

Taq1A and DARPP polymorphisms are associated with worse working memory updating in high-BMI individuals

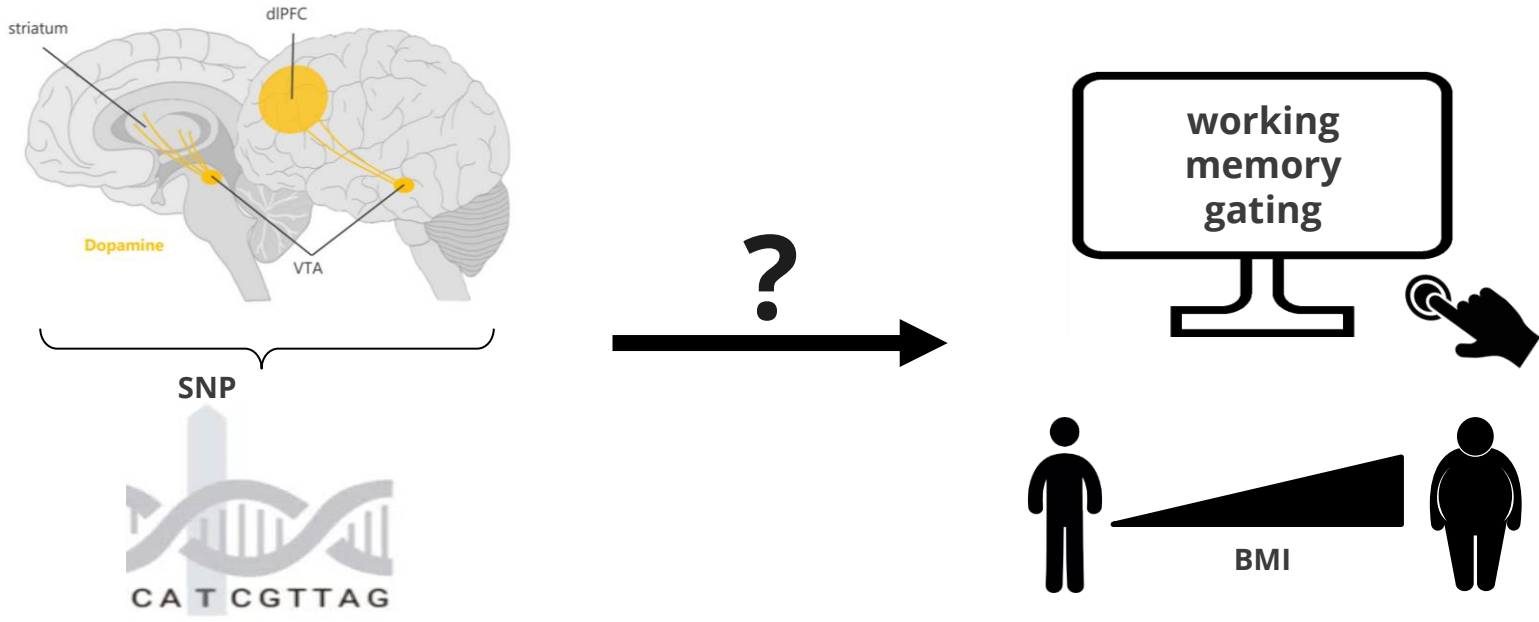
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Introduction



Working memory (WM) gating requires stable maintenance and flexible updating of information. These processes are implemented via dopamine-dependent signaling in the dorsolateral prefrontal cortex (PFC) and striatum [1,2]. Two single nucleotide polymorphisms (SNPs), COMT (rs4680) and Taq1A (rs1800497), have been associated with dopamine in PFC and striatum respectively, as well as with working memory functioning [3,4,5]. Furthermore, altered dopamine transmission has been observed in individuals with high BMI [6]. It remains unclear, however, if and how these two SNPs interactively influence working memory gating, depending on BMI.

Research questions:

Do COMT and Taq1A interact to foster differential performance in WM gating, depending on BMI?

Exploratory:
Do other proxies of dopamine modulate WM gating, depending on BMI?

Methods

Participants

- combined data set from 3 studies (fMRI: GREYDT, BEDOB; EEG: WORMCRI),
- final N = 318 (152 females; mean Age = 26.93 years (SD = 6.79, min = 12.17, max = 49.75); mean BMI = 26.40 kg/m² (SD = 6.37, min = 17.51, max = 45.54).
- all physically and mentally healthy, right-handed

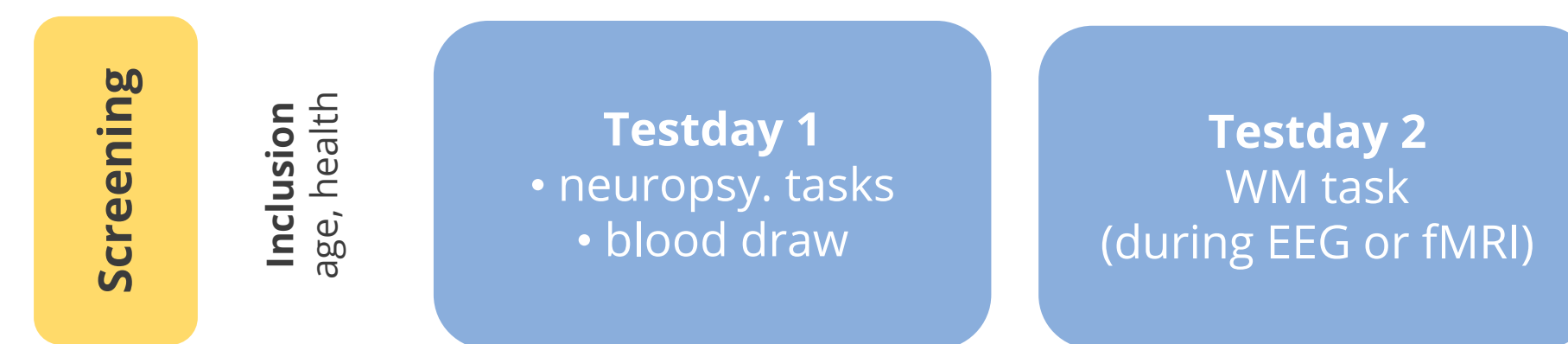
Proxies for Dopamine Differences

- **Taq1A:** A1+ associated with less D2 Receptors in Striatum
- **COMT:** met/met associated with more dopamine in PFC

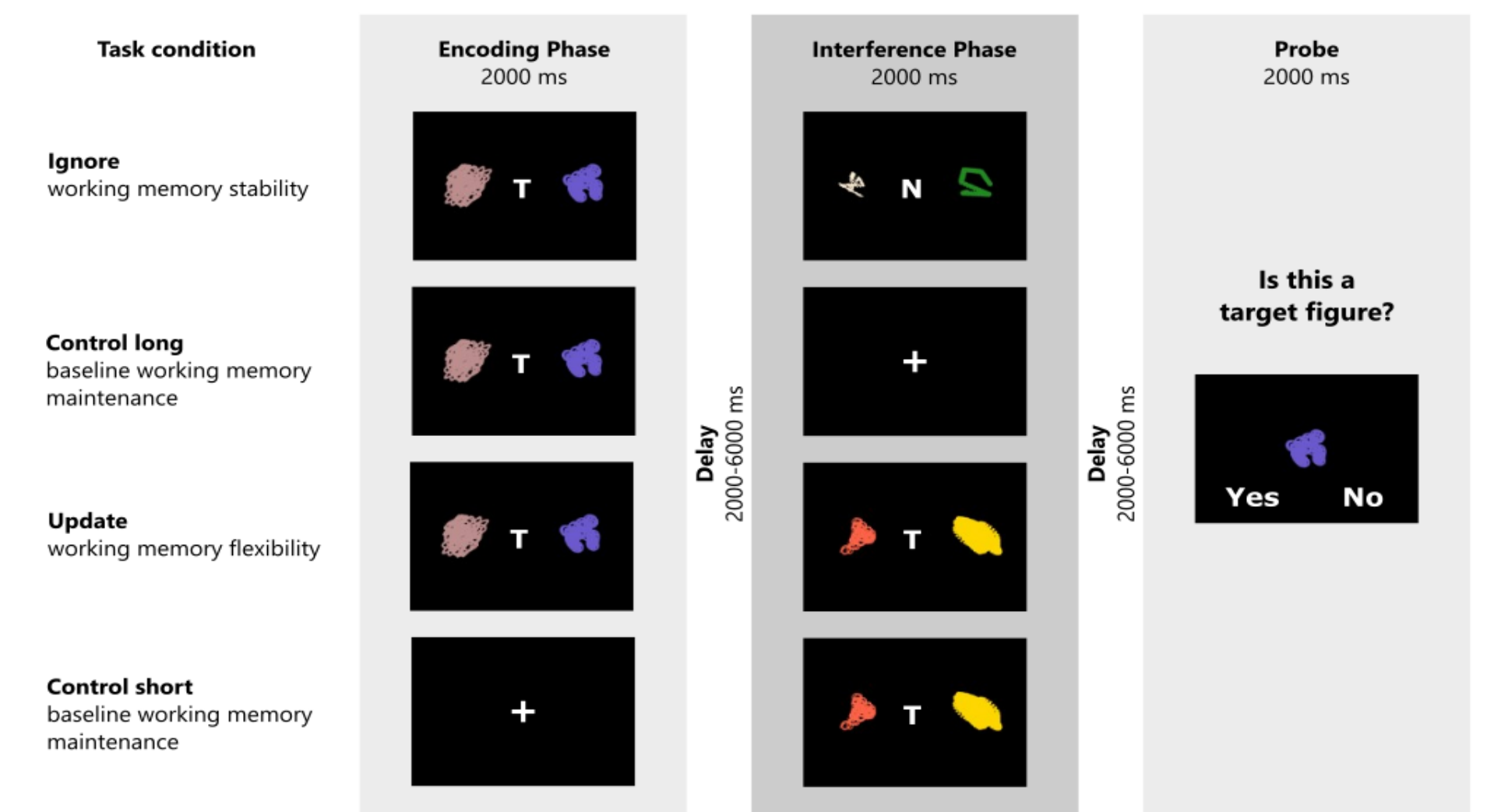
exploratory:

- **DARPP:** A/A associated with enhanced striatal D1 efficacy
- **C957T:** T/T higher D2 receptor availability (PFC & Striatum)
- **ratio of phenylalanine and tyrosine to large neutral amino acids:** proxy for endogenous dopamine levels

Study Design



Working memory gating task



Analysis

trial based analysis: logistic regression for accuracy (correct vs. incorrect)
 model comparison approach to find best fitting, least complex model

$$(1) \text{ accuracy} \sim \text{COMT} * \text{Taq1A} * \text{condition} * z\text{BMI} + z\text{tiredness} + z\text{IQ} + z\text{concentration} + \text{gender} + (1 | \text{subject})$$

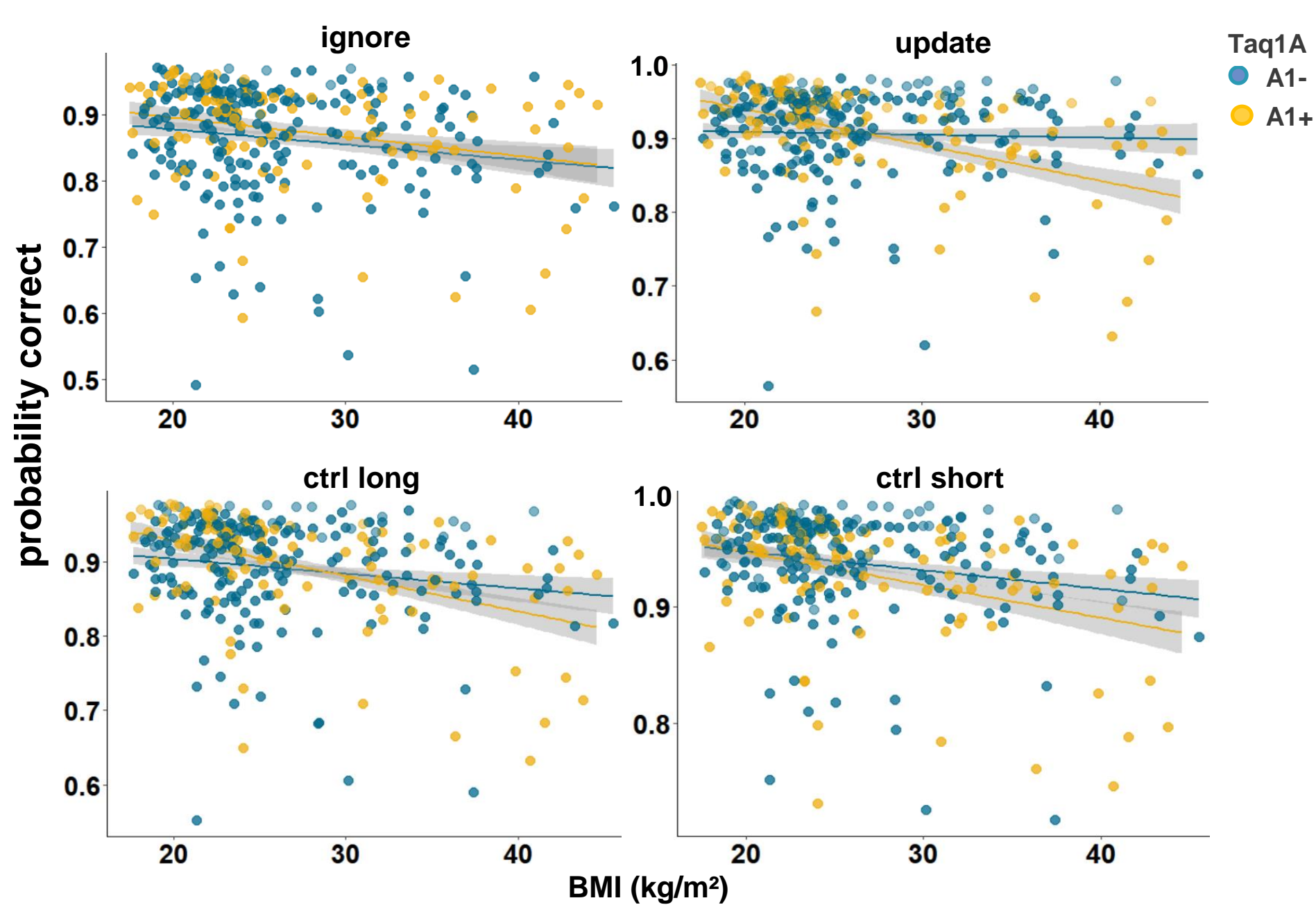
exploratory (corrected for multiple comparisons, p-value time 5)
 (2-5) $\text{accuracy} \sim \text{SNP} * \text{condition} * z\text{BMI} + z\text{IQ} + z\text{tiredness} + z\text{concentration} + \text{gender} + (1 | \text{subject})$

$$(6) \text{ accuracy} \sim \text{amino acid ratio} * \text{condition} * z\text{BMI} + z\text{IQ} + z\text{concentration} + \text{gender} + (1 | \text{subject})$$

Results

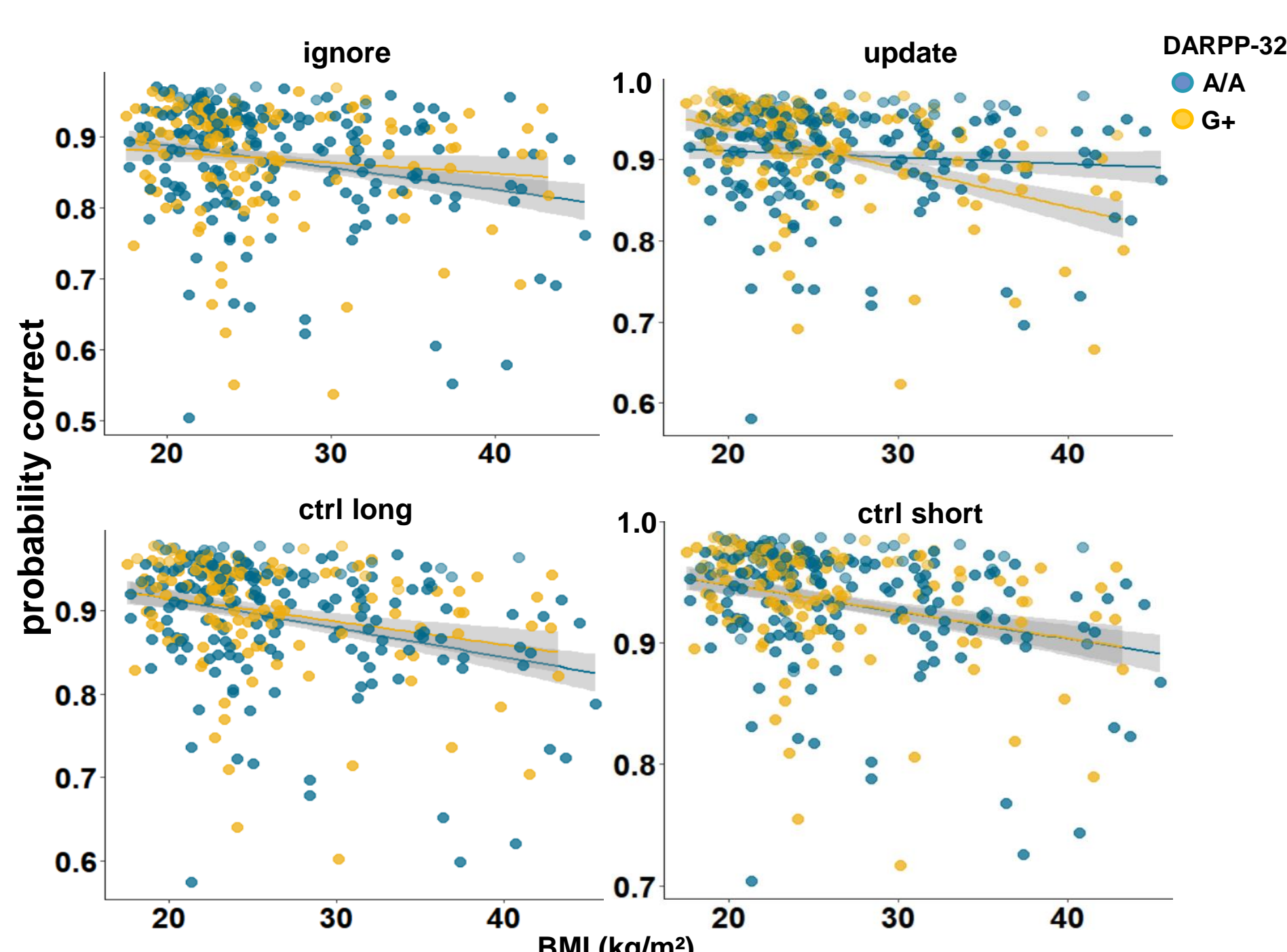
No gene-gene interaction, but BMI-dependent effect of Taq1A on working memory updating

model 1: $p_{\text{COMT} * \text{Taq1A} * \text{condition} * \text{BMI}} = 0.106$; $p_{\text{Taq1A} * \text{BMI} * \text{condition}} < 0.000$
 Posthoc: $p_{\text{Taq1A} * \text{BMI}}$ for update = 0.001; $p_{\text{Taq1A} * \text{BMI}}$ all other conditions > 0.05



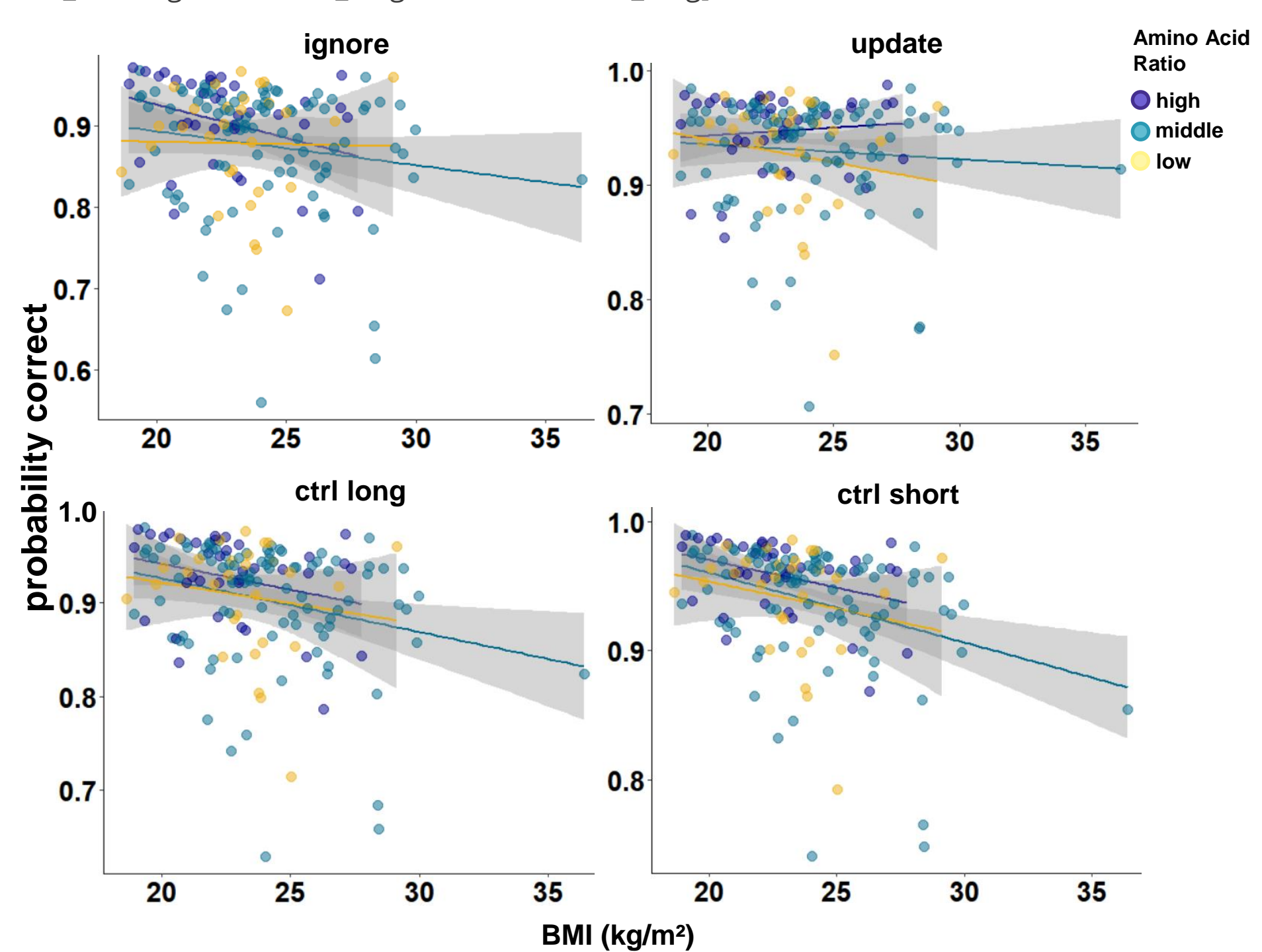
BMI-dependent effect of DARPP on updating of working memory contents

model 2: $p_{\text{DARPP} * \text{BMI} * \text{condition}} = 0.001$
 Posthoc: $p_{\text{DARPP} * \text{BMI}}$ for update = 0.011; $p_{\text{DARPP} * \text{BMI}}$ all other conditions > 0.189



Amino acid ratio interacts with BMI to foster differential ignoring vs. updating

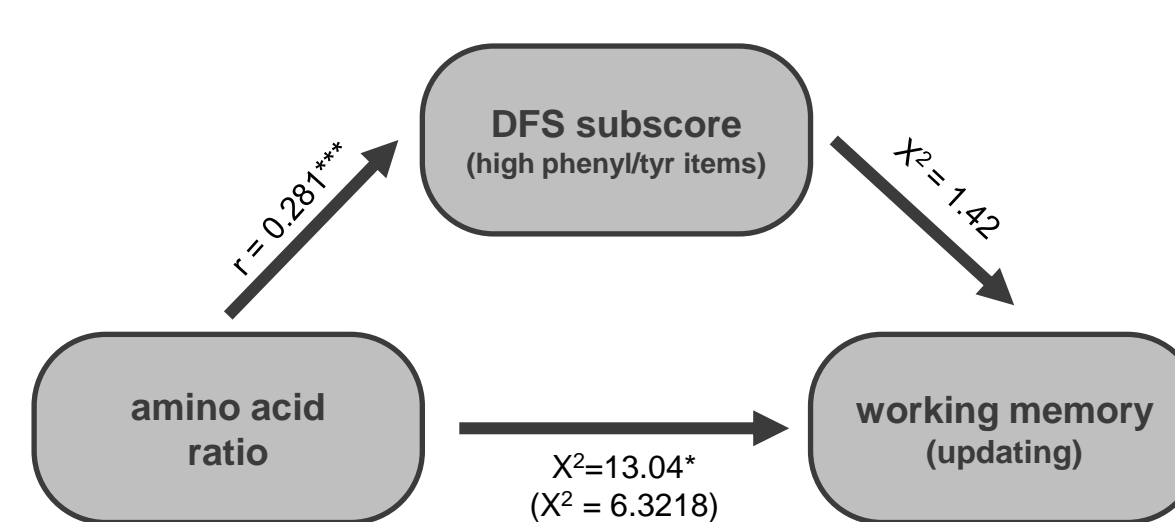
model 6: $p_{\text{AAratio} * \text{BMI} * \text{condition}} = 0.023$
 Posthoc: $p_{\text{update vs. ignore}} = 0.011$; all other comparisons [update vs. ctrl_short; ignore vs. ctrl_long; ctrl_short vs. ctrl_long] $p > 0.226$



Association between amino acid ratio and BMI-dependent ignore/update is mediated by food intake

additional mediation analysis: subscore of items high in phenylalanine and tyrosine from the Dietary Fat and Sugar Questionnaire.

- $p_{\text{subscore}} \sim \text{AAratio} < 0.001$ ($r = -0.281$, 95CI = -0.4180702 - 0.1307529)
- subscore as a covariate in model 6
 → $p_{\text{AAratio} * \text{BMI} * \text{condition}} = 0.097$, indicating a mediation.



The standardized coefficient between amino acid ratio and working memory, controlling for DFS subscore, is in parentheses. *** $p < 0.001$; * $p < 0.05$

Discussion

High BMI combined with an "disadvantageous" genotype is associated with worse updating. Specifically, SNPs associated with striatal dopamine transmission are at play here

- complies with evidence suggesting that high BMI is associated with impaired dopamine transmission within the striatum [6]
- first study to show that specifically updating of WM is affected in individuals with high BMI who also possess a disadvantageous genotype

We found no significant associations between COMT or C957T and BMI-dependent working memory gating

- both SNPs are (also) related to PFC dopamine [3,7]
- in line with that there are no ignore-related effects
- emphasizes that specifically updating/striatal effects are at play

Blood amino acid ratio, which is likely to be influenced by intake of foods high in phenylalanine and tyrosine, can affect BMI-dependent WM gating performance

- suggests that a diet high in phenylalanine and tyrosine could rescue "bad" updating
 - highly speculative!
 - needs specifically designed studies, with explicit measures targeted to quantify food intake properly

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[1] Durstewitz, D., & Seamans, J. K. (2008). The dual-state theory of prefrontal cortex dopamine function with relevance to catechol-O-methyltransferase genotypes and schizophrenia. *Biological psychiatry*, 64(9), 739-749.

[2] O'Reilly, R. C., & Frank, M. J. (2006). Making working memory work: a computational model of learning in the prefrontal cortex and basal ganglia. *Neural computation*, 18(2), 283-328.

[3] Chen, J., Lipska, B. K., Halim, N., Ma, Q. D., Matsumoto, M., Melhem, S., Kolachana, B. S., Hyde, T. M., Herman, M. M., Apud, J., Egan, M. F., Kleinman, J. E., & Weinberger, D. R. (2004). Functional analysis of genetic variation in catechol-O-methyltransferase (COMT): effects on mRNA, protein, and enzyme activity in postmortem human brain. *American Journal of Human Genetics*, 75(5), 807-821.

[4] Jönsson, E. G., Nöthen, M. M., Grünhage, F., Farde, L., Nakashima, Y.,

Propping, P., & Sedvall, G. C. (1999). Polymorphisms in the dopamine D2 receptor gene and their relationships to striatal dopamine receptor density of healthy volunteers. *Molecular Psychiatry*, 4(3), 290-296.

[5] Berryhill, M. E., Wiener, M., Stephens, J. A., Lohoff, F. W., & Coslett, H. B. (2013). COMT and ANKK1-Taq1a genetic polymorphisms influence visual working memory. *PLoS One*, 8(1), e5862.

[6] Horstmann, A., Fenske, W. K., & Hankir, M. K. (2015). Argument for a non-linear relationship between severity of human obesity and dopaminergic tone. *Obesity Reviews*, 16(10), 821-830.

[7] Hirvonen, M., Lumme, V., Hirvonen, J., Pesonen, U., Nagren, K., Vahlberg, T. (2009). C957T polymorphism of the human dopamine D2 receptor gene predicts extrastriatal dopamine receptor availability in vivo. *Prog. Neuropsychopharmacol. Biol. Psychiatry* 33 (4) 630-636.