

# Normalized Amygdala fMRI Response during Emotional Processing as Marker of Recurrence Severity of MDD in the UK Biobank

Jerke van den Berg<sup>1,2</sup>, Henk Marquering<sup>1,2</sup>, Henricus Ruhé<sup>3</sup>, Liesbeth Reneman<sup>2</sup>, Matthan Caan<sup>1</sup>

<sup>1</sup>Biomedical Engineering and Physics, <sup>2</sup>Radiology and Nuclear Medicine, Amsterdam UMC, Amsterdam, Netherlands, <sup>3</sup>Department of Psychiatry, Radboud UMC, Nijmegen, Netherlands.

## INTRODUCTION

- Studies have consistently shown increased amygdala activity measured in BOLD response in Major Depressive Disorder (MDD) during emotional tasks involving facial stimuli, namely the Hariri task<sup>1</sup>.
- Such increase was not found by Tamm et al.<sup>2</sup> in the UK Biobank (UKBB), one of the largest population studies to date, collecting the median BOLD response of faces-shapes contrast masked to the amygdala (see Figure 1).
- Tamm's analysis<sup>2</sup> assessed associations with concurrent depressive symptoms, focusing on state rather than trait factors of depression.
- Normative Modeling is an emerging new tool to combat heterogeneity in neuroimaging (see Figure 2).

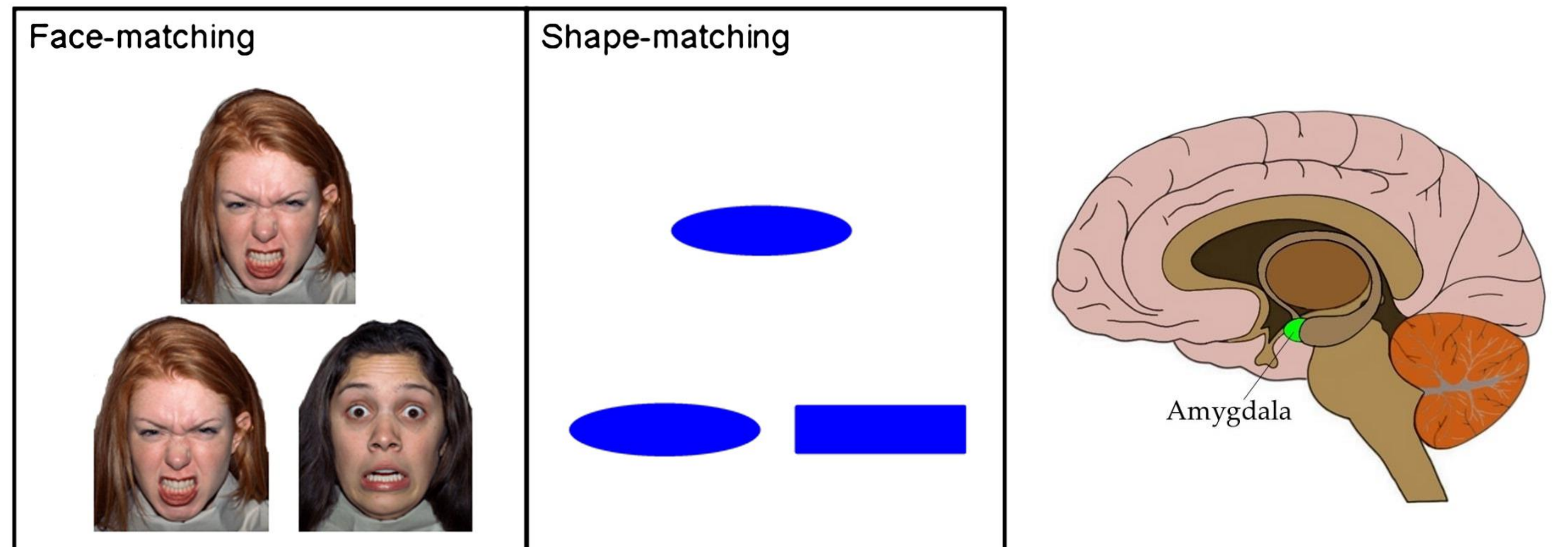


Figure 1 – The left shows examples of the Hariri task, consisting of alternating Face-matching and Shape-matching blocks. The median BOLD response is taken from the contrast between these two and masked to the amygdala (shown on the right).

## AIM OF STUDY

- Analyse and distinguish **state** from **trait** associations of MDD in a multilevel analysis of the BOLD response in the Amygdala measured during the Hariri task, to find out whether the BOLD-contrast signal associates with longterm recurrence-severity of MDD or current onset.
- Regress out associated covariates (age and sex) and determine whether the BOLD response in this experiment is a potential biomarker for MDD.

## METHODS

- Using **Normative Modeling**, we calculate deviation scores of depressed individuals relative to healthy controls, and regress out unwanted effects induced from age and sex. For determining the normative range, a sample of 6442 healthy individuals is used (separated from testing) to fit a Bayesian Linear Regression model.
- Using questionnaire data, we determine patient history of depressive episodes, categorized by single episode, moderate recurrence (2-5 episodes) and severe recurrence (5+ episodes). To determine remission/occurrence of an episode on day of scanning, the Recent Depressive Symptoms (RDS) is used by summing up the score of four questions closely related to the PHQ-9.

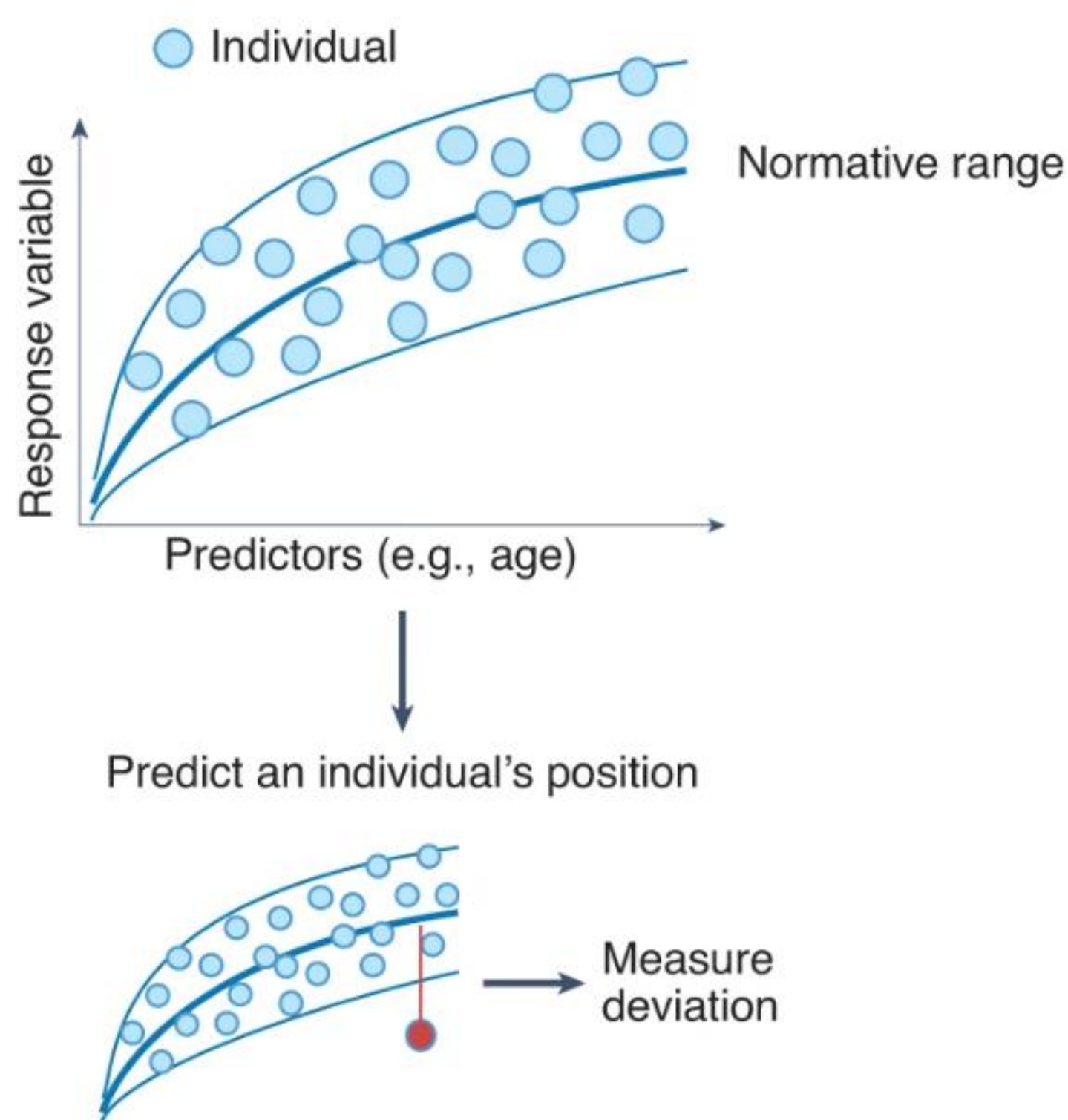


Figure 2 – Normative Modeling for individual prediction.

## RESULTS

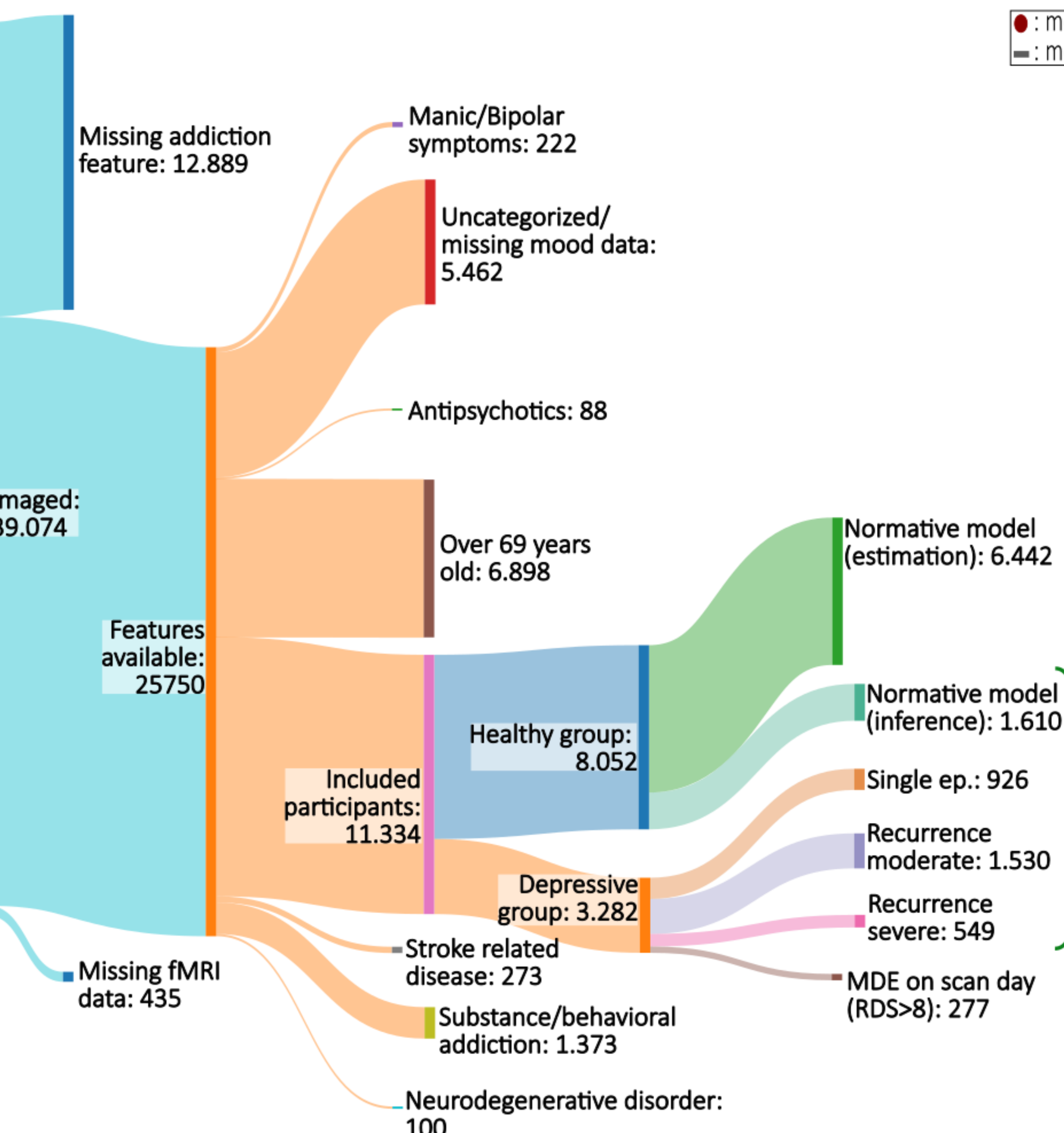


Figure 3 – Inclusion from UKBB, with median BOLD response of healthy, single episode, moderate- and severe recurrent MDD (in remission; RDS<9) participants from left to right respectively. Fisher's test is reported above, with a bar representing the pairwise significance post-hoc test between healthy controls and severe recurrent MDD.

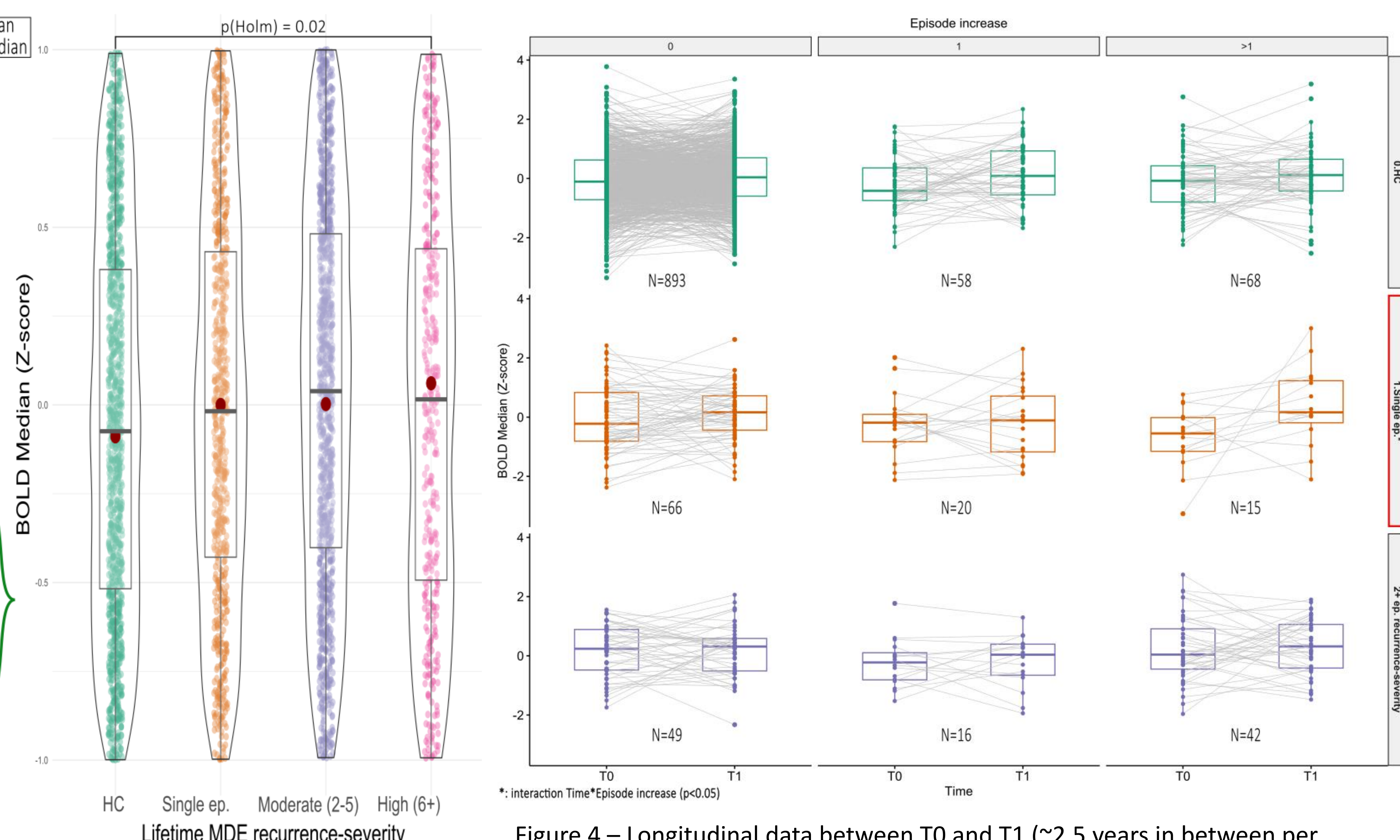


Figure 4 – Longitudinal data between T0 and T1 (~2.5 years in between per participant), divided into recurrence-severity at T0 (rows) and relative increase of episodes at T1 (columns).

## DISCUSSION AND CONCLUSION

- Normative Modeling **increased predictive power** of the amygdala BOLD feature as marker for MDD recurrence, without inherently 'training' for it.
- Population studies introduce more heterogeneity both in labels and neurological data; Normative Modeling is a step to overcome this.
- No effect was found in symptom-severity**, in line with Tamm et al.<sup>2</sup>
- A significant increase in the BOLD signal was found within **severe recurrence** of MDD (5+ episodes), as well as in longitudinal analysis in the **advancement of single episode to further recurrence** of MDD episodes.

A significant effect in the recurrence-severity stages was found ( $p=0.02$ ,  $\eta^2=0.002$ ), with post-hoc analysis indicating a significant difference between healthy controls and **severe recurrence** (Holm-adjusted  $p=0.02$ ) (Fig. 3). A significant interaction time \* episode increase ( $p=0.02$ ,  $\eta^2=0.03$ ) within the **single episode** group was found, highlighted in red (Fig. 4).