Article: 0105

Topic: S44 - Symposium 46: Parental psychopathology and children's developmental trajectories: data from prospective cohort studies

Impact of Maternal Prenatal Risk Factors On Oxytocin Methylation, and On Occurrence of Conduct Disorders During Adolescence

E.D. Barker<sup>1</sup>, C.A.M. Charlotte<sup>1</sup>

<sup>1</sup>King's College London, Institute of Psychiatry, London, United Kingdom

Youth with high callous-unemotional traits (CU) are at risk for early-onset and persistent conduct problems. Research suggests that

there may be different developmental pathways to CU (genetic/constitutional vs environmental), and that the absence or presence

of co-occurring internalizing problems is a key marker. However, it is unclear whether such a distinction is valid. Intermediate

phenotypes such as DNA methylation, an epigenetic modification regulating gene expression, may help to clarify etiological

pathways. This is the first study to examine prospective inter-relationships between environmental risk (prenatal/postnatal) and

DNA methylation (birth, age 7 and 9) in the prediction of CU (age 13), for youth low vs high in internalizing problems. We focused

on DNA methylation in the vicinity of the oxytocin receptor (OXTR) gene as it has been previously implicated in CU. Participants

were 84 youth with early-onset and persistent conduct problems drawn from the Avon Longitudinal Study of Parents and Children.

For youth with low internalizing problems (46%), we found that (i) OXTR methylation at birth associated with higher CU (age 13) as

well as decreased experience of victimization during childhood (birth-age 9), (ii) higher prenatal parental risks (maternal

psychopathology, criminal behaviors, substance use) associated with higher OXTR methylation at birth and (iii) OXTR methylation

levels were more stable across time (birth–age 9). In contrast, for youth with high internalizing problems, CU were associated with

prenatal risks of an interpersonal nature (that is, intimate partner violence, family conflict) but not OXTR methylation. Findings

support the existence of distinct developmental pathways to CU.