Genetic Components in Dental Fear and Anxiety – Preliminary GWAS Results

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Introduction:

The results of a recent genome-wide association study (GWAS) showed 3 significant associations between dental fear and anxiety and genetic loci, nearby genes NTSR1, DMRTA1, and FAM84A (Zhou et al. 2022). NTSR1 mediates the functions of neurotensin, NTSR1-deficient mice may have emotional disorders involving fear memory (Yamauchi et al. 2007). DMRTA1 is a candidate gene located within a known quantitative trait locus for contextual fear memory in mice (Carhuatanta et al. 2014). FAM84A encodes neurologic sensory protein 1, a candidate gene for schizophrenia (Sundararajan et al. 2018). Also, genes SLC6A4 (5-HTTLPR) and MC1R have been associated with dental fear and anxiety. 5-HTTLPR is a polymorphism of the serotonin transporter encoding gene. Disturbed function of serotine may result in the development of various anxious symptoms (Franceski et al. 2018). MC1R variant alleles have predicted higher levels of dental fear and fear of pain (Randall et al. 2016).

Objectives:

The aim was to identify genes associated with dental fear and anxiety in Northern Finland Birth Cohort1966 (NFBC1966).

Materials and methods:

- The study population was the NFBC1966, which consisted of 95.6% of the children born in the two northernmost provinces of Finland in 1966 (N=12,058).
- > Dental fear and anxiety was measured with Modified Dental Anxiety Scale (MDAS) when the participants were 46-year-olds.
- > The phenotype used in a genome-wide association study was dental fear and anxiety.
- > MDAS was used as natural logarithmic conversion for sum of MDAS, where higher values indicated higher anxiety.
- \succ The GWAS (n=1,440) was performed with the additive models using SNP test v2.5.1 and the frequentist association models.
- \blacktriangleright Associations at p-value <1x10⁻⁶ were considered suggestive and those at p-value $<5x10^{-8}$ genome-wide significant.



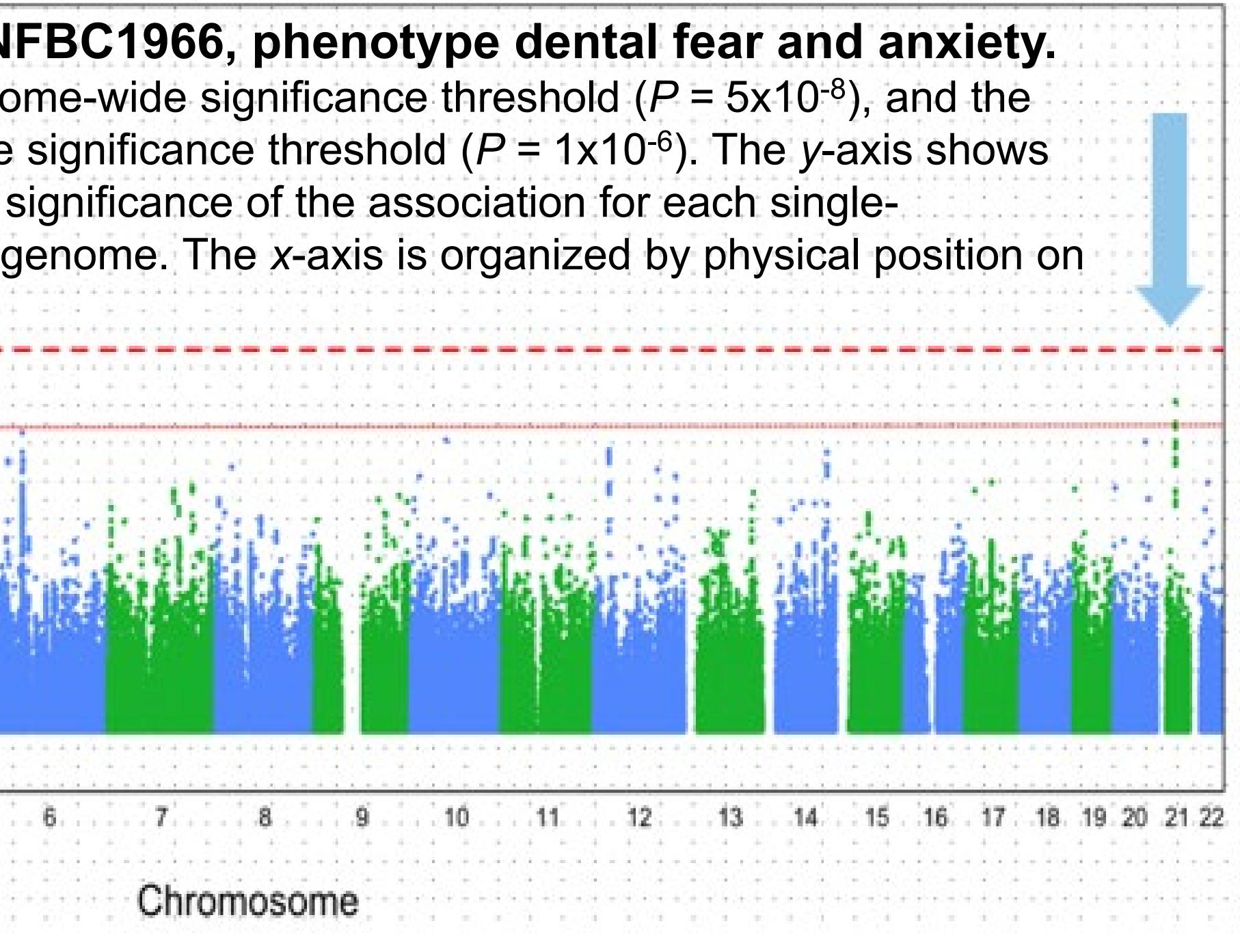
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¹² Figure 1. Manhattan plots for NFBC1966, phenotype dental fear and anxiety. The red dotted line indicates the genome-wide significance threshold ($P = 5 \times 10^{-8}$), and the red solid line indicates the suggestive significance threshold ($P = 1 \times 10^{-6}$). The y-axis shows the -log10-transformed P, indicating significance of the association for each singlenucleotide polymorphism across the genome. The x-axis is organized by physical position on each chromosome.

Conclusion:

Genes associated with dental fear and anxiety are poorly known. The results of this genome-wide association study suggest a possible connection between dental fear and anxiety and the neurotransmitter or neuroendocrine system. This result needs to be replicated in other study populations.

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Results:

SNP rs58988229, p=3.6x10⁻⁷. ADAMTS1 (ADAM Metallopeptidase with Tyrosine Rich 1).

Contacts:

The analysis revealed one suggestive association for dental fear and anxiety situated in chromosome 21, lead

This SNP is intergenic and situated near genes

Thrombospondin Type 1) and CYYR1 (Cysteine and

The ADAMTS1 gene has been suggested in mice to connect with inhibitory neurotransmitter system (GABA), leading to altered brain function related to anxiety in the hippocampus (Kurumaji & Nishikawa 2012). > The CYYR1 gene has been predicted to encode a

membrane protein that is an integral component of membranes (www.genecards.org) and relate to neuroendocrine system (Vitale et al.2007).

Both ADAMTS1 and CYYR1 represent biologically plausible candidate genes for dental fear and anxiety. \succ In this relatively small population, we were unable to confirm the previously found GWAS results between genetic loci and dental fear and anxiety.

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