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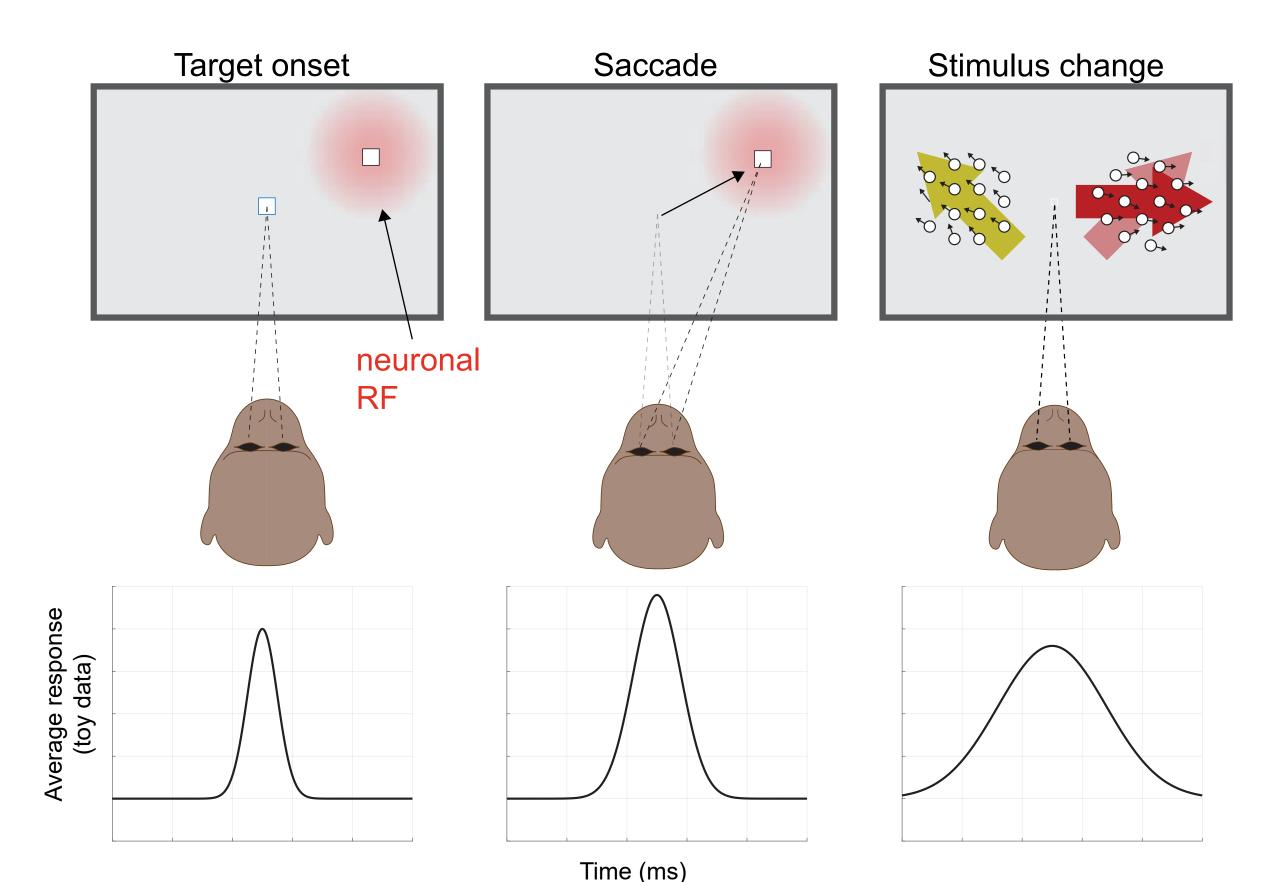
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Introduction / Background

- The primate superior colliculus (SC) responds stimulus onset(s), saccade(s) and subtle stimulus changes with phasic increases in activity.
- Do SC activity patterns discriminate these events or does interpretation of SC output rely on other structures?
- Here we present a method for quantifying and comparing population encoding of task variables using partial least squares discriminant analysis (PLS-DA).

How are visual and behavioral events encoded in the SC?

 SC responds to diverse events with transient increases in average activity.

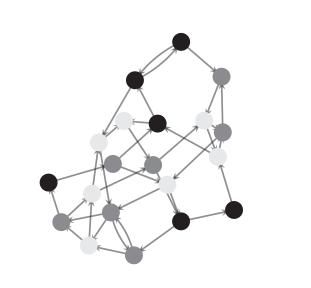


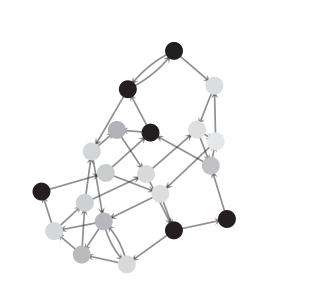
Do underlying activity patterns discriminate events?

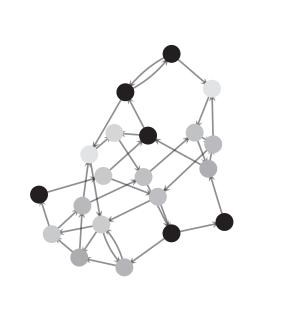
Similar activation patterns do not discriminate events.

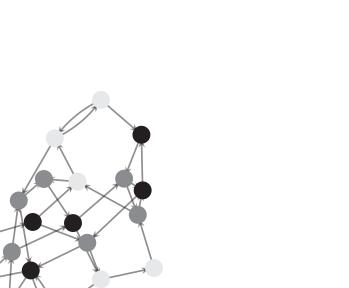
Distinct activation patterns do

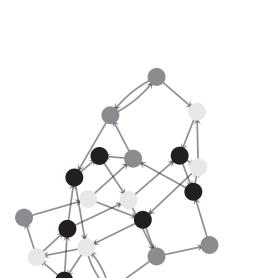
discriminate events.







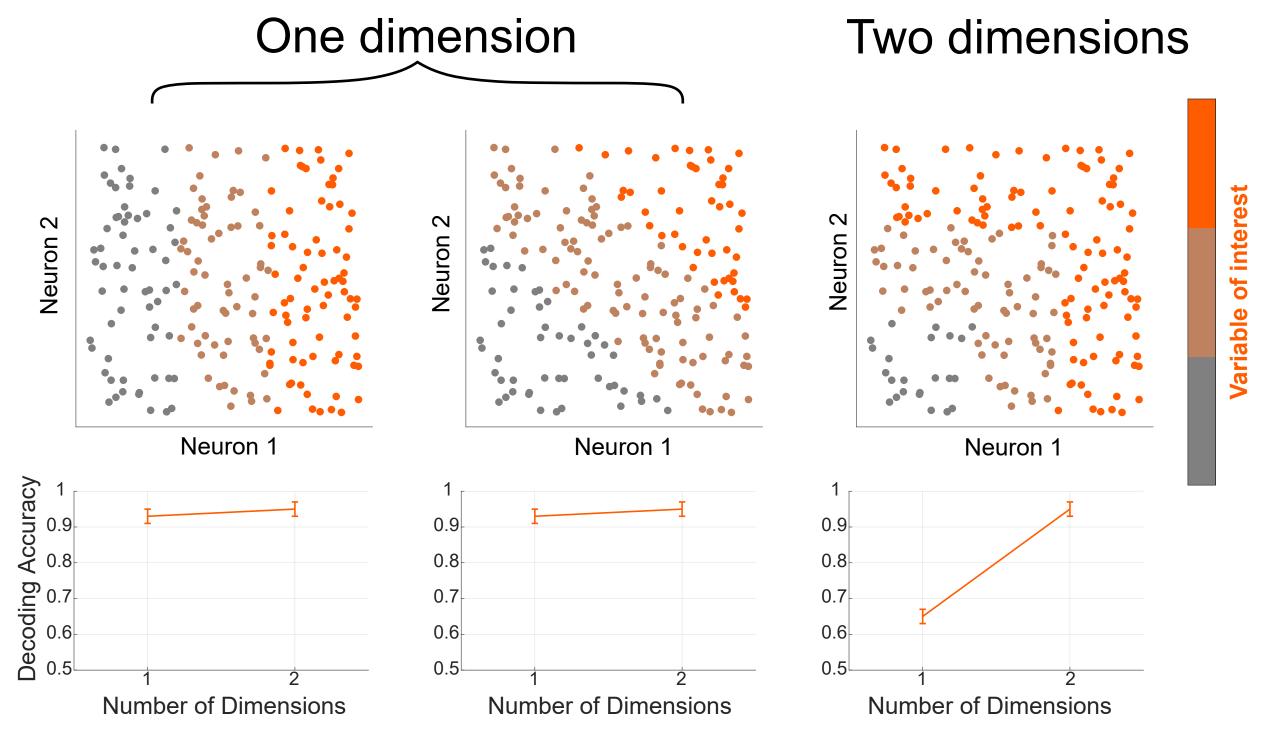




Quantifying population encoding patterns: a 2D toy example

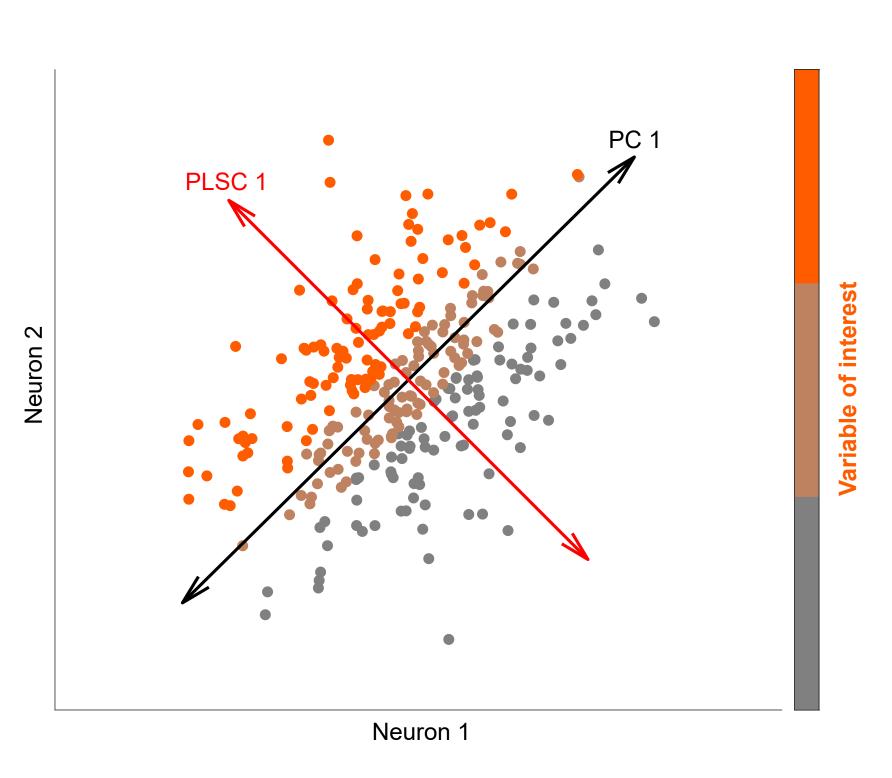
 A variable of interest may be encoded in one or two dimensions of a 2D population.

 How many dimensions of activity are needed to reach the decoding accuracy available in the full population?



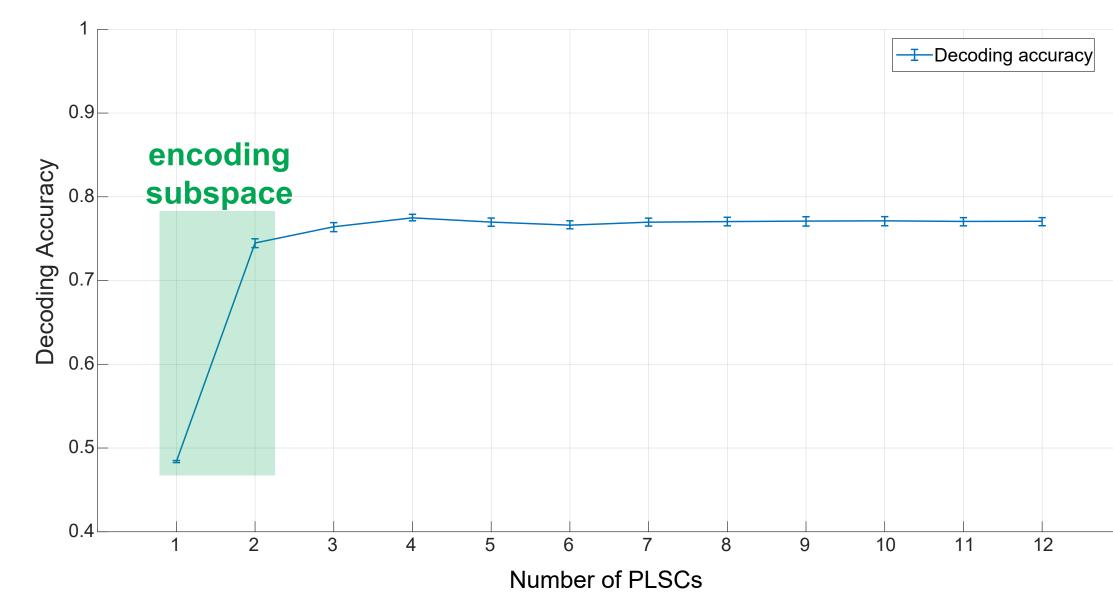
How can we intelligently select a minimal subset of encoding dimensions?

Partial least squares discriminant analysis (PLS-DA)



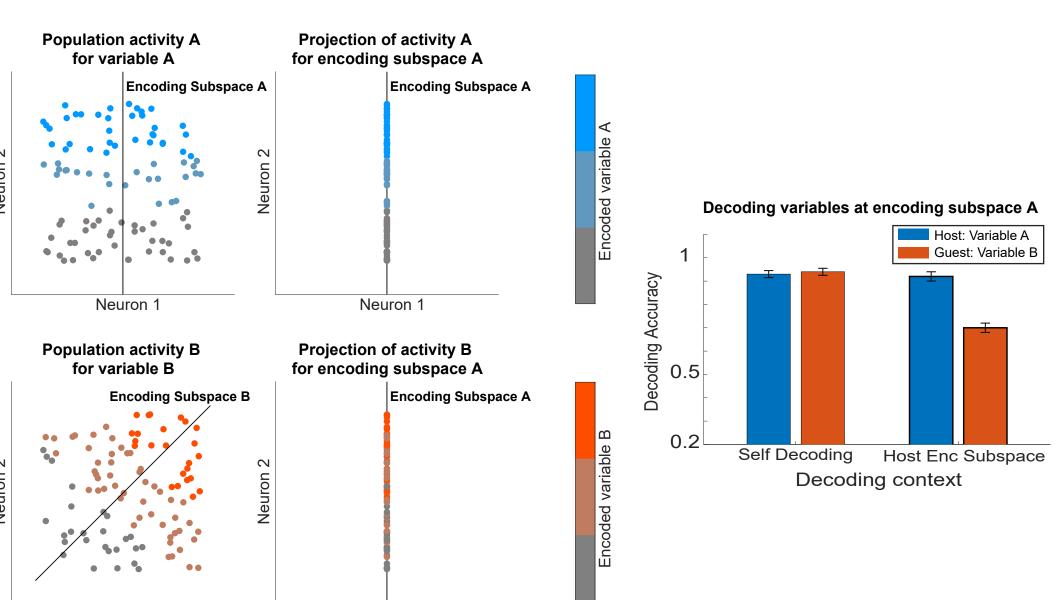
- PLS-DA is an iterative, supervised method that identifies components (dimensions) of the input that maximize covariance between input features and class labels.
- The identified dimensions capture the most relevant information for **prediction** while **reducing dimensionality**.
- Difference from principal component analysis (PCA): PCA maximizes variance in the data without considering class labels.

Identifying an encoding subspace using PLS-DA



- Individual labeled trials are used.
- Decoding is performed while incrementally increasing the count of PLS components (PLSCs).
- The encoding subspace is the subset of PLS components where decoding accuracy plateaus.

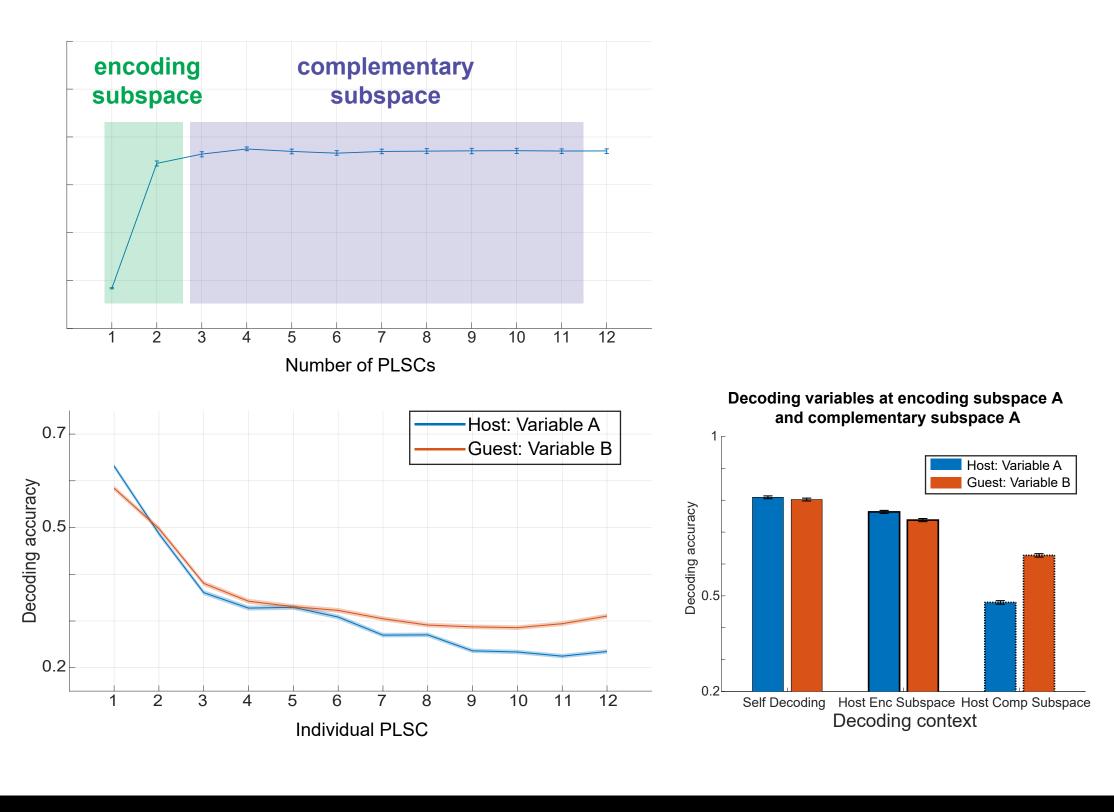
Comparing encoding of two variables using an identified subspace



- We identify the encoding subspace of variable A (the host), called encoding subspace A.
- Host: Variable A Guest: Variable B as the guest.

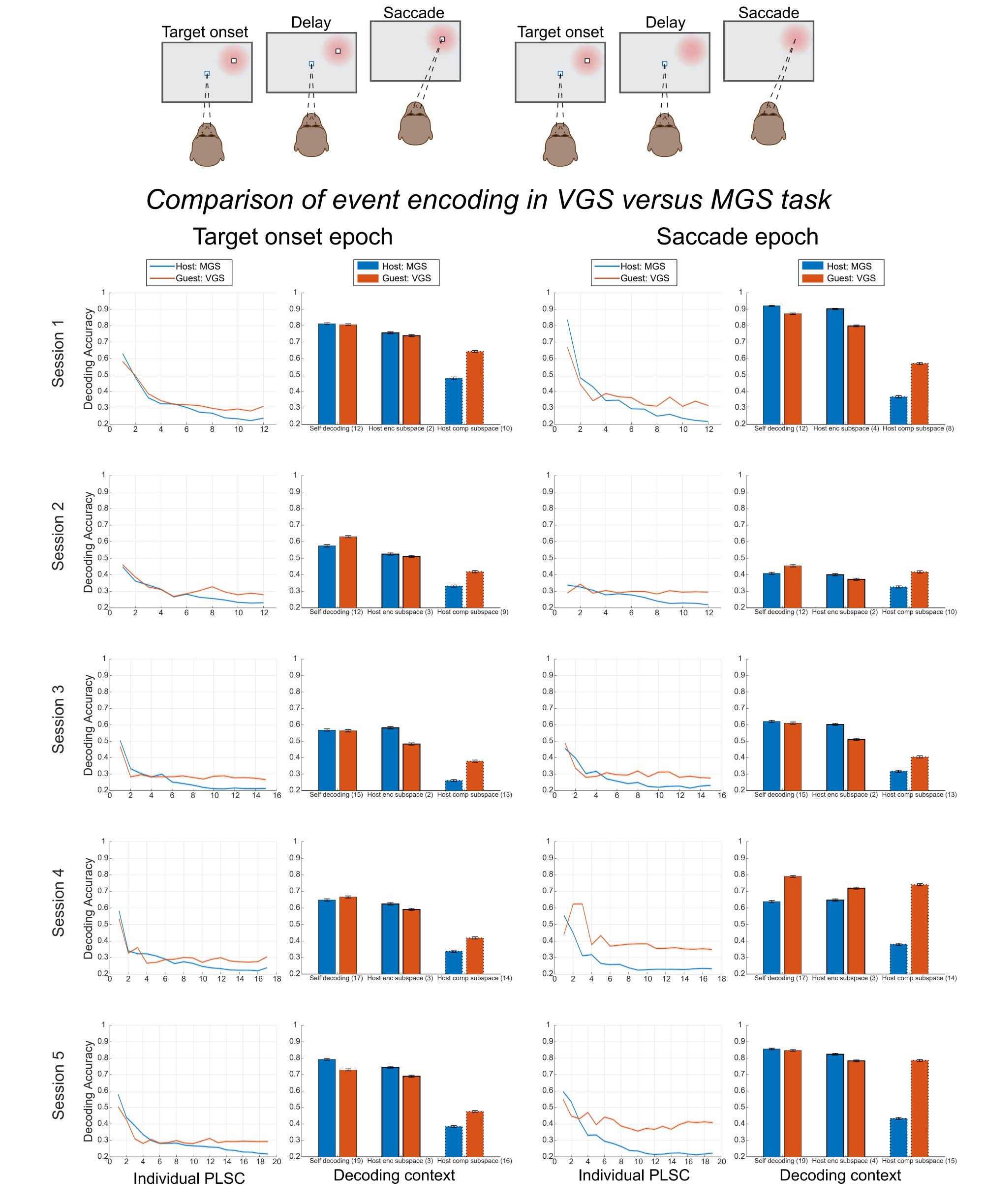
 We define Variable B as the guest.
 - Population activity for variables A and B is projected onto encoding subspace A.
 - Variables A and B are decoded within encoding subspace A to estimate the similarity or difference in the neural population's encoding of each variable.

What about the other dimensions?



- Components not in the encoding subspace form a complementary subspace.
- Is the complementary subspace a "null" space? Or does it contain information?
- For a detailed view of how information is distributed across dimensions, we can decode from individual dimensions.

Visually guided saccade (VGS) and memory-guided saccade (MGS) tasks



- **Target location** was decoded from a target onset epoch (+50 to +200 ms) and a saccade epoch (-25 to +75 ms). Locations varied between sessions, but each session used a single set of 4 locations for both VGS and MGS tasks.
- Decoding accuracy in the first few dimensions are most consistent, but more differences emerge in the later dimensions.
- Encoding differences between tasks divergence more prominently in the saccade epoch compared to the target onset epoch.
- We suspect variations across sessions reflect functional differences in recorded neuronal populations.

Conclusion

- We propose using the "encoding subspace" to investigate neuronal population coding.
- PLS-DA offers a principled approach to defining the encoding subspace.
- Superior colliculus neuronal populations exhibit distinct encoding patterns for the same target locations across different task contexts.