Brain-wide mapping reveals that engrams for a single memory are distributed across multiple brain regions

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01.

Important Terms

Important Terms

cFOS+

regulate genes that are switched on or repressed depending on influences from external stimuli.

Memory Engram

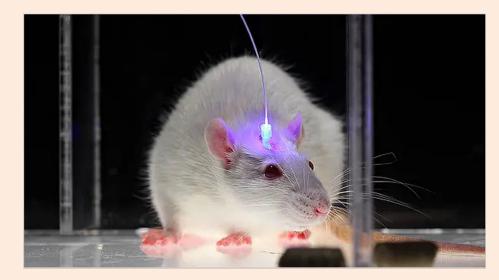
the enduring physical or chemical changes that occur in brain networks upon learning representing acquired memory information

4-hydroxytamoxifen (4-OHT)

4-OHT is a compound that can specific activate these ERT2 receptors (estrogen receptor) and initiate the express of the Credependent tdTomato stain.

Optogenetics

Scientists genetically engineer neurons to produce lightsensitive proteins known as opsins. Opsins permit the transmembrane movement of ions (causing an action potential) so scientists are able to use light to control specific neuron firing.



O2. Previous Research

Previous Studies

Previously Identified Engram Regions

- Hippocampal subfields
- Amygdala subregions
- Retrosplenial cortex
- Prefrontal cortex

"Indeed, early limited studies supported [Semon's engram concept] for contextual fear memory. While these findings enhanced our understanding of engram-based memory storage, a thorough mapping of a unified engram complex for a specific memory has been a challenging endeavor."

Previous Studies

- Optogenetic stimulation of hippocampal engram activates fear memory recall.
 - [PMC free article] [PubMed] [Google Scholar]
- Wiring and molecular features of prefrontal ensembles representing distinct experiences.
 - [PMC free article] [PubMed] [Google Scholar]
- Distinct neural circuits for the formation and retrieval of episodic memories.
 - [PMC free article] [PubMed] [Google Scholar]
- Memory retrieval by activating engram cells in mouse models of early Alzheimer's disease.
 - <u>PMC free article</u>] [PubMed] [Google Scholar]
- Bidirectional switch of the valence associated with hippocampal contextual memory engram.
 - [PMC free article] [PubMed] [Google Scholar]

03. Hypothesis

Hypothesis

A specific memory is stored in functionally connected engram cell ensembles that are widely distributed across multiple brain regions and are activated by natural recall cues. This is not an original hypothesis but part of Richard Semon and Donald Hebb's theory of synaptic plasticity.



Evidence Supporting Hypothesis:

- Gene expression is altered by widespread, behaviorally-defined neural circuits.
- Distributed brain regions involved in memory formation have been identified.
- Engram cells have been identified in hippocampal subfields and the amygdala.

O4. Test Groups

Test Groups

Group 1: Home Cage Group

Mice that received 4-hydroxytamoxifen (4-OHT)

and stayed in their home cage

Group 2: CFC Group

Mice that received 4-OHT and contextual fear

training and then went back into their home cage

Group 3: Recall group

Mice that received 4-OHT and contextual fear

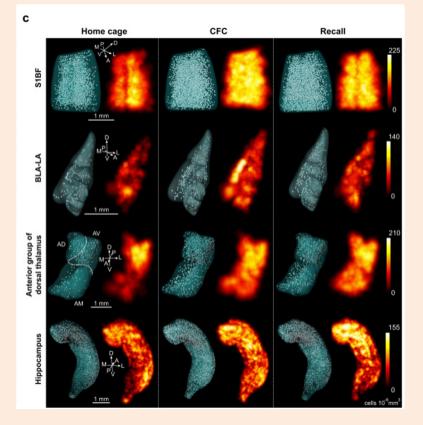
training and then after CFC were tested for fear memory recall.



05. Methods

Methods

- The brains of the mice were processed using the SHIELD technique to preserve endogenous tdTomato fluorescence and the tissue structure.
- They created a list of engram-containing regions of the brain. From this data they were able to form an engram index.
- To verify their engram index, the scientists found the overlap between tdTomato and recall-activated cFos+ neurons.
- They used optogenetics (control neurons through light) to test memory recall.
 - Dorsal CA1 and BLA engram cells were used as positive controls because they show significant memory recall when optogenetically activated.





New Engram Regions Discovered

- LDTg emotional arousal under adverse conditions
- Anterior hypothalamic nucleus (AHN) expression of conditioned fear behaviors
- RE Thalamus contextual memories, specifically discerning similar environments (supports the idea that thalamic ensembles don't solely play a passive relay role)
- Ventrolateral orbital area (VLO) long-term memory retention
- AM thalamus context-specific, high valence memories
- Periaqueductal gray (PAG) freezing behavior
- Paraventricular (PVT) Thalamus arousal, stress, emotional memory, and motivation.

*Note: many of these functions are just possibilities and have not been fully confirmed yet.

Engram Index

The scientists devised an engram index that ranks the possible engram cell complexes. This index is signified by two parts:

- Calculation of the average number of cFos+ neurons during memory encoding as compared to home cage.
- 2. Calculation of difference between activation levels in memory encoding and recall.

	(10 -0 1)			
ngram	index = $\log_{10} \left(\frac{ \bar{\mu}_{CFC} - \bar{\mu}_{HC} }{ \bar{\mu}_{CFC} - \bar{\mu}_{HC} } \right)$			
	$ \bar{\mu}_{CFC} - \bar{\mu}_{RE} $			
		с		
-		C	-	
Engram index			Engram index	
1.50	Midbrain reticular nucleus		0.34	Nucleus of reuniens
1.38	Laterodorsal tegmental nucleus	*	0.34	Posterior amygdalar nucleus
1.36	Presubiculum		0.33	Ventral posteromedial nucleus of the thalamus
1.33	Fields of Forel		0.32	Primary auditory area
1.25	Cuneiform nucleus		0.32	Lateral visual area
1.02	Basolateral amygdalar nucleus, anterior part		0.32	Posterior pretectal nucleus
0.97	Substantia innominata		0.31	Taenia tecta, ventral part
0.95	Nucleus of the lateral olfactory tract, body		0.31	Subparafascicular nucleus
0.95	Nucleus of the lateral olfactory tract, molecular layer		0.31	Parataenial nucleus
0.87	Medial pretectal area		0.30	Central amygdalar nucleus, medial part
0.84	Tegmental reticular nucleus		0.29	Ventral group of the dorsal thalamus
0.82	Nucleus of the lateral lemniscus		0.29	Globus pallidus, external segment
0.76	Dentate gyrus		0.28	Ventromedial hypothalamic nucleus
0.74	Ectorhinal area	*	0.28	Paraventricular nucleus of the thalamus
0.71	Tuberomammillary nucleus, ventral part		0.28	Cerebral nuclei
0.69	Olivary pretectal nucleus		0.26	Postrhinal area
0.65	Superior colliculus, deep white layer		0.26	Epithalamus
0.65	Paraventricular hypothalamic nucleus, descending division		0.26	Medial amygdalar nucleus
0.65	Supraoptic nucleus		0.26	Medial habenula
0.61	Basomedial amygdalar nucleus		0.26	Striatum Brain stem
0.56	Field CA3 Magnocellular reticular nucleus		0.25	Cortical subplate
0.55	Basolateral amygdalar nucleus		0.25	Tuberal nucleus
0.54	Claustrum	*	0.25	Anterior hypothalamic nucleus
0.52	Lateral habenula		0.25	Diagonal band nucleus
0.50	Tuberomammillary nucleus		0.25	Paraventricular hypothalamic nucleus
0.49	Field CA1	*	0.24	Orbital area, ventrolateral part
0.49	Frontal pole, layer 6a		0.24	Cortical amygdalar area, posterior part
0.48	Lateral hypothalamic area		0.24	Piriform area
0.47	Inferior colliculus		0.24	Infralimbic area
0.47	Posterior hypothalamic nucleus		0.24	Hypothalamic medial zone
0.46	Hippocampo-amygdalar transition area		0.23	Nucleus accumbens
0.45	Ventral premammillary nucleus		0.23	Basolateral amygdalar nucleus, ventral part
0.45	Subparafascicular nucleus, magnocellular part		0.23	Striatum-like amygdalar nuclei
0.45	Superior central nucleus raphe		0.23	Entorhinal area, lateral part
0.45	Interanterodorsal nucleus of the thalamus		0.22	Basomedial amygdalar nucleus, posterior part
0.45	Trapezoid body		0.22	Posterior parietal association areas
0.45	Pontine central gray		0.22	Fundus of striatum
0.44	Dorsal motor nucleus of the vagus nerve			Caudoputamen
0.43	Midbrain		0.21	Posterior intralaminar thalamic nucleus Dorsal peduncular area
0.43	Olfactory tubercle		0.21	Postsubiculum
0.42	Intercalated amygdalar nucleus		0.21	Prelimbic area
0.41	Mammillary body		0.20	Orbital area, medial part
0.41	Pedunculopontine nucleus		0.20	Orbital area, lateral part
0.41	Endopiriform nucleus, dorsal part Pallidum		0.19	Retrosplenial area, dorsal part
0.40	Periaqueductal gray		0.18	Agranular insular area, ventral part
0.39	Anteromedial thalamic nucleus		0.18	Retrohippocampal region
0.38	Piriform-amygdalar area		0.17	Dorsal cochlear nucleus
0.37	Lateral septal nucleus, ventral part		0.16	Thalamus
0.37	Prosubiculum		0.16	Anterior cingulate area, dorsal part
0.37	Magnocellular nucleus		0.16	Anterior cingulate area, ventral part
0.37	Temporal association areas		0.15	Perirhinal area
0.37	Endopiriform nucleus, ventral part		0.14	Entorhinal area, medial part
0.37	Medial septal nucleus		0.13	Thalamus, polymodal association cortex related
0.36	Basolateral amygdalar nucleus, posterior part		0.13	Midline group of the dorsal thalamus
0.36	Anteromedial visual area	*	0.13	Mediodorsal nucleus of thalamus
0.35	Interanteromedial nucleus of the thalamus			
0.35	Anterior amvodalar area			

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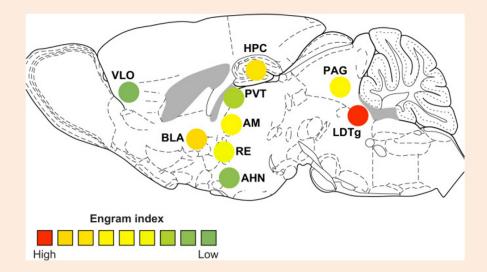
0.34

Anteroventral nucleus of thalamus (continued on top of panel c)

Engram Index

Brain regions that have an increase in activation levels during encoding as compared to home cage and don't have much difference between encoding and recall activation levels are placed highest on the engram scale.

To the scientists surprise, they found some regions with a higher recall activation than compared to memory encoding. They believe that this is because of some **unknown recall enhancing process.**



O7. Discussion

Summary

The engram index is based on the concept that engrams are held by neuronal ensembles that are activated by learning and are reactivated to support recall. The rain-wide engram mapping experiments revealed that certain brain regions exhibit higher recall activation as compared to CFC memory encoding. While there have been previous studies on brain mapping, this study permits a more complete map that shows the different roles that different parts of the brain play in memory and provides more support for the hypothesis.



Future Research

Future studies can, using the extensive brain mapping of this study, generate a more extensive map of engram cell ensembles that includes the identification of their functional connectivity as well as the mnemonic functions of individual ensembles.

Although this study provides evidence supporting the concept that a memory is stored in a functionally connected engram ensembles' complex distributed broadly across the brain, consistent with Semon's unified engram complex hypothesis, it would be interesting to see studies that either contradict, or prove a different hypothesis.

As mentioned earlier, the engram index was flawed so future research or replication could rectify that.



Thank You