

Associations of parental history of diabetes mellitus with the offspring's incidence is modified by offspring's body weight, findings from a Japanese work-site based cohort

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INTRODUCTIONS

- Family history may reflect complex relationships between genetic factors and environmental conditions that are important for developing diabetes.
- A greater risk from maternal history of diabetes compare to paternal history has been reported in some, but not in all studies.
- Cross-sectional report from the same population of our cohort study found that maternal history of hypertension, diabetes and dyslipidemia were more strongly associated with off-spring's prevalence of metabolic syndrome than paternal history.¹
- Proposed explanations included following:
 - \succ Mutations of mitochondrial DNA;
 - \succ Intrauterine environment;
 - \succ Behavior influence of the mother;
- It has been hypothesized that interactions between genetic- and environmental factors are of specific importance for the development of type 2 diabetes (T2D).
- Previous study indicated that obesity may significantly modify the association between parental history of diabetes and risk of new incidence.²

OBJECTIVES

In the current worksite-based Japanese cohort study:

- (1) The difference in the risk of T2D between individuals with paternal- and maternal history of diabetes was investigated;
- 2 The interaction effect of overweight on this association was tested.

METHODS

Participants:

5,471 workers (4,299 men; 1,172 wom **Exclusions:**

- History of diabetes (n = 430)
- 2 Baseline fasting blood glucose equal to 126 mg/dL or self-re under treatment of T2D (n =
- Body mass index (BMI), sex, s (3) drinking status, physical activ available (n = 374)
- n = 4,446 (3,492 men; 954 women)

Follow-ups: from May 2002 until April 20

Diagnosis of T2D:

- 1 Fasting glucose measured in the annua check-up first exceeded 126 mg/dL;
- 2 Self-reported in the questionnaires sen years during the follow-up. And confirm physician in charge for the date of diag

Other data collections including:

- 1 Parental history of diabetes and cates None, Father-only, Mother-only or Bot
- ② BMI was calculated from the weight a measured during the annual health ch from 2002 through the year of 2011, value was applied for primary analysis utilizing the formula (weight, kg) \div (h Overweight was defined as BMI higher to 25 kg/m²;
- Smoking status (current, past, never)
- 4 Alcohol consumption (times/week);
- 5 Physical activity was defined as partic moderate or vigorous leisure-time exe least 12 days/month or more than 360 in total (yes or no);

	METHODS cont.	<u>RES</u>	SULI
nen)	 Statistical analysis: Baseline descriptive statistics of the participants were reported as mean ± standard deviation (SD) and groups were compared using analysis of variance or ² test; 	 During the median cases of T2D (227 n Multivariable adjust 1,000 person-year; 	nen,
e higher or eported 221) smoking and vity not	 Incidence rate of T2D during the follow-up was evaluated by Poisson regression adjusted for age, sex, smoking status, alcohol consumption, BMI and physical activity; Multivariable-adjusted hazard ratios (HRs) and 	Figure 1: Hazard ratio and 9	95% CI
)11	95% confidence intervals (95% CIs) were estimated including the same above covariates by Cox proportional hazard model taking the category of "None" for parental history as the reference group in the total sample analysis;	4	
al health	 Another analyses was performed: HRs of T2D incidence according to the presence of paternal or maternal history of diabetes; 	∘- Neither Fathe	r-only
nt every 2 med by the gnosis.	 Stratified by overweight. Normal weight (n = 3499) Total sample analysis (n = 4,446) Overweight (n = 947) 	IR [*] HR (95% CI) IF - 6.7 1 5 Maternal	eight. al weigh R [*] 5.2
gorized as: th; and height neck-ups	 Likelihood ratio test was used to test the interaction effect of overweight. <u>RESULTS</u> 	+ 11.8 (1.20 - 2.67) - 6.6 1 5 Paternal	3.2 5.4 5.2 (
the baseline s, by	Table 1. Means (SD) and percentages of the participants at baseline, Aichi, 2002.	 Parental histories both) of diabetes 	•
neight, m)²; r or equal	Neither Father Mother Both P value n 3,832 373 216 25 Age (SD), years 47.7 (7.1) 45.9 (6.8) 48.4 (6.8) 46.1 (6.5) < 0.001	T2D incidence in r workers in Japan.	midd
; cipation of	Men, %79.571.373.680.0< 0.001Body mass index (SD), kg/m²22.9 (2.8)23.4 (2.9)23.5 (2.8)23.4 (2.8)< 0.001	② Maternal history of T2D in normal wei was observed in of	of dia ight
ercise for at 0 min/moth	Presence of physical activity, % 55.6 52.0 58.8 56.0 0.42 Alcohol consumed everyday, % 30.8 24.3 28.2 32.0 0.27	 parental history. ③ Being Overweight of T2D in subjects 	-

92.2

geometric mean, (95% Cl) (91.9 - 92.6) (91.2 - 93.3) (92.1 - 94.9) (91.3 - 100.2) 0.11

93.5

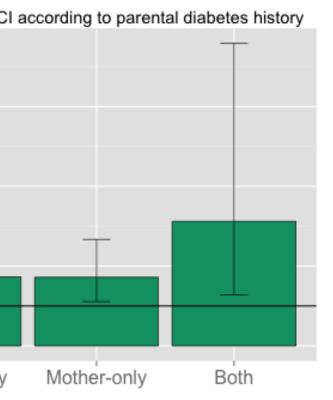
95.7

92.3

Fasting glucose,

TS cont.

.9 years follow-up, 277 50 women) developed; incidence rate: 7.9 per



cording to parental history of

ght (n = 3,499)	Overweight (n = 947)		P for	
HR (95% CI)	IR*	HR (95% CI)	interaction	
1	14.2	1	0.017	
2.57 (1.61 - 4.12)	12.8	0.86 (0.40 - 1.86)		
1	12.9	1		
1.59 (0.97 - 2.60)	22.3	1.92 (1.19 - 3.08)	0.485	

her-only, mother-only, or e positively associated with dle-aged male and female

abetes increased the risk of subjects to a degree that veight subjects without

nificantly increased the risk h paternal diabetes history.

DISCUSSIONS

Potential explanations:

- > Genomic imprinting, regulatory regions of certain genes are differentially methylated and expressed depending on whether the gene is inherited from the mother or father and may also interact with offspring's BMI.³
- \succ Maternal effects on the intrauterine environment on fetal development (including perinatal nutrition, and metabolism), and epigenetic modifications in oocytes (both nuclear and mitochondrial DNA methylation) may contribute to the maternal transmission of diabetes in normal weight subjects.

Limitations:

- Self-reported information might had recall bias
- Baseline information were not updated (BMI, parental DM might change during the follow-up), further confirmation needed.
- Incidence ascertainment was through annual health check-up and self-reported. Validation was done by reviewing the medical records with consent (38%).
- We do not have the information about age of diagnosis of the subjects parents.

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