1	Interactions of genetic and environmental risk factors with respect to body fat mass in
2	children: Results from the ALSPAC study
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31 **Running Head:** Genetic and environmental determinants of obesity

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33 Key words: ALSPAC; child; fat mass; genetics; overweight; risk factors

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37	Abbreviations:	BMI- body mass index
38		ALSPAC-Avon Longitudinal Study of Parents and Children
39		GWA-genome-wide association
40		DXA-dual energy X-ray absorptiometry
41		FMI-fat mass index
42		CI-confidence interval
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49 **Abstract:**

50 **Background:** Genetic risk factors for childhood obesity were found to have greater effects on 51 children with a higher body fat mass. Similarly environmental and lifestyle risk factors for 52 childhood obesity were found to have a stronger effect at high body mass index (BMI) 53 percentiles. We hypothesized that these findings might reflect gene-environment interactions 54 with respect to the development of overweight.

55 **Methods:** We analysed data of 2 346 children from the Avon Longitudinal Study of Parents 56 and Children (ALSPAC), using quantile regression with body fat mass index (FMI) for 57 children at the age of 9 years as outcome variable. We assessed interactions of an "obesity-58 risk-allele-score" with environmental and nutritional factors.

Results: There was no evidence of interactions between the obesity-risk-allele score and the environmental variables except for maternal overweight. However, we found a clear interaction with respect to intake of mono- and polyunsaturated fatty acids at the age of 7. In children with low intake, genetic risk was associated with increasing effect sizes by FMI percentile.

64 **Conclusions:** Our results suggest an interaction between a low dietary content of unsaturated 65 fatty acids and genetic risk factors for overweight on FMI. This effect is likely to be stronger 66 in children with higher FMI. Apart from maternal overweight, which might also reflect 67 unknown genetic factors, we found no evidence for interactions of genetic disposition with 68 other environmental or nutritional factors.

70 Introduction:

Genetic factors are likely to determine the risk of overweight in children. Evidence comes 71 originally from adoption and twin studies ^{1, 2} and has been confirmed in recent genome-wide 72 association (GWA) studies, in which a number of risk alleles for overweight / obesity have 73 been identified ³⁻⁷. Combining such genetic variants in a risk score appears to be an 74 appropriate measurement of an individual's genetic predisposition for overweight ^{8, 9}. In a 75 76 previous study, we found that such a genetic risk score was associated with differential effect sizes depending on children's body composition, with greatest effects on higher body mass 77 index (BMI) percentiles ¹⁰. 78

Although the association of overweight / obesity with specific genetic predisposition is therefore well-established, the underlying mechanisms are still largely unknown. Recent studies suggest that interaction with environmental risk factors may be important ¹¹⁻¹⁴. This might also help to explain why we had previously found similar patterns of associations (i.e., different effect sizes by children's BMI) for environmental risk factors ^{15, 16}.

We therefore hypothesized that potential interactions with environmental risk factors might strengthen the effect of the "obesity-risk-allele score" in the upper body fat mass percentiles. In order to answer this question, we assessed interactions between the mentioned "obesityrisk-allele score" and a number of priming, life-style and nutritional factors with respect to body fat mass in children at primary school age using quantile regression.

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90 Methods:

91 Study population and data sources

92 The Avon Longitudinal Study of Parents and Children (ALSPAC) is an ongoing longitudinal birth cohort study that has been described in more detail elsewhere ¹⁷. A total of 14 541 93 94 pregnant women living in the former Avon Health Authority (South-West England) with an estimated date of delivery between April 1991 and December 1992 were enrolled, resulting in 95 96 a cohort of 13 971 children at the age of 1 year. Information about demographic data, lifestyle 97 habits, disease history etc. was collected using self-administered questionnaires, data extraction from medical notes, and linkage to routine information systems and at research 98 99 clinics. Dietary data were collected at the age of 7 years with the use of 3-day unweighed diet diaries ¹⁸. Ethical approval for the study was obtained from the ALSPAC Law and Ethics 100 101 Committee and Local Research Ethics Committees.

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103 **Outcome and explanatory variables**

104 Childhood height and weight were measured to the nearest 0.1 cm and 0.1 kg, respectively, at 105 dedicated ALSPAC Focus clinics by a trained research team. Fat mass was assessed at the 9-106 year visit using whole body dual energy X-ray absorptiometry (DXA) measurements (Prodigy 107 scanner, Lunar Radiation Corp, Madison, Wisconsin, US). We calculated fat mass index 108 (FMI) for each child from DXA measurements at age 9 y by dividing fat mass (kg) by height 109 squared (m^2).

In accordance with our previous studies ^{8, 10}, we calculated an "obesity-risk-allele score" by counting the total number of obesity risk alleles across the following eight genetic variants with known associations with BMI in children ³⁻⁶: rs9939609 (in/near to FTO), rs17782313 (MC4R), rs6548238 (TMEM18), rs10938397 (GNPDA2), rs368794 (KCTD15), rs2568958 (NEGR1), rs925946 (BDNF) and rs7647305 (ETV5). In doing so only individuals with

115 complete genotype data at all eight variants were included in the analyses and only one 116 variant at each locus was chosen.

117 We selected environmental variables if they had shown a distribution dependent effect on children's BMI^{19, 20} or by a priori considerations. These were dichotomized in order to 118 119 compare FMI distributions in exposed and non-exposed children. Maternal smoking during 120 pregnancy and exclusive formula feeding are established priming factors for childhood overweight ^{19, 20}. As factors associated with life-style, we chose low physical activity (defined 121 122 as child goes to special classes or clubs for some activity (e.g. dancing, judo, sports) less than 123 once a week), high TV consumption (defined as more than 2 hours per day), both at the age 124 of 9 y, and low parental education (i.e. neither father nor mother achieved O-level). Maternal 125 overweight defined as a BMI of at least 25 kg/m² was used as a proxy for both environmental and unknown genetic risk factors. Finally, we defined the following nutritional factors 126 127 extracted from the 7-year food dietary records and dichotomized them as suggested by the Institute of Medicine in 2005 ²¹: high caloric intake (> 1 700 kcal / day in females; > 1 900 128 129 kcal / day in males), low intake of protein (< 10% of daily caloric intake), high intake of 130 carbohydrates (> 60% of daily caloric intake), high intake of saturated fatty acids (> 20% of 131 daily caloric intake) and low intake of mono- and polyunsaturated fatty acids (< 20% of daily 132 caloric intake).

Our analyses were restricted to white European children. In addition to all singletons, we included one randomly selected child from each mother with more than one child in the study $(n = 7 \ 146 \ children)$ in order to avoid potential intercorrelation in close family members. In total, the dataset contained $n = 4 \ 616$ observations with full information on FMI at the age of 9 y, of which $n = 2 \ 346$ contained full information on all environmental variables (see **Table** 138 1). 139

140 Statistical analysis

Quantile regression is a statistical technique which has been applied in a wide research area ²²⁻ 141 142 ²⁵. While traditional mean regression estimates the conditional mean of the outcome 143 distribution, quantile regression estimates the conditional quantiles, i.e. sample percentiles, 144 e.g. the 0.9 quantile / 90th percentile. In case of a binary risk factor the quantile regression 145 coefficient represents the difference of the particular conditional percentile in the estimated 146 outcome distribution, e.g. FMI, in subjects that are exposed or not exposed, whereas all other 147 explanatory variables remain constant. Thus, quantile regression gives a more complete 148 picture of the response distribution than mean regression. Since it does not rely on 149 distributional assumptions, it is even more adequate than mean regression when the outcome 150 distribution is skewed, as in our case for FMI.

We calculated quantile regression models with the 3rd, 10th, 20th, ..., 90th and 97th percentiles 151 152 of FMI as outcomes and the obesity-risk-allele score, maternal smoking during pregnancy, 153 formula feeding, low physical activity, high TV consumption and low parental education as 154 explanatory variables. We did not include the nutritional variables in these models, as these 155 were not independent from each other (since the proportions of daily caloric intake were expected to sum up to 100% for each observation). These variables were assessed separately 156 157 in additional models considering the other environmental factors as potential confounders. 158 Bootstrap methods were used to calculate 95 % confidence intervals (CIs) for the quantile regression effect estimates. We adjusted all models for sex, age and height at the age of 9 y. If 159 160 a specific predictor showed a distribution dependent association with FMI, we additionally 161 calculated separate models that included an interaction term between the obesity-risk-allele

162 score and this factor adjusted for the other potential predictors. Thereby, an effect of the risk 163 allele score was estimated separately for each level of the binary risk factors. If a significant 164 interaction between the obesity-risk-allele-score and a specific risk factor was found, and if 165 the risk factor was significant at the 90th percentile, we stratified our analyses by exposure to 166 this factor.

167 All calculations were carried out with the statistical software R 2.14.2 (<u>http://cran.r-</u> 168 project.org), using the *quantreg* package.

169 **Results:**

The children analyzed had a mean FMI of 4.1 kg/m² and a median FMI of 3.6 kg/m² at 9 years
of age, whereby the distribution was right-skewed (**Table 1**). The prevalence of overweight
(including obesity) according to IOTF criteria ²⁶ based on BMI was 18.2 %.

In the first model without interactions, increasing effect sizes by FMI percentile were found with respect to the obesity-risk-allele-score, maternal overweight, high TV consumption, maternal smoking during pregnancy, low parental education and low protein intake as well as decreasing effect sizes for low intake of unsaturated fatty acids (**Table 2,3**). For example, the estimated regression coefficients of maternal overweight were 0.46 [95% CI: 0.33; 0.59] at the 10th percentile, 0.99 [0.84; 1.14] at the median, and 2.46 [2.07; 2.85] at the 90th percentile.

180 Interaction with the obesity-risk-allele-score

In the second model, there was no evidence for interactions between the obesity-risk-allele score and the environmental variables except for maternal overweight at the 90th percentile of FMI (data not shown). Stratified analyses suggested that exposition to maternal overweight increased the effect of the genetic factors in the upper FMI percentiles, but not in the lower parts of the distribution (**Figure 1**).

With respect to the nutritional factors, the only significant interaction with the obesity-riskallele-score was found with respect to intake of mono- and polyunsaturated fatty acids. In children with low intake, genetic risk was associated with continuously increasing effect sizes by FMI percentile, while in non-exposed children no such pattern occurred (**Figure 2**).

191 **Discussion:**

We found no evidence for a potential interaction of genetic disposition for overweight with a number of established environmental risk factors such as high TV consumption, low physical activity, maternal smoking during pregnancy, formula feeding or low parental education as well as for high caloric intake, high intake of saturated fatty acids, low protein intake or high intake of carbohydrates. However, there was a weak interaction with maternal overweight and a clear interaction with intake of unsaturated fatty acids.

198 The interaction of genetic disposition with unsaturated fatty acids is particularly interesting. We were able to demonstrate that the effect of genetic disposition was weight status 199 200 dependent in children with low intake of unsaturated fatty acids, while there was virtually no 201 effect in children with appropriate intake. This finding might potentially indicate that a 202 genetic disposition for overweight might only have an effect in subjects who have a relatively 203 low intake of unsaturated fatty acids (which might e. g. due to a low proportion of vegetarian 204 food or fish), while an appropriate intake seemed to be protective against the effects of 205 genetic obesity risk factors. Interestingly, this finding would confirm the result of a recent 206 study which showed an interaction between the ratio of polyunsaturated and saturated acids in the diet with the FTO gene ²⁷: A allele carriers with an intake ratio of lower than 0.43 had a 207 208 higher risk for becoming obese than TT carriers irrespective of their intake ratio. However, 209 this finding was based on a relatively small dataset (n=354). With respect to other genetic settings, similar findings had been reported before ^{28, 29}. 210

Our results did not confirm, however, other observational studies reporting interactions between genetic disposition and intake of carbohydrates ^{30, 31}, total fat ^{28, 29, 32, 33} or total energy ^{34, 35}. Unfortunately, we were not able to assess whether this was due to lack of statistical power, as, to our knowledge, no methods exist for power estimation in quantileregression.

Maternal overweight may represent the outcome of the effects of nutrition, life-style and genetic disposition and is therefore likely to be a risk factor for overweight especially in younger children. Therefore, its observed interaction with genetic disposition for overweight may at least partially reflect the effect of polyunsaturated fatty acids mentioned above. An alternative explanation might be that the effects of identified genetic risk factors are modified by other (unknown) obesity risk genes which may also be part of the impact of maternal overweight on offspring's body composition.

The prospective design of the ALSPAC data set constitutes a strength of our analyses because reverse causation is not likely to be an issue. Furthermore, the use of quantile regression enabled us to examine differential effect sizes of explanatory variables and their interactions with respect to FMI in children.

227 As previous evidence with respect to our findings is relatively scarce, confirmation of these 228 results in another dataset would be desirable to exclude potential chance findings. 229 Unfortunately, we had no access to other datasets in which both detailed genotyping and 230 assessment of food habits during childhood had been collected. Further, our dataset was not 231 big enough to allow dividing it into a "training" and a "validation" sample. Another potential 232 weakness of our study might constitute in the fact that diet diary data allowing for detailed 233 assessment of caloric intake etc. were collected at the age of 7 years, while fat mass was 234 measured about two years later. However, this approach has been used previously on these data and yielded plausible results ¹⁸. Furthermore, there was at least reasonable agreement 235

between general food patterns at the ages of 7 and 9 years, as assessed by food frequency
 questionnaires ³⁶.

In conclusion, our analyses suggest potential interactions between a diet with a low content of unsaturated fatty acids and genetic risk factors for overweight on FMI. This effect is likely to be stronger in children with higher FMI. Apart from maternal overweight, which might also reflect unknown genetic factors, we found no evidence for interactions of genetic disposition with other environmental or nutritional factors.

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The authors' responsibilities were as follows: AB developed the study hypothesis, performed data management and contributed to the first and final draft of the paper together with RvK. CR was responsible for statistical analyses and contributed to the first and final draft. ARN contributed to the final draft of the manuscript. NF contributed to the interpretation of the results and to subsequent drafts of the manuscript. KS made suggestions with respect to data analysis and variable selection.

Ethics approval

This study was conducted with the approval of the ALSPAC Ethics and Law Committee.

Competing interests

None of the authors had a conflict of interest.

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Variable	Mean (SD) / n (%)
Fat mass index [kg/m ²]	4.1 (2.2)
Overweight children *	n = 428 (18.2 %)
Obese children *	n = 83 (3.5 %)
Age [years]	9.8 (0.3)
Sex (female)	n = 1,146 (48.8 %)
Maternal overweight	n = 448 (19.1 %)
High TV consumption	n = 781 (33.3 %)
Low parental education	n = 1,016 (43.3 %)
Maternal smoking during pregnancy	n = 963 (41.0 %)
Exclusive formula feeding	n = 389 (16.6 %)
Low physical activity	n = 918 (39.1 %)
High caloric intake	n = 848 (36.1 %)
High intake of saturated fatty acids	n = 70 (3.0 %)
Low intake of unsaturated fatty acids	n = 1,934 (82.4 %)
Low protein intake	n = 126 (5.4 %)
High intake of carbohydrates	n = 333 (14.2 %)

TABLE 1. Characteristics of the study population (n = 2,346).

*	classified	using	IOTF	cut-off	values	(20)
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TABLE 2: Regression coefficients (±1.96 s.e. values) of risk factors as estimated by quantile regression at specific percentiles p. The effects of the

factors obesity-risk-allele score and the environmental factors adjusted by age, height and sex were estimated.

Risk factor	3р	10p	30p	50p	70p	90p	97p
obesity-risk-allele-score	$0.04 (\pm 0.02)$	$0.03 (\pm 0.02)$	$0.09 (\pm 0.02)$	0.13 (± 0.02)	0.23 (± 0.03)	$0.30 (\pm 0.06)$	$0.32 (\pm 0.07)$
maternal BMI > 25	$0.22 (\pm 0.09)$	0.46 (± 0.13)	0.74 (± 0.10)	$0.99 (\pm 0.15)$	1.53 (± 0.22)	$2.46 (\pm 0.39)$	2.17 (± 0.33)
high TV consumption	$0.19 (\pm 0.07)$	$0.19 (\pm 0.07)$	0.21 (± 0.08)	0.30 (± 0.10)	$0.30 (\pm 0.13)$	0.62 (± 0.24)	$0.72 (\pm 0.32)$
smoking during pregnancy	$0.10 (\pm 0.07)$	$0.09 (\pm 0.07)$	0.21 (± 0.06)	$0.29 (\pm 0.09)$	0.21 (± 0.12)	0.23 (± 0.24)	$0.20 (\pm 0.30)$
formula feeding	0.04 (± 0.13)	$0.22 (\pm 0.08)$	$0.25~(\pm 0.08)$	0.24 (± 0.13)	0.33 (± 0.18)	0.17 (± 0.25)	$0.07 (\pm 0.35)$
low physical activity	$0.04 (\pm 0.07)$	$-0.03 (\pm 0.07)$	$0.08~(\pm 0.07)$	$0.03 (\pm 0.09)$	$-0.04 (\pm 0.12)$	$-0.02 (\pm 0.22)$	0.13 (± 0.29)
low parental education	-0.01 (± 0.06)	$-0.09 (\pm 0.06)$	-0.01 (± 0.07)	0.03 (± 0.10)	0.15 (± 0.13)	$0.26 (\pm 0.22)$	0.35 (± 0.29)

TABLE 3: Regression coefficients (±1.96 s.e. values) of risk factors as estimated by quantile regression at specific percentiles p. The effect of each

nutrition factor was estimated separately considering the obesity-risk-allele score and the environmental factors adjusted by age, height and sex.

Risk factor	3 p	10p	30p	50p	70p	90p	97p
high caloric intake	$0.12 (\pm 0.07)$	$0.15 (\pm 0.07)$	$0.09~(\pm 0.07)$	0.21 (± 0.11)	0.23 (± 0.13)	$0.03 (\pm 0.23)$	0.04 (± 0.32)
high intake of saturated fatty acids	$0.40 (\pm 0.18)$	0.27 (± 0.11)	0.01 (± 0.17)	-0.18 (± 0.38)	0.35 (± 0.39)	$-0.20 (\pm 0.69)$	1.52 (± 1.22)
low intake of unsaturated fatty acids	0.02 (± 0.10)	$-0.02 (\pm 0.09)$	-0.09 (± 0.10)	-0.16 (± 0.13)	-0.37 (± 0.20)	-0.61 (± 0.27)	$-0.59 (\pm 0.39)$
low protein intake	-0.07 (± 0.09)	-0.18 (± 0.14)	$0.02 (\pm 0.17)$	$0.20 (\pm 0.24)$	$0.50 (\pm 0.28)$	0.38 (± 0.46)	0.51 (± 0.48)
high intake of carbohydrates	$-0.07 (\pm 0.08)$	-0.13 (± 0.10)	$-0.09 (\pm 0.07)$	-0.33 (± 0.10)	-0.17 (± 0.20)	$-0.22 (\pm 0.29)$	-0.19 (± 0.51)

FIGURE 1: Point estimates and 95% confidence bounds (grey areas) for increase in fat mass index (FMI) at 9 years per obesity-risk-allele a) in interaction with maternal overweight, b) in children of overweight mothers and c) in children of non-overweight mothers.



FIGURE 2: Point estimates and 95% confidence bounds (grey areas) for increase in fat mass index (FMI) at 9 years per obesity-risk-allele a) in interaction with low intake of mono- and polyunsaturated fatty acids, b) in children with low intake of mono- and polyunsaturated fatty acids.

