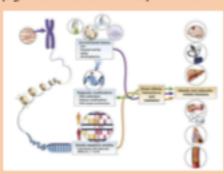
A genome-wide methylation study of body fat traits in the Norfolk Island Isolate

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Introduction

- · Obesity is a complex disorder that increases the risk of many endstage diseases (e.g. diabetes, heart disease).
- Obesity is influenced by both genetic and environmental factors.
- Epigenetic variation may mediate the interaction of these factors





Study Objectives

The aim of this research is to identify epigenetic markers associated with body fat traits in a healthy cohort. In this poster, the results from cross-sectional study of all samples is provided.

Methods

The study was performed on the cohort of 48 healthy adults with following methods:

- Body fat traits:
 - body fat percentage (BF)
- Waist circumference (WC)
- Body mass index (BMI)
- Waist-hip ratio (WHR)
- Hip circumference (HC)
- Weight (W)
- Genome-wide DNA methylation measured using Illumina 450K from whole blood samples
- Statistical analysis was performed using custom R scripts including the MINFI package to identify differentially methylated positions (DMPs) associated with each trait.

Results

Table 1: Significant DMPs associated with body fat traits

Trait	СрБ	Chr	Positio n	Gene	beta	P-valuet	Region		
PC1	cg04946709	16	5978 9030	GOT2-CDH8	t	3.19 x 10 ⁻⁸	Body		
BW	og04946709	16	5978 9030	GOT2-CDH8	t	5.83 x 10 ⁻⁸	Body		
нс	og22455090	5	8982 5803	LYSMD3	Ť	1 x 10-7	TSS 1500		
PC2	og22538323	7	27702700	HIBADH	t	1.17 × 10 ⁻⁷	TSS 200		
нс	og16379671	12	131647826	ADGRD1	t	2.02 x 10 ⁻⁷			
виш	og26576937	20	2739938	EBF4	t	2.08 x 10 ⁻⁷	3'UTR		
† adjusted for covariates									

BW = Body Weight HC= Hip Circumference

BMI = Body Mass Index

Table 2: Pathways associated with body fat traits

10	Name	P-value	P _{n·n} ·		Total				
P00012	Cadherin signalling pathway	3.23 x 10 ⁻¹⁰	6.76 x 10 ⁻⁷	list 28	Genes 159				
P00057	Wnt signalling pathway	7.44 x 10°	7.78 x 10°	38	305				
9 46 59 8	Thyroid hormone signalling pathway	5.1 x 10°s	2.82 x 10 ⁻²	16	116				
692234	PI3K-Akt signalling pathway	5.39 x 10·s	2.82 x 10-z	32	342				
1268881	Constitutive Signalling by AKT1 E17K in Cancer	7.75 x 10 ⁻⁶	3.25 x 10 ⁻²	7	25				
M295	Genes related to PIP3 signalling in cardiac myocytes	1.58 x 10 ⁻⁴	4.79 x 10 ⁻²	11	67				
	Rap1 signalling pathway	1.6 x 10 ⁻⁴	4.79 x 10 ⁻²	22	210				
Benjamini-Hochberg adjusted P-value									

Conclusions

By investigating a healthy cohort of individuals, we have identified:

- Five significant DMPs, which mapped to GOT2-APOOP5-CDH8, LYSMD3, HIBADH, ADGRD1 (GPR133)-LINC01257 and EBF4 genes
- Two significant pathways (Cadherin signalling and Wnt signalling) associated with body fat traits

Overall, this study provides new insights into epigenetically-influenced mechanisms underlying body fat variation in a healthy cohort, which could be considered targets for studying obesity in future studies

