

# Building a hypothesis

Cognitive analysis

Hypothesis testing

Functional  
decomposition  
on

$$A = \{f_{attn}, f_{ment}, \dots, f\{\cdot\}\}$$

$$P_{JA} := \{f_{attn}, f_{ment}\} \subset A$$

Temporal  
dependance

$$P(f, t) = \beta_1 + \beta_2 JA + \beta_3 pJA + \beta_4 JA \cdot pJA$$

Biological  
signature

$$f_{ment} \mapsto P_\beta(TPJ)$$

$$f_{attn} \mapsto NC_{ERP}$$

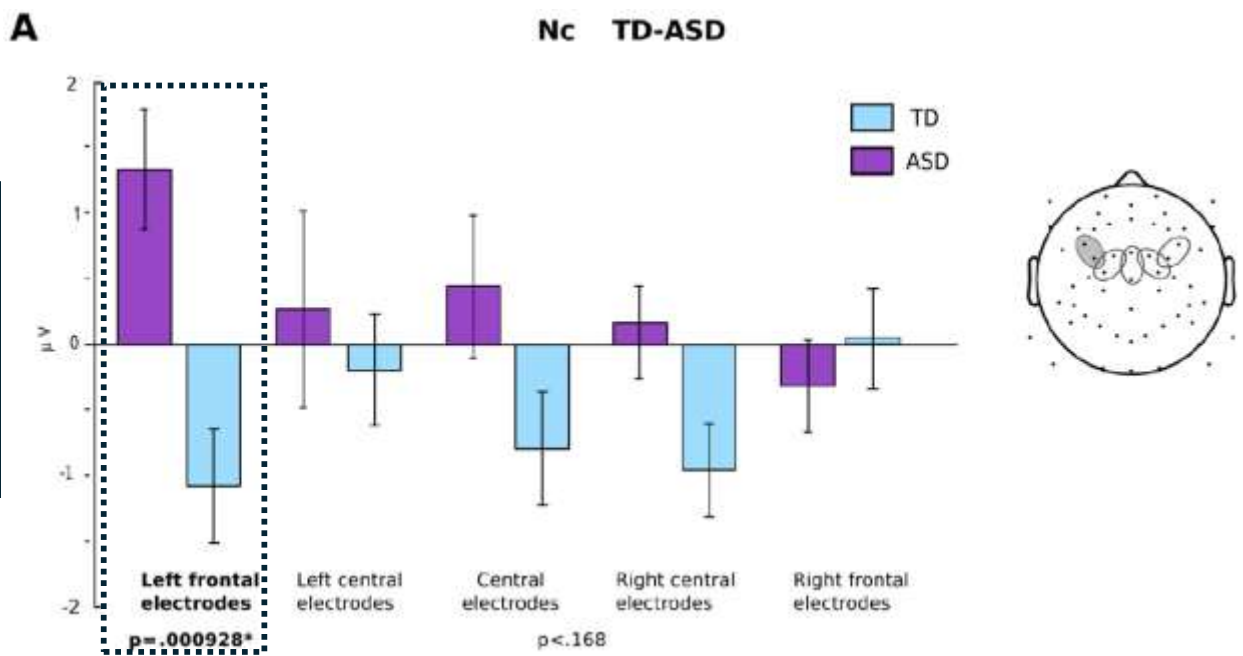
Dedicated  
regressor

$$P(f, t) = \beta_1 + \beta_2 JA_{first} + \beta_3 JA$$

# Positive controls, and first differences

Biological signature

$f_{ment} \mapsto P_{\beta}(TPJ)$   
 $f_{attn} \mapsto NC_{ERP} \uparrow ASD \downarrow TD$



Behavior was similar between TD and ASD

nJA trials (from valid trials <sup>b</sup> )	52.7% ± 22.5 SD	61.8% ± 28.3 SD	.1
JA trials (from valid trials <sup>b</sup> )	47.3% ± 22.5 SD	38.2% ± 28.3 SD	.13

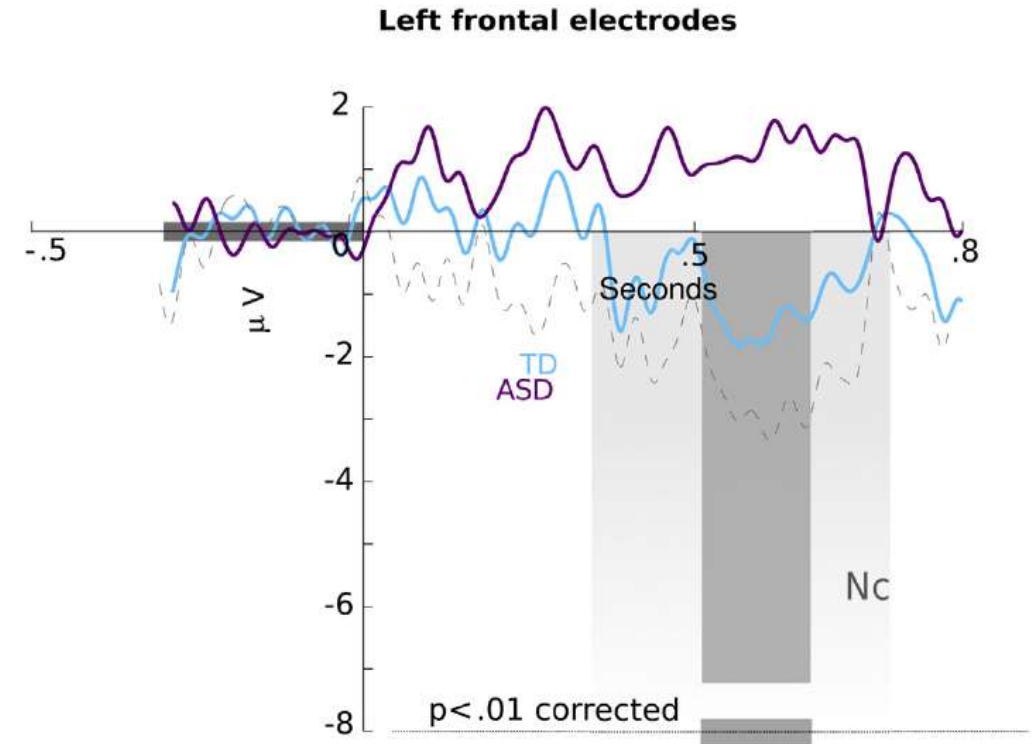
ERP signal differs precisely  
at Nc component

Biological  
signature

$$f_{ment} \mapsto P_{\beta}(TPJ)$$

$$f_{attn} \mapsto NC_{ERP} \uparrow ASD \downarrow TD$$

**B**



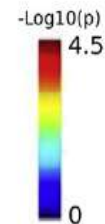
# Source localization of Nc component differences

Biological  
signature

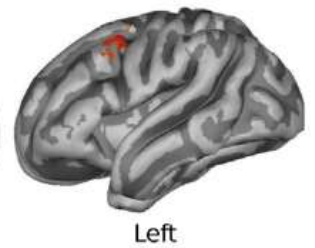
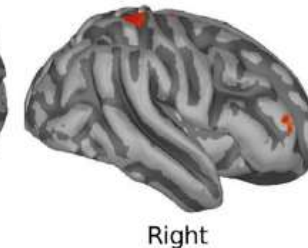
$$f_{ment} \mapsto P_{\beta}(TPJ)$$

$$f_{attn} \mapsto Nc_{ERP} \mapsto \textit{Source}$$

C



TD-ASD  
500-680ms

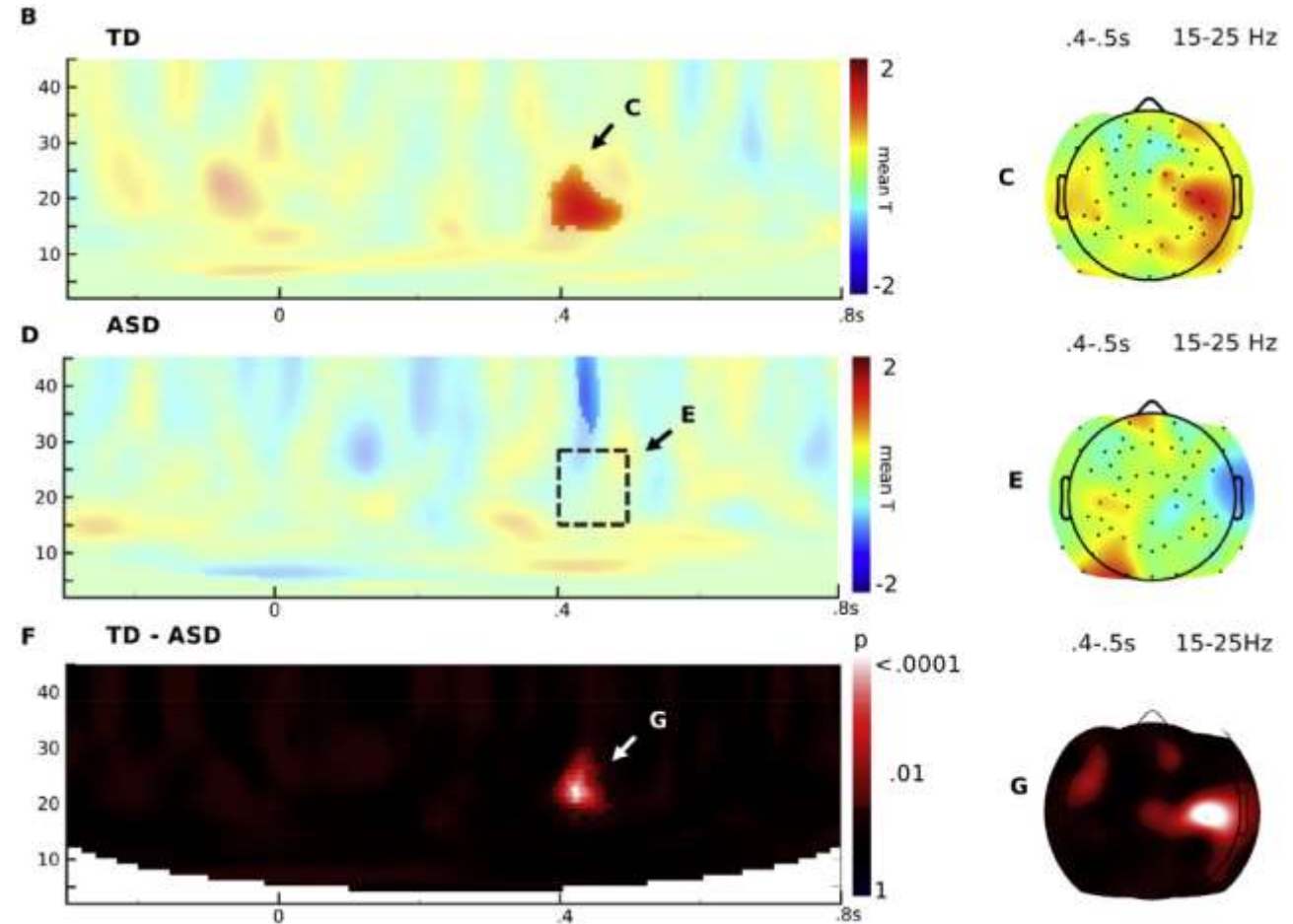


# Beta power modulation in JA

Biological  
signature

$$f_{ment} \mapsto P_{\beta}(TPJ)$$

$$f_{attn} \mapsto NC_{ERP} \mapsto \textit{Source}$$



# Source localization of beta power modulation differences

Biological  
signature

$f_{ment} \mapsto P_{\beta}(TPJ) \mapsto \textit{Source}$

$f_{attn} \mapsto NC_{ERP} \mapsto \textit{Source}$

