# Genomic variants correlated with cognitive behavioral development in the Canadian Healthy Infant Longitudinal Development (CHILD) study

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#### **BACKGROUND**

Sleep disordered breathing (SDB), a collective term for chronic conditions including habitual snoring and obstructive sleep apnea, affects up to 10% of children between 2 and 8 years old. Earlier studies have shown that SDB in children is associated with neurobehavioral functions related to executive functioning, behavior development and attention deficit hyperactivity disorder (ADHD) [1]. We hypothesize that preschool SDB may share a common genetic predisposition with neurobehavioral functions during early childhood.

## **METHODS**

In this study, genomics data were ascertained from the Canadian Healthy Infant Longitudinal Development (CHILD) study using the Illumina HumanCore Exome BeadChip. A total of 2048 Caucasian subjects had available SDB variables derived from parent-reported sleep related breathing disorder subscale (age 5) in addition to cognitive behaviour assessments (i.e. Child Behavior Checklist (CBCL) internalizing and externalizing scores, which are associated with anxiety and aggressive behaviours, respectively). We selected 108 loci for a candidate gene analysis of both SDB and CBCL scores that included variants previously associated with schizophrenia from the psychiatric genomic consortium (PGC) genome wide association study [2]. In addition to main genetic effects, we investigated the potential for genetic interactions with exposures such as exclusive breastfeeding until 3 months.

#### **RESULTS**

Single variant association of CBCL externalizing score identified 14 significant variants located at chromosome 6q12 (p <  $3.63 \times 10^{-5}$ ). In addition, we identified 2 variants significantly associated with SDB at chromosome 6p22.1 (p =  $1.8 \times 10^{-5}$ ). Moreover, we identified an interaction effect between genetic variants at chromosome 16q21 and exclusive breastfeeding at 3 months for CBCL externalizing score.

### CONCLUSION

Our study identified that genetic variants associated with schizophrenia in adults may contribute to cognitive behavioural traits and SDB among children during early childhood. These results suggest a common genetic predisposition that can be detected early in childhood and is

modifiable by environmental exposures such as breastfeeding. On-going analyses include genetic risk score analysis and gene-set association tests of rare variants. Furthermore, we will explore gene-environmental interactions using additional exposures such as parental SDB, sleep duration, apnea-hypopnea index, sleep habits, and physical activity.

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