing under the blue light-off condition. This blue light-induced freezing was prevented when MEC-Va fibers in the PFC were inhibited during CFC on Day-1 (Fig. 1T, U, fig. S4). Using transsynaptic retrograde tracing combined with the activity-dependent cell labelling, we confirmed that the PFC engram cells generated by CFC receive monosynaptic input from MEC-Va cells (Fig. 1V–X, fig. S5).

To examine whether PFC engram cells are also reactivated by the conditioned context (rather than by blue light) at a recent and remote time point, we targeted injections of AAV₉-TRE:

human histone H2B-green fluorescent protein (H2B-GFP) to the PFC of c-fos:tTA transgenic mice (Fig. 2A). The mice underwent CFC on Day-1 and then re-exposed to the conditioned (Context-A) or an unconditioned (Context-B) context on Day-2 or Day-13 (Fig. 2B). Cells activated by CFC were labeled with H2B-GFP and the cells activated by the context test with a c-Fos antibody, and we calculated the proportion of double labeled cells (Fig. 2A–B, fig. S6B). H2B-GFP⁺ cells (PFC engram cells) were preferentially reactivated in Context-A on Day-13 but not Day-2, compared to the H2B-GFP negative cells (Fig. 2C). There was no difference in c-Fos expression between H2B-GFP positive and negative cells when mice were tested in Context-B (Fig. 2C). We also found that the spine density of the PFC engram cells on Day-12 was significantly higher than on Day-2 (Fig. 2D, E, fig. S7), in line with previous findings of a positive correlation between dendritic spine density on memory engram cells and memory expression triggered by natural recall cues (24–26).

To test whether PFC engram cells are necessary for memory recall by natural cues, we bilaterally targeted injections of AAV₉-c-fos:tTA and AAV₉-TRE:ArchT-eGFP (Fig. 2D, F) and optic fibres to the PFC of WT mice and labeled the PFC engram cells that were activated by CFC with ArchT while the mice were OFF-Dox (Fig. 2F). Cell body inhibition of the PFC engram cells by green light during retrieval did not affect recent memory (Day-2), however at the remote time-point (Day-12) memory retrieval was disrupted compared to the green light-off condition (Fig. 2F).

To further investigate the characteristics of PFC engram cells we monitored transient calcium (Ca²⁺) events in PFC cells *in vivo*. WT mice were injected with AAV₅-Syn:GCaMP6f into the PFC and implanted with a micro GRIN lens targeting the PFC (Fig. 2G-I, fig. S8) (22, 27). On Day-1 mice were first exposed to Context-B followed by CFC in Context-A. Mice were then re-exposed to both contexts in the same order on Day-2 and Day-15 (Fig. 2J). The averaged frequency of Ca^{2+} events in PFC cells did not significantly change in either a time- or context-dependent manner (fig. S9B). However a small but significant difference was revealed in the cumulative distribution curves of a rate difference index (assessing context selectivity; see Methods) between Day-1 conditioning and Day-15 recall, and between Day-2 recall and Day-15 recall (fig. S9C). PFC cells did not appear to discriminate between the two contexts on Day-1 prior to footshock presentation (Fig. 2K, L). However, after footshock presentation, about 11% of cells showed a significant increase in Ca²⁺ transients (shock responding cells; SR cells) (Fig. 2K, L). The remaining ~89% of PFC cells did not respond to the shocks (shock non-responding cells; SNR cells). The SR cells were less active than SNR cells during exposure to Context-B and Context-A on Day-1 prior to footshock presentation (Fig. 2L-N, fig. S9D). During recall, the transient Ca²⁺ activity of SR cells in Context-A (conditioned context) was significantly higher compared to that in Context-B on Day-15 but not on Day-1 or Day-2, whereas the frequency of Ca²⁺ transient events in SNR cells remained constant, irrespective of context (Fig. 2M, N). This produced a significant rate difference index of Ca²⁺ activity for the conditioned context between the SR and SNR cells on Day-15 but not on Day-1 (excluding the shock delivery period) or Day-2 (Fig. 2O). These results combined with c-Fos activation data (Fig. 1M–O) suggest that the SR cells may be the PFC memory engram cells, given that the generation of the PFC engram cells requires both context exposure and footshocks.

Our calcium imaging data suggest that footshock stimulus input into the PFC is crucial for the generation of PFC engram cells. Because the BLA integrates footshock information arriving from the thalamus (28) and projects to the PFC (fig. S5I, S10), we optogenetically inhibited the pathway from BLA to PFC during CFC (Fig. 2P). Optogenetic inhibition of BLA terminals in the PFC during CFC disrupted the generation of PFC engram cells (Fig. 2Q). The terminal inhibition during CFC also inhibited remote memory formation (Fig. 2R).

To test whether the HPC engram cells play a crucial role in the functional maturation of PFC engram cells during the system consolidation process, we bilaterally targeted injection of AAV9-TRE:tetanus toxin light chain (TeTX) or AAV9-TRE:eYFP (as a control) to the hippocampal dentate gyrus (DG) of c-fos:tTA transgenic mice (Fig. 3A). When the mice were subjected to CFC, DG engram cells were labeled with TeTX. DG engram cell labeling with TeTX caused a robust inhibition of DG engram cell output, as revealed by greatly reduced immunoreactivity of vesicle-associated membrane protein 2 (VAMP2), which is essential for activity-dependent neurotransmitter release from presynaptic terminals (13), within the stratum lucidum (s.l.) in hippocampal CA3 in OFF-Dox mice compared to ON-Dox mice (Fig. 3B, C). Optogenetic activation of DG engram cells with ChR2 failed to produce the increase in c-Fos⁺ cells in CA3 of TeTX-expressing mice that was observed in control mice compared to home cage controls (Fig. 3D). TeTX expression in HPC engram cells inhibited the reactivation of PFC engram cells during exposure to the conditioned context 12 days after CFC compared to the eYFP control group (Fig. 3E, F). TeTX expression also blocked the increase of dendritic spine density of PFC engram cells compared to the eYFP group (Fig. 3G). In vivo calcium imaging revealed that TeTX expression in HPC engram cells after CFC blocked the increase of the context discrimination index in SR cells in the PFC (Fig. 3H, fig. S11).

To investigate the post-consolidation fate of HPC engram cells we crossed c-fos:tTA TG mice with TRE:H2B-GFP TG mice (29), subjected them to CFC, and then re-exposed them to the conditioned (Context-A) or an unconditioned (Context-B) context on Day-2 or Day-13 (Fig. 3I–K). DG engram cells were preferentially reactivated in Context-A on Day-2 but not Day-13, compared to the non-engram cells (Fig. 3L). No difference was observed in the activation of DG engram and non-engram cells by Context-B (Fig. 3L). We were unable to maintain labeled DG engram cells with ChR2 beyond 12 days with injection of AAV₉-TRE:ChR2-mCherry. To extend this technical limit we targeted injections of AAV_{1,5,8,9}-TRE:CCre, AAV_{1,5,8,9}-TRE:NCre and AAV₅-EF1a:ChR2-mCherry to the DG of c-fos:tTA transgenic mice (Fig. 3M). We could thus extend viable labelling by a few days (Fig. 3N). The spine density of DG engram cells on Day-15 was significantly reduced compared to Day-5 (Fig. 30, fig. S12). On both Day-5 and Day-15, optogenetic activation of DG engram cells induced freezing behaviour (Fig. 3P, Q).

We finally investigated the role of MEC-Va projections to the BLA in recent and remote memory (Fig. 4A, fig. S2A). Inhibition of MEC-Va terminals in the BLA during CFC disrupted contextual fear memory formation. Retrieval was impaired at all time-points tested (Fig. 4B). When terminal inhibition was restricted to retrieval, recent memory tested on Day-2 and Day-8 was impaired, but remote memory retrieval on Day-15 and Day-22 was unaffected (Fig. 3C). In contrast, inhibition of PFC engram cell terminals in the BLA did not

impair memory retrieval on Day-2 but did impair memory retrieval on Day-12 (Fig. 4D, E). To investigate whether the BLA fear memory engram cells formed on Day-1 are maintained and used for PFC engram-dependent remote memory recall, we subjected the double transgenic mice (Fig. 4F, I) to CFC and re-exposed them to the conditioned context at recent or remote time-points (Fig. 4G). BLA engram cells were reactivated equally well by the conditioned context at both recent and remote time-points (Fig. 4H). Similarly, BLA cells activated by recent recall were reactivated again equally well by re-exposure to the conditioned context at recent and remote time-points (Fig. 4J, K).

Here, we found that PFC memory engram cells for CFC were rapidly formed during Day-1 training by virtue of inputs from both MEC-Va and BLA, but they were not retrievable with natural recall cues. The immature PFC engram cells functionally, structurally, and physiologically matured during the subsequent few weeks and this process required inputs from HPC engram cells presumably through MEC-Va. In contrast to their formation on Day-1, retrieval of the PFC engram at a remote time did not require MEC-Va input. HPC engram cells formed during training became silent with time; they are not retrieved on Day-14 by natural recall cues but are still re-activatable optogenetically for recall. However, fear memory BLA engrams formed during training are functionally maintained even after the consolidation-mediated switch in recall circuits (Fig. 4L).

Our model (Fig. 4L) introduces the concept that the prefrontal memory engram is already generated, albeit in an immature form, on Day-1 of training via inputs from both HPC-EC and BLA (Fig. 1). The standard model (1, 2, 4, 6, 7, 11) hypothesized that remote memory is formed in the cortex by a slow transfer of hippocampal memory. In contrast, in our study the role of the hippocampus in cortical memory is for the rapid generation of immature engram cells in the PFC during training and for the subsequent functional maturation of these preexisting engram cells (Fig. 3). The immature PFC engram may correspond to the cortical "tagging" speculated in an earlier study (14). Whereas in a previous study the BLA is crucial for both recent and remote fear memory expression (30), our results demonstrated an overlapping set of BLA engram cells for both recent and remote fear memory retrieval which were quickly formed during training (Fig. 4). However, the source of input into the BLA engrams for retrieval shifts from MEC-Va at recent time-points to the PFC engram at remote time-points (Fig. 4L). The route through which contextual stimuli activate the mature PFC engram is unknown. Most likely the information processed in a variety of sensory cortices reaches the PFC via the thalamus (31). Supporting this idea, PFC engram cells receive monosynaptic input from both the medial-dorsal and anteromedial thalamus (fig. S5).

Our finding of the lasting hippocampal engrams (Fig. 3Q) is consistent with multiple trace theory (5, 11). However, at the post-consolidation stage, the hippocampal engrams were not activatable by natural recall cues, but with optogenetic stimulation. A similar state of hippocampal engrams has previously been observed in anisomycin-induced amnesia (24) and in mouse models of early Alzheimer's disease (26), and the early (Day-2) PFC engram cells showed a similar property (Fig. 1S, 2C). Although we did not determine how long after encoding this "silent state" of the hippocampal engram lasts, we speculate that the hippocampal engram will eventually lose the original memory information (29, 32, 33).

Alternatively, the silent engram cells may still participate in the successful remote recall of discrete episodic details (5, 11).

As in previous studies (18, 20, 29), we observed that training resulted in wide-spread neuronal activation in the neocortex, including the ACC and RSC. However, whereas the activation of PFC neurons is crucial for formation of remote memory, MEC-Va input into the cACC or RSC is dispensable for this process. For remote memory, the PFC may thus have a distinctive role compared to other neocortical areas in integrating multiple sensory information stored in various cortical areas (11). Finally, our data show that the remote memory expressed by the PFC engram is conditioned-context specific, suggesting that it is episodic-like.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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References and Notes

- 1. Marr D. Philos Trans R Soc Lond B Biol Sci. Jul 1.1971 262:23. [PubMed: 4399412]
- 2. Squire LR. Science. Jun 27.1986 232:1612. [PubMed: 3086978]
- 3. Kim JJ, Fanselow MS. Science. May 1.1992 256:675. [PubMed: 1585183]
- 4. McClelland JL, McNaughton BL, O'Reilly RC. Psychol Rev. Jul. 1995 102:419. [PubMed: 7624455]
- 5. Nadel L, Moscovitch M. Curr Opin Neurobiol. Apr.1997 7:217. [PubMed: 9142752]
- 6. Tse D, et al. Science. Apr 06.2007 316:76. [PubMed: 17412951]
- 7. McClelland JL. J Exp Psychol Gen. Nov.2013 142:1190. [PubMed: 23978185]
- 8. Buzsaki G. Cereb Cortex. Mar-Apr;1996 6:81. [PubMed: 8670641]
- 9. Siapas AG, Wilson MA. Neuron. Nov.1998 21:1123. [PubMed: 9856467]
- 10. Wiltgen BJ, Brown RA, Talton LE, Silva AJ. Neuron. Sep 30.2004 44:101. [PubMed: 15450163]
- 11. Frankland PW, Bontempi B. Nat Rev Neurosci. Feb.2005 6:119. [PubMed: 15685217]
- 12. Preston AR, Eichenbaum H. Curr Biol. Sep 9.2013 23:R764. [PubMed: 24028960]
- Nakashiba T, Buhl DL, McHugh TJ, Tonegawa S. Neuron. Jun 25.2009 62:781. [PubMed: 19555647]
- 14. Lesburgueres E, et al. Science. Feb 18.2011 331:924. [PubMed: 21330548]
- 15. Zelikowsky M, Bissiere S, Fanselow MS. J Neurosci. Mar 07.2012 32:3393. [PubMed: 22399761]
- Reijmers LG, Perkins BL, Matsuo N, Mayford M. Science. Aug 31.2007 317:1230. [PubMed: 17761885]
- 17. Liu X, et al. Nature. Mar 22.2012 484:381. [PubMed: 22441246]
- 18. Tonegawa S, Liu X, Ramirez S, Redondo R. Neuron. Sep 2.2015 87:918. [PubMed: 26335640]
- 19. Kitamura T, et al. Science. Feb 21.2014 343:896. [PubMed: 24457215]
- 20. Ye L, et al. Cell. Jun 16.2016 165:1776. [PubMed: 27238022]

- 21. Deisseroth K. Nat Neurosci. Sep.2015 18:1213. [PubMed: 26308982]
- 22. Ziv Y, et al. Nat Neurosci. Mar.2013 16:264. [PubMed: 23396101]
- 23. Surmeli G, et al. Neuron. Dec 2.2015 88:1040. [PubMed: 26606996]
- Ryan TJ, Roy DS, Pignatelli M, Arons A, Tonegawa S. Science. May 29.2015 348:1007. [PubMed: 26023136]
- 25. Hayashi-Takagi A, et al. Nature. Sep 17.2015 525:333. [PubMed: 26352471]
- 26. Roy DS, et al. Nature. Mar 24.2016 531:508. [PubMed: 26982728]
- 27. Kitamura T, et al. Neuron. Sep 23.2015 87:1317. [PubMed: 26402611]
- 28. Pellman BA, Kim JJ. Trends Neurosci. Jun.2016 39:420. [PubMed: 27130660]
- Tayler KK, Tanaka KZ, Reijmers LG, Wiltgen BJ. Curr Biol. Jan 21.2013 23:99. [PubMed: 23246402]
- 30. Maren S, Aharonov G, Fanselow MS. Behav Neurosci. Aug.1996 110:718. [PubMed: 8864263]
- Do-Monte FH, Quinones-Laracuente K, Quirk GJ. Nature. Mar 26.2015 519:460. [PubMed: 25600268]
- 32. Denny CA, et al. Neuron. Jul 02.2014 83:189. [PubMed: 24991962]
- 33. Kitamura T, et al. Cell. Nov 13.2009 139:814. [PubMed: 19914173]
- 34. Tumbar T, et al. Science. Jan 16.2004 303:359. [PubMed: 14671312]
- Okuyama T, Kitamura T, Roy DS, Itohara S, Tonegawa S. Science. Sep 30.2016 353:1536. [PubMed: 27708103]
- 36. Miyamichi K, et al. Nature. Apr 14.2011 472:191. [PubMed: 21179085]
- 37. Ramirez S, et al. Science. Jul 26.2013 341:387. [PubMed: 23888038]
- 38. Ramirez S, et al. Nature. Jun 18.2015 522:335. [PubMed: 26085274]
- 39. Yu CR, et al. Neuron. May 27.2004 42:553. [PubMed: 15157418]
- 40. Hirrlinger J, et al. PLoS One. 2009; 4:e4286. [PubMed: 19172189]
- 41. Shima Y, et al. Elife. Mar 21.2016 5:e13503. [PubMed: 26999799]
- Wickersham IR, Finke S, Conzelmann KK, Callaway EM. Nat Methods. Jan.2007 4:47. [PubMed: 17179932]
- 43. Kohara K, et al. Nat Neurosci. Feb.2014 17:269. [PubMed: 24336151]
- 44. Goosens KA, Maren S. Learn Mem. May-Jun;2001 8:148. [PubMed: 11390634]
- 45. Kim J, Pignatelli M, Xu S, Itohara S, Tonegawa S. Nat Neurosci. Oct 17.2016
- 46. Kitamura T, Macdonald CJ, Tonegawa S. Learn Mem. Sep.2015 22:438. [PubMed: 26286654]
- 47. Yokoyama M, Matsuo N. Front Behav Neurosci. 2016; 10:218. [PubMed: 27872586]
- 48. Redondo RL, et al. Nature. Sep 18.2014 513:426. [PubMed: 25162525]
- Restivo L, Vetere G, Bontempi B, Ammassari-Teule M. J Neurosci. Jun 24.2009 29:8206. [PubMed: 19553460]
- 50. Pinto L, Dan Y. Neuron. Jul 15.2015 87:437. [PubMed: 26143660]
- 51. Sun C, et al. Proc Natl Acad Sci U S A. Jul 28.2015 112:9466. [PubMed: 26170279]

One Sentence Summary

We discovered the neocortical remote memory engram cells, and the neural circuit mechanisms responsible for their generation and maturation.

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Fig. 1. MECVa input to PFC during conditioning is crucial for generation of PFC engram cells (A) CTB injection into PFC. Sagittal section of MEC with CTB-labeled cells (red) (B) Coronal sections of PFC with MECVa axons expressing eYFP (green). rACC; rostral ACC, PL; prelimbic cortex. (C, D) Sagittal section of MEC with CTB-labeled cells (red) and immunostained with anti-PCP4 (green) and anti-NeuN (blue). PCP4 is a marker for layer III and Vb cells in MEC. CTB injection into BLA. (E, J) Viral injections and optic fiber implantations. (F, G) Time courses of freezing during recall tests. Green light was shone into the PFC during conditioning (F) or testing (G). N presents number of animals. (H) CTB injection into caudal ACC (cACC). Sagittal section of MEC with CTB-labeled cells (red). (I) Coronal sections of ACC with MECVa axons (green). (K, L) Time courses of freezing during recall tests. Green light was shone into the ACC during conditioning (K) or testing (L). (M) Experimental schedule. (N) Coronal section of PFC with anti-c-Fos (green). (O) Percentages of c-Fos⁺ cells in PFC of homecage (HC), context exposure (CTX), CFC with eYFP, and CFC with eArchT group. (P) Virus-mediated engram cell labelling with ChR2. (Q) Coronal section of PFC with ChR2-mCherry (red). (R, T) Experimental schedule. (S, U) Averaged freezing for light-off and light-on epochs. (V) Retrograde trans-synaptic labeling with activity-dependent cell labeling. (W) Sagittal section of MEC with rabies virus-specific mCherry (red). (X) Distribution of mCherry⁺ cells in the MEC (n=212 mCherry⁺ cells, n presents number of cells.). *P < 0.05 by unpaired t-test compared to eYFP (F, G, K, L), oneway ANOVA with Tukey-Kramer test (O) and paired t-test (S, U). Error bars mean \pm s.e.m.



Fig. 2. PFC engram cells mature with time

(A) PFC engram cell labelling with H2B-GFP. Coronal sections of PFC with H2B-GFP (green), anti-c-Fos (red). Marked cells are double-positive. (B) Experimental schedule. (C) Percentages of c-Fos⁺ cells in H2B-GFP⁺ and H2B-GFP⁻ cells in PFC. (D) PFC engram cell labelling with ArchT. (E) Images showing dendritic spines from PFC engram cells. Cumulative probability of the spine density of PFC engram. (F) Experimental schedule. Averaged freezing for light-off and light-on epoch during recall test. (G, H) Viral injections and GRIN lens implantation. (I) Stacked image acquired through the microendoscope over

10 mins of imaging in PFC. (J) Experimental schedule. (K) Raster plots of Ca^{2+} event in shock non-responding (SNR) cells and shock responding (SR) cells in PFC (showing 12 example cells). (L-N) Average Ca^{2+} event frequency of SNR cells and SR cells on Day-1, Day-2 and Day-3. (O) Average rate difference index of Ca^{2+} activity. (P) Viral injections and optic fiber implantations. Coronal sections of PFC visualizing BLA axons (green). (Q) Percentages of c-Fos⁺ cells in PFC of homecage (HC), shock only (Shock), CFC with eYFP, and CFC with eArchT, groups. (R) Time courses of freezing during recall tests. *P < 0.05 by unpaired t-test (C, O, R), KS test (E), paired t-test (F, M, N), one-way ANOVA with Tukey-Kramer test (Q). Error bars mean \pm s.e.m.



Fig. 3. HPC engram cells support the maturation of PFC engram cells, while HPC engram cells become silent with time

(A) DG engram cell labelling with TeTX. (**B**, **C**) Sagittal sections of HPC with anti-VAMP2 (red). (**D**) Percentages of c-Fos⁺ cells in hippocampal CA3 of homecage, blue-light-ON mice with eYFP or with TeTX. (**E**) Experimental schedule. (**F**) Percentages of c-Fos⁺ cells in H2B-GFP⁺ and H2B-GFP⁻ cells in the PFC of eYFP- and TeTX-expressing mice. (**G**) Images showing dendritic spines from PFC engram. Cumulative probability of the spine density of PFC engram of eYFP- and TeTX-expressing mice. (**H**) Experimental schedule, average Ca²⁺ event frequency of SNR cells and SR cells under TeTX-expressing condition. (**I**) Transgenic strategy of DG engram cell labelling with H2B-GFP. (**J**) Coronal sections of DG with H2B-GFP (green), anti-c-Fos (red). Marked cells are double-positive. (**K**, **P**) Experimental schedule. (**L**) Percentages of c-Fos⁺ cells in H2B-GFP⁺ and H2B-GFP⁻ cells in the DG. (**M**) Long-term DG engram cell labelling with ChR2. (**N**) Coronal sections of DG with ChR2-mCherry (red). (**O**) Images showing dendritic spines from DG engram. Cumulative probability of the spine density of DG engram. (**Q**) Averaged freezing by blue light stimulation for light-off and light-on epochs. *P < 0.05 by one-way ANOVA with

Tukey-Kramer test (D), unpaired t-test (F, L), KS test (G, O), paired t-test (H, Q). Error bars mean \pm s.e.m.



Fig. 4. BLA engram cells are maintained throughout consolidation but with a switch of the recall circuit

(A) Viral injections and optic fiber implantations. Coronal sections of BLA with MECVa axons expressing eYFP (green). (**B**, **C**) Time courses of freezing during recall tests. Green light was shone into the BLA during conditioning (**B**) or testing periods (**C**). (**D**) Viral injections and optic fiber implantations. Coronal sections of BLA visualizing axons of PFC engram cells (green). (**E**) Averaged freezing for green light-off and light-on epochs during recall test. (**F**, **I**) BLA engram cell labelling with H2B-GFP. (**G**, **J**) Experimental schedules. (**H**, **K**) Percentages of double labeling with c-Fos⁺ and H2B-GFP⁺ in the BLA compared to the calculated chance level. (**L**) A new model for systems consolidation of memory. *P < 0.05 by unpaired t-test (**B**, **C**, **H**, **K**) or by paired t-test (**E**). Error bars mean \pm s.e.m.