# 20<sup>th</sup> annual Congress of the

# **EUROPEAN COLLEGE OF SPORT SCIENCE**

24<sup>th</sup> - 27<sup>th</sup> June 2015, Malmö – Sweden **BOOK OF ABSTRACTS** 

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Radmann, A., Hedenborg, S., Tsolakidis, E.

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#### DEVELOPMENT OF THE ATHLETES RECRUITING SYSTEM IN CYCLIC SPORTS BASED ON INDIVIDUAL MORPHOFUNC-TIONAL AND GENETICS PARAMETERS

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INTRODUCTION Modern approaches to the sportsmen recruitment and individualization of the training process should take into account both genetic and environmental factors contribution to the making of the world-class athlete. The aim of this study was to develop a genetic monitoring program to determine the individual profile for athletes reserve in cyclic sports. METHODS The study group comprised 250 swimmers, both males and females, aged 10-19. We evaluated the morpho-functional state of swimmers using morphological and functional suitability scales (Davydov V.Y. et al.). Genotyping and frequency of 5HTT, ACE, AGT, PPARA, PPARD, PPARGC1A, and TFAM polymorphisms were obtained by site-specific PCR amplification that was performed on genomic DNA isolated from buccal epithelial

cells (Akhmetov I.I.), RESULTS The morphofunctional state evaluation showed that most of the athletes had 'average' (66.7% total: 62.2% in airls; 69.7% in boys) and 'above average' morphophysiological level (21.6% total; 24.4%; in airls; 21.6% in boys). 9.0% of the study group had 'below the average' level (8.9% in girls; 9.1% in boys), and only 2.7% had 'high level' (4.5%; and 1.5%). The prevalence of the following aenotypes was discovered: AGT CC (77.78%±5.66% in boys; 73.17%±6.92% in girls), PPARA GG (65.38%±6.60%; 70%±7.25%) TFAM GG (59.62%±6.80%; 53.66%±7.79%), PPARD TT (58.82%±6.90%; 70%±7.25%). These genotype frequencies were significantly higher than in controls. There was also significant difference of the distribution of the ACE ID in boys (24.53%±5.91%) and girls (46.34%±7.79%), and ACE Il in study group (22.64%±6.15% in boys vs. 14.63%±5.52% in girls); the frequency of the TFAM GC was also significantly higher in girls (31.71%±7.27% vs. 13.46%±4.73%). We also observed the trend in PPARD TT genotype distribution (70%±7.25% in girls vs. 58.82%±6.90% in boys). DISCUSSION Conducted anthropological examination allowed to develop the model parameters of morphological and functional parameters of the athletes. We analyzed the morphophysiological developmental level based of these characteristics and evaluated maximum potential taking into account so called 'testosterone peak'. The most promising athletes were observed in groups of 14 and 15 years (airls) and 11 and 15 years (boys). On the basis of previous studies we selected 5 candidate genes and studied a common genetic profile of the swimmers. As a result, the basic polymorphic genes set for subsequent screening study was developed. Thus, AGT C, PPARA G, PPARD T and TFAM G alleles can be considered as relevant genetic markers for swimmers. After initial screening examination of these genetic markers further molecular genetic testing for the 13 gene polymorphism will be carried out. References Akhmetov II Molecular genetics of sports: a monograph. – M.: Soviet Sport, 2009. – 268 p. (RUS) Davydov VY Selection and orientation of swimmers in terms of physique in the long-term training (theoretical and practical aspects): monograph / Davydov VY, Avdienko VB. (2013). – M.: Soviet Sport, 2013. – 384 p. (RUS)

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