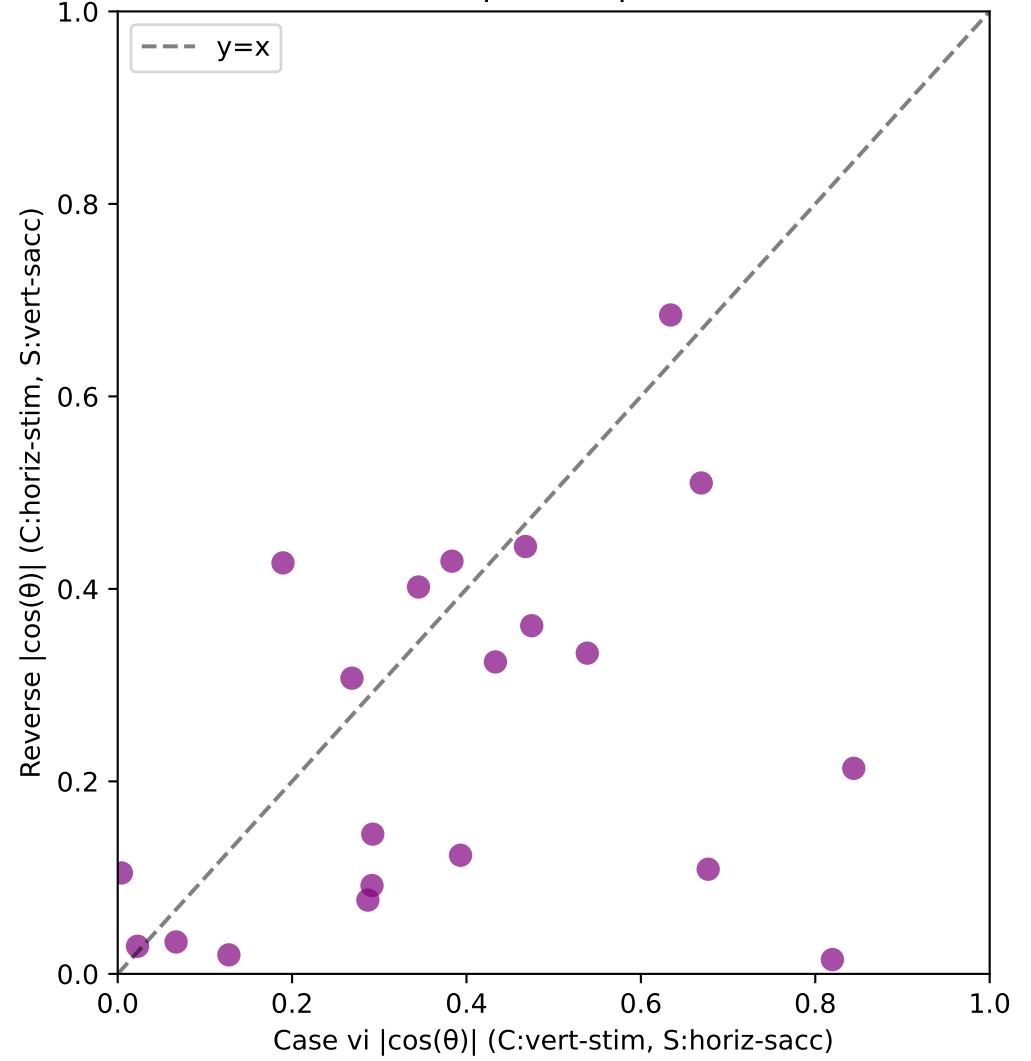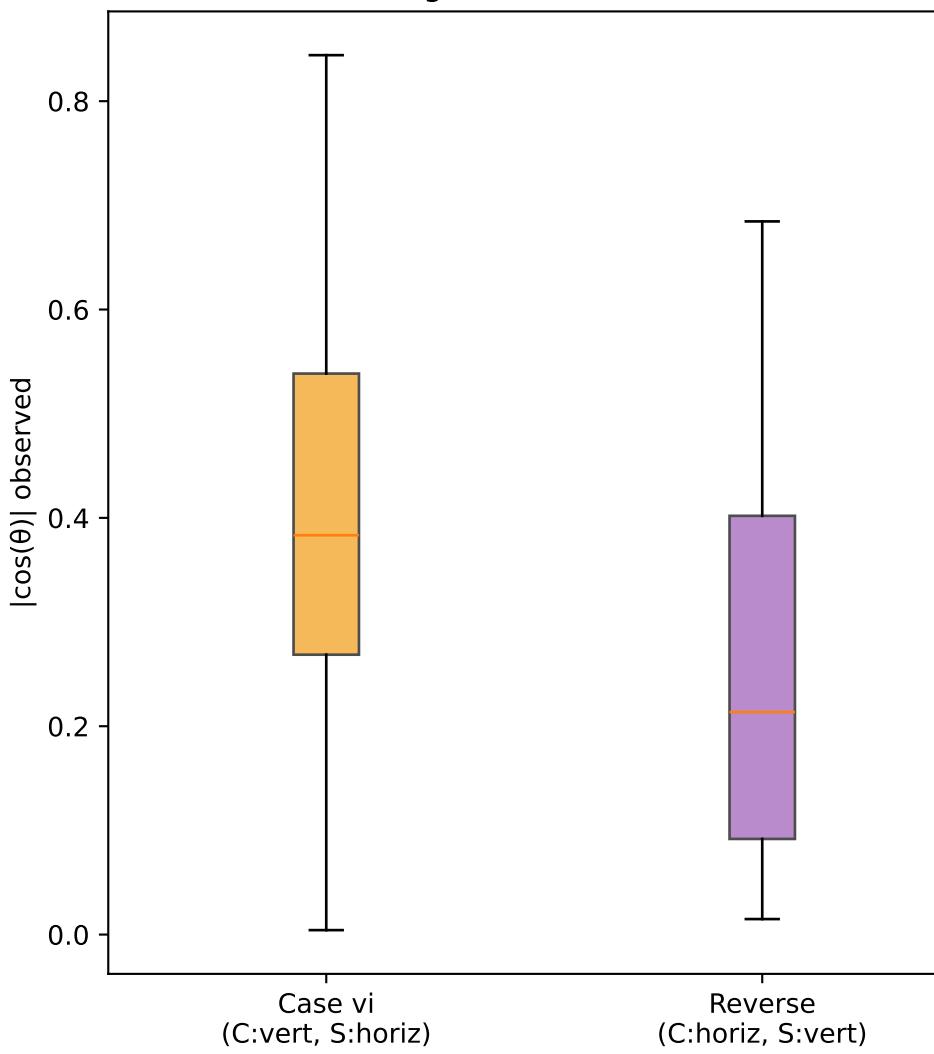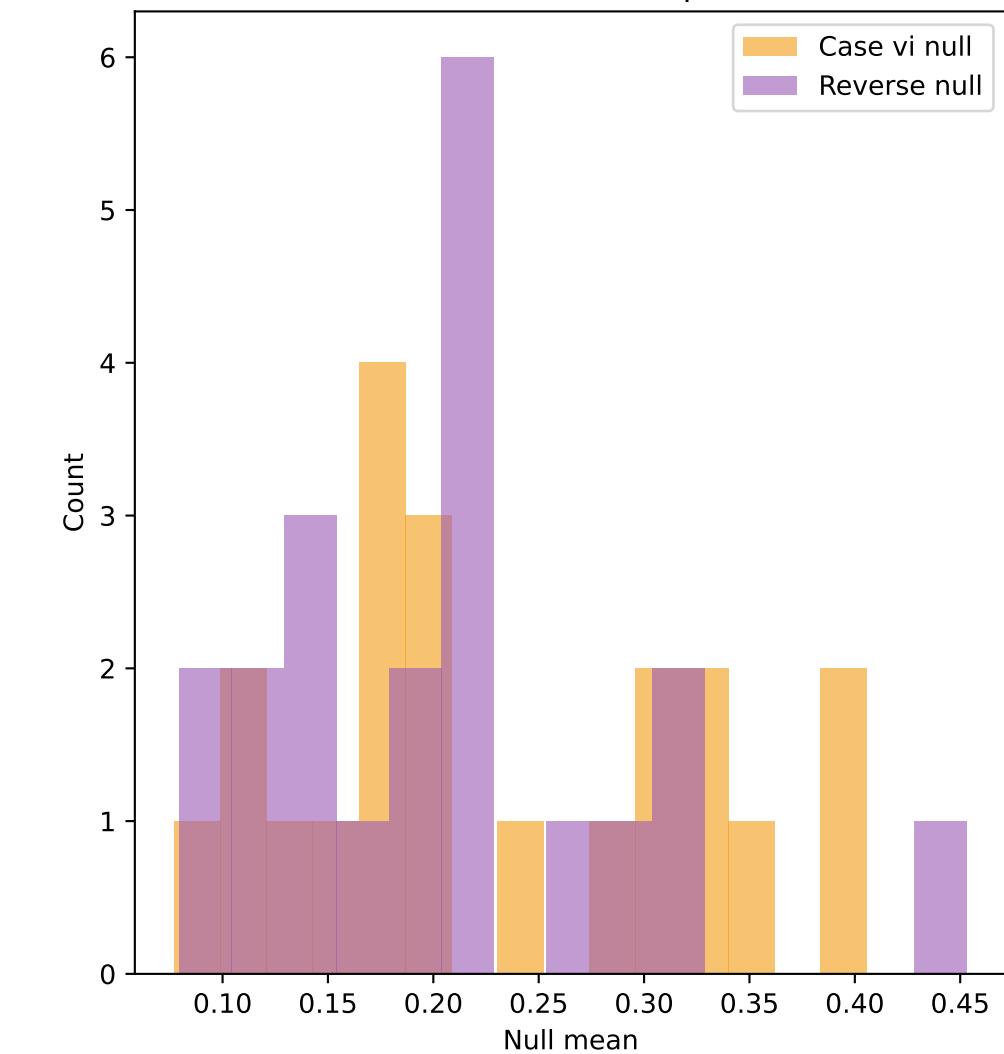**Case vi vs Reverse Case**
r=0.343, paired-t p=0.0163
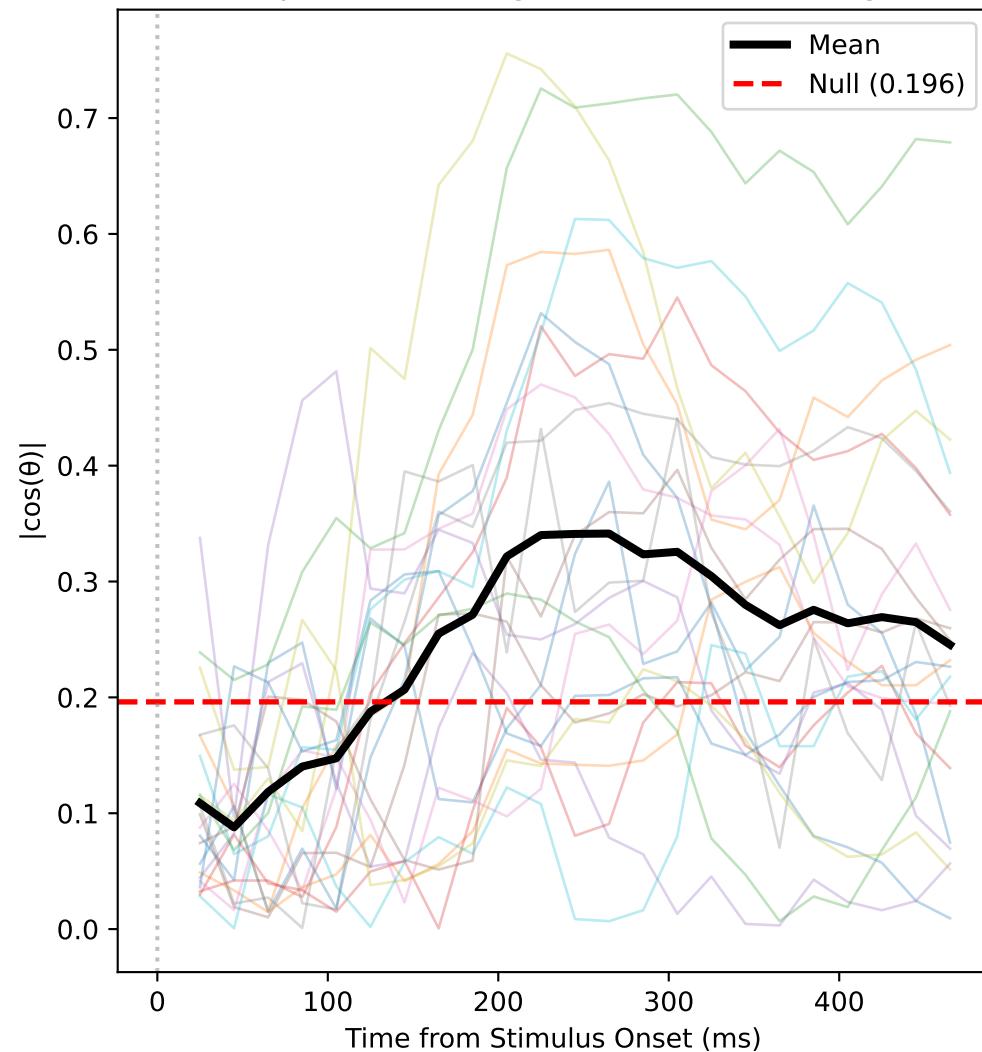
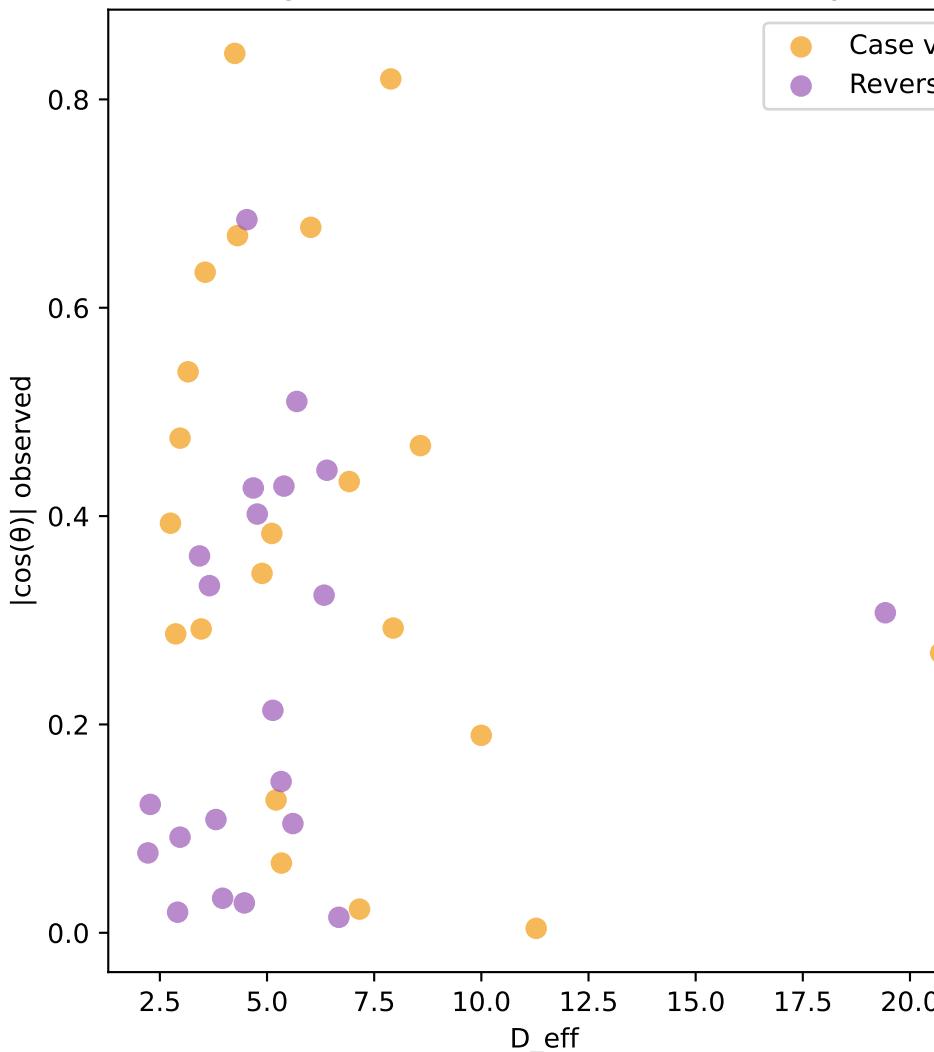**Observed Alignment: Case vi vs Reverse**

Null Distribution Comparison

Temporal Cross-Alignment (Case vi config)

Alignment vs Effective Dimensionality

Alignment Relative to Null