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Aerobic Fitness Does Not Modify the Effect of FTO Variation on Body Composition Traits

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Abstract

Purpose: Poor physical fitness and obesity are risk factors for all cause morbidity and mortality. We aimed to clarify whether common genetic variants of key energy intake determinants in leptin (LEP), leptin receptor (LEPR), and fat mass and obesity-associated (FTO) are associated with aerobic and neuromuscular performance, and whether aerobic fitness can alter the effect of these genotypes on body composition.

Methods: 846 healthy Finnish males of Caucasian origin were genotyped for FTO (rs8050136), LEP (rs7799039) and LEPR (rs8179183 and rs1137101) single nucleotide polymorphisms (SNPs), and studied for associations with maximal oxygen consumption, body fat percent, serum leptin levels, waist circumference and maximal force of leg extensor muscles.

Results: Genotype AA of the FTO SNP rs8050136 associated with higher BMI and greater waist circumference compared to the genotype CC. In general linear model, no significant interaction for FTO genotype-relative VO2max (mL·kg⁻¹·min⁻¹) or FTO genotype-absolute VO2max (L·min⁻¹) on BMI or waist circumference was found. Main effects of aerobic performance on body composition traits were significant (p<0.001). Logistic regression modelling found no significant interaction between aerobic fitness and FTO genotype. LEP SNP rs7799039, LEPR SNPs rs8179183 and rs1137101 did not associate with any of the measured variables, and no significant interactions of LEP or LEPR genotype with aerobic fitness were observed. In addition, none of the studied SNPs associated with aerobic or neuromuscular performance.

Conclusions: Aerobic fitness may not modify the effect of FTO variation on body composition traits. However, relative aerobic capacity associates with lower BMI and waist circumference regardless of the FTO genotype. FTO, LEP and LEPR genotypes unlikely associate with physical performance.

Introduction

Obesity and low physical fitness frequently associate with each other, and are individual risk factors for many pathological conditions and cardiovascular mortality [1,2]. Body composition and physical performance are outcomes of cumulative effects of common and rare variants in a large number of genes with environmental and gene-gene interactions [3]. The effects of behavioural factors on human performance, including physical activity and dietary habits have been widely studied, yet knowledge on genetic background is sparse. The heritability estimates for obesity range from 40% to 70% [3], from 20 to 40% for aerobic performance [4,5], and approximately 60% for muscle force [6]. Single nucleotide polymorphisms (SNPs) represent the most common type of genetic variation in the human genome [7]. However, it is not known whether genetic variants in genes encoding fat and obesity-associated (FTO), leptin (LEP), and leptin receptor (LEPR) associate with physical performance, and whether fitness level modifies the risk for obesity associated with these gene variants.

Genome-wide association studies have identified the FTO gene as the first susceptibility locus for common obesity [8,9]. Minor allele of this variant is associated with increased risk for obesity and elevated body weight [8,9]. In a recent meta-analysis, physical activity level was shown to modify the relationship between the FTO risk variant and body weight and risk of obesity [10]. However, effects of FTO genotype and its interaction with aerobic fitness on body composition have not been reported so far. Despite the fact that FTO is under extensive research, the role and function of the FTO gene product remains incompletely understood.

Leptin is a peptide hormone secreted mainly from white adipose tissue. It regulates hunger, body temperature and energy metabolism, and has been used as a biomarker of energy deficiency [11,12]. Recent studies have shown that obese individuals are in fact leptin resistant [12], and that physical exercise may restore leptin sensitivity [13]. Associations of leptin...
The present study investigated the association of selected SNPs in *FTO*, *LEP* and *LEPR* with body composition, neuromuscular and cardiorespiratory performance and plasma leptin levels in 846 healthy male subjects. In addition, genotype-aerobic fitness interactions on body composition traits were studied. Our hypothesis was that these common variants are associated with human performance, body composition and health-related risk factors. Furthermore, we hypothesized that aerobic fitness modifies the relationship between the genetic variants and body composition. The information this study provides may be used to identify those individuals having higher health risks and tendency for poor physical fitness, and for better understanding of the effect of genetic factors on human physical performance.

**Materials and Methods**

**Study Subjects**

The subjects of Caucasian origin were 846 healthy male Finnish volunteers with mean age (SD) of 25±5 years. All subjects were informed on the purpose of the study, gave a written informed consent and underwent medical examination prior to the tests. This study was approved by the ethics committee of the University of Jyväskylä and the hospital district of Central Finland. Anthropometric data and blood samples were collected after an overnight fast. The subjects consumed light breakfast 1 to 2 hours before the exercise tests.

**Physical Performance and Body Composition**

Height, weight and waist circumference were recorded, and body mass index (BMI, kg·m⁻²) was calculated. Body fat and lean mass percentage were recorded by using the eight-polar bioimpedance method with multifrequency current (InBody 720; Biospace Company, Seoul, Korea). Bioimpedance was recorded after an overnight fast and with at least one day off from any intensive physical activity.

Maximal isometric force production of the leg extensor muscles was measured with a dynamometer (Department of Biology of Physical Activity, University of Jyväskylä, Finland). The test was performed in sitting position with 107-degree knee angle. The subjects were supervised to generate maximal force as fast as possible and maintain this force for 3 seconds. The data were analysed with a 16-bit AD converter (CED power 1401, Cambridge Electronic Design Ltd, England) and Signal (2.16) software.

Aerobic fitness was assessed by maximal bicycle ergometer test (Ergoline 800 S, Ergoselect 100 K, Ergoselect 200 K, Bitz, Germany) as previously described [22]. The initial workload was 50 W with 25 W increase on 2-minute intervals until exhaustion. Heart rate was monitored throughout the test (Polar T-31; Polar Vantage, Kempele, Finland). The analyzed variables were maximum heart rate, maximal workload and maximal oxygen consumption (mL·kg⁻¹·min⁻¹) estimated by software (Mifit4/ Fitware, Finland) [(11.016 · maximum work load) x (1·body weight⁻²) +7.0]. The test was terminated when the subject could not maintain the required cycling speed (60-90 rpm). Physical activity was assessed with international physical activity questionnaire (IPAQ) [23].

**Blood Samples and Genotyping**

The fasting blood samples were collected and analysed immediately with a hemacytometer (Sysmex Co., Kobe, Japan). Plasma was separated from the whole blood and stored at -80°C until analysis. Plasma leptin concentrations were assayed by commercial ELISA according to the manufacturer's instructions (Quantikine, R&D Systems, Minneapolis, MN, USA). Assay specifications were as follows: sensitivity limit 7.8 pg·mL⁻¹, and maximum intra- and interassay CV% 3.3% and 5.4%, respectively.

The SNP genotyping was performed using allele-specific PCR assays. Briefly, genomic DNA was first isolated from the blood mononuclear cells using QIAamp DNA Blood kit (Qiagen, Hilden, Germany). Next, 50 nanograms of the DNA was amplified with Brilliant QPCR Master Mix (Stratagene, La Jolla, CA, USA) and allele-specific SNP assays on a MX3000P Real-time PCR System (Stratagene). For *FTO* (rs8050136) and *LEPR* SNP Lys656Asn (rs8179183), the commercially available TaqMan SNP genotyping assays were used (Applied Biosystems, Foster City, CA, USA), and for *LEP* (rs7799039) and *LEPR* SNP Gln223Arg (rs1137101), the following molecular beacons SNP genotyping assays were used: LEP SNP forward primer 5′-CCTGAATTTCCTCATGAGAC-3′ and reverse primer 5′-TGCAACATCTCAGCACCTATG-3′, and the molecular beacons 5′-FAM/HEX-CGGACGTGGAGTAATTTTCC(A/G)-BHQ1; LEPR SNP forward primer 5′-TCACGGACACTCTCCGTATG-3′ and reverse primer 5′-TTATGGGCTGAACTGACAT-3′, and the molecular beacons 5′-FAM/HEX- CGGACGTGGAATTTTCC(A/G)TCACCTCGCCG-3′.

**Statistics**

Calculations were performed with SPSS software (Illinois, Chicago, USA) by using one-way ANCOVA or non-parametric tests, when appropriate. Age, smoking and earlier physical activity were set as covariates. Smoking (smoker or non-smoker) and physical activity (vigorous physical activity more than 3 times per week) were set as dichotomous variables, whereas age VO2max, BMI, waist circumference were set as continuous variables.

Genotype-VO2max interaction on BMI, waist circumference and fat percent were analyzed by using general linear model. The general linear model included main effects terms for earlier physical activity, smoking, age, genotype, VO2max and genotype-VO2max interaction. Physical activity and smoking were set as dichotomous variables, and VO2max, age and body composition traits were set as continuous variables. Genotype was set as discrete variable in every analysis.

Additionally, the *FTO* genotype and *FTO* genotype-VO2max interaction related odds for overweight (BMI over 25 kg·m⁻²) and abdominal obesity (waist circumference over 90 cm) were analyzed by logistic regression. Recessive model was applied in analysis (AA genotype vs. CA and CC genotype). Age, smoking and physical activity were set as covariates. The subjects were divided into quartiles according to VO2max. The 75% percentile limits for VO2max were 3.7 L·min⁻¹ and 46.9 mL·kg⁻¹·min⁻¹. The subjects in the highest 25% quartile were defined as high VO2max, and compared with the 75% percentile of subjects (VO2max <3.7 L·min⁻¹ or <46.9 mL·kg⁻¹·min⁻¹).

Data are presented as mean ± standard deviation unless otherwise stated. Statistical significance was set at p<0.0125 to account for multiple testing. Bonferroni correction was applied to
post-hoc comparisons. Statistical power and sample sizes were estimated with SISA web calculator [24]. A 90% level was chosen for power calculations, and the number of subjects needed to demonstrate 1 S.D. difference in continuous parameters was estimated.

Results

All SNPs conformed to Hardy-Weinberg’s equilibrium. None of the SNPs were associated with earlier physical activity estimated with IPAQ.

In ANCOVA analysis, the FTO SNP rs8050136 associated with BMI and waist circumference (Table 1). Genotype AA carriers had significantly higher BMI and greater waist circumference compared to the genotype CC carriers. In general linear model, the main effects of relative and absolute VO$_2$max on BMI and waist circumference were significant ($p<0.001$). However, no significant FTO genotype-relative VO$_2$max (mL·kg$^{-1}$·min$^{-1}$) interaction on BMI ($p=0.081$) or waist circumference ($p=0.093$) was found. In addition, the test result was insignificant when FTO genotype-absolute VO$_2$max (L·min$^{-1}$) interaction on BMI ($p=0.081$) or waist circumference ($p=0.093$) were analyzed (Figures 1–4). In logistic regression, genotype AA increased odds for waist circumference over 90 cm. However, the interaction of genotype AA with high VO$_2$max was insignificant (Table 2). No association was found with aerobic fitness (VO$_2$max expressed as L·min$^{-1}$ or mL·kg$^{-1}$·min$^{-1}$) (Table 1).

In ANCOVA analysis, LEP SNP rs7799039 did not associate with any variable (Table 3). No interactions were observed between aerobic fitness and genotype. LEPR SNP rs1137101 did not associate with BMI, body fat, leptin levels or physical performance in ANCOVA (Table 3), and no aerobic performance-genotype interactions were observed in the general linear model. In ANCOVA analysis, LEPR Lys656Asn SNP rs8179183

Table 1. Showing the results of ANCOVA–analysis according to FTO genotype.

<table>
<thead>
<tr>
<th>FTO SNP rs8050136 (N = 773)</th>
<th>Genotype</th>
<th>CC</th>
<th>CA</th>
<th>AA</th>
</tr>
</thead>
<tbody>
<tr>
<td>N</td>
<td>269</td>
<td>380</td>
<td>124</td>
<td></td>
</tr>
<tr>
<td>BMI (kg·m$^{-2}$)</td>
<td>24.5±3.6</td>
<td>24.8±3.7</td>
<td>25.7±4.3*</td>
<td></td>
</tr>
<tr>
<td>Waist circumference (cm)</td>
<td>85.8±10.3</td>
<td>86.0±9.7</td>
<td>88.8±12.2#</td>
<td></td>
</tr>
<tr>
<td>Fat per cent (%)</td>
<td>17.5±7.2</td>
<td>17.9±7.0</td>
<td>19.4±8.0</td>
<td></td>
</tr>
<tr>
<td>VO$_2$max (mL·kg$^{-1}$·min$^{-1}$)</td>
<td>42.0±8.3</td>
<td>41.8±7.8</td>
<td>40.2±8.6</td>
<td></td>
</tr>
<tr>
<td>VO$_2$max (L·min$^{-1}$)</td>
<td>3.29±0.56</td>
<td>3.30±0.58</td>
<td>3.30±0.62</td>
<td></td>
</tr>
<tr>
<td>Maximal working capacity (Watts)</td>
<td>244.2±44.7</td>
<td>245.0±46.5</td>
<td>242.7±49.1</td>
<td></td>
</tr>
<tr>
<td>Maximal force of leg extensors (N)</td>
<td>2927±872</td>
<td>2950±816</td>
<td>2942±992</td>
<td></td>
</tr>
<tr>
<td>Plasma leptin (pg·mL$^{-1}$)</td>
<td>3598±3657</td>
<td>3702±3780</td>
<td>4289±4278</td>
<td></td>
</tr>
</tbody>
</table>

Earlier physical activity, smoking and age were set as covariates. Bonferroni correction was accounted for multiple post-hoc comparisons. BMI: $p=0.007$ for main effect, *$p=0.005$ between the genotype groups CC and AA.

Waist circumference: $p=0.012$ for main effect, #$p=0.012$ between the genotype groups CC and AA.

Maximal working capacity: $p=0.007$ for main effect, #$p=0.005$ between the genotype groups CC and AA.

Maximal force of leg extensors: $p=0.012$ for main effect, #$p=0.012$ between the genotype groups CC and AA.

Plasma leptin: $p=0.012$ for main effect, #$p=0.012$ between the genotype groups CC and AA.

Figure 1. Linear regression analysis. The main effect of VO$_2$max (mL·kg$^{-1}$·min$^{-1}$) on BMI was significant ($p<0.001$). The main effects of FTO genotype and FTO genotype-VO$_2$max (mL·kg$^{-1}$·min$^{-1}$) interaction were not significant ($p=0.029$ and $p=0.081$, respectively).
doi:10.1371/journal.pone.0051635.g001
did not associate with any variable (Table 3). In addition, no interactions were observed between genotype and aerobic fitness in the general linear model.

**Discussion**

The main finding of the present study was that aerobic fitness does not modify the effect of *FTO* variation on BMI or waist circumference. However, relative aerobic capacity did associate

**Table 2. Showing results of the logistic regression analysis.**

<table>
<thead>
<tr>
<th></th>
<th>OR</th>
<th>95% C.I.</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>BMI over 25 kg m(^{-2}) adjusted for age, smoking, physical activity and relative aerobic fitness (mL kg(^{-1}) min(^{-1})</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Genotype AA</td>
<td>1.51</td>
<td>0.96–2.37</td>
<td>0.073</td>
</tr>
<tr>
<td>VO(_{2\text{max}}) (&gt;46.9 mL kg(^{-1}) min(^{-1}))</td>
<td>0.15</td>
<td>0.09–0.25</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Genotype AA by VO(_{2\text{max}}) (&gt;46.9 mL kg(^{-1}) min(^{-1}))</td>
<td>1.31</td>
<td>0.43–3.96</td>
<td>0.639</td>
</tr>
<tr>
<td><strong>BMI over 25 kg m(^{-2}) adjusted for age, smoking, physical activity and absolute aerobic fitness (L min(^{-1}))</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Genotype AA</td>
<td>1.61</td>
<td>1.02–2.53</td>
<td>0.038</td>
</tr>
<tr>
<td>VO(_{2\text{max}}) (&gt;3.7 L min(^{-1}))</td>
<td>1.10</td>
<td>0.75–1.63</td>
<td>0.626</td>
</tr>
<tr>
<td>Genotype AA by VO(_{2\text{max}}) (&gt;3.7 L min(^{-1}))</td>
<td>1.02</td>
<td>0.41–2.58</td>
<td>0.959</td>
</tr>
<tr>
<td><strong>Waist circumference over 90 cm adjusted for age, smoking, physical activity and relative aerobic fitness (mL kg(^{-1}) min(^{-1}))</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Genotype AA</td>
<td>1.92</td>
<td>1.22–3.03</td>
<td>0.005</td>
</tr>
<tr>
<td>VO(_{2\text{max}}) (&gt;46.9 mL kg(^{-1}) min(^{-1}))</td>
<td>0.08</td>
<td>0.04–0.18</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Genotype AA by VO(_{2\text{max}}) (&gt;46.9 mL kg(^{-1}) min(^{-1}))</td>
<td>1.07</td>
<td>0.19–5.94</td>
<td>0.941</td>
</tr>
<tr>
<td><strong>Waist circumference over 90 cm adjusted for age, smoking, physical activity and absolute aerobic fitness (L min(^{-1}))</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Genotype AA</td>
<td>1.96</td>
<td>1.22–3.15</td>
<td>0.005</td>
</tr>
<tr>
<td>VO(_{2\text{max}}) (&gt;3.7 L min(^{-1}))</td>
<td>1.26</td>
<td>0.81–1.97</td>
<td>0.299</td>
</tr>
<tr>
<td>Genotype AA by VO(_{2\text{max}}) (&gt;3.7 L min(^{-1}))</td>
<td>0.88</td>
<td>0.33–2.40</td>
<td>0.884</td>
</tr>
</tbody>
</table>

OR and 95% C.I. for the genotype AA (vs. genotype CA and CC) of the FTO rs8050136 according to overweight and abdominal obesity.

doi:10.1371/journal.pone.0051635.t002
Figure 3. **Linear regression analysis.** The main effect of VO$_2$max (mL·kg$^{-1}$·min$^{-1}$) on waist circumference was significant (p<0.001). The main effects of FTO genotype and FTO genotype-VO$_2$max (mL·kg$^{-1}$·min$^{-1}$) interaction were not significant (p = 0.040 and p = 0.093, respectively).
doi:10.1371/journal.pone.0051635.g003

Figure 4. **Linear regression analysis.** The main effect of VO$_2$max (L·min$^{-1}$) on waist circumference was significant (p<0.001). The main effects of FTO genotype and FTO genotype-VO$_2$max (L·min$^{-1}$) interaction were not significant (p = 0.521 and p = 0.262, respectively).
doi:10.1371/journal.pone.0051635.g004
with lower BMI and waist circumference regardless of the FTO genotype. We also report that the LEP promoter -2548 SNP and the LEPR Lys656Asn SNP did not associate with BMI, waist circumference or body fat percent.

Our observations that genotype AA of the FTO variant associated with higher BMI and greater waist circumference are in line with a recent meta-analysis [25]. Our novel finding, however, is that aerobic fitness does not modify the effect of FTO variation on BMI and waist circumference. In another recent meta-analysis, the level of physical activity was shown to modify the relationship between the FTO risk variant and body weight and risk of obesity [10]. Nevertheless, these studies are not fully comparable to ours because at present, this is the first study to report objectively measured aerobic fitness - FTO-interaction. The questionnaire-based assessment of physical activity may have some reporting bias [26]. In addition, the present findings may be explained by the fact that apart from physical activity, other factors may affect aerobic fitness. Approximately 20 to 40% of VO2max has been estimated to be heritable, and other factors, such as physical activity, intake of carbohydrates, smoking, body weight, blood haemoglobin levels, age and presence of chronic disease account for the remaining [4, 5]. In addition, the present and earlier studies differ in study design because in our study, all subjects were young males. Nevertheless, in the present study, aerobic fitness was associated with body composition traits regardless of the FTO genotype. Despite lack of FTO genotype-aerobic fitness interaction, the genotype AA carriers are more susceptible to obesity and may benefit from increased aerobic fitness to a greater extent due to its favourable effects on the regulation of body weight. In the present study, FTO variant was not found to associate with aerobic fitness, which is in line with previous studies [27, 28]. It can be speculated that low physical fitness combined with increased body weight in the allele A carriers of the FTO variant may further increase risk for lifestyle-associated disease, such as cardiovascular events [29]. This would in turn highlight the importance of physical activity to attenuate risk for obesity. Furthermore, the studied FTO variant did not associate with plasma leptin levels, which in line with other studies [30, 31].

In the present study, the leptin promoter SNP rs7799039 did not associate with BMI, which is supported by earlier reports [17, 18]. It has been speculated that lower leptin levels in allele G carriers of this SNP would be an underlying mechanism for increased adipose tissue [32]. However, in the present study, leptin levels were similar between the genotype groups. Furthermore, no association with aerobic fitness, neuromuscular performance, or interaction with VO2max was found.

The LEPR Gln223Arg genotype showed no association with cardio-respiratory fitness, which is supported by our previous report [19]. Furthermore, in agreement with other studies [19, 33], no differences in leptin levels between the genotype groups were found, although controversial findings do exist [15, 34, 35]. It should be noted, however, that in many published studies, interpretation of the results is limited due to small sample size, differences in ethnicity of the subjects, and other factors related to the study settings.

The LEPR Lys656Asn SNP did not associate with BMI, waist circumference or body fat percent, and these findings are also supported by others [36-38]. Moreover, subjects of the present study were not morbidly obese, and small prevalence of the minor Asn allele may also complicate evaluation of the effect of this SNP on body weight and body composition. Therefore, no significant genotype-VO2max interactions were found.

<table>
<thead>
<tr>
<th>Genotype</th>
<th>LEP2548SNP</th>
<th>LEPRGln223Arg</th>
<th>FTO-2548SNP</th>
<th>Fat per cent %</th>
<th>VO2max (ml min^-1 kg^-1)</th>
<th>Maximal working capacity (Watts)</th>
<th>Maximal force of leg extensors (N)</th>
<th>Plasma leptin (pg mL^-1)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GG</td>
<td>N = 176</td>
<td>N = 153</td>
<td>N = 175</td>
<td>9.7 ± 6.3</td>
<td>2.6 ± 0.5</td>
<td>2.0 ± 0.3</td>
<td>837 ± 293</td>
<td>46.1 ± 137</td>
</tr>
<tr>
<td>GA</td>
<td>N = 549</td>
<td>N = 472</td>
<td>N = 545</td>
<td>10.8 ± 6.9</td>
<td>2.7 ± 0.5</td>
<td>2.1 ± 0.4</td>
<td>863 ± 293</td>
<td>43.7 ± 137</td>
</tr>
<tr>
<td>AA</td>
<td>N = 12</td>
<td>N = 12</td>
<td>N = 13</td>
<td>9.3 ± 6.1</td>
<td>2.4 ± 0.4</td>
<td>1.9 ± 0.3</td>
<td>787 ± 293</td>
<td>41.1 ± 137</td>
</tr>
</tbody>
</table>

Table 3: Showing the results of ANCOVA – analysis according to LEP and LEPR genotypes.
There are certain limitations to the present study. First, the sample size was moderate for genetic association analyses. The relatively large confidence intervals may bias identification of a true interaction in logistic regression analysis. In addition, all subjects were males, which may affect generalization of the results to other populations. Our study design was cross-sectional, which may increase risk for selection bias. However, earlier physical activity status was assessed with IPAQ survey, and no associations with pre-study physical activity and genotypes were found. The bioimpedance method is both sensitive and specific for analysis of body fat mass, but may have limitations for accurate evaluation of muscle mass [39]. The method for assessing aerobic fitness was indirect, whereas direct gas exchange analysis would provide greater accuracy.

In conclusion, aerobic fitness may not modify the effect of FTO variation on body composition traits. However, relative aerobic capacity had a favourable effect on BMI and waist circumference, regardless of the FTO genotype. It is unlikely that FTO, LEP and LEPR genotypes associate with physical performance. These findings can be applied in sports medicine to decrease risk for diseases related to sedentary lifestyle and obesity, where improved aerobic fitness is beneficial regardless of the FTO-related predisposition to obesity.

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Author Contributions

Conceived and designed the experiments: MS KH HK JL MA. Performed the experiments: AH JL MA. Analyzed the data: AH JL NO. Contributed reagents/materials/analysis tools: AH JL MA. Wrote the paper: AH JL NO MA.

References

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FTO Genetic Variation and Aerobic Fitness
