

Introduction

- Excitation-inhibition (E-I) balance is essential for central aspects of cortical functioning, such as the dynamic stability of activity and efficient coding of the information¹.
- Adolescence involves substantial macro- and microscale changes in the brain, including maturation of the E-I balance²⁻³.
- Currently available evidence on the *in vivo* maturation of the E-I balance in humans is limited, as the invasive methods used in animal studies are not feasible in humans.
- Biophysical network modeling (BNM) of the brain is a promising technique that can bridge different scales of investigation at a whole-

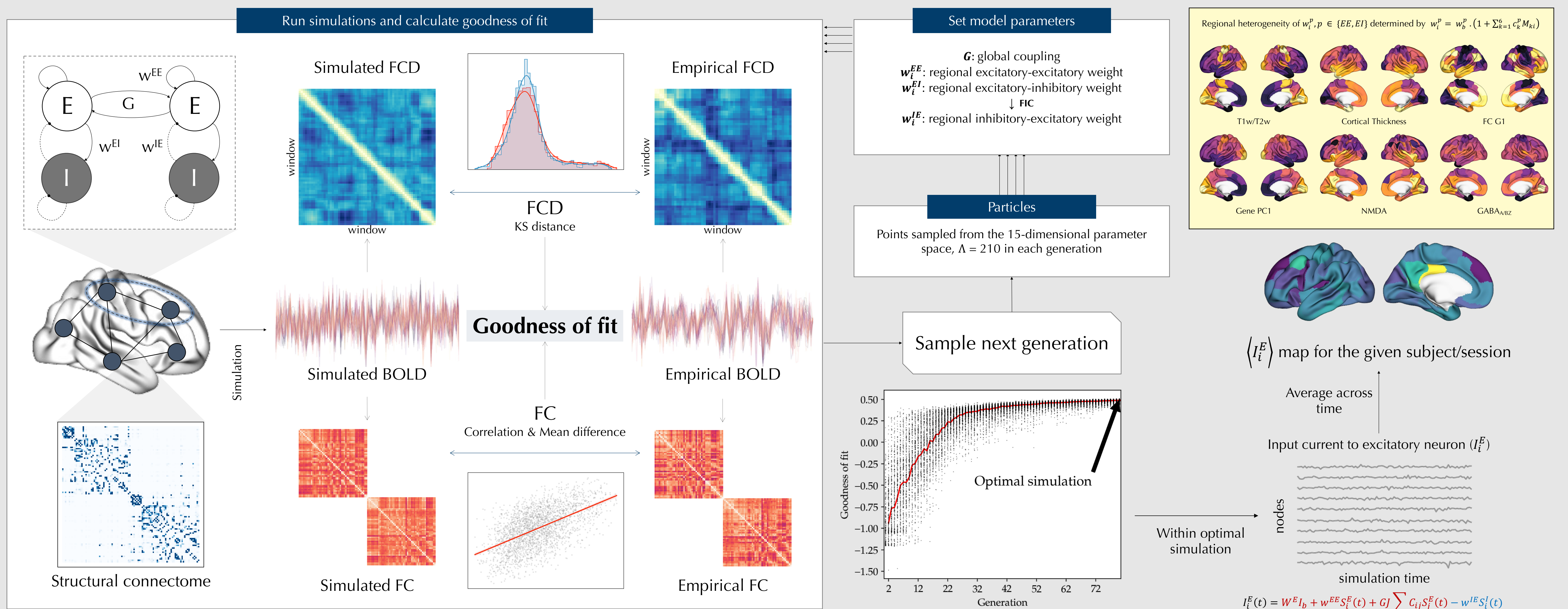
cortical level, and provides a tool to derive mechanistic inferences about the E-I balance based on the observed (empirical) *in vivo* imaging data at the macroscale⁴⁻⁵.

- Recently, this approach was applied to group-averaged cross-sectional data of 29 age groups of adolescents and revealed widespread relative decrease of inhibition across the cortex, most prominently in the sensorimotor areas⁶.
- However, an individualized *in vivo* estimation of E-I balance is needed for more precise characterization of E-I maturation based on subject-specific simulations, and enables assessing longitudinal changes.

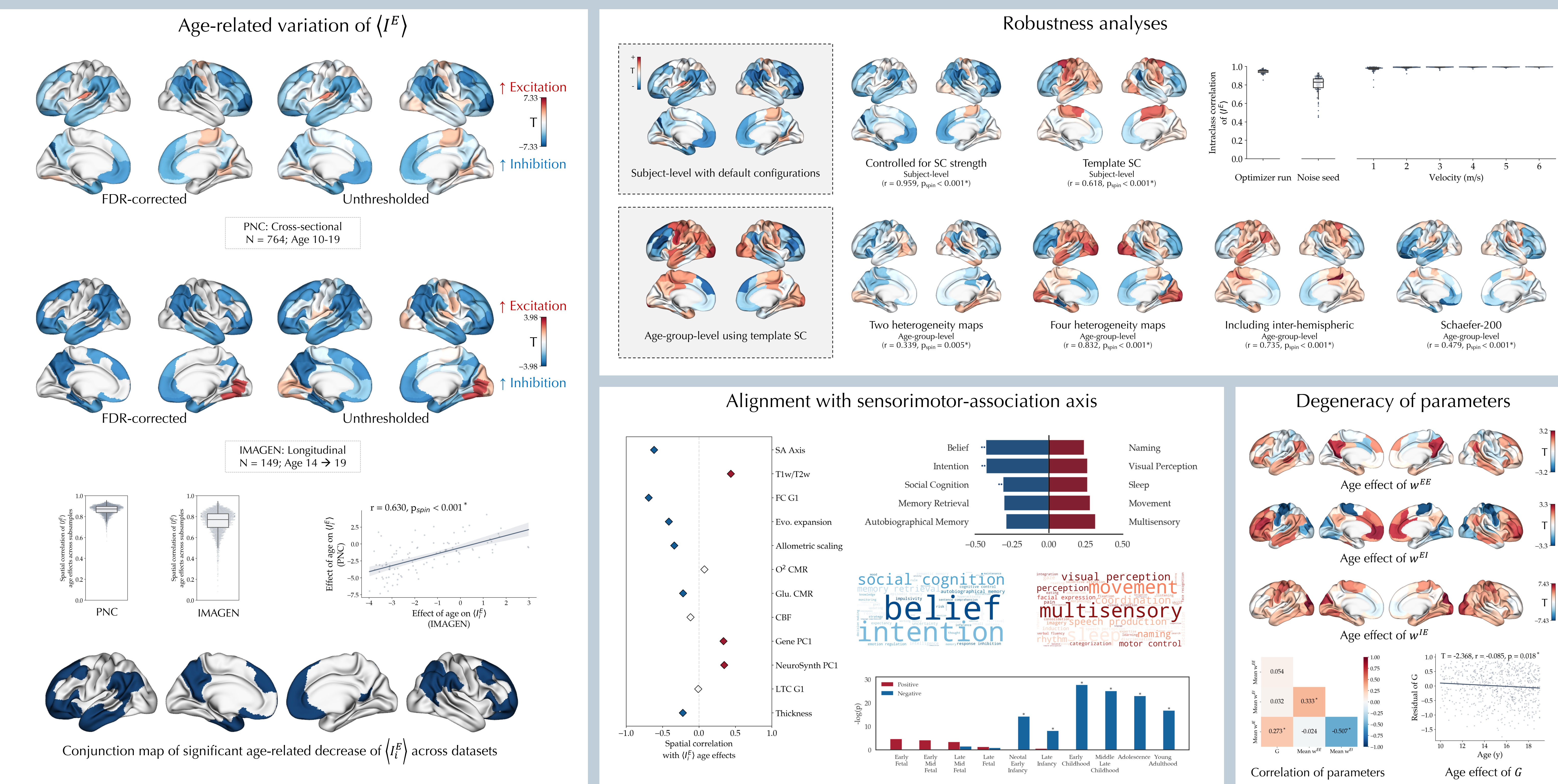
AIM

How does excitation-inhibition balance mature during human adolescence at an individual level, cross-sectionally and longitudinally?

Methods



Results



Take Home

Cross-sectional and longitudinal data showed reliable and robust patterns of E-I balance maturation with:

- Shift towards \uparrow inhibition / \downarrow excitation in association areas
- Lack of change or shift towards \downarrow inhibition / \uparrow excitation in sensorimotor areas

The spatial pattern of differential E-I balance maturation across cortical areas co-aligns with the proposed sensorimotor-association axis of neurodevelopment, similar to several other cortical features².

There are important methodological considerations with using BNMs for the estimation of E-I balance. For example, inter-dependency of model parameters and the difficulty of interpreting their concurrent changes make them unsuitable for this purpose.

References:

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