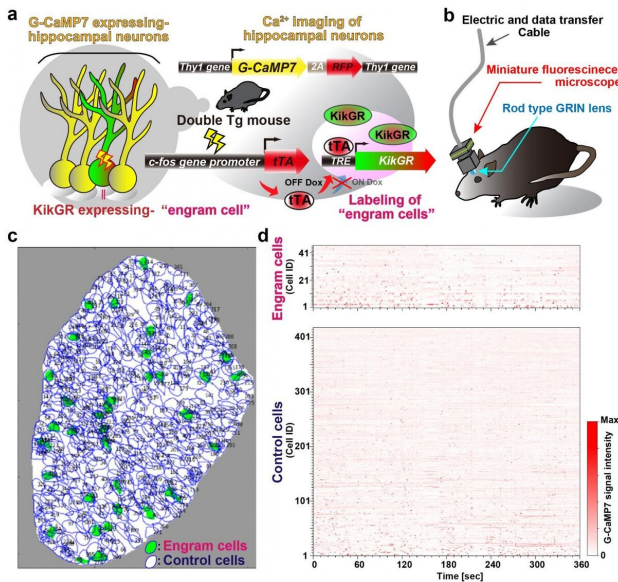


Discovery of neuronal ensemble activities orchestrated to represent one memory

5 September 2019



New optogenetic system to reveal neuronal activity of engram and non-ensemble by light. a. Engram cells can be specifically targeted in c-fos-tTA mice. Thy1-G-CaMP7 is useful for Ca²⁺ imaging of hippocampal CA1 neurons. Sander symbol, indicating neuronal activation state. b. Image of observation of fluorescence in mouse brain with miniature fluorescent microscope. c. Representative image of all engram cells and non-ensemble control cells detected from a single animal. d. Normalized calcium transients corresponding to neuronal activity for all engram cells (top) and non-ensemble cells (bottom) during experience of novel episode (displayed over time). Credit: University of Toyama

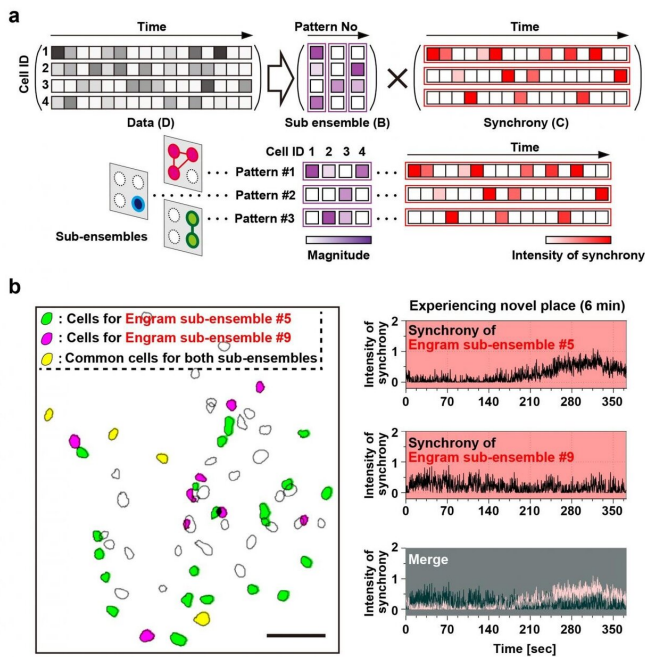
The brain stores memories through a neuronal ensemble, termed engram cells. A unique system was established to transfer neuronal population activity into light with discrimination between engram and non-ensemble cells using fluorescence proteins. By using this system, it is revealed that engram sub-ensembles represent distinct pieces of information, which are then orchestrated to constitute an entire memory. In addition, some sub-

ensembles preferentially reappear during post-learning sleep, and these replayed sub-ensembles are more likely to be reactivated during retrieval.

A Japanese research group supervised by Dr. Noriaki Ohkawa (Lecturer) and Professor Kaoru Inokuchi of the University of Toyama succeeded in establishing a system to investigate characteristic activity of cell ensembles acquiring memory and visualized ways for representation and consolidation of memory experienced novel episodes in the brain.

We are exposed to many episodic events and then memorize their information throughout our life. This kind of memory, episodic memory, is processed in several brain regions, and one of the regions is the hippocampus. It is believed that, in the hippocampus, one specific episodic memory is stored within and retrieved from a neuronal **ensemble** composed of **neurons**, termed engram cells, that are activated during learning. Indeed, activation or inhibition of engram cell ensembles induces or inhibits memory retrieval, respectively; thus, engram cell ensembles represent the physiological manifestation of a specific memory trace.

However, one episodic memory is composed of several components of the episode, and each component should be encoded by specific substrates, e.g. engram sub-ensembles. Nevertheless, it had not been clear how activity in these engram cells was assembled to represent the corresponding event because of technical limitations, and these mean that it is difficult to distinguish the population activity of engram cells from that of non-ensemble cells.



Episodic memory of experience of novel space is constructed by several sub-ensembles within a single engram population. a. Diagrammatic representation of NMF analysis; colored bars indicate the relative activity of each neuron in the detected pattern (shown as the magnitude) (left), and the relative intensity of the synchronous activity of this pattern across time (right). b. Example of two detected engram sub-ensembles. Distribution map of detected engram cells (left panel). Green and red magenta filled contours indicate neurons assigned to two different ensembles (corresponding to patterns #5 and #9, respectively), and yellow filled contours denote cells shared between patterns. Scale bar, 100 μ m. The temporal pattern of synchronous activities of patterns #5 and #9 during experiencing novel space (top and middle right) and their overlay (bottom right). Credit: University of Toyama

To address how one [episodic memory](#) is represented and consolidated in engram cell ensembles, one must visualize the activity of engram and non-engram cells. Engram cells can be specifically targeted in c-fos-tTA mice because the neural activity associated with memory formation induces c-fos gene expression, which in turn induces activity-dependent tTA expression under the control of the c-fos promoter. In the absence of doxycycline, tTA can bind to the tetracycline responsive element (TRE), enabling downstream

expression of the TRE-dependent transgene (Figure 1a).

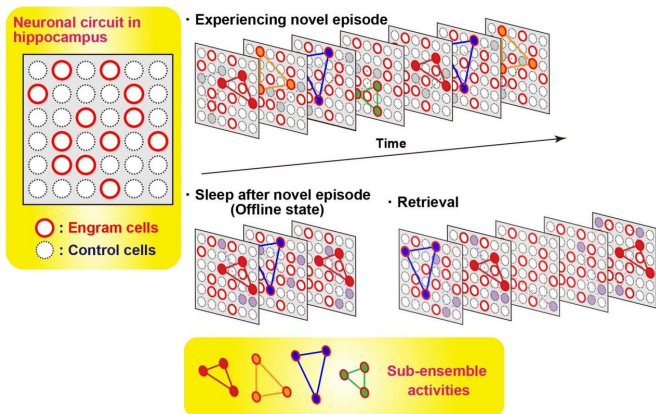
When a neuron activates, Ca²⁺ flows into their soma. Thy1-G-CaMP7 mice express a Ca²⁺ indicator, G-CaMP7, in pyramidal neurons of hippocampal CA1 in the mice. Thus, neuronal activity is transferred into G-CaMP7 fluorescence, called Ca²⁺ imaging. We developed a technique that combines a head-mounted, miniature fluorescent microscope, with Thy1-G-CaMP7/c-fos-tTA double transgenic mice. The hippocampal CA1 region in double transgenic mice was injected with LV expressing a fluorescent protein, Kikume Green Red (KikGR), under the control of TRE (Figure 1a and 1b). Using this approach, engram cells can be identified with KikGR, and the Ca²⁺ signals corresponding to the activity of engram cells and non-engram cells can be tracked during the experience of a novel episodic event (Figure 1c and 1d).

The data indicated that population activity of engram cell ensembles exhibited the characteristic trait of highly repetitive activity during novel episodic events. To address components of one memory, it was proposed to deconstruct population activity into sub-ensemble groups. Non-negative matrix factorization (NMF) decomposes population activity into a time series of coactivated neuronal ensembles (Fig. 2a). Each sub-ensemble is composed of different cells, which are spatially intermingled (Fig. 2b left), to make their synchronous activity even among the group of engram cells associated with a single event (Fig. 2b right). These results suggest that the total information of one event is structured into sub-ensemble ensembles.

In order to measure the activity of engram cells across different memory processing stages, recording of Ca²⁺ transients from novel experience through post-experience sleep to retrieval was conducted. Around 40% of the total number of sub-ensembles that appeared in a novel experience in engrams reactivated during post-experience sleep and then preferentially reappeared in retrieval sessions, while almost none of sub-ensembles in non-engrams did not show this feature. Thus, engram sub-ensembles formed during a novel

experience and that were reactivated during sleep sessions were mostly reactivated during the retrieval session (Fig. 3). By contrast, most non- engram ensembles that were activated during the novel experience were not reactivated in the later sessions.

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Diagrammatic model showing orchestration of ensembles of engram cells across memory processing. Many sub-ensembles express during experiencing novel episode. A part of sub-ensembles of engram cells re-appear during post-episode experience sleep. This set of sub-ensembles appeared during sleep are more likely observed during memory retrieval. In contrast, control, non- engram, cells almost do not show this feature. Credit: University of Toyama

The findings reported in this study demonstrate that engram cells possess synchronous activity, formed by several sub-ensembles in the engram population. Only in engram [cells](#) does this synchronous activity survive through post-learning sleep sessions that contribute to the consolidation process. The present work sheds light on the relationship between ensemble activities and coding principles in [learning](#) and [memory](#).

More information: Khaled Ghandour et al, Orchestrated ensemble activities constitute a hippocampal memory engram, *Nature Communications* (2019). [DOI: 10.1038/s41467-019-10683-2](https://doi.org/10.1038/s41467-019-10683-2)

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