c fe e ce e ie .c

JOINT EVENT 10th International Conference on

&

2nd International Conference on

June 12-13, 2017 Rome, Italy

Tatyana Bairova, Ekaterina A Sheneman, Kseniia D levleva and Lubov V Rychkova 6FLHQWLÄF FHQWHU IRU IDPLO\ KHDOWK DQG KXPDQ UHSURGXFWLRQ SUREOHPV 5XVVLDQ)HGI

Statement of the Problem: Among the causes of obesity an important role is played by the heredity. $\,$ e FTO is one of the genes associated with obesity and body mass fat. And polymorphism rs9939609 of this gene located in the $\,$ rst intron (the code is W=A/T) appears the most signi $\,$ cant.

Methodology & eoretical Orientation: e study included total of 128 Caucasian female adolescents (average age 15.86 ± 1.02) living in Eastern Siberia (Irkutsk region, Russia). SDS BMI, % of body mass fat was measured. 59 girls were included in group with normal weight (SDS BMI 0.84 ± 0.55) and 69 girls were included in group with overweight and obesity (SDS BMI 2.52 ± 0.72). Genomic DNA was extracted from EDTA-treated whole blood by commercial kits (DNA-Sorb-B, AmpliSens, Russia). Genotyping of the *FTO* rs9939609 was performed using polymerase chain reaction in real time with DT-Prime cycler (DNA-technology, Russia). Statistical analysis was performed by so "STATISTICA8.0".

Findings: A-allele frequency was 40% in control group and 49% was in group with overweight and obesity (=0.223). Comparisons of SDS BMI and percent of body mass fat stratied by *FTO rs9939609* genotypes are shown in table 1. We found the signicant increase of SDS BMI in carriers of A-allele in group with overweight and obesity. ere is no association in group with normal weight.

Conclusion & Signi cance:

Volume 7, Issue 3 (Suppl)