

SUPPLEMENTARY DATA

Online Supplementary Material

Insulin-like growth factor binding protein 2 (IGFBP-2) and the risk of developing type 2 diabetes

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Supplementary Note 1

Analysis of cis regulatory elements of *IGFBP2* gene

For the identification of putative regulatory elements located within or in close proximity to the *IGFBP-2* gene, hepatic H3K27ac, H3K4me3, H3K9ac histone marks known to be associated with active promoters were downloaded from the ENCODE-Project. BAM-files were converted to bed files using BEDTools and then filtered for chromosomes. Bed-files were loaded into R (version 3.4.3) by use of Gviz (version 1.22.3), rtracklayer (version 1.38.3) and chipseq (version 1.28.0) packages. Chip-Seq scores were calculated within a size of 100 base pairs and plotted with the plot Tracks function. The resulting tracks of the single tissues were overlapped with help of Gimp (version 2.8).

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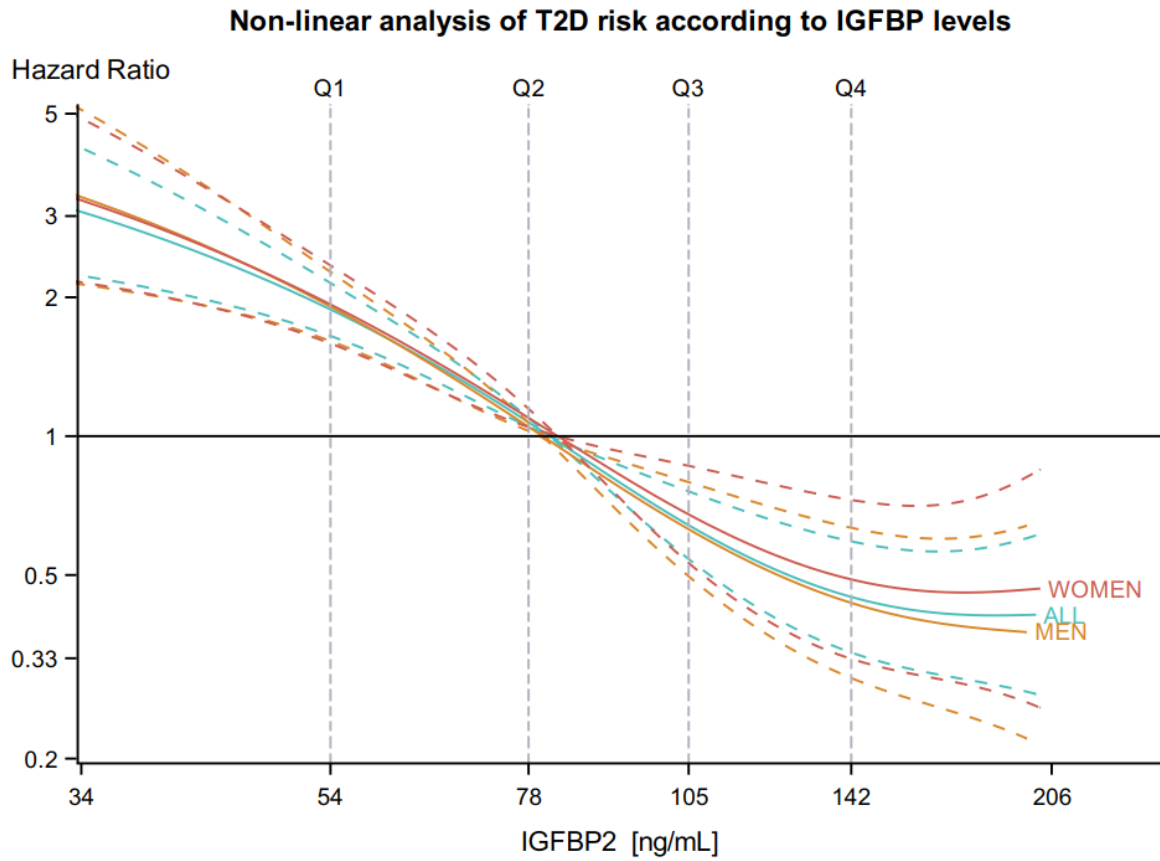
Supplementary Table 1. Baseline characteristics in cases and controls of the nested case-control sample with methylation data.

All (n=580)	Controls n=290	Cases n=290
IGFBP-2 [ng/ml]	100 (67, 142)	60 (44, 87)
Gender (% female)	48	48
Age at baseline [y]	56 (49, 60)	56 (49, 61)
BMI [kg/m ²]	26 (24, 28)	30 (27, 33)
Waist [cm]	89 (81, 95)	100 (92, 107)
Smoking Status [%]		
never smoker	47	33
ex-smoker, < 20 u./d.	23	26
ex-smoker, ≥ 20 u./d.	11	19
smoker, < 20 u./d.	13	11
smoker, ≥ 20 u./d.	7	12
Education [%]		
vocational training or lower technical college	37	47
university	24	22
university	39	31
Alcohol [g/day]	9.7 (3.3, 22.3)	8.2 (2.5, 19.6)
FLI [%]	37 (13, 62)	79 (54, 90)
IGF-1 [ng/ml]	157 (126, 197)	153 (125, 194)
IGFBP-3 [μg/ml]	3 (2.7, 3.5)	3.1 (2.7, 3.5)
Triglyceride [mg/dl]	117 (84, 168)	168 (123, 241)
Fetuin [μg/ml]	257 (223, 302)	279 (236, 317)
GPT [U/L]	22 (16, 30)	29.5 (21, 42)
gamma-GT [U/L]	20 (13, 36)	34 (21, 53)
Adiponectin [μg/ml]	7.6 (5.4, 9.8)	5.3 (4, 7.5)
Glucose [mg/dl]	103 (95, 113)	118 (105, 140)
HbA1c [%]	5.4 (5.1, 5.7)	6.1 (5.7, 6.7)

Comparison of the baseline characteristics between randomly selected incident type 2 diabetes cases and 1:1 matched controls; matching criteria were age (± 6 months), sex, and fasting time (<3h, 3h to <6h, and ≥ 6 h before blood draw), time of day of blood sampling (± 2 h), and season at blood sampling.

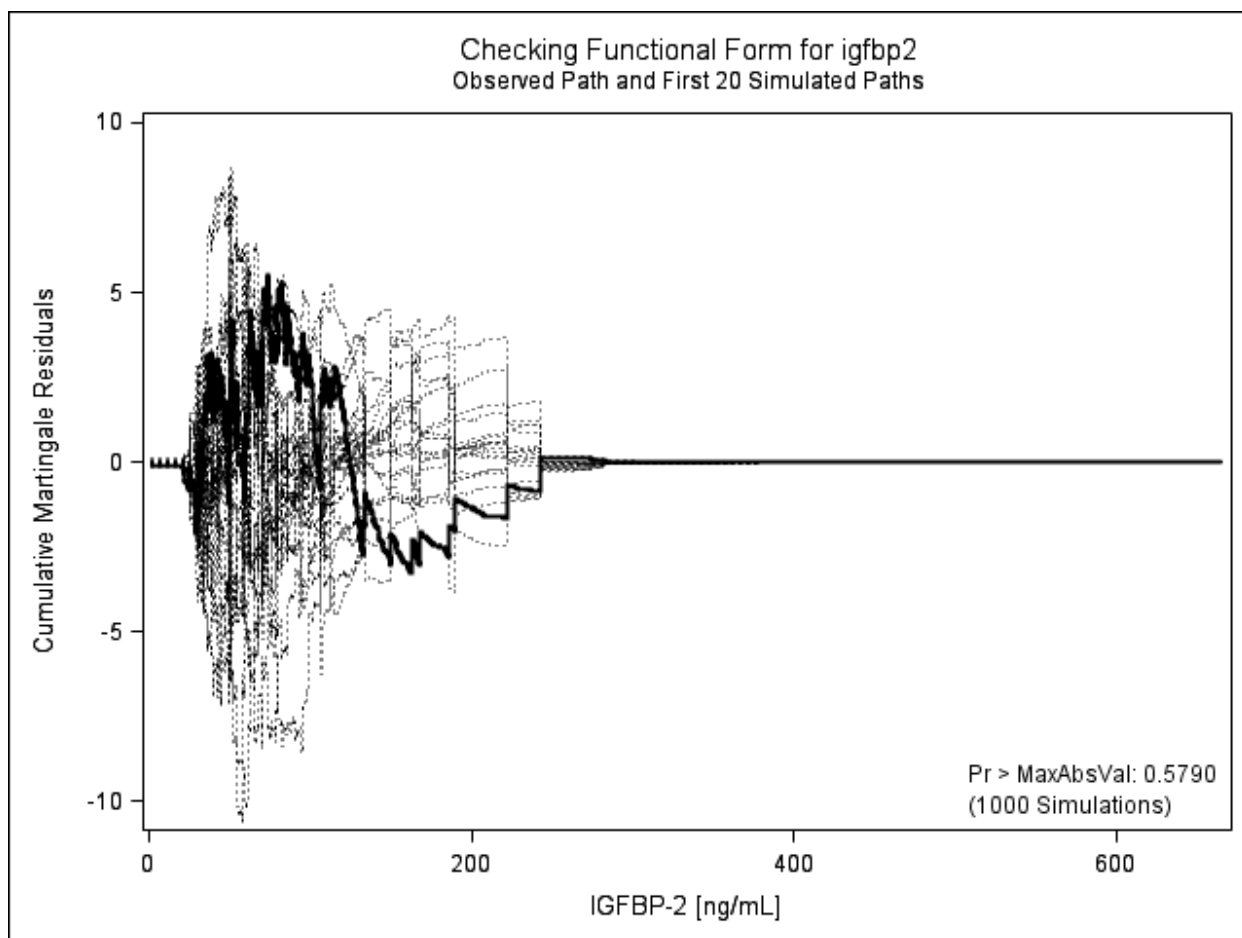
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Supplementary Figure 1. Restricted cubic spline analysis of the non-linear relation between circulating IGFBP2 concentrations and type 2 diabetes (Q1-Q4: Quintiles of the IGFBP2 distribution)



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Supplemental Figure 2. Testing potential interactions with follow-up time



Supremum Test for Functional Form

<i>Variable</i>	<i>Replications</i>	<i>Seed</i>	<i>Pr > MaxAbsVal</i>
igfbp2	1000	48680156	0.579

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Supplemental Figure 3. Checking proportional hazards assumption

