

MRC

Brain Network
Dynamics Unit

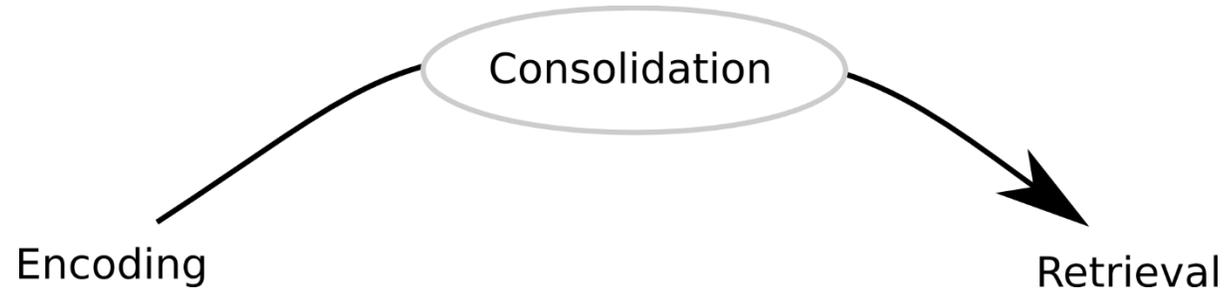


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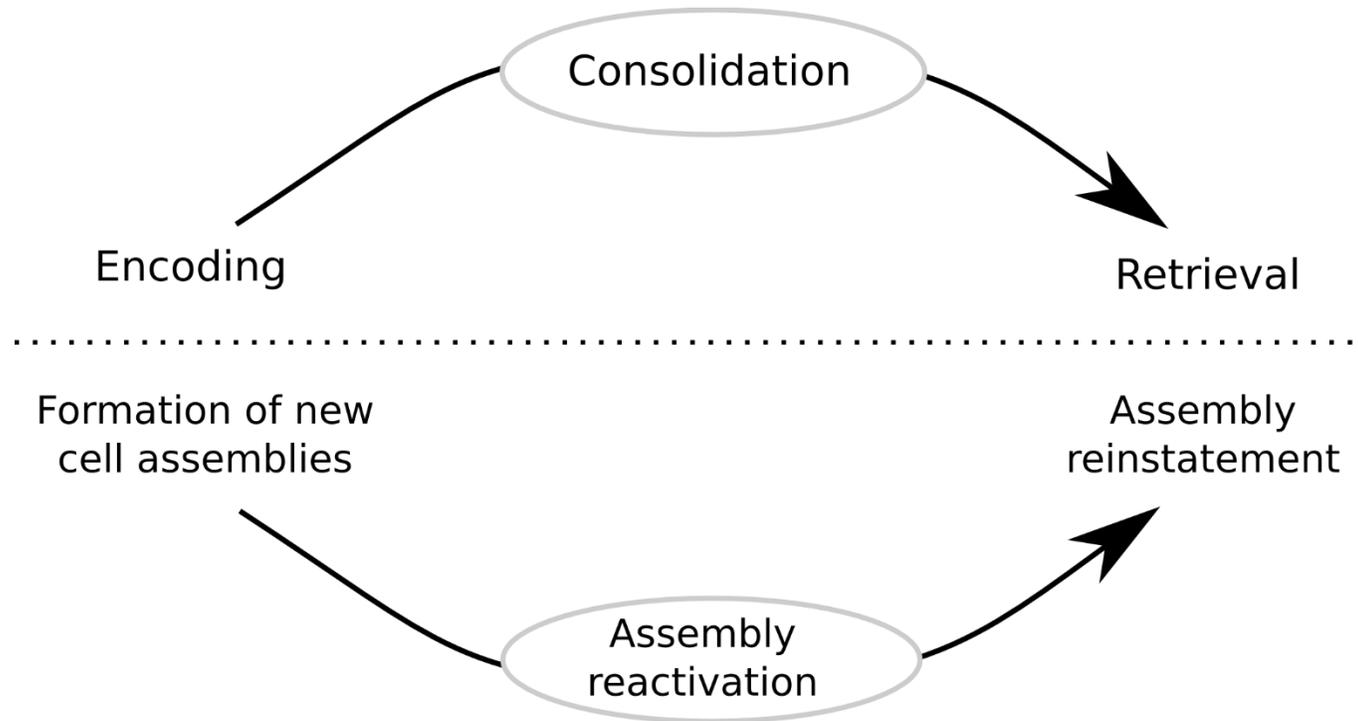
Hippocampal reactivation stabilizes recently formed cell assembly patterns

Gido van de Ven

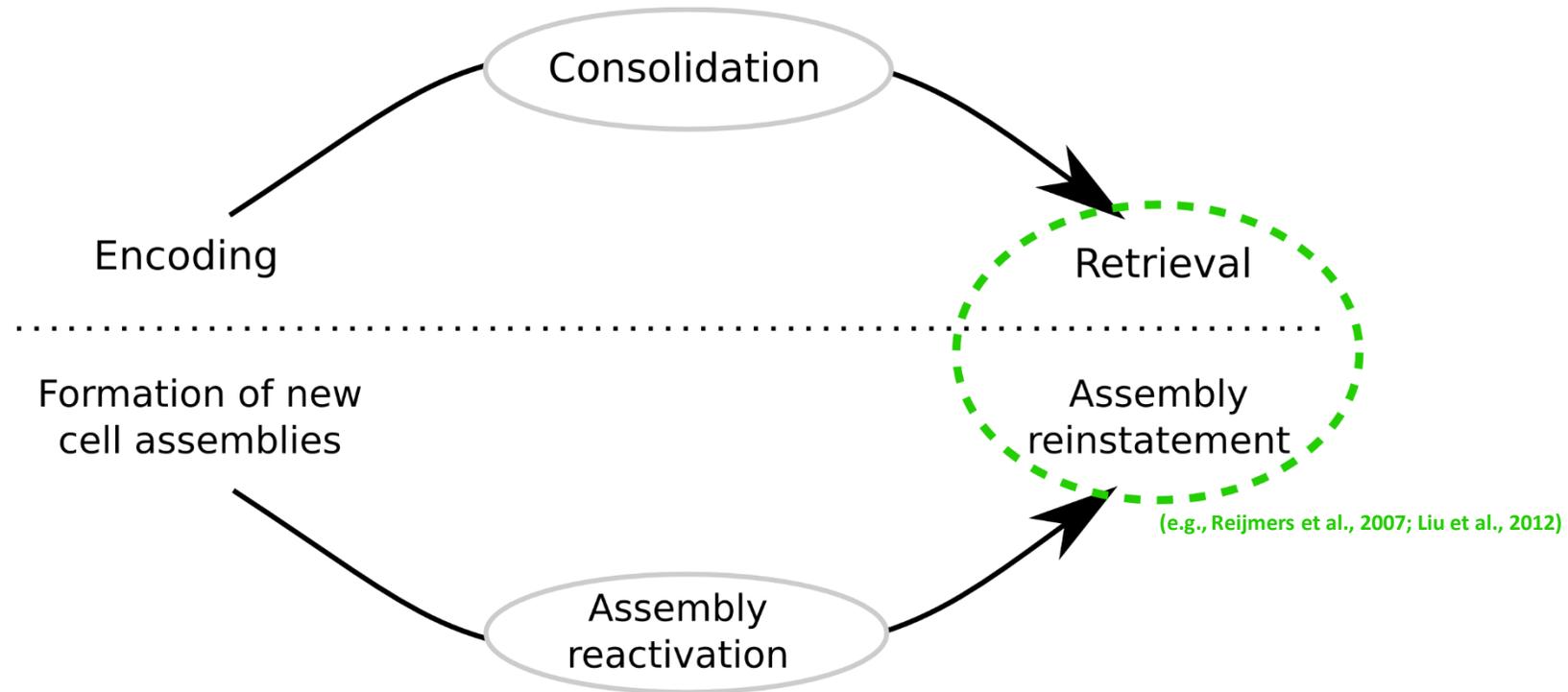
Memory-consolidation



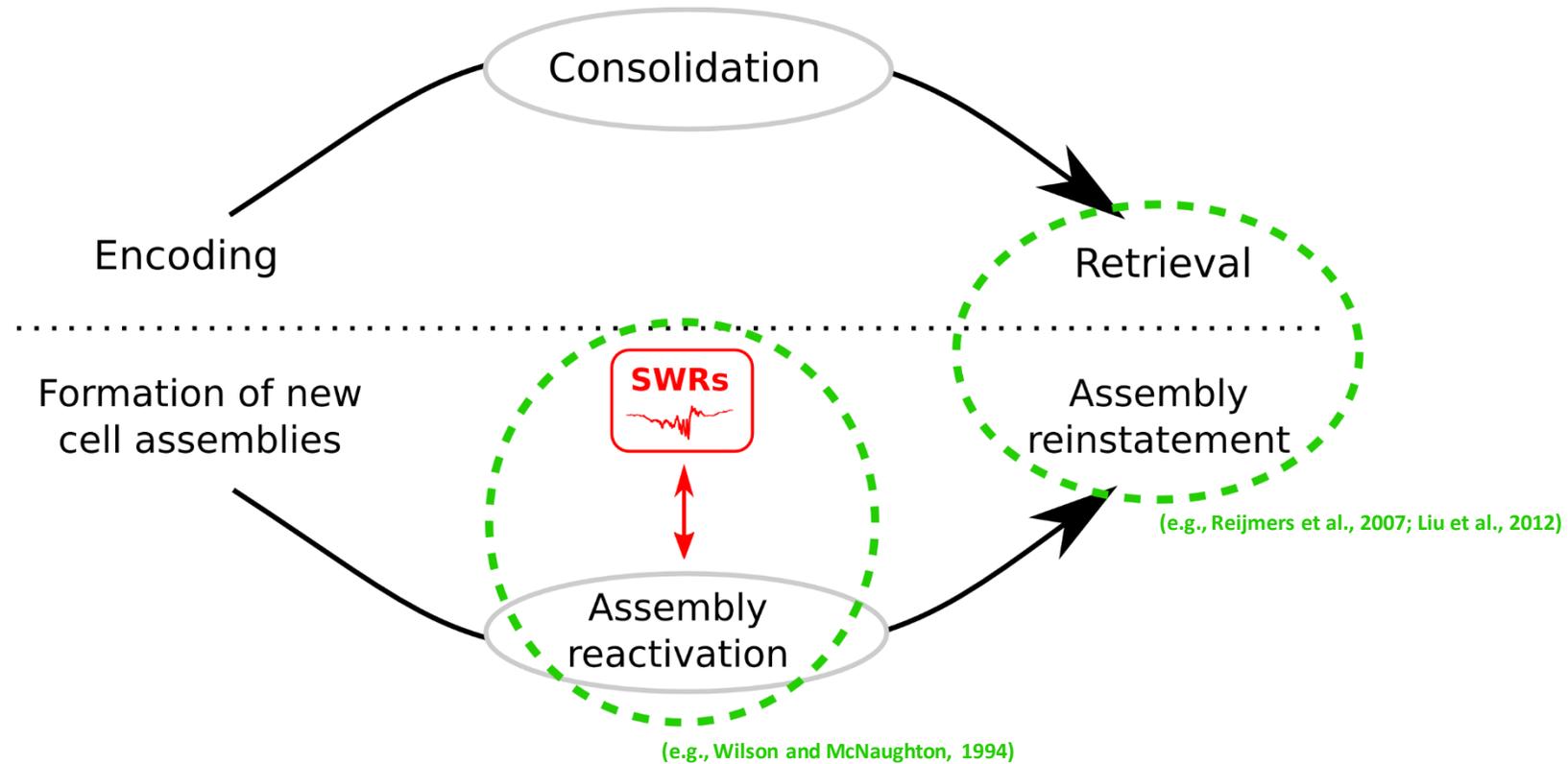
Memory-consolidation: cell assembly / reactivation hypothesis



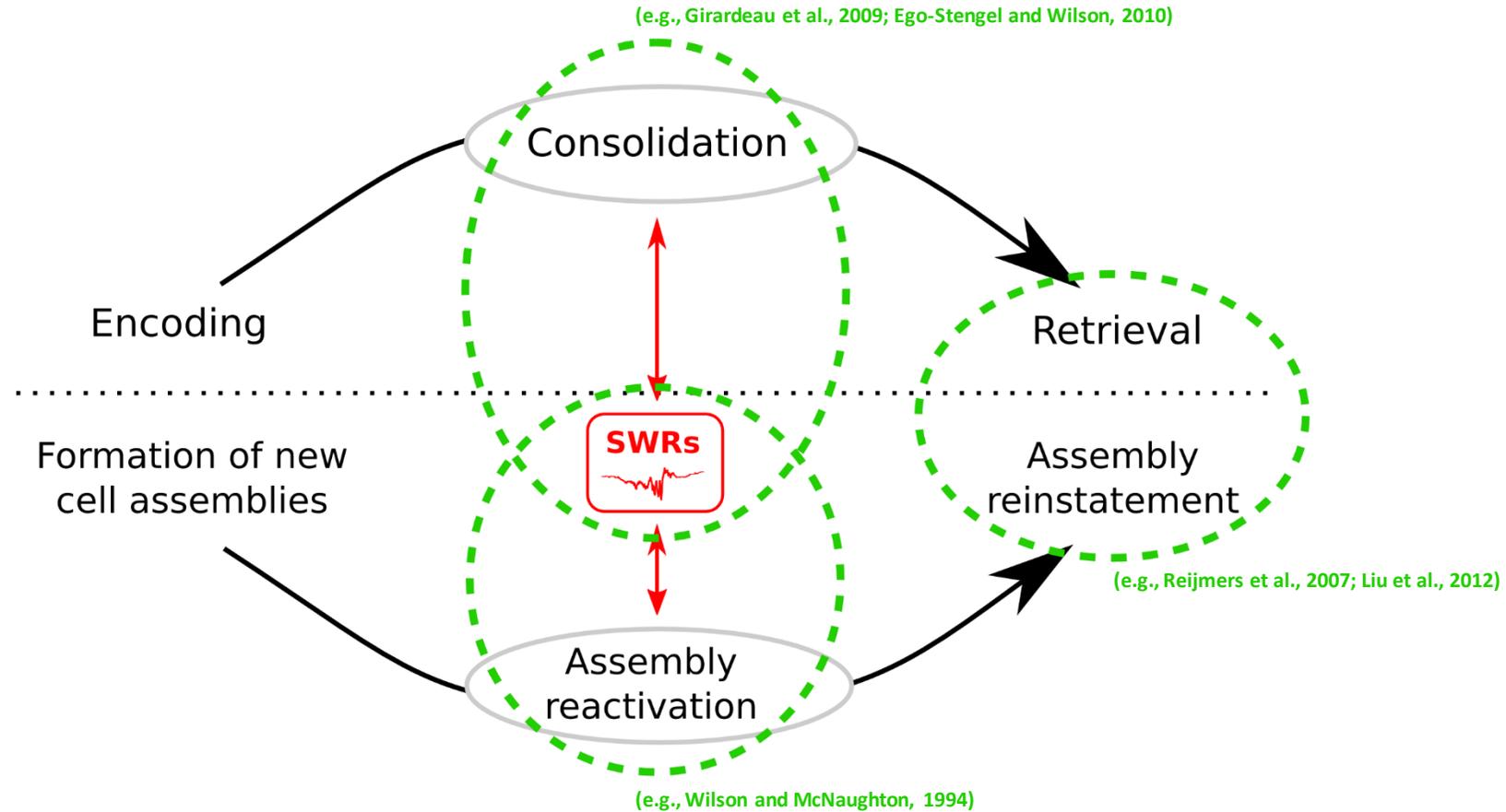
Memory-consolidation: cell assembly / reactivation hypothesis



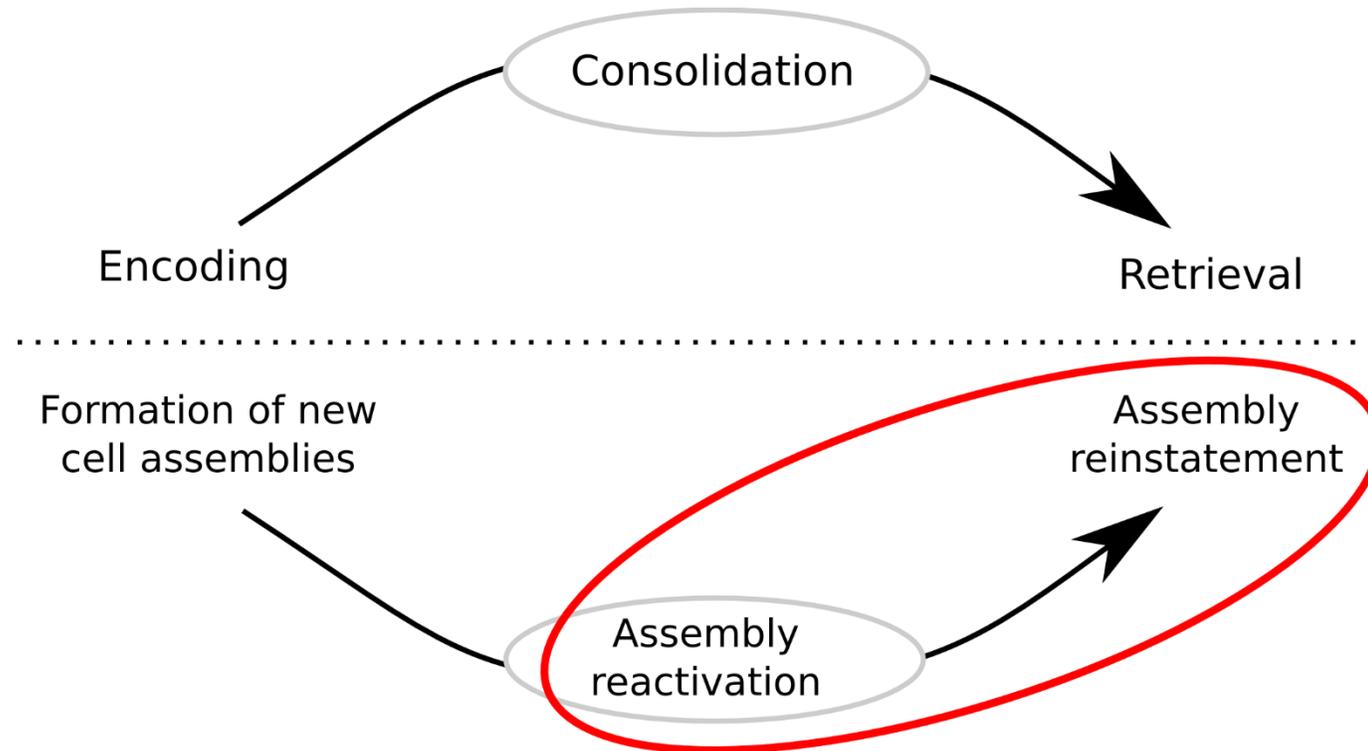
Memory-consolidation: cell assembly / reactivation hypothesis



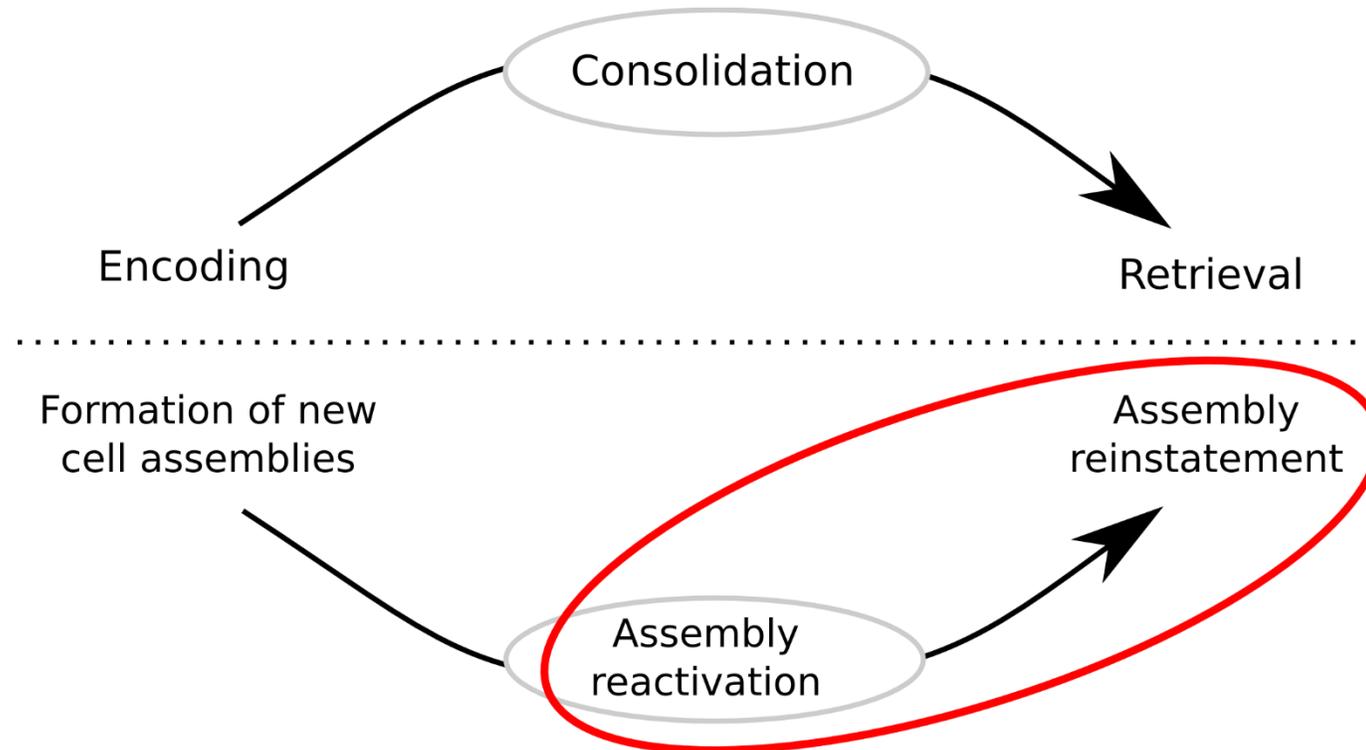
Memory-consolidation: cell assembly / reactivation hypothesis



Memory-consolidation: cell assembly / reactivation hypothesis



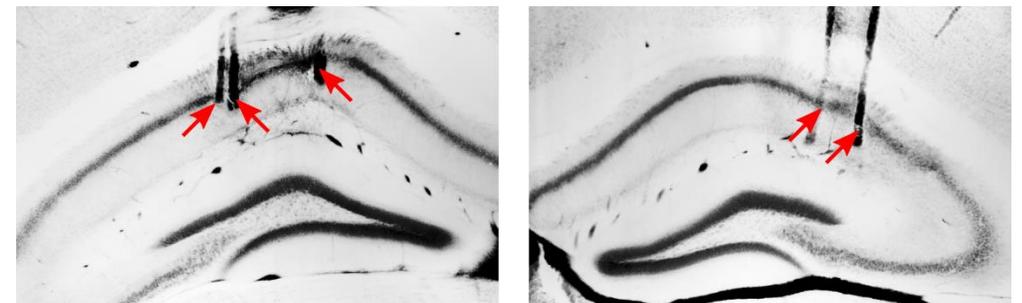
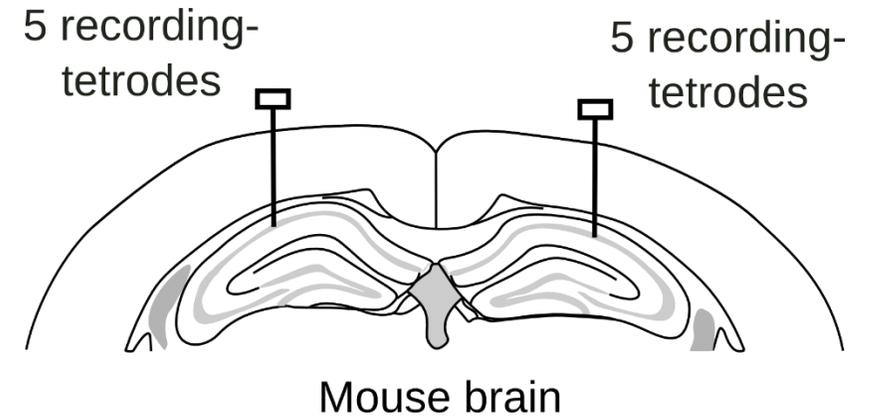
Memory-consolidation: cell assembly / reactivation hypothesis



2 challenges: → Identification & tracking of “memory-representing” cell assemblies
→ Selective disruption of reactivation

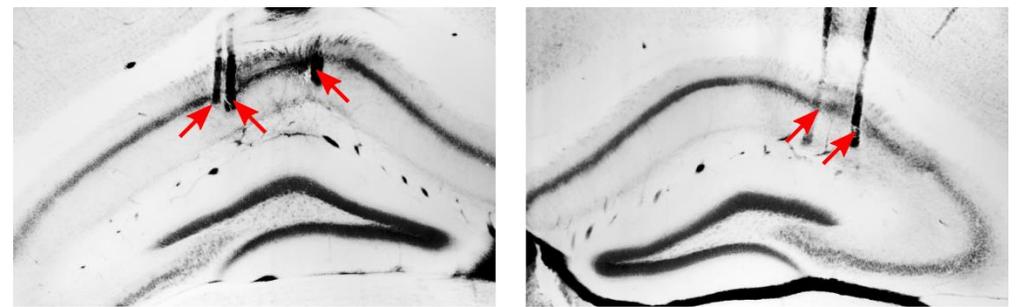
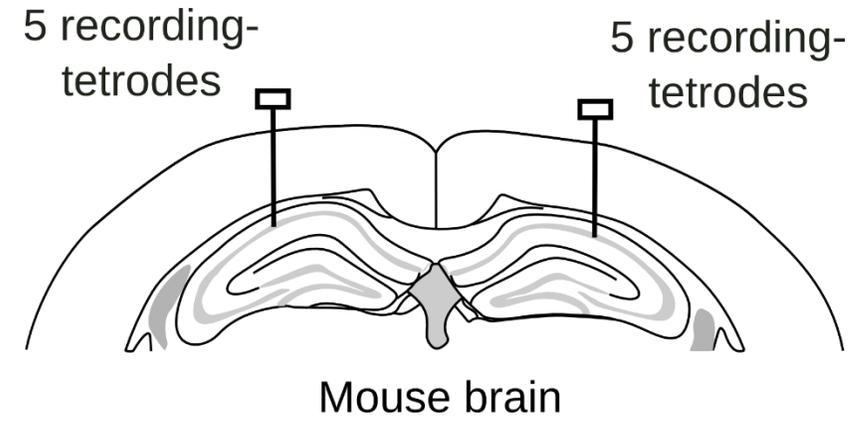
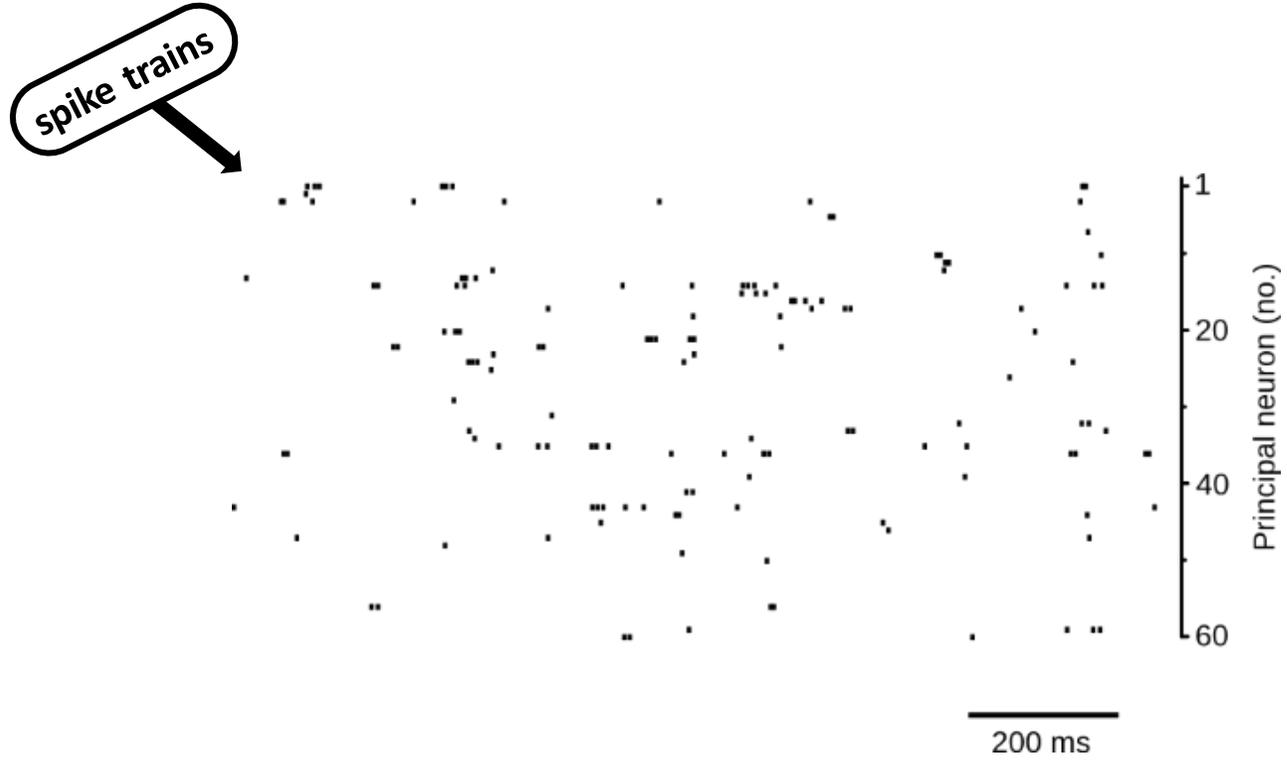
Identification of “cell assemblies”

= **challenge 1**



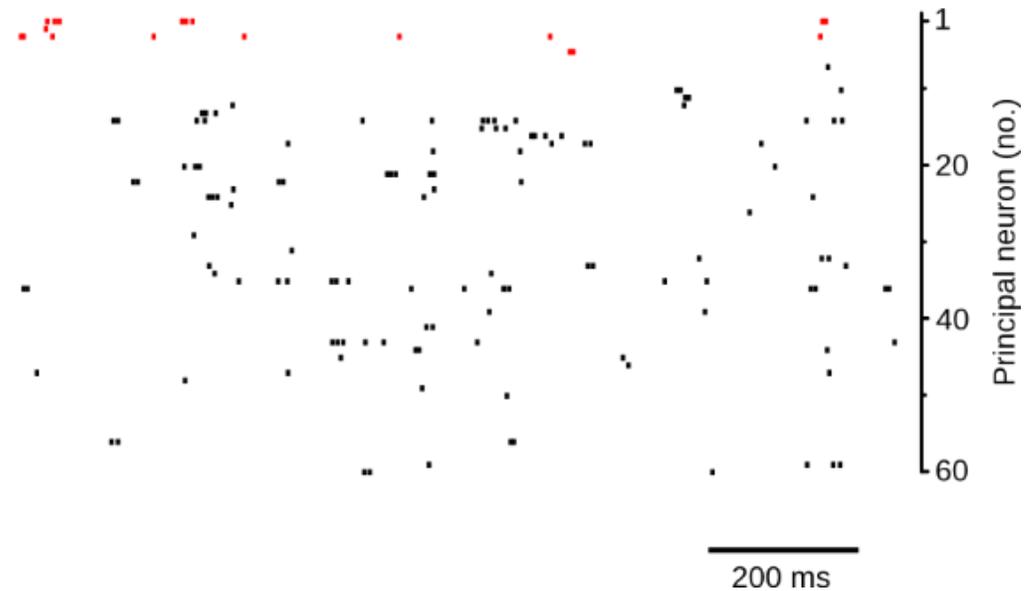
Identification of “cell assemblies”

= **challenge 1**

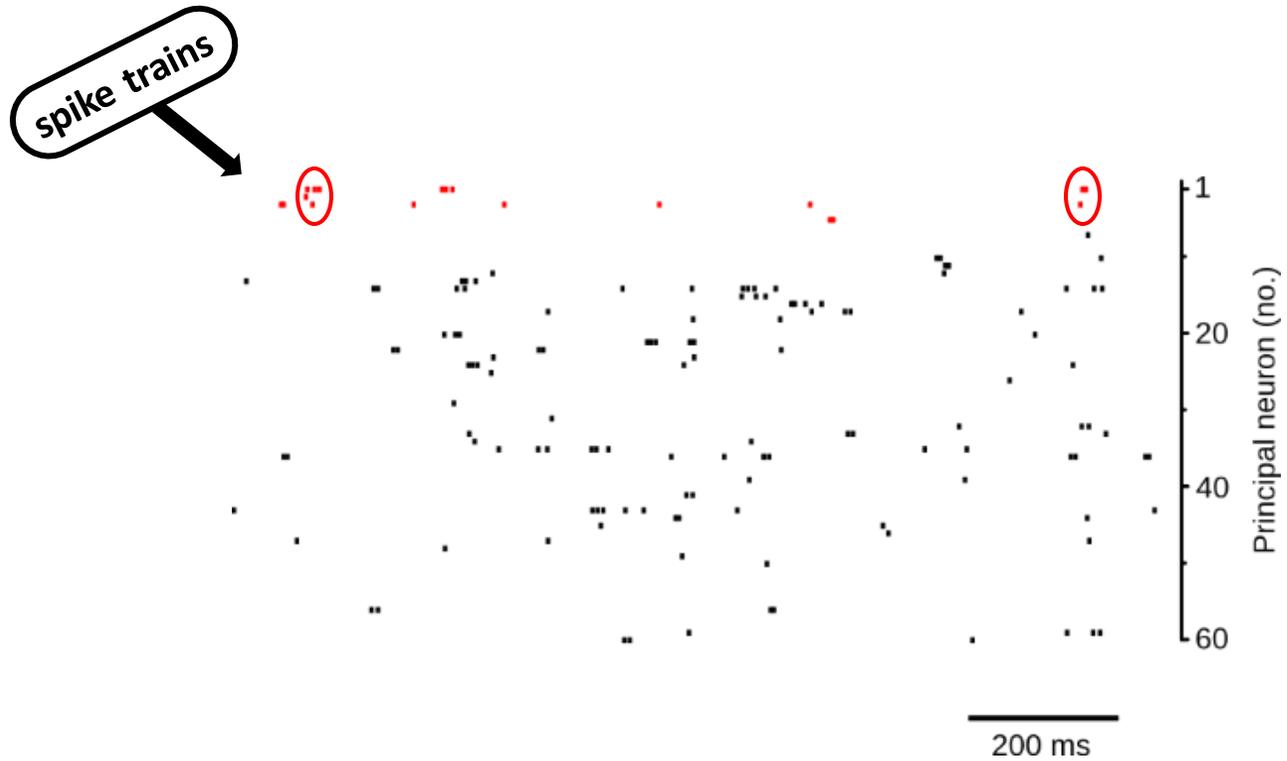


Identification of “cell assemblies”

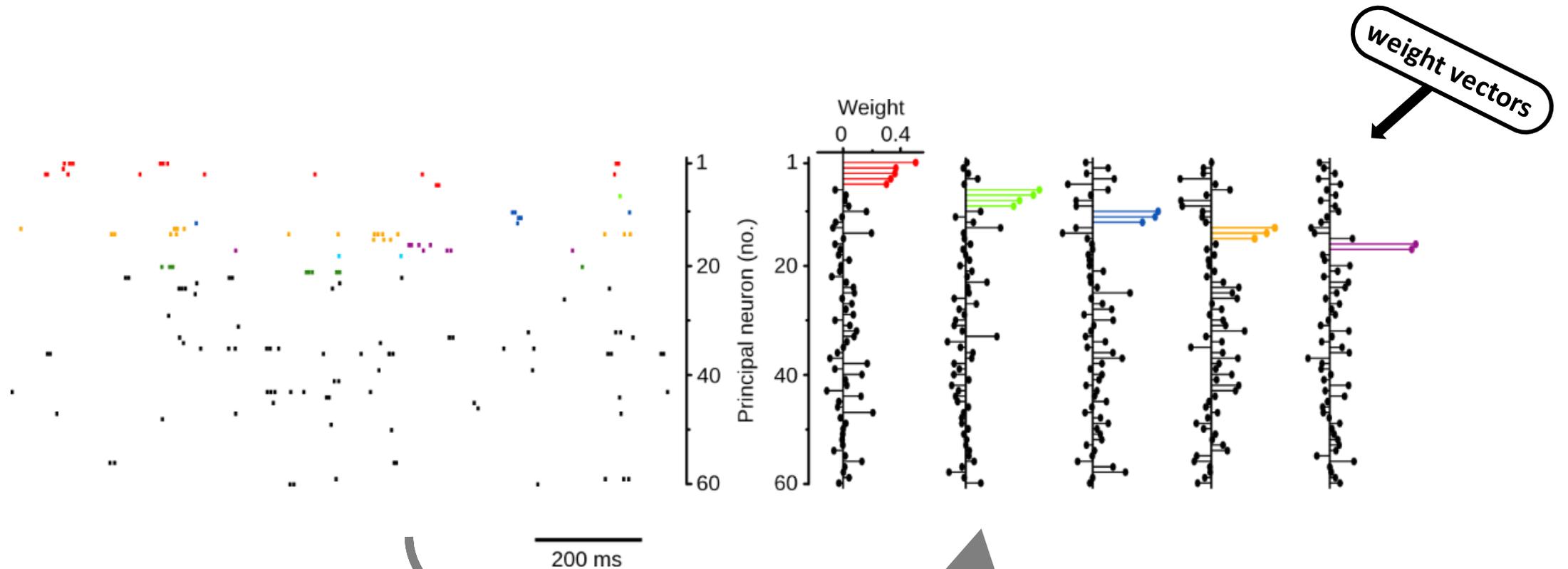
spike trains



Identification of “cell assemblies”

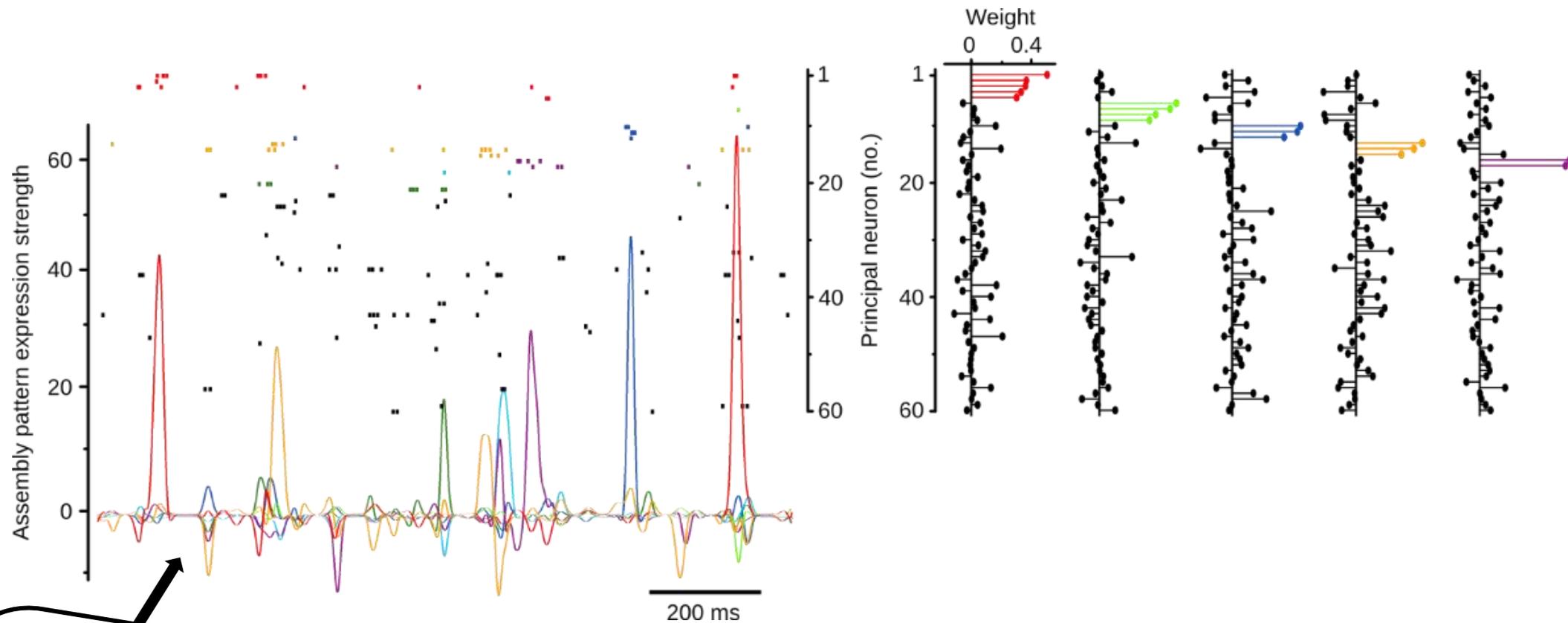


Identification of “cell assemblies”



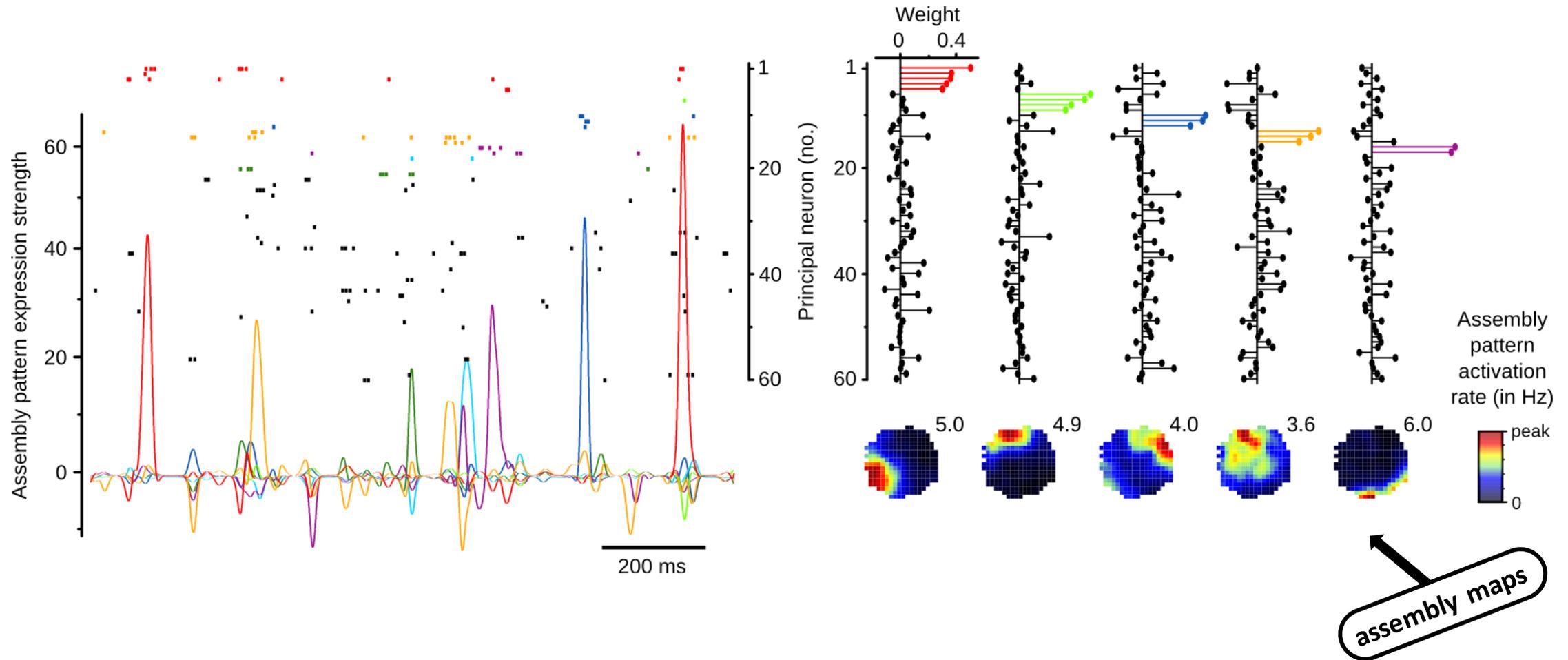
*Assembly-detection method
based on PCA and ICA*

Identification of “cell assemblies”



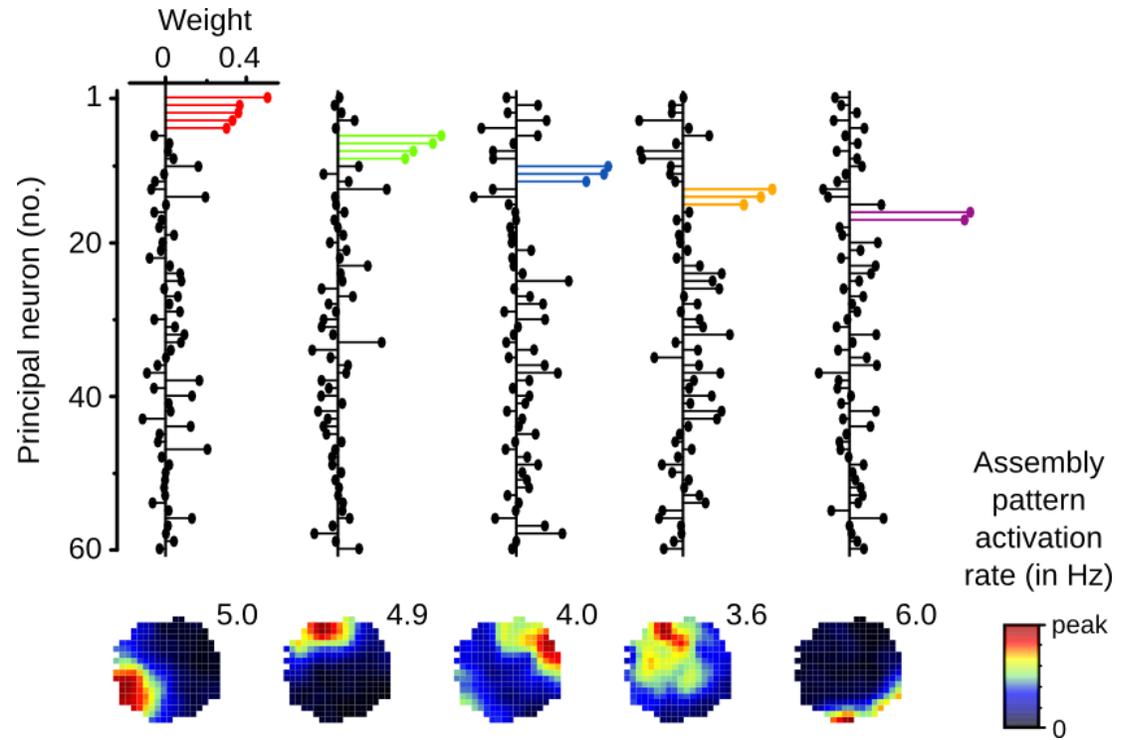
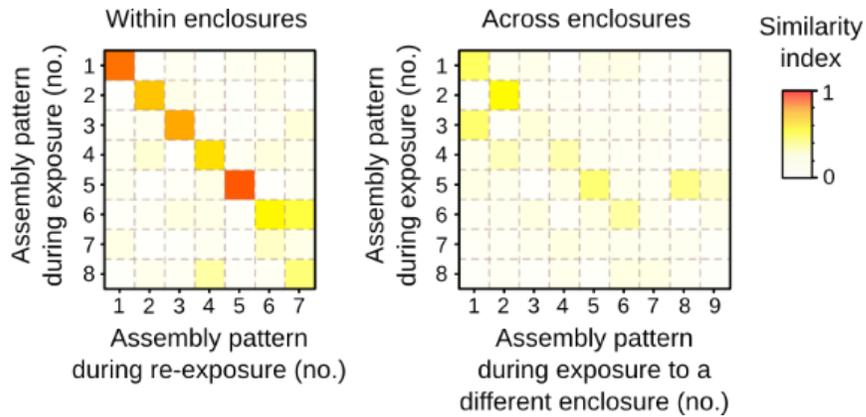
assembly pattern expression-strength tracked over time

Identification of “cell assemblies”



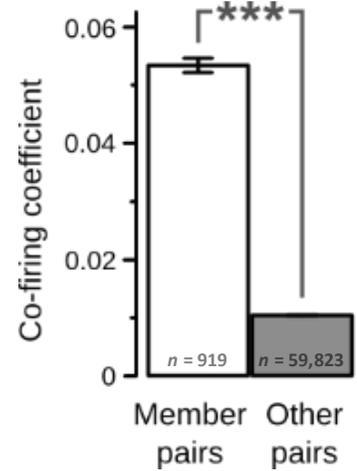
Identification of “cell assemblies”

assembly patterns are environment-specific

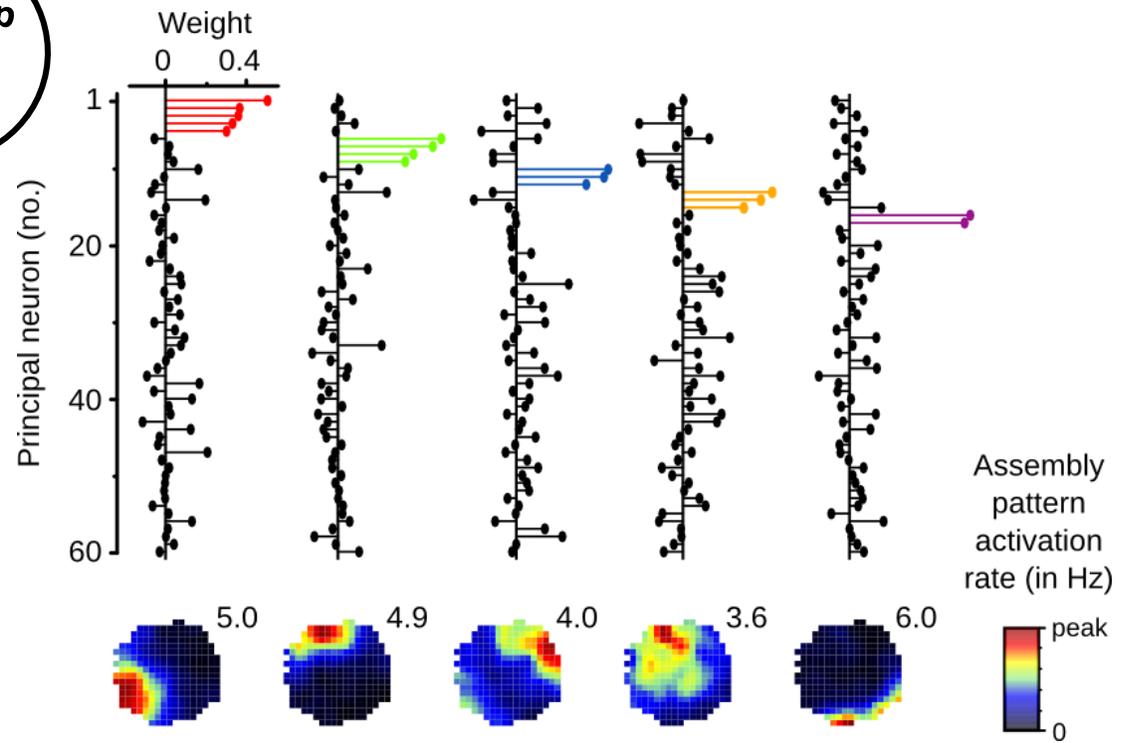
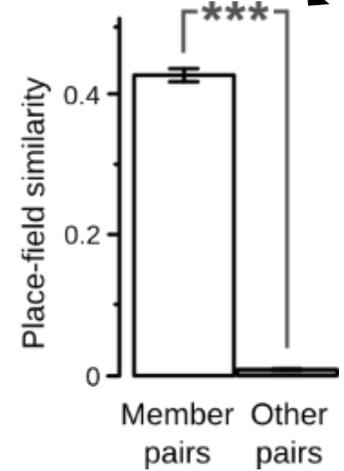


Identification of “cell assemblies”

assembly patterns group together co-active neurons

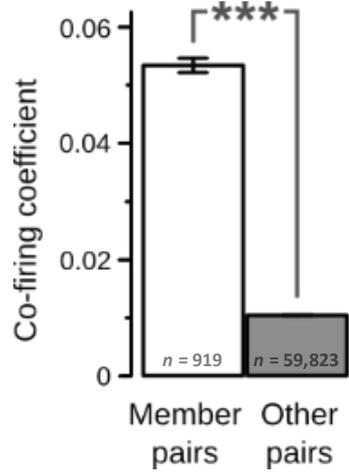


assembly patterns group together neurons with overlapping place fields

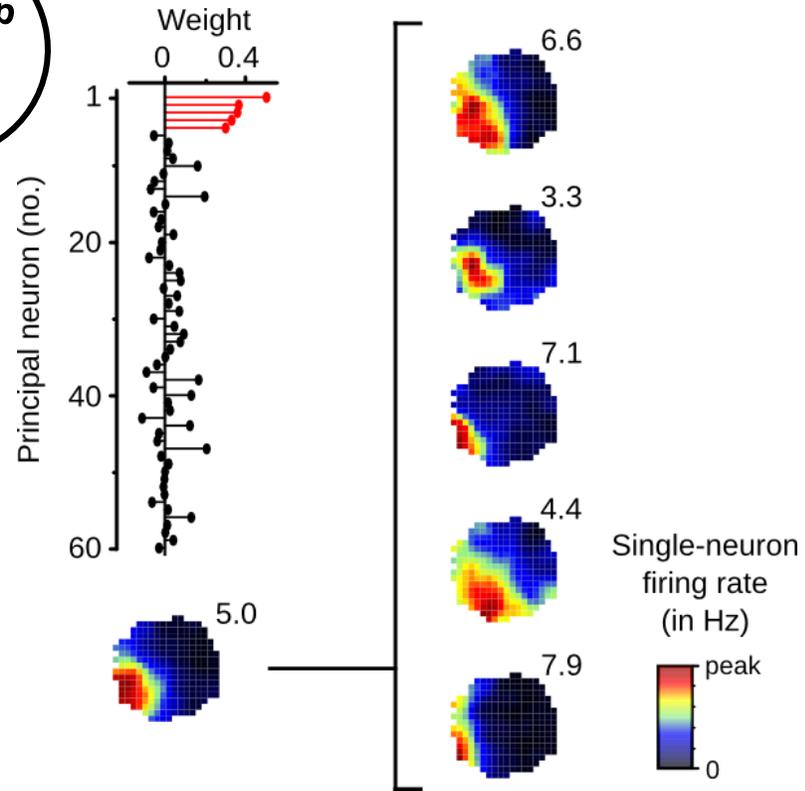
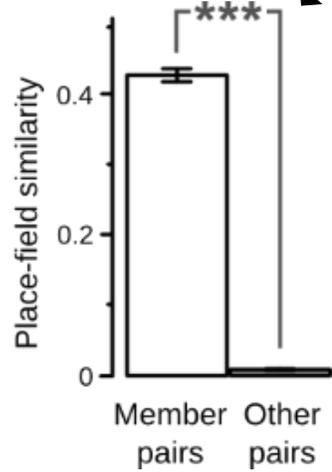


Identification of “cell assemblies”

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assembly patterns group together neurons with overlapping place fields



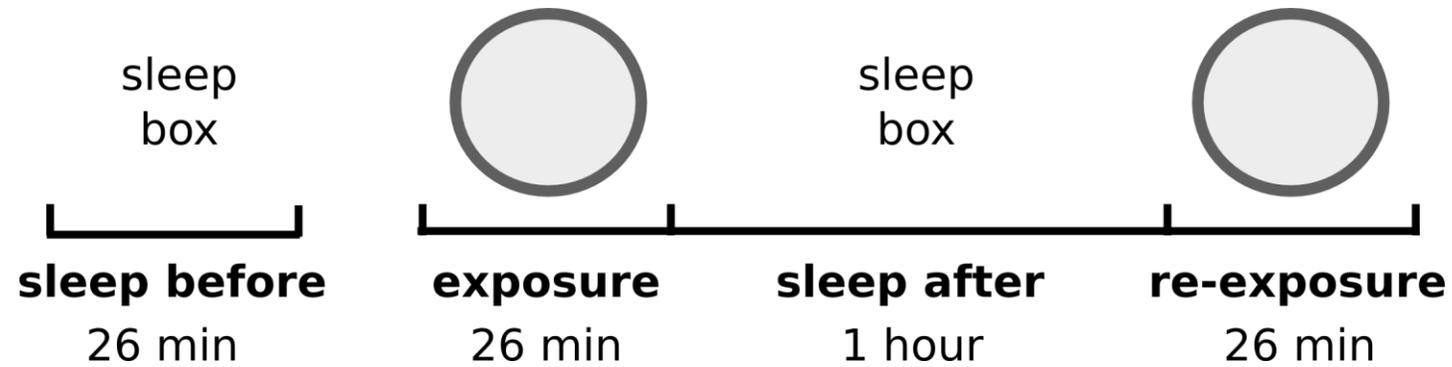
Identification of “cell assemblies” - summary

Based on short-time scale interactions (25 ms) in tetrode recordings of hippocampal principal neurons, an unsupervised statistical framework based on PCA and ICA **detects** and **tracks** cell assembly patterns that:

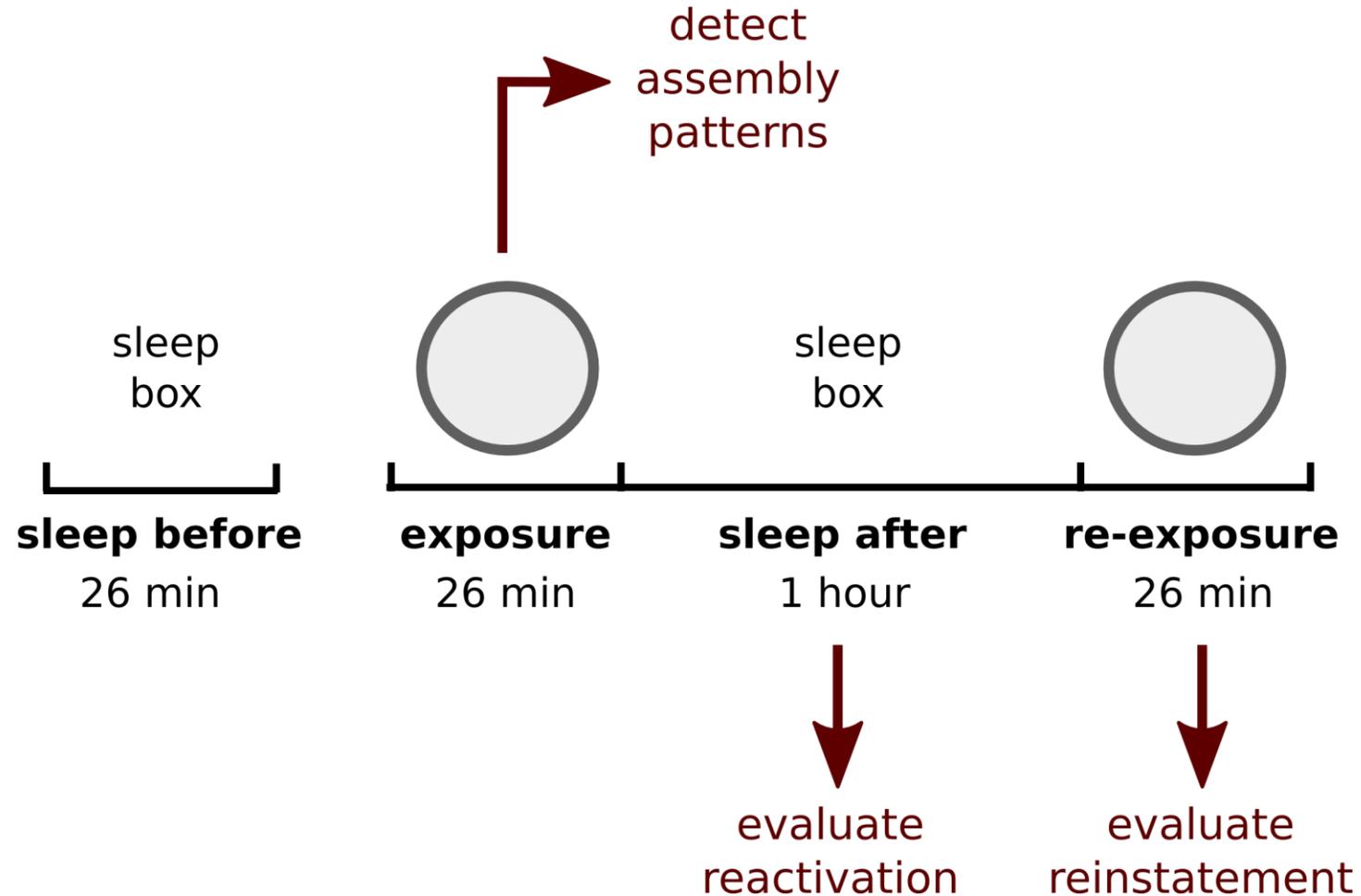
- are spatially selective;
- are environment-specific;
- bind together co-active neurons; and
- bind together neurons with overlapping spatial tuning.

→ **challenge 1**: Identification & tracking of “memory-representing” cell assemblies

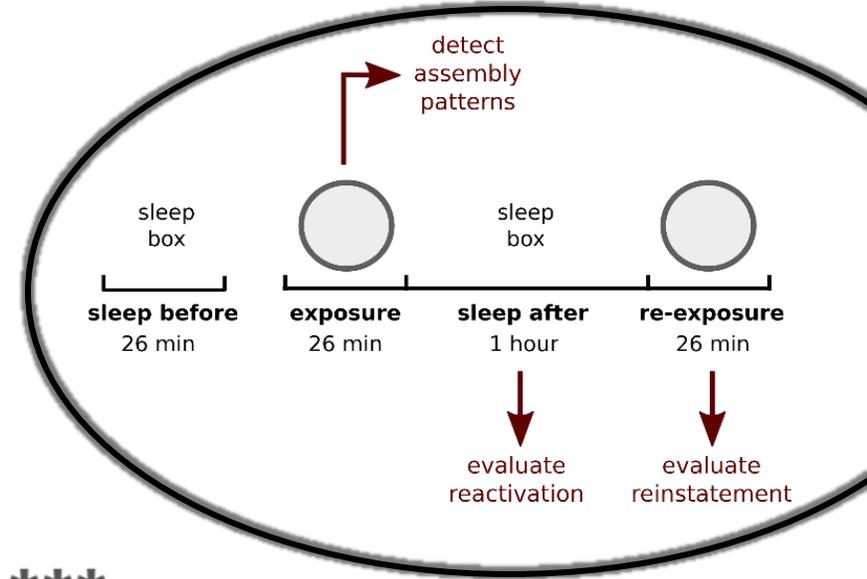
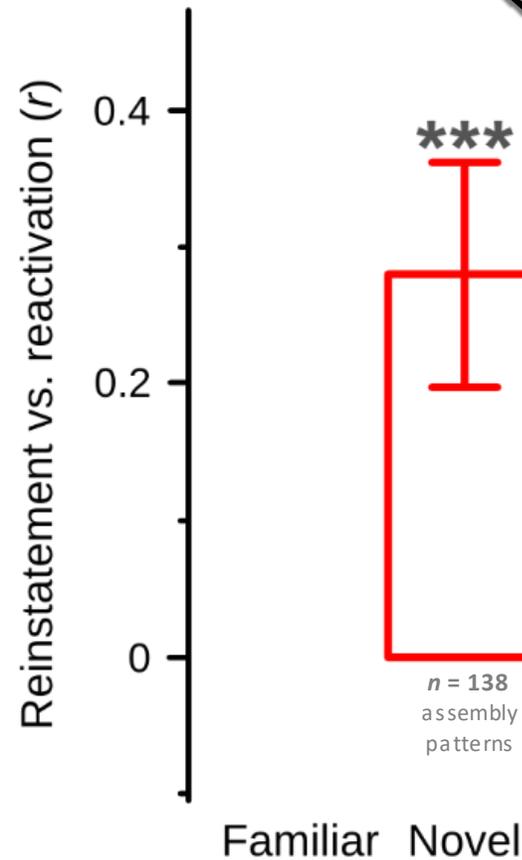
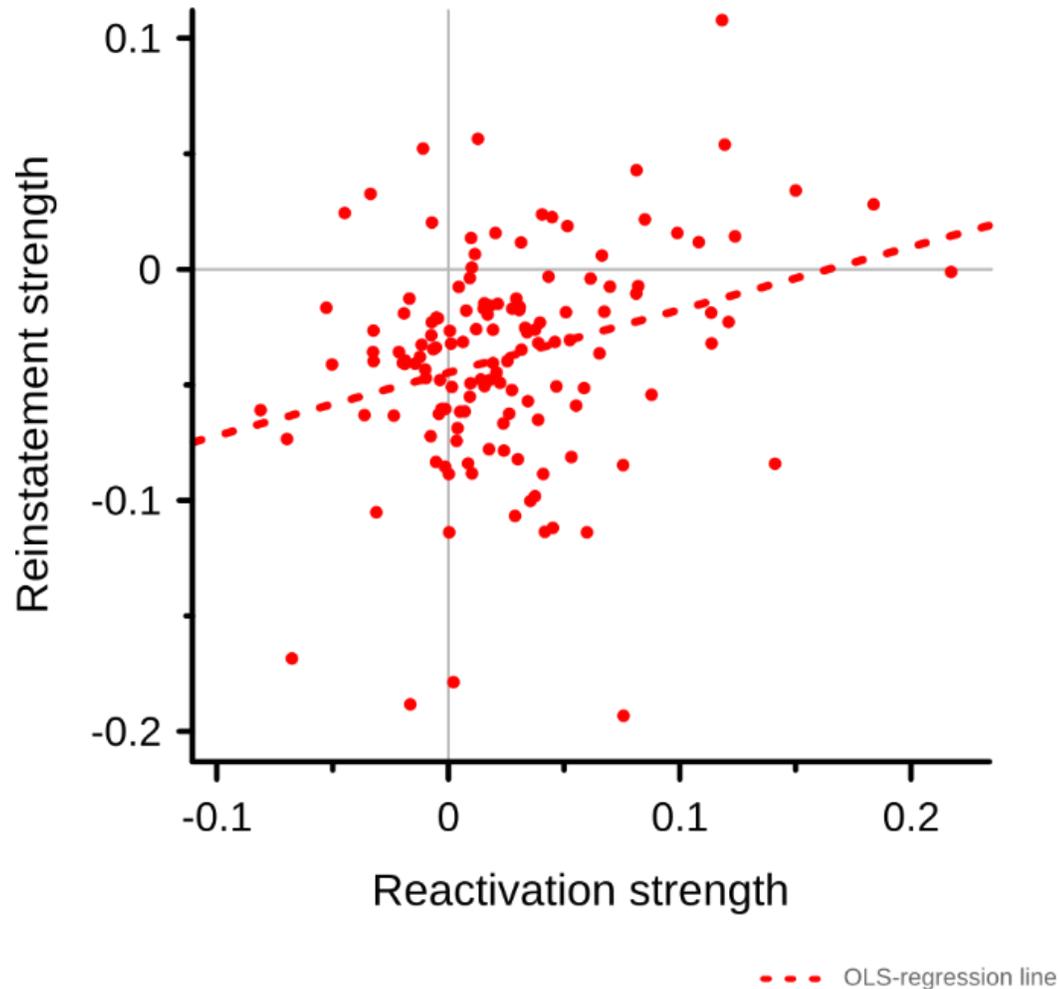
Experimental protocol



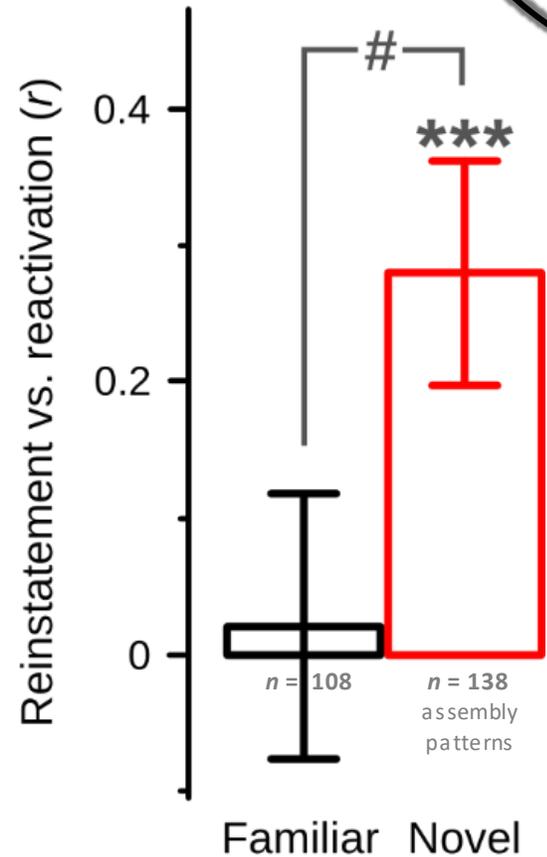
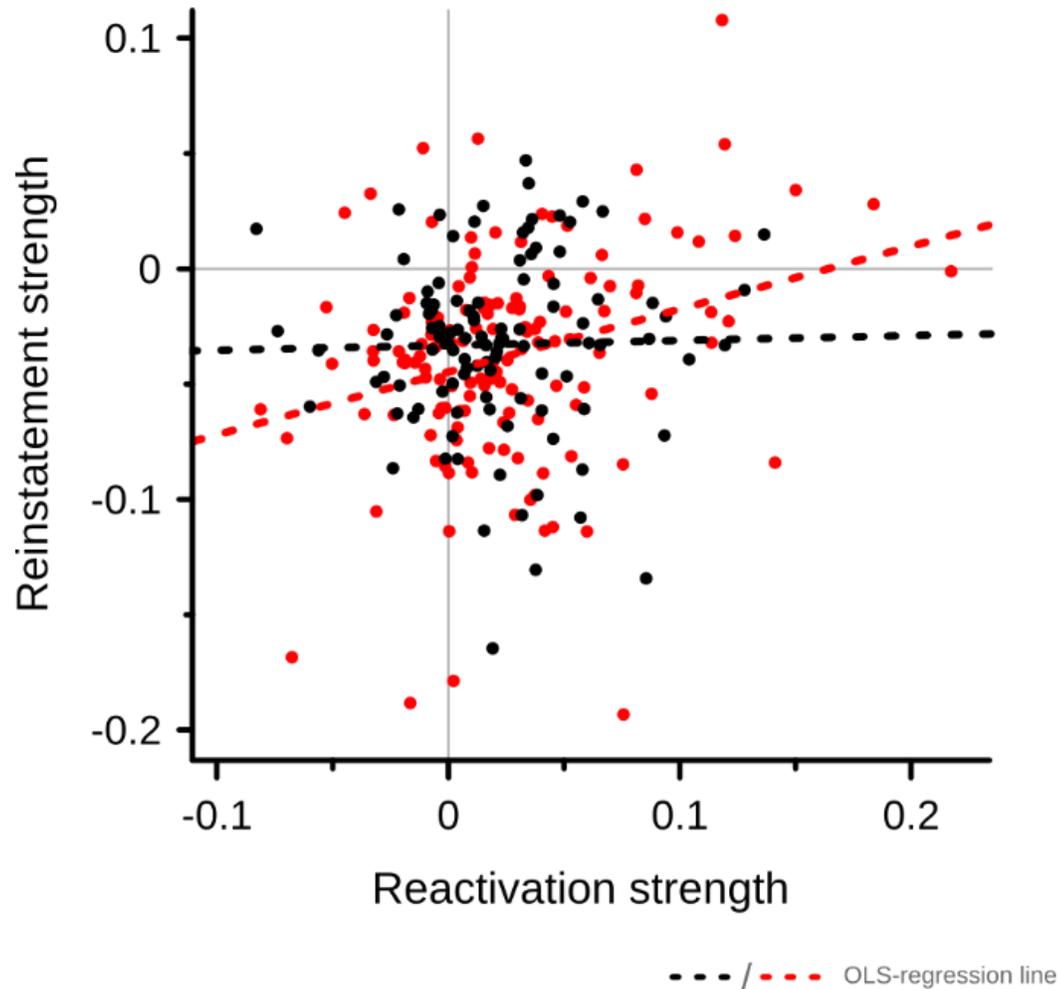
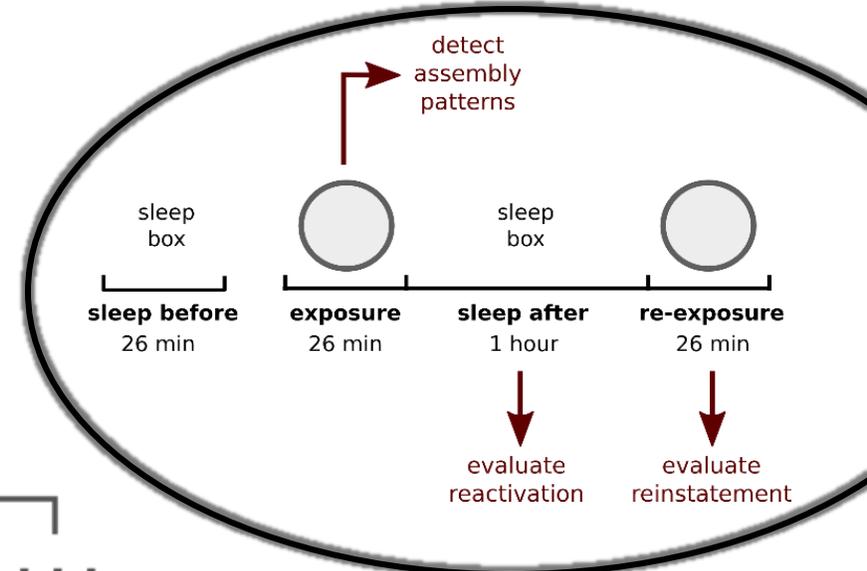
Experimental protocol - (*correlation*)



An assembly pattern's reactivation predicts it subsequent reinstatement



An assembly pattern's reactivation predicts its subsequent reinstatement

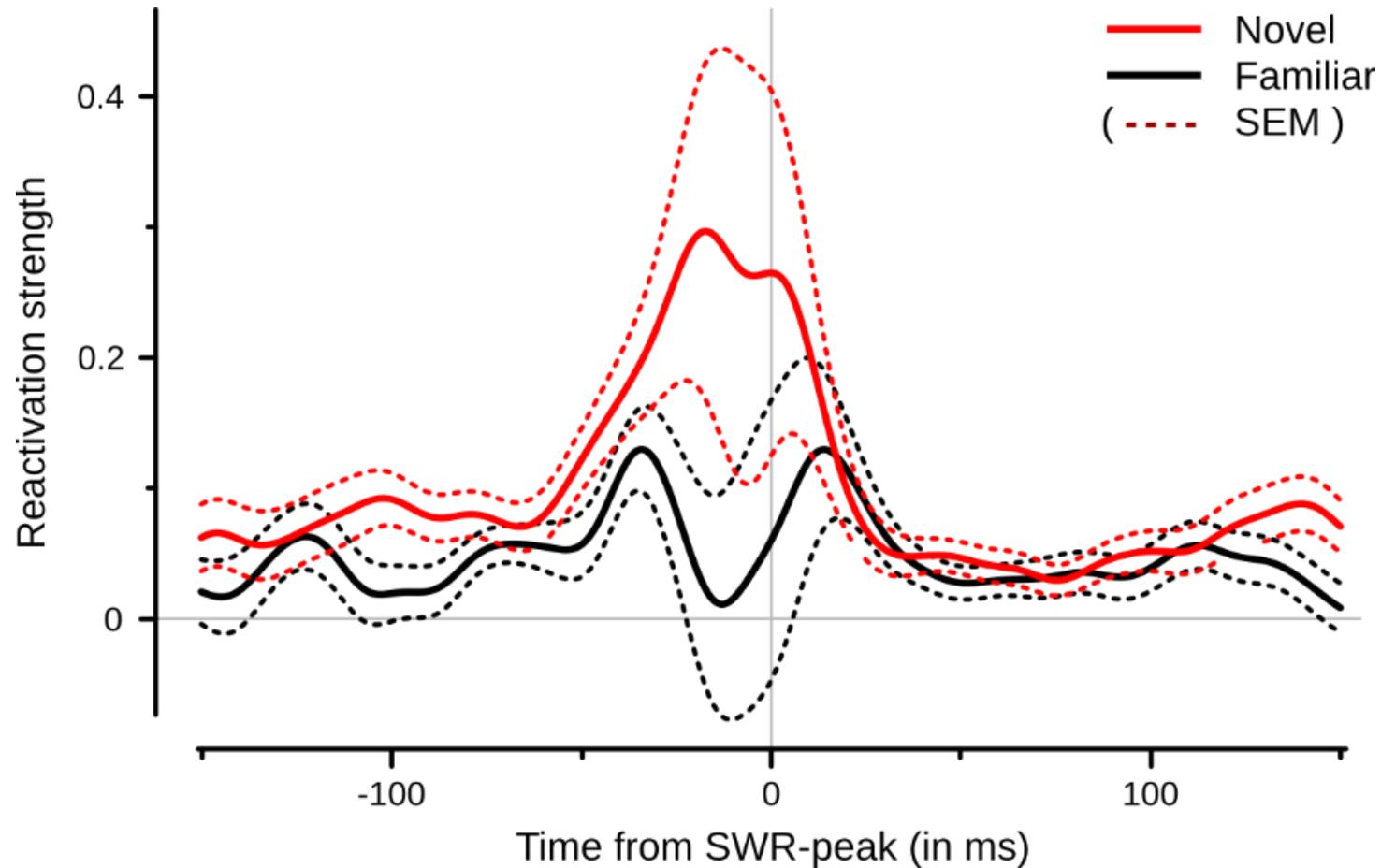


Selective disruption of reactivation?

= **challenge 2**

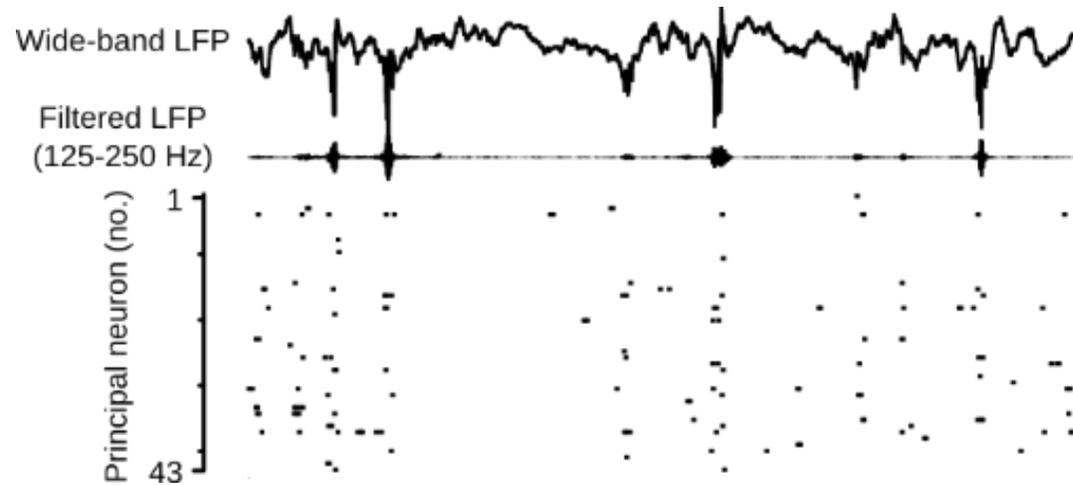
Selective disruption of reactivation?

= challenge 2

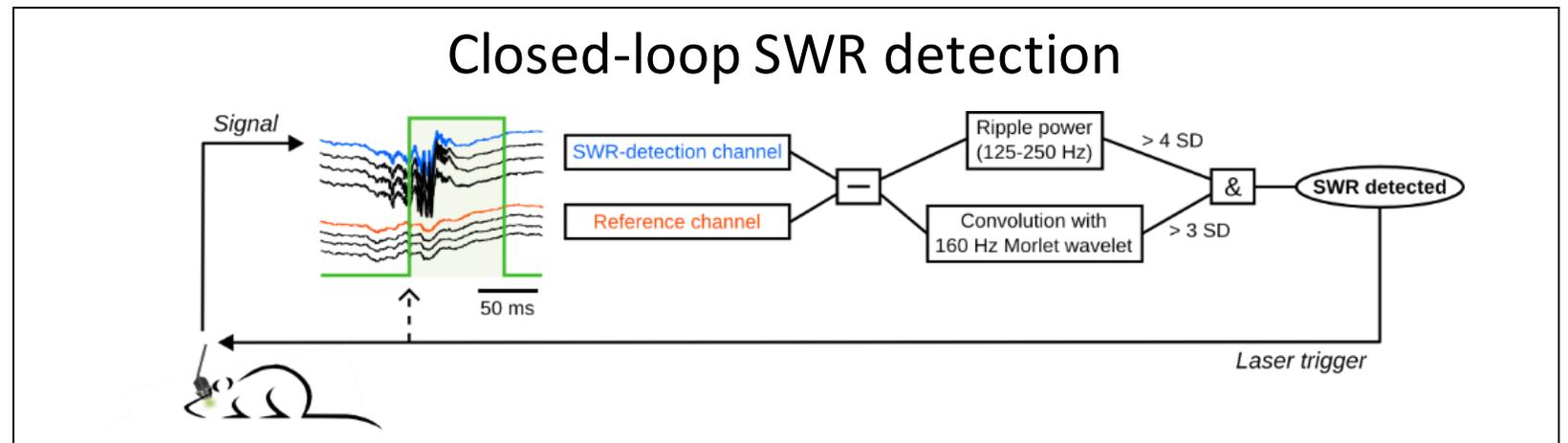
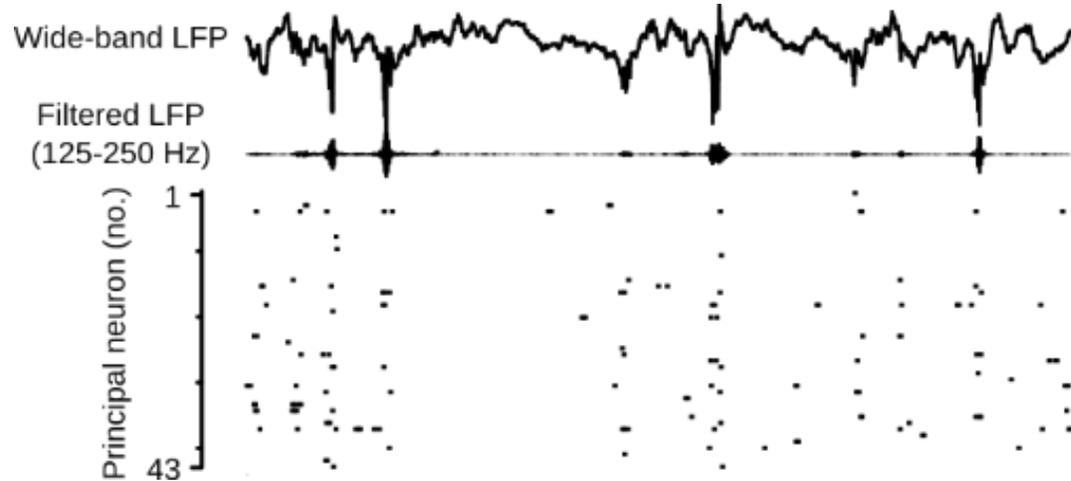


Novel: $n = 139$ assembly-patterns
Familiar: $n = 108$ assembly-patterns
(based on 43 recording-blocks from 8 mice)

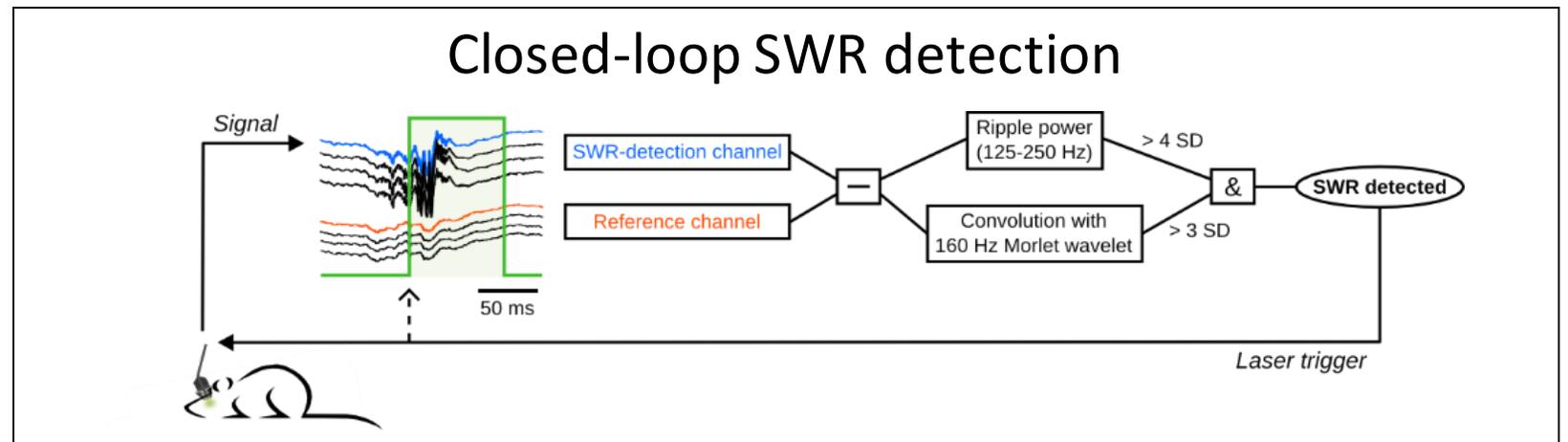
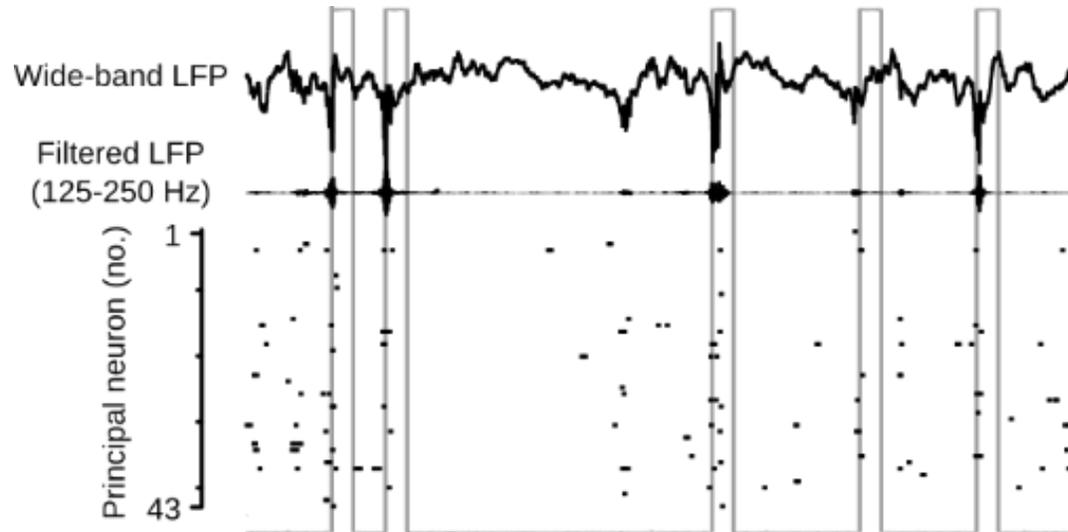
Selective disruption of reactivation: *optogenetic SWR silencing*



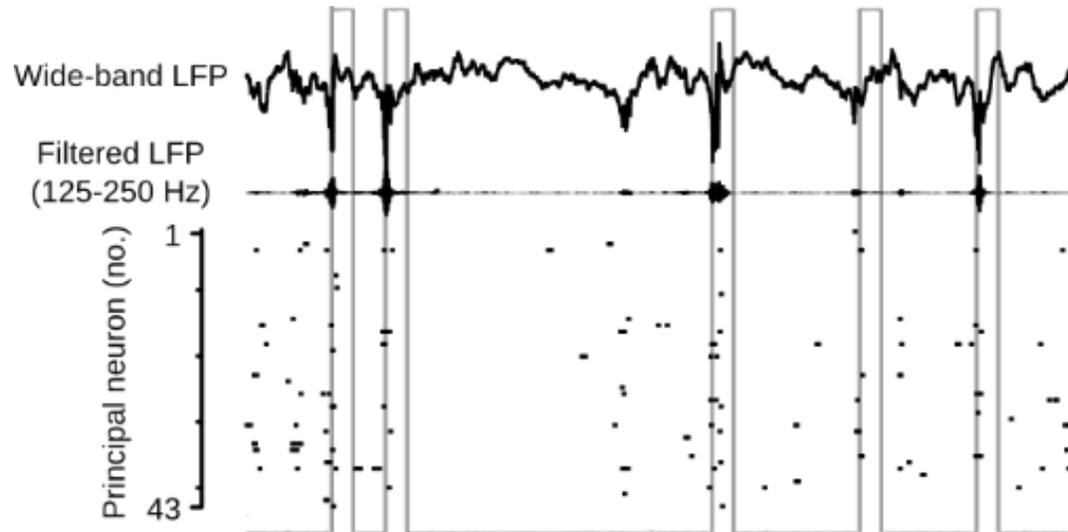
Selective disruption of reactivation: *optogenetic SWR silencing*



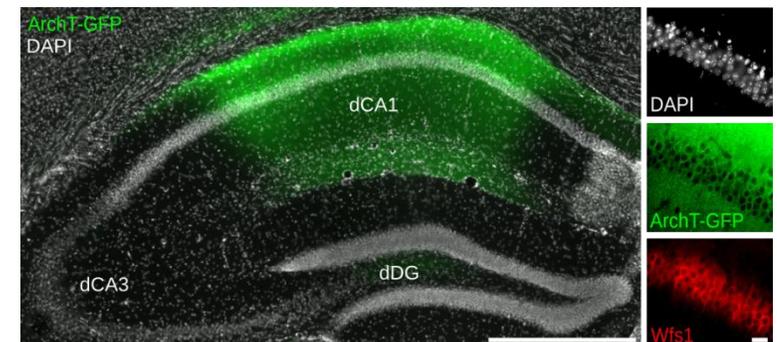
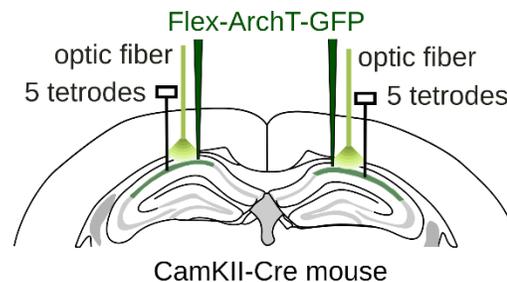
Selective disruption of reactivation: *optogenetic SWR silencing*



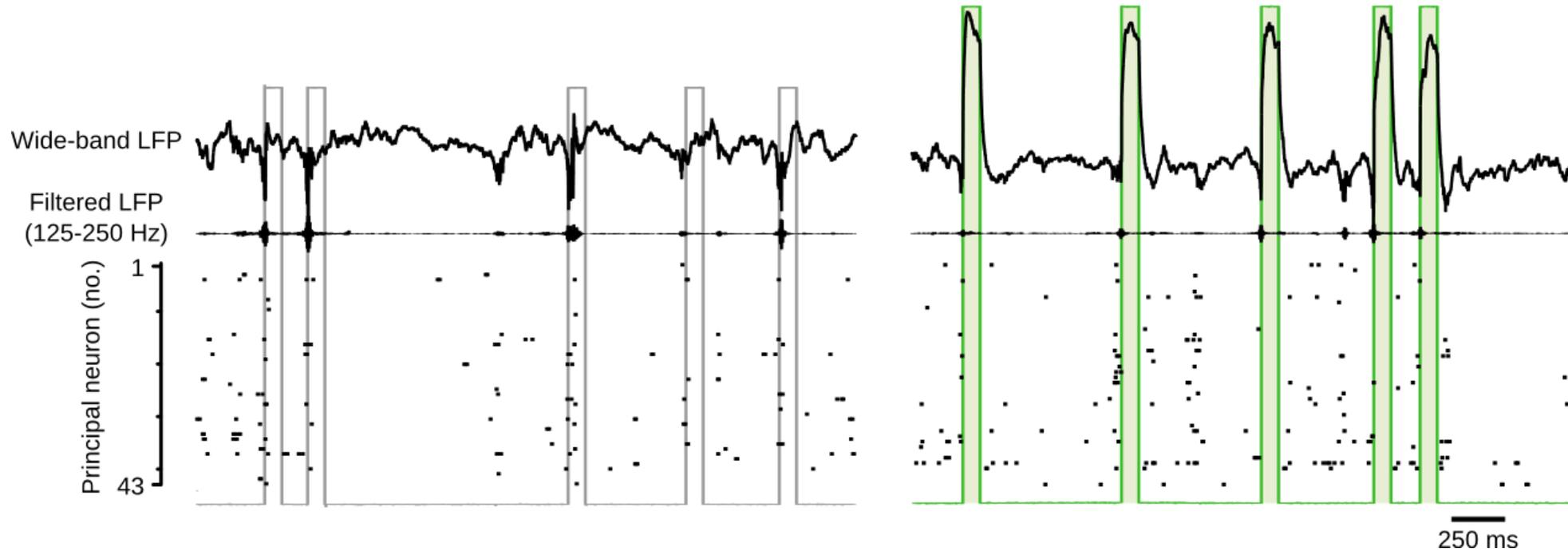
Selective disruption of reactivation: *optogenetic SWR silencing*



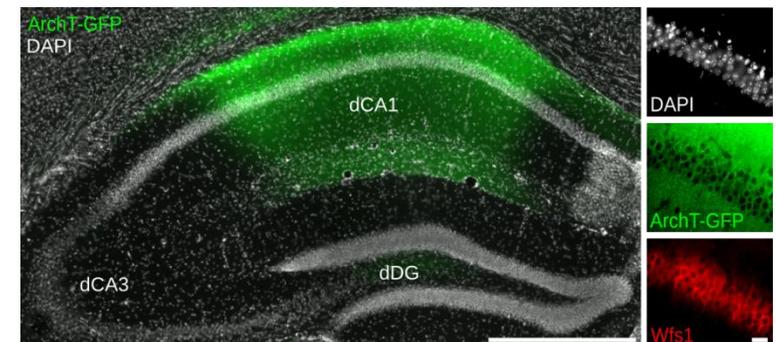
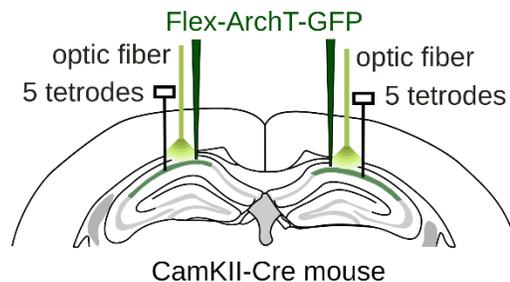
Optogenetic silencing of CamKII-positive cells using ArchT



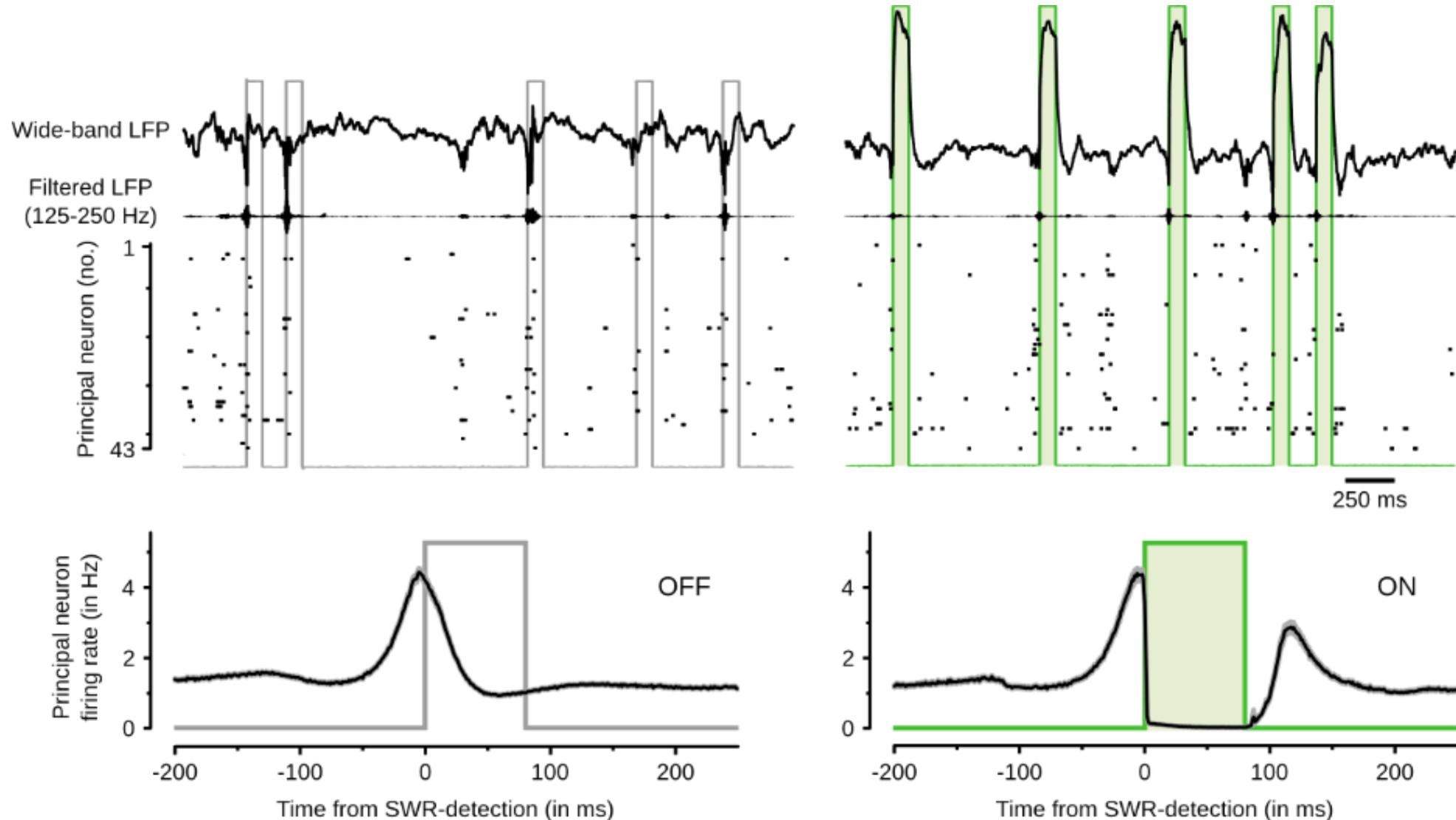
Selective disruption of reactivation: *optogenetic SWR silencing*



Optogenetic silencing of CamKII-positive cells using ArchT

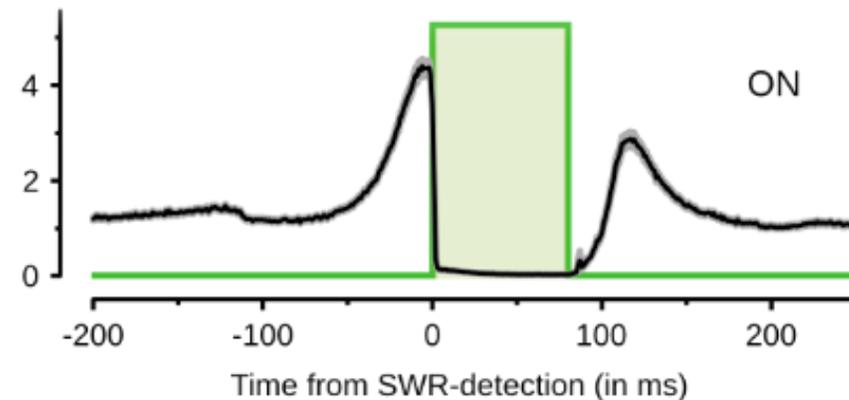
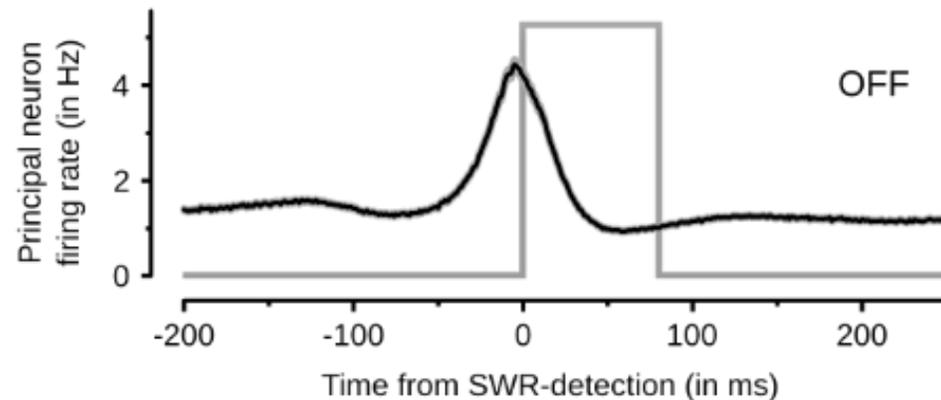
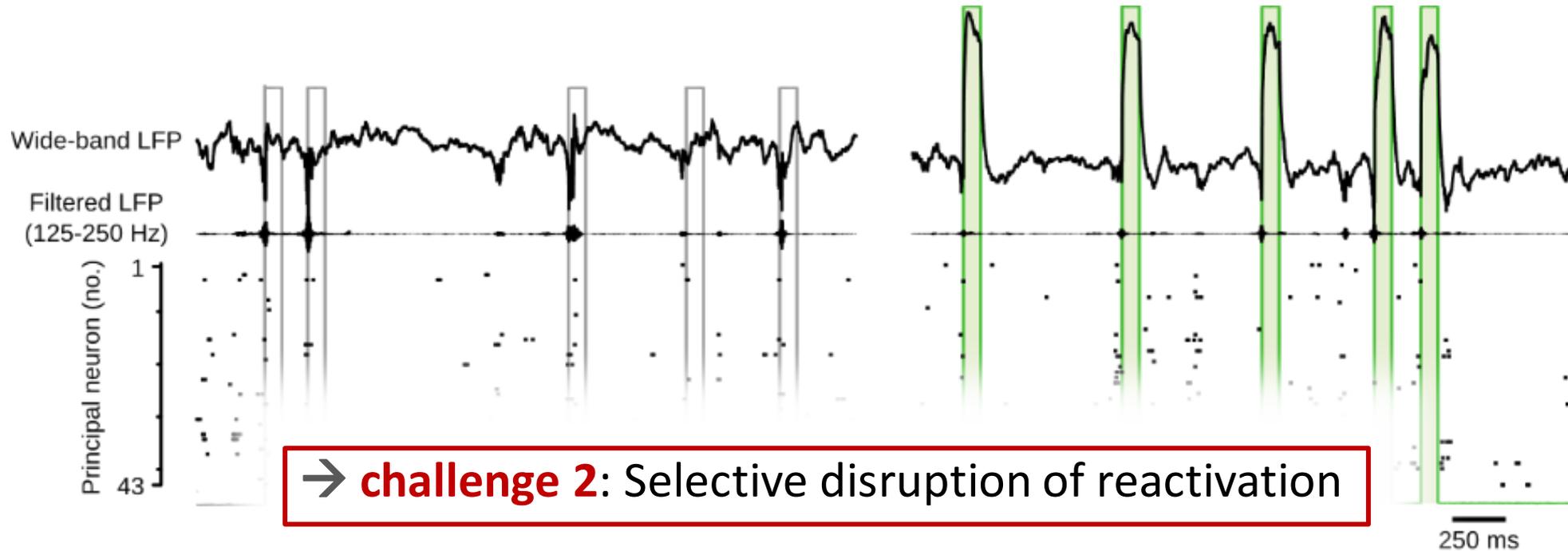


Selective disruption of reactivation: *optogenetic SWR silencing*



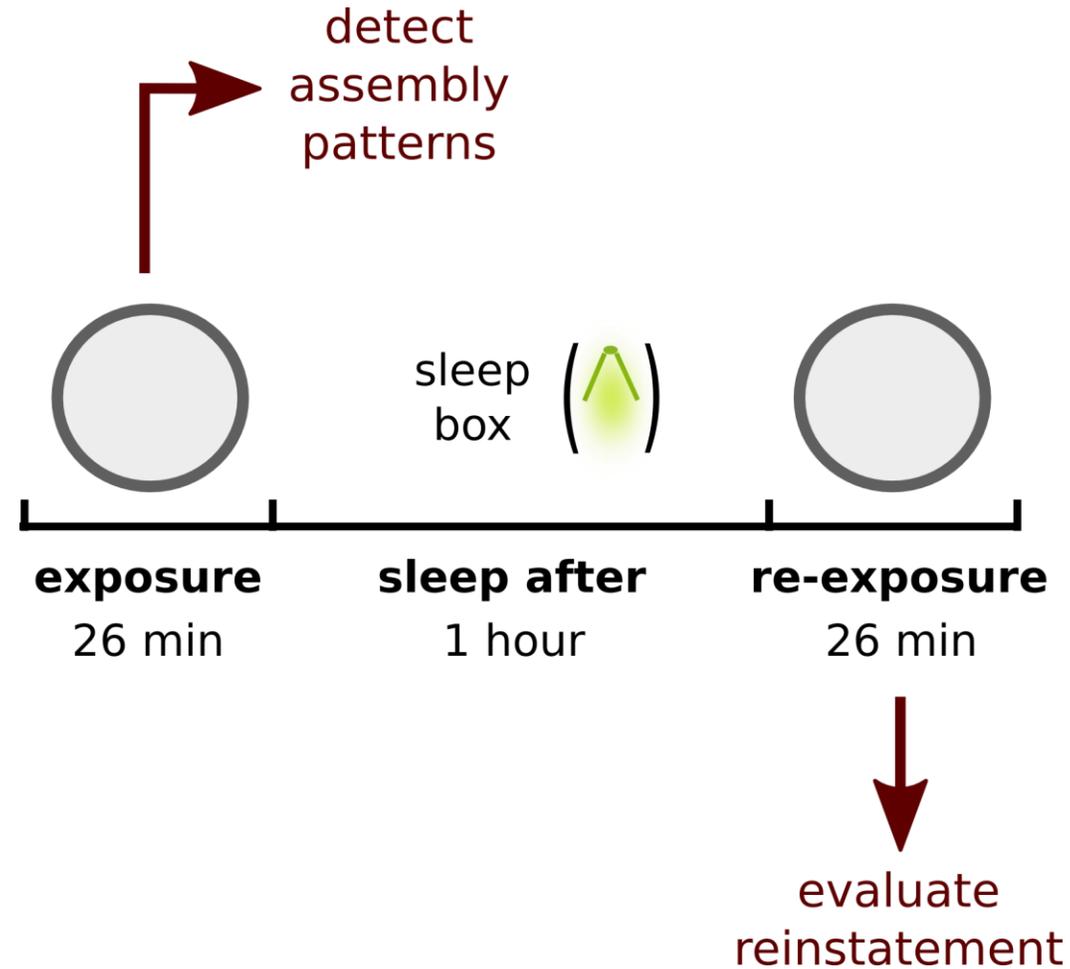
OFF: $n = 1,988$ neurons (from 43 sessions)
ON: $n = 1,527$ neurons (from 37 sessions)

Selective disruption of reactivation: *optogenetic SWR silencing*

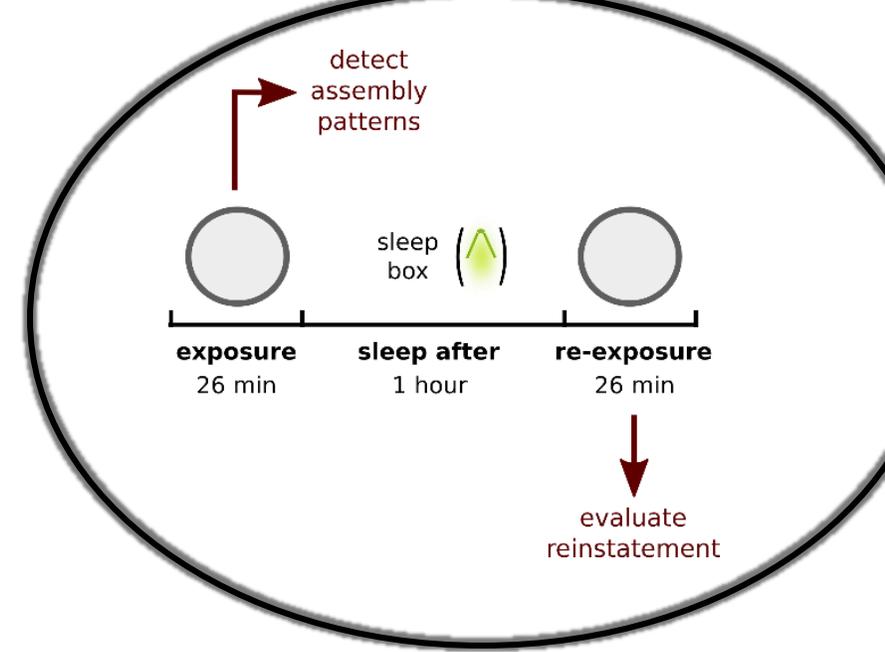
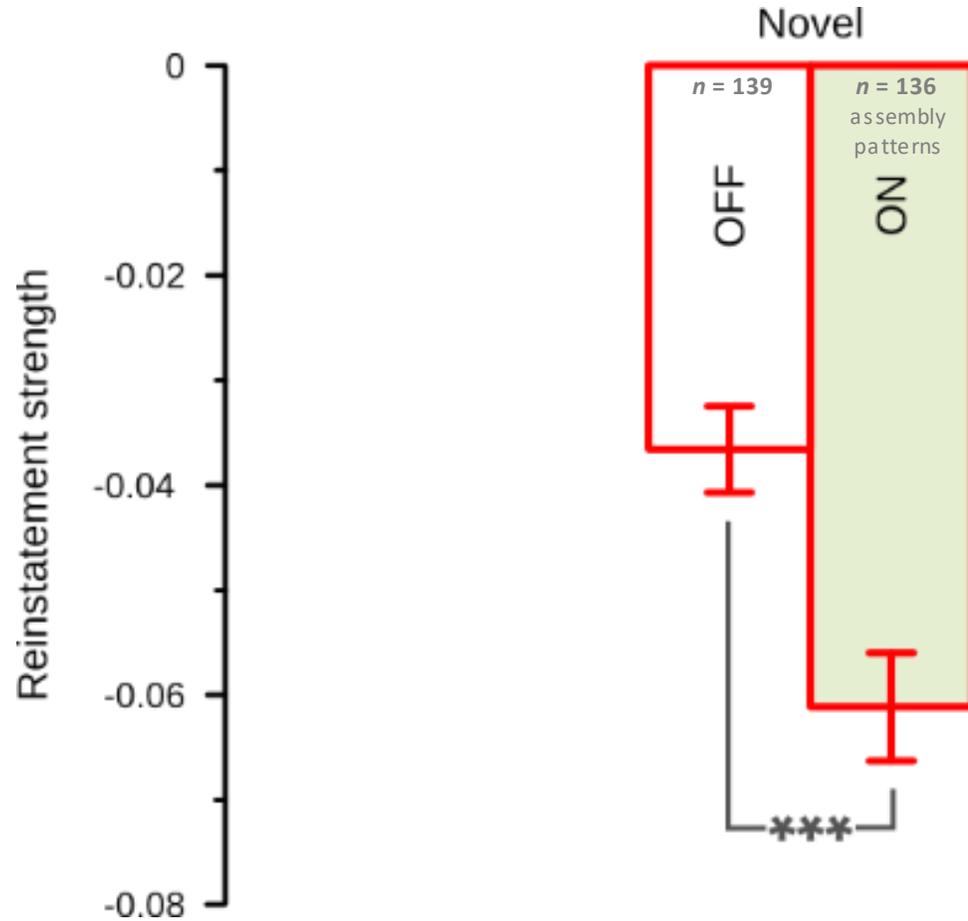


OFF: $n = 1,988$ neurons (from 43 sessions)
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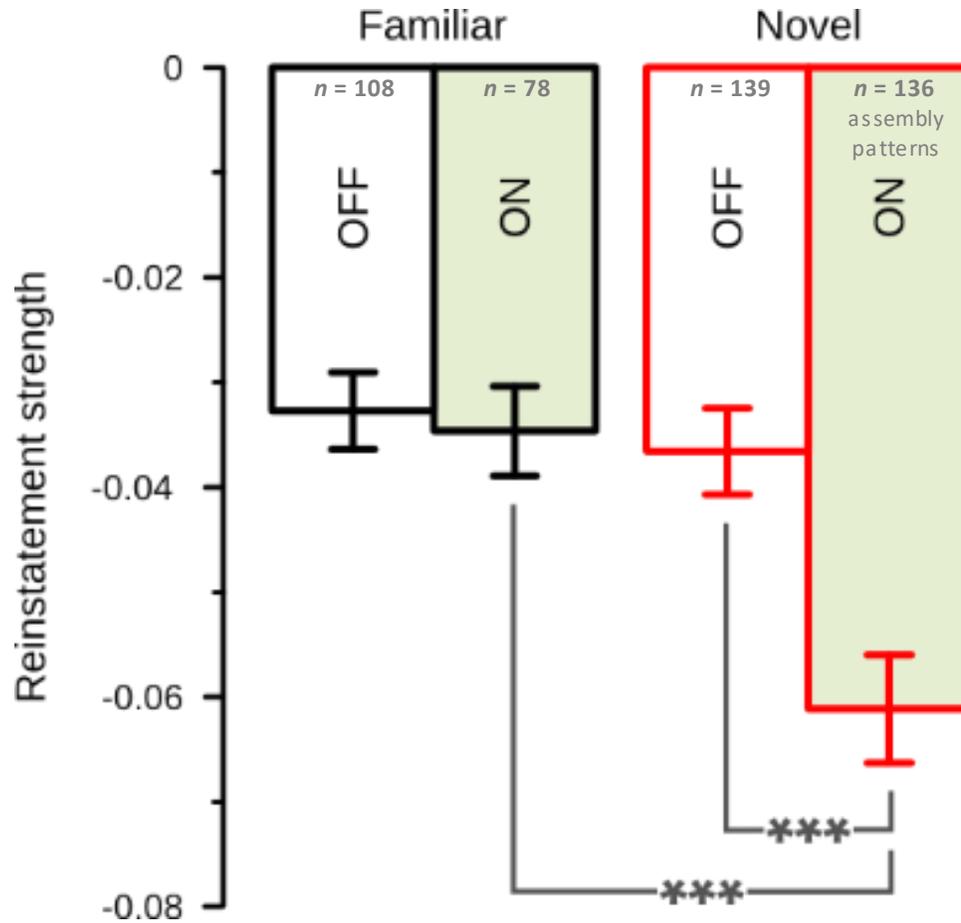
Experimental protocol - (*causation*)



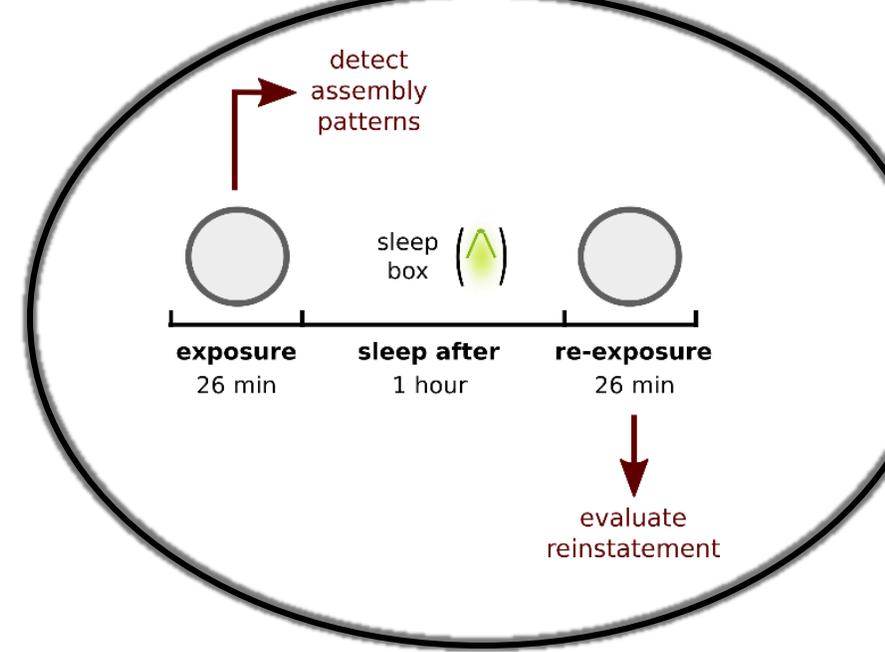
SWR-silencing impairs assembly pattern reinstatement



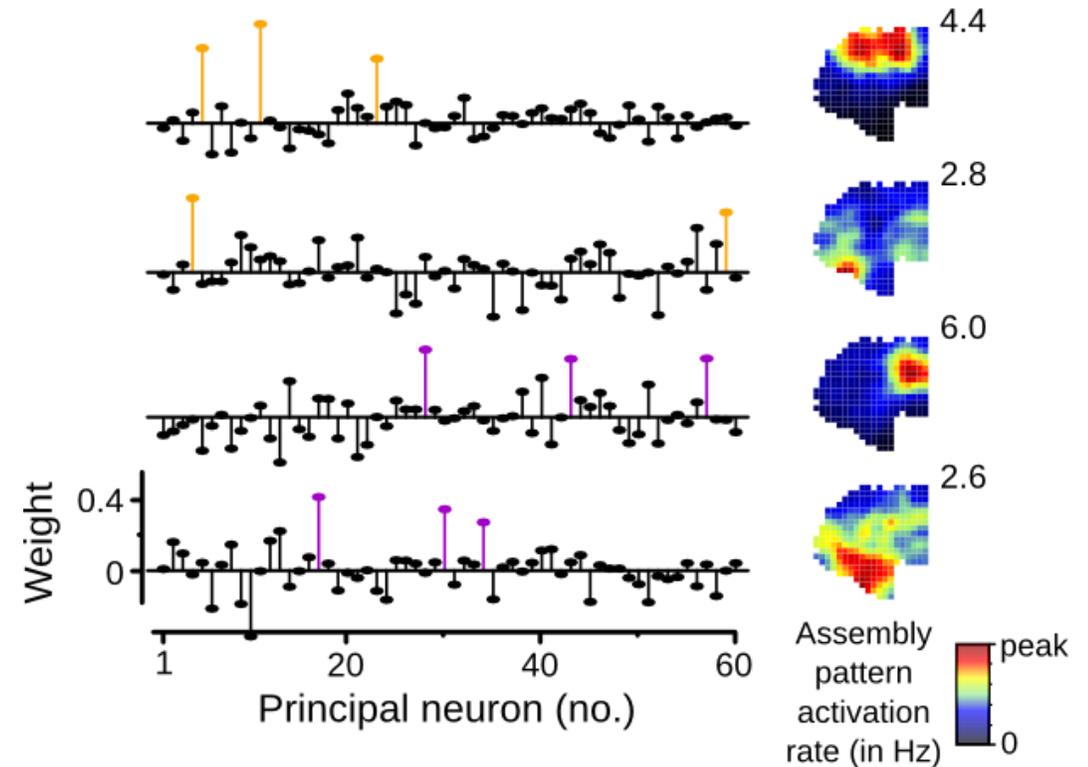
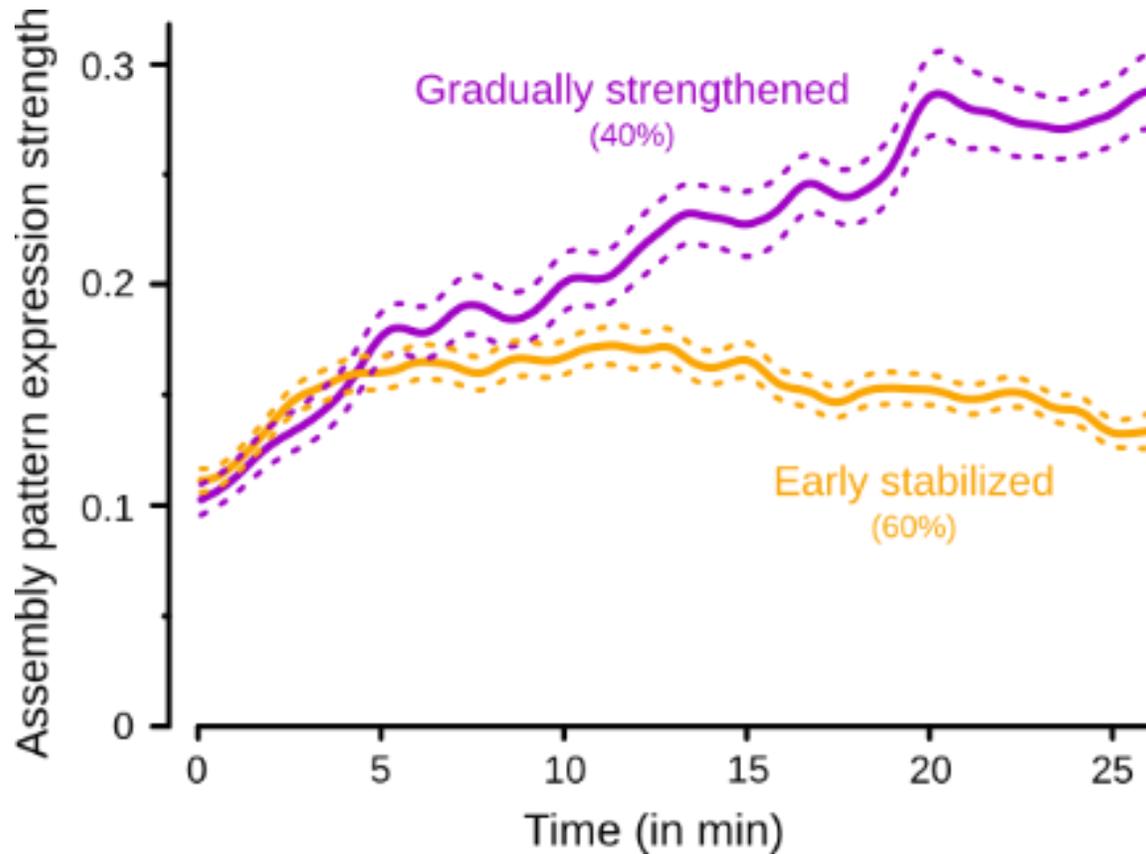
SWR-silencing impairs assembly pattern reinstatement



interaction SWR-silencing x enclosure type:
 $F(1,318) = 5.05, P < 0.05$

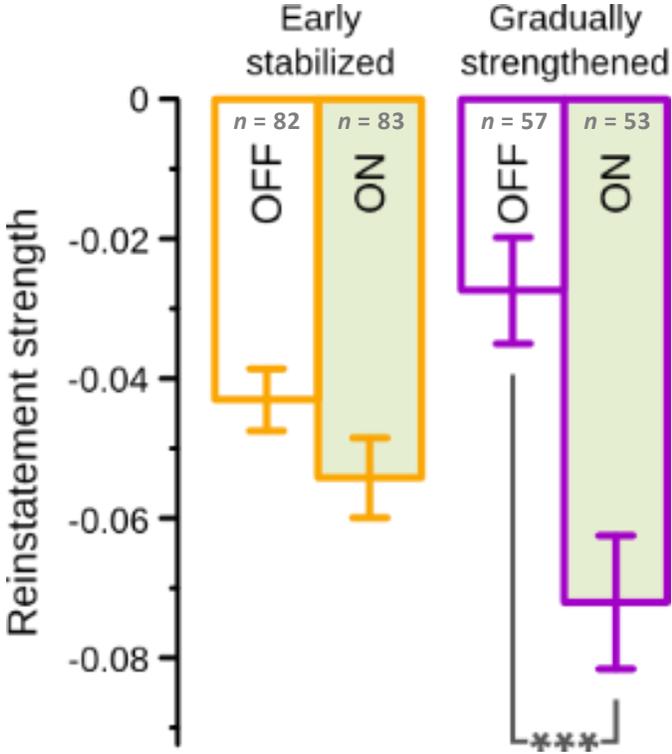
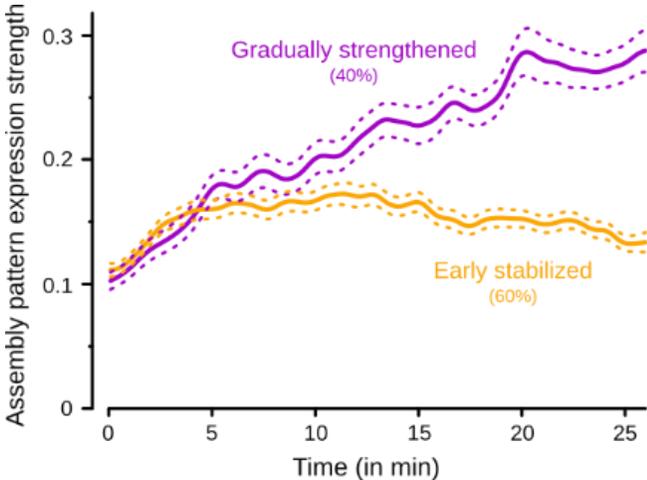


Only stability of gradually strengthened patterns requires offline reactivation



Gradually strengthened: $n = 134$ assembly-patterns
Early stabilized: $n = 201$ assembly-patterns
(based on 50 recording-blocks from 8 mice)

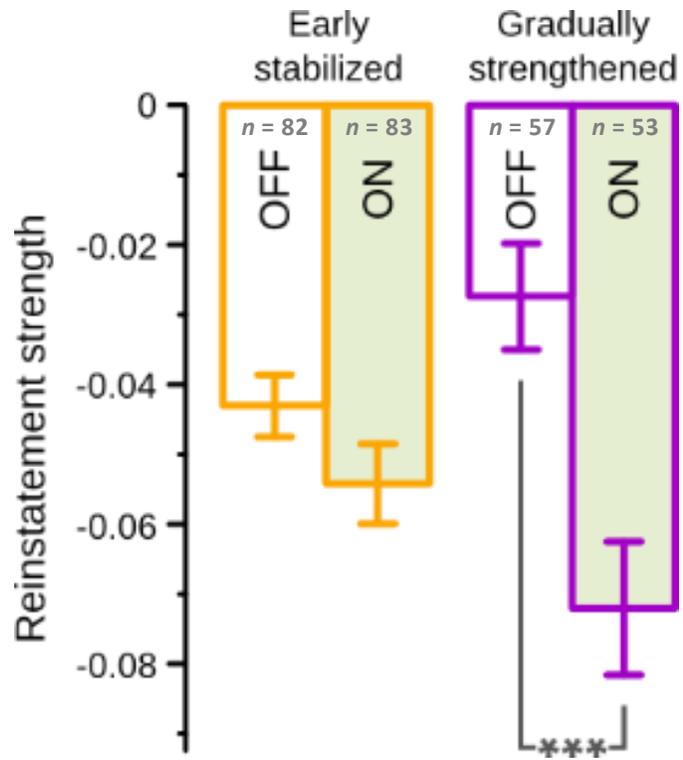
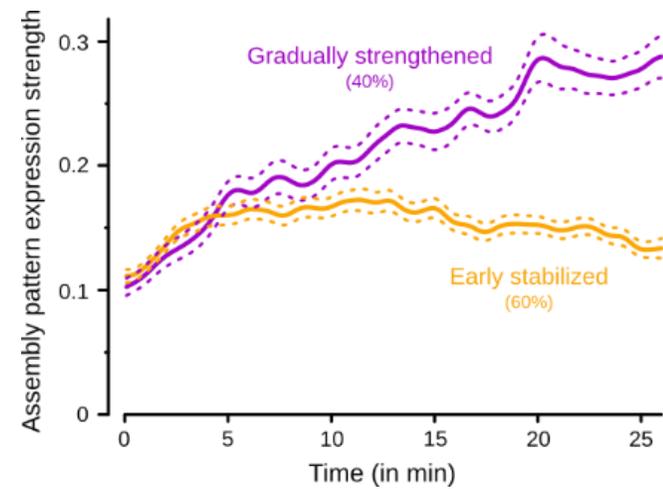
Only stability of gradually strengthened patterns requires offline reactivation



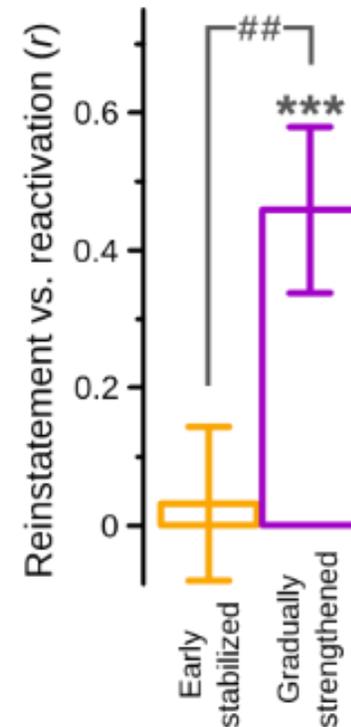
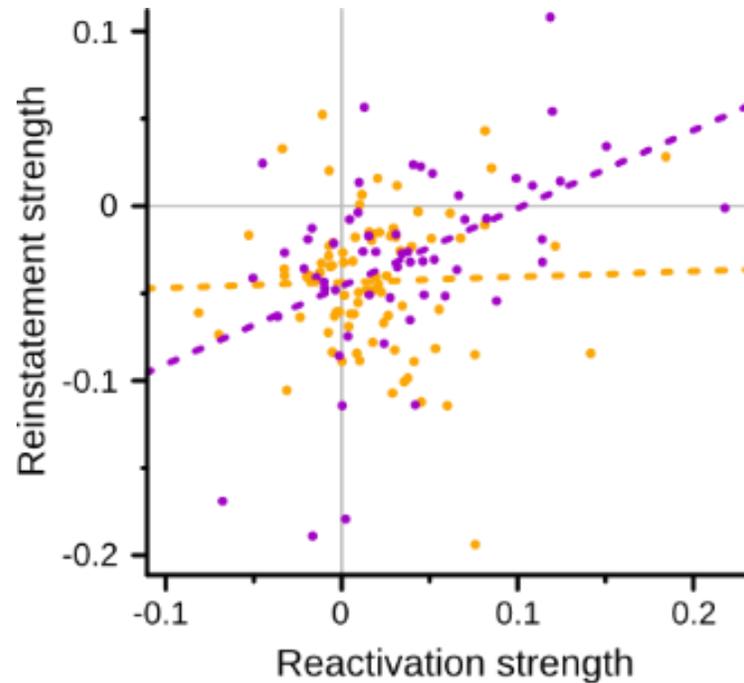
interaction SWR-silencing x pattern type:
 $F(1,271) = 6.28, P < 0.05$

(based on 50 recording-blocks from 8 mice)

Only stability of gradually strengthened patterns requires offline reactivation



interaction SWR-silencing x pattern type:
 $F(1,271) = 6.28, P < 0.05$



One-sentence summary

The stability of “Hebbian-like” cell assembly patterns, which are gradually strengthened during their initial expression, depends on their offline reactivation.

Acknowledgements

Dupret lab

David Dupret

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Mohamady El-Gaby

Stephen McHugh

Visualizing tetrode tracks

Ben Micklem

Linda Katona

Research support

Jane Janson

Liz Norman

Lisa Conyers

Katharine Whitworth

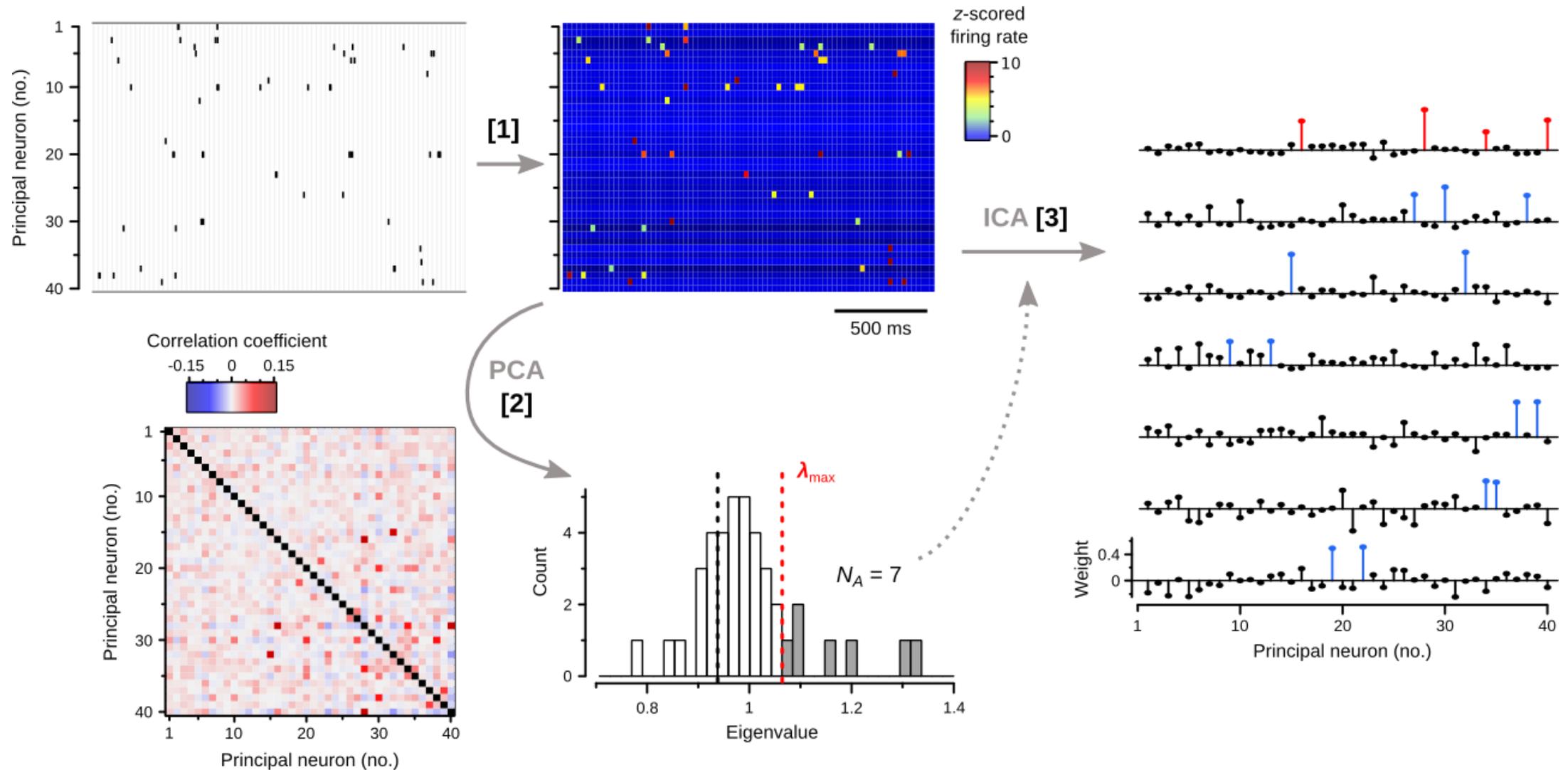
Kristina Wagner

Heidelberg

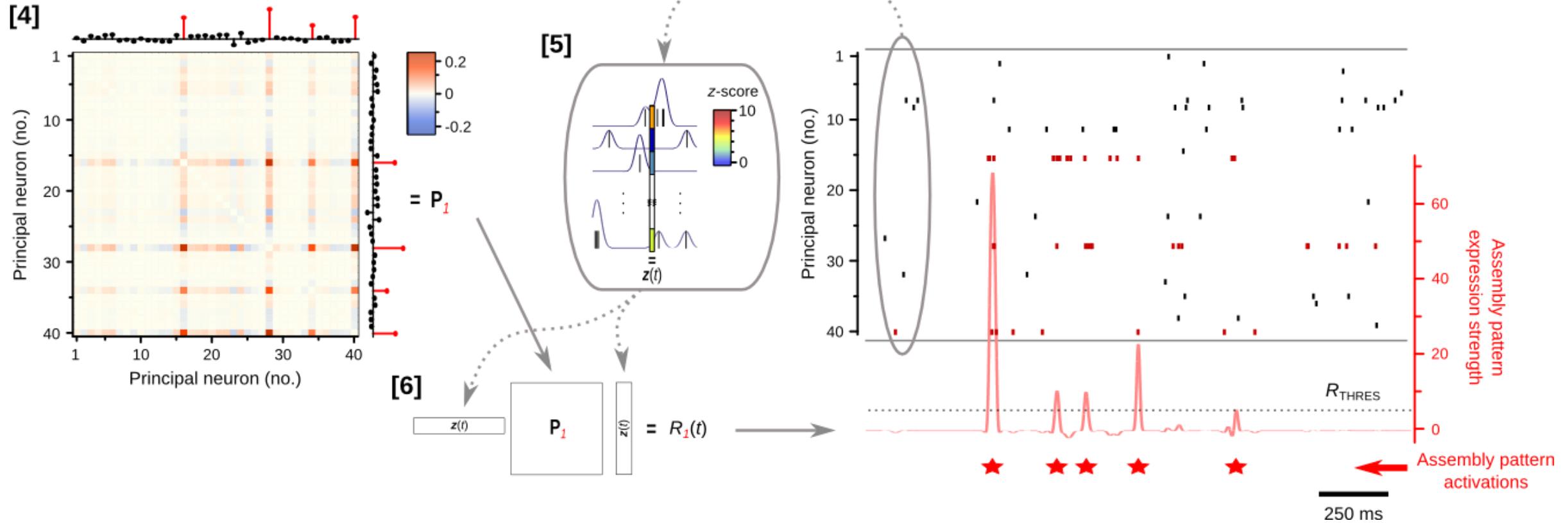
Kevin Allen (real-time SWR detection)



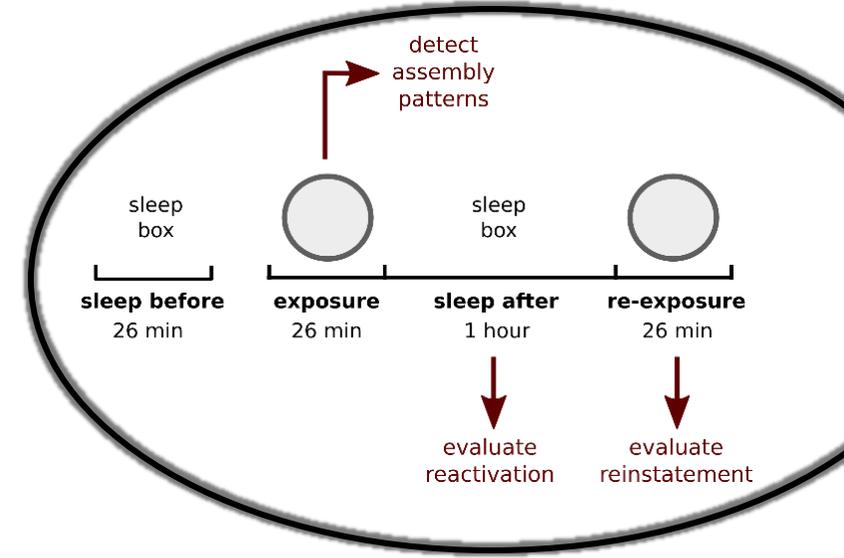
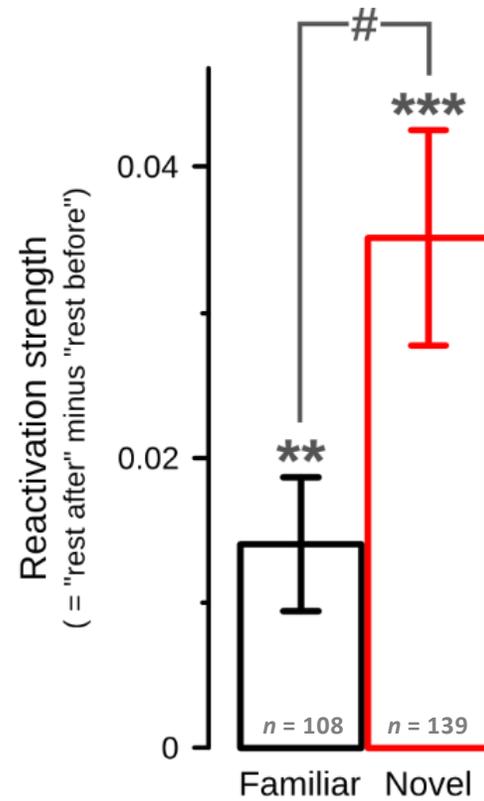
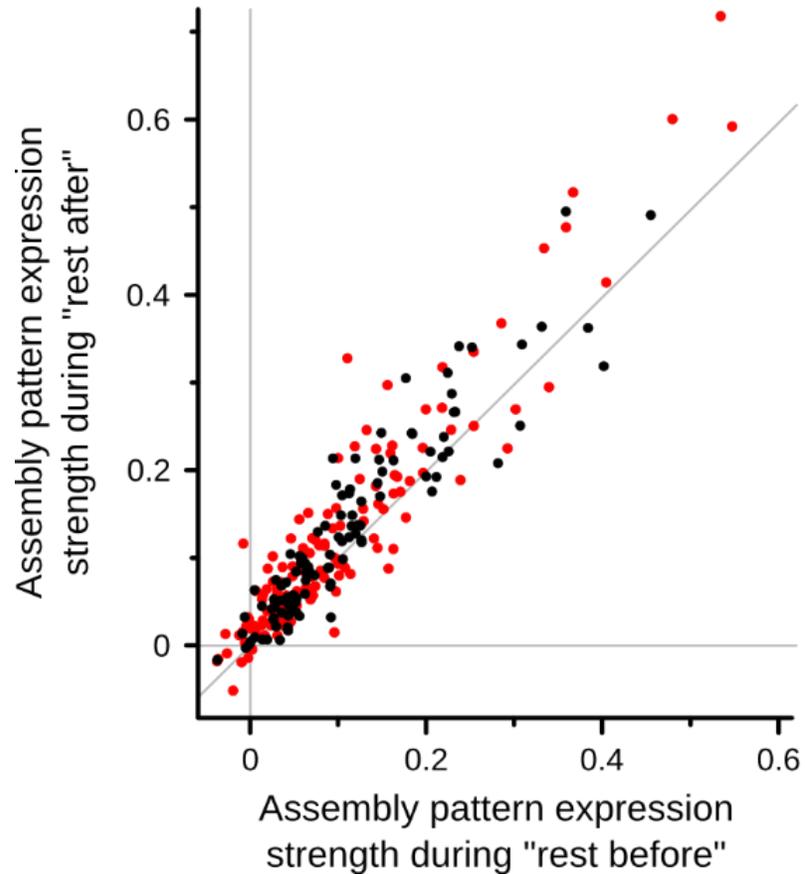
Assembly pattern identification



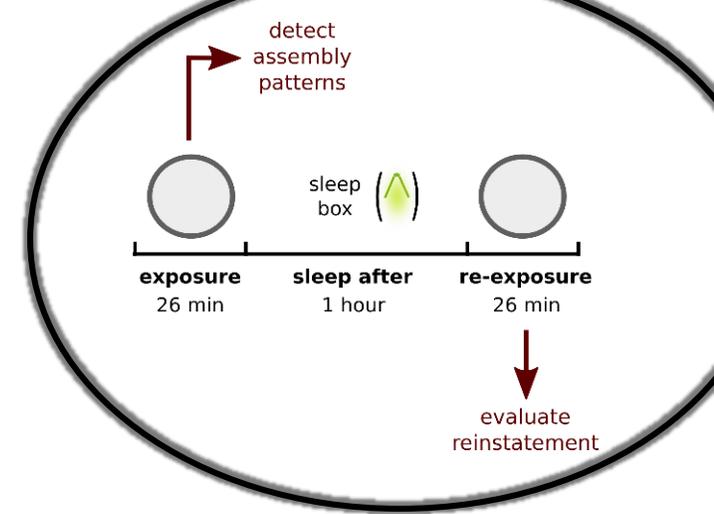
Assembly pattern tracking



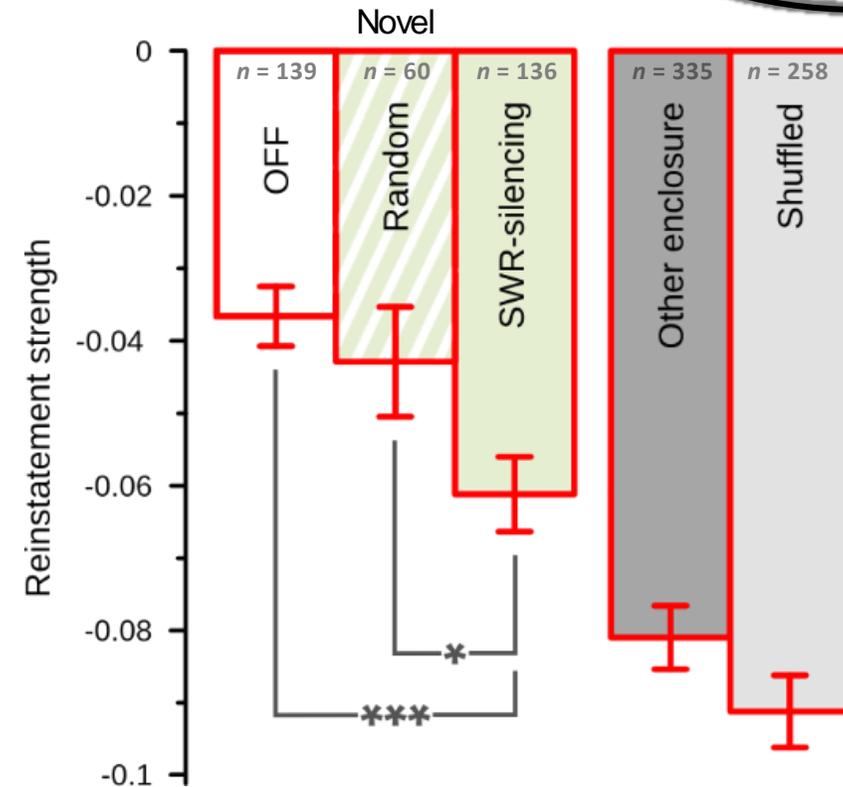
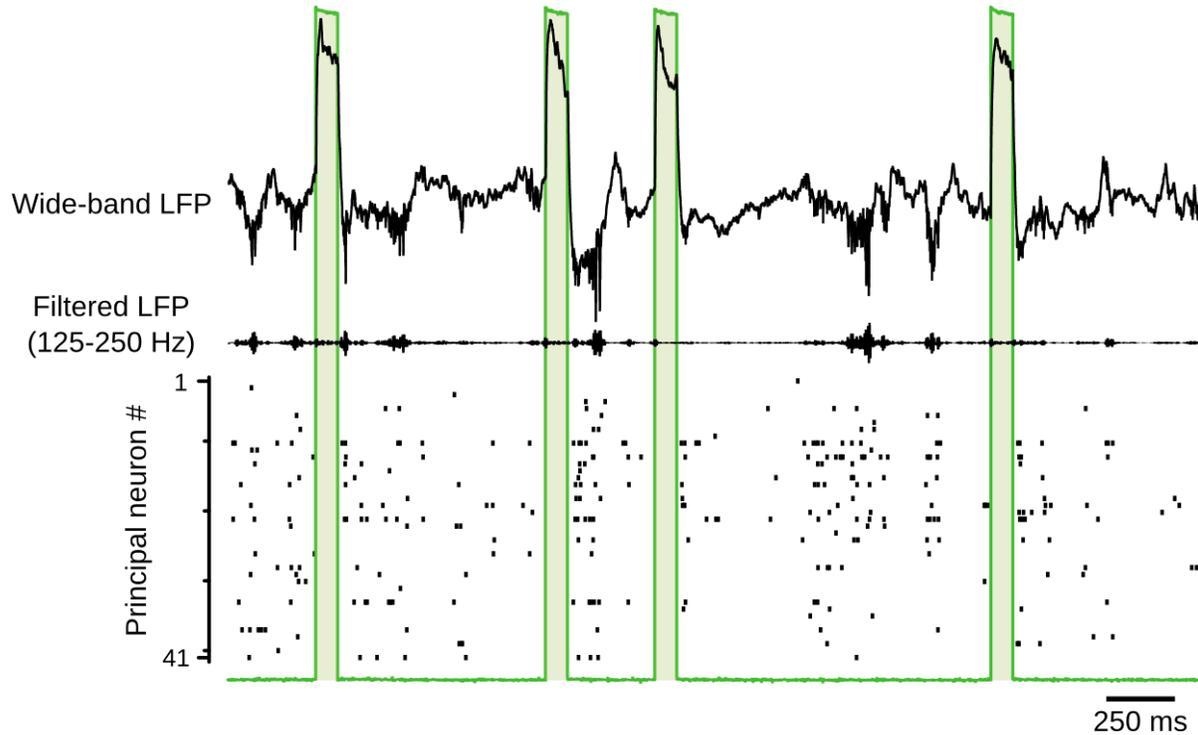
Assembly pattern reactivation



Control experiment: random silencing

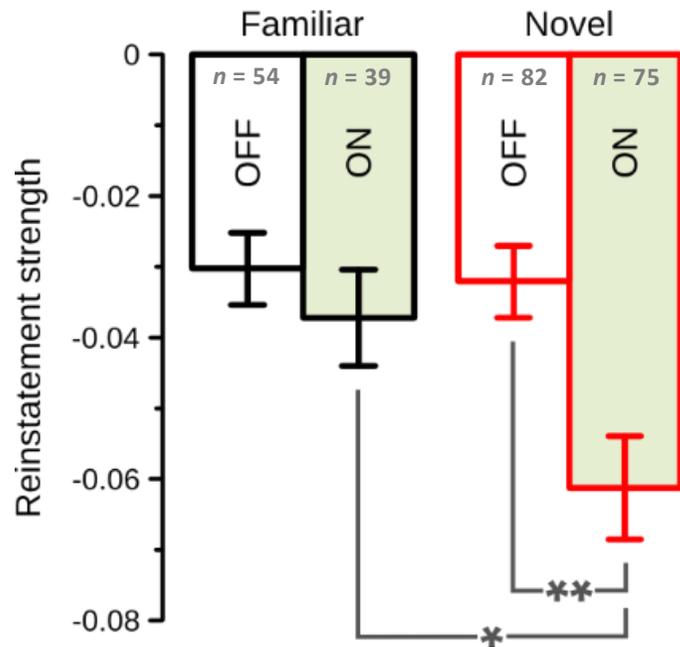


Random silencing control experiment

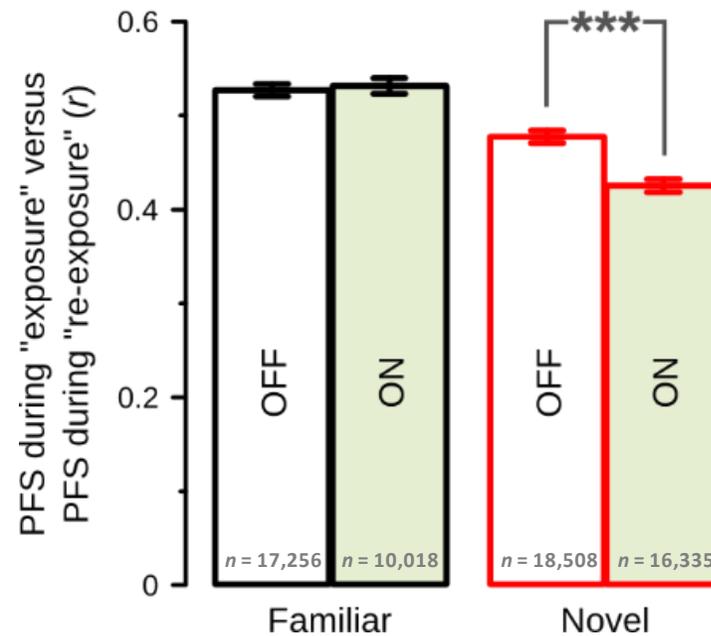


Effect of SWR-silencing: control analyses

"Across-tetrodes only" analysis



Neuron-pair analysis



Single-neuron analysis

