# FREQUENCY OF THE IGF A/G rs7136446 POLYMORPHISM AND ATHLETIC PERFORMANCE

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#### ABSTRACT

Previous studies have shown that carrying the minor T allele of the IGF C-1245T polymorphism was associated with higher circulating IGF-I levels, greater muscle mass and improved power athletic performance. The aim of the present study was to assess the frequency distribution of another IGF-I single nucleotide polymorphism (SNP), the A/G rs7136446, among Israeli athletes. The IGF A/G rs7136446 polymorphism was determined in 185 short (n=72) and long-distance (n=113) runners, 94 short (n=44) and long-distance (n=50) swimmers, 54 weight-lifters and 111 controls. There were no significant differences in GG carriers, previously described as associated with higher maximal force production, between the athletes and controls. The only statistical significant difference in GG carriers was found between the sprinters (24%) and weight-lifters (9%; p<0.05). Although a single polymorphism cannot determine an athlete's ability to succeed or fail in sports, the present findings suggest a potential importance of IGF-I polymorphisms mainly to power sports and in particular to speed sport performance.

Keywords: genetic profile; athletic performance; speed sport events

#### INTRODUCTION

Directing children to focus in a single sport before adolescence is discouraged. Despite that, an increasing number of children do specialize in specific sport at an early age, and even compete in "elite" level. Moreover, several Olympic sports initiate their selection processes in an attempt to identify future champions even before elementary school [13]. Therefore, it seems that one of the key factors for competitive sports success is identification of a sport event that best matches the athlete's ability at a proper age [32]. Recent efforts to identify relevant genes for human athletic excellence are difficult, mainly because each gene makes only a small contribution to the overall heritability. However, genetic polymorphism has been suggested as an additional, sometimes very important, piece in the puzzle of sport selection. The idea that genetic hormonal polymorphism may also be used for prediction of sports excellence has emerged in recent years, and is only in its origins.

It is now well established that changes in insulin-like growth factor-I (IGF-I) play a pivotal role in the muscle adaptation to exercise [1, 9, 11, 14, 15, 25, 28]. Previous studies in pre and late pubertal girls have shown that both functional (i.e. maximal oxygen consumption - VO<sub>2max</sub>) and structural (i.e. thigh muscle volume determined by magnetic resonance images) indices of fitness were correlated with serum IGF-I levels [10, 12]. Therefore, the majority of studies on hormonal genetic polymorphism and athletic success focused on the IGF-I system. Interestingly, very few studies examined the variation of IGF-I polymorphism in professional athletes. C-1245T (rs35767) is a genetic variation in the promoter region of the IGF-I gene. The minor allele T is associated with higher circulating IGF-I levels [7, 26, 27]. The frequency of the IGF-I promoter polymorphism was significantly greater among athletes compared to non-active controls with a considerably higher sequence frequency among strength athletes compared to team-sport athletes [20]. Higher VO<sub>2max</sub> level was found among Polish athlete's carriers of the i88 and i94 allele tandem repeats of the IGF gene polymorphism, while carriers of the i86 allele had lower VO<sub>2max</sub> levels [16]. Consistent with the previous results, we demonstrated a higher frequency of the T/T polymorphism among Israeli athletes compared to non-existence among controls [4]. While half of the T/T polymorphism carriers were endurance athletes and half power athletes, endurance T/T carriers were of national level, but the power athletes were top-level athletes (international and Olympic). This suggested that the IGF-I T/T polymorphism possible association with greater muscle mass is more important to power sport excellence.

The aim of the present study was to assess the frequency distribution of another IGF-I single nucleotide polymorphism (SNP), the A/G, rs7136446, among Israeli athletes. It was previously demonstrated that GG homozygotes of the IGF-I rs7136446 SNP showed higher maximal force production compared with the other genotypes, and logistic regression analysis showed that presence of the A allele decreased the odds for higher maximal force production. Therefore, we hypothesized that the prevalence of the GG IGF-I rs7136446 SNP would be higher among the athletes and mainly among the power athletes.

## MATERIALS AND METHODS

### Participants

Two hundred and thirty-nine track and field athletes (182 males and 57 females, age 17-55 years) and 94 swimmers (61 males and 33 females, age 16-49 years) participated in the study. Track and field athletes were assigned to three main sub-groups according to their event speciality during their athletic career: 1) long-distance runners (major event: 5000 m to marathon run, n=113), 2) short-distance runners (100-200 m sprinters and jumpers, n=72), and 3) weight lifters (major event: Snatch, Clean and jerk, n=54). Swimmers were assigned to two groups according to their main swimming event during their swimming career: 1) long-distance swimmers (major event: 400-1500 m swim, n=50), and 2) short-distance swimmers (major event: 50-100 m swim, n=44). All athletes were ranked among the top alltime Israeli results in their event and had competed at national and/or international level on a regular basis. The control group consisted of 113 (92 males and 21 females, age 20-29 years) non-athletic healthy individuals who were not engaged in competitive sport. Characteristics of the athletes and controls are presented in Table 1.

Group	n	Main event	M/F	Top/ National level	Age (Mean±SD, range)	
Endurance runners	113	5000m-marathon	92/21	48/65	32.4 ±9.5 (17–55)	
Sprinters/Jumpers	72	100–200m, jumps	49/23	28/44	30.5 ±13.7 (17–71)	
Weight lifters	54	Snatch, Clean & Jerk	41/13	34/20	28.7 <u>+</u> 9.4 (17–47)	
Long distance swimmers	50	400–1500m	32/18	20/30	25.3 ±9.3 (16–48)	
Short distance swimmers	44	50–100m	29/15	10/34	23.1 ±7.6 (16–49)	
Controls	111	_	77/34	-	26.4 ±5.8 (19–29)	

The study was approved by the Institutional Review Board of the Hillel Yaffe Medical Center, Hadera, Israel, according to the Declaration of Helsinki. Written informed consent was obtained from each participant before the commencement of the study.

### Genotyping

Genomic DNA was extracted from peripheral EDTA treated anti-coagulated blood using a standard protocol, described elsewhere [24]. Briefly, Lyses buffer (5 mL) containing 100 mM of tris HCl, pH = 8.5 + 0.5 M of EDTA + 10% SDS + 5 M of NaCl 40 mL added D.D.W. to 1000 mL was added to the white cell solution. Proteinase K (200 µg) was added to the solution. The tubes were incubated at 50°C in a shaker incubator for overnight for digestion. After digestion, one volume of ethanol ( $-20^{\circ}$ C) was added to the lysate and the samples were swirled until precipitation was completed (20–30 min, until it became completely transparent). The DNA was recovered by lifting the aggregated precipitate from the solution, using a disposable tip. Excess liquid was dabbed off and the DNA was dispersed in a pre-labelled eppendorf tube. The DNA was washed in TE solution. Genotypes were determined using the TaqMan allelic discrimination assay. The Assay-by-Design service (http://www.thermofisher.com) was used to set up a TaqMan allelic discrimination assay for the IGF1 A/G (rs7136446).

Primer sequences were forward: AATTGGTTACCTGCTACATTGA, reverse: G GAGTTAACGCATCTCCTTACTG. Probe sequences were for -174G/C forward: VIC- CGCGTAGTCGAGCG, reverse: FAM-CGCTCGCTGCCCTAAGTGCT. The PCR reaction mixture included 5ng genomic DNA, 0.125µl TaqMan assay (40\*, ABI), 2.5µl Master mix (ABI) and 2.375µl water. PCR was performed in 96 well PCR plates in an ABI 7300 PCR system (Applied Biosystems Inc., Foster City, CA, USA) and consisted of initial denaturation for 5min at 95°C, and 40 cycles with denaturation of 15s at 95°C and annealing and extension for 60s at 63°C. Results were analyzed by the ABI TaqMan 7900HT using the sequence detection system 2.22 software (Applied Biosystems Inc.).

# Statistical analysis

The SPSS statistical package, version 20.0, was used to perform all statistical analyses (SPSS, Chicago, IL, USA). A  $\chi^2$ -test was applied to confirm that the observed genotype frequencies were within the Hardy-Weinberg equilibrium and to compare allele and genotype frequencies between athletes and controls, as well as between athletes from different sports and between different competitive groups (e.g. short versus long-distance). If observed or

expected values included a cell with a value of 5, we used a Fisher's exact test to compare allele and genotype frequencies.

### RESULTS

The prevalence of the A/G IGF-I rs7136446 single nucleotide polymorphism among Israeli long and short distance runners and swimmers and among weight lifters is shown in Table 2 and Figure 1. Genotype subtype did not differ by age or sex. In addition, IGF-I A/G genotype distribution was in agreement with the Hardy–Weinberg equilibrium within all groups (p > 0.05).

		Genotype frequencies					Allele frequencies				
	n	AA		AG		GG		A – allele		G – allele	
		Ν	%	Ν	%	Ν	%	Ν	%	Ν	%
Control	111	45	41%	50	45%	16	14%	140	63%	82	37%
Endurance Runners											
Top level	48	18	38%	23	48%	7	15%	59	61%	37	39%
National level	65	28	43%	26	40%	11	17%	82	63%	48	37%
Total	113	46	41%	49	43%	18	16%	141	62%	85	38%
Sprinters and Jumpers											
Top level	28	9	32%	13	46%	6	21%	31	55%	25	45%
National level	44	15	34%	18	41%	11	25%	48	55%	40	45%
Total	72	24	33%	31	43%	17	24%	79	55%	65	45%
Weight Lifters											
Top level	34	15	44%	15	44%	4	12%	45	66%	23	34%
National level	20	5	25%	14	70%	1	5%	24	60%	16	40%
Total	54	20	37%	29	54%	5	9%	69	64%	39	36%
Short distance s	wimme	rs									
Top level	10	4	40%	5	50%	1	10%	13	65%	7	35%
National level	34	11	32%	18	53%	5	15%	40	59%	28	41%
Total	44	15	34%	23	52%	6	14%	53	60%	35	40%
Long distance swimmers											
Top level	20	7	35%	12	60%	1	5%	26	65%	14	35%
National level	30	11	37%	11	37%	8	27%	33	55%	27	45%
Total	50	18	36%	23	46%	9	18%	59	59%	41	41%

**Table 2.** The prevalence of the A/G IGF-I rs7136446 polymorphism among Israeli long and short distance runners and swimmers and among weight lifters.

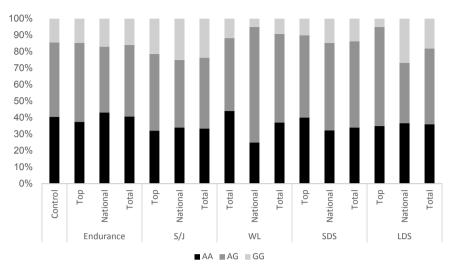


Figure 1. The prevalence of the A/G IGF-I rs7136446 polymorphism among national and top level Israeli long and short distance runners and swimmers and among weight lifters.

*Note*:  $\chi^2(1) = 4.41$ , p = 0.036, for GG genotype frequency, sprinters vs. weight lifters.

The higher prevalence of IGF-I rs7136446 GG genotype was found among sprinters and jumpers (21%, 25%, and 24% for top-, national-level and total, respectively); while the lower GG genotype frequencies were found among weight lifters (12%, 5%, and 9% for top-, national-level and total, respectively, p<0.05).

IGF-I rs7136446 GG genotype frequencies among endurance runners, swimmers and controls vary between 10 to 18%, and were similar to the prevalence found among healthy Finnish males [17]. There were no significant differences in GG genotype carriers between short and long-distance runners and controls, between short and long-distance swimmers and controls, and between weight-lifters and controls. Also, there was no difference in IGF-I rs7136446 GG genotype frequencies between top and national level for all groups.

#### DISCUSSION

We examined the prevalence of the A/G IGF-I rs7136446 polymorphism among Israeli long and short distance runners and swimmers and among weight lifters. In contrast to our hypothesis, there were no significant differences in GG carriers between athletes (runners – short and long-distance, swimmers – short and long-distance, and weight-lifters) and controls. However, interestingly, the prevalence of GG carriers was significantly greater among sprinters compared to weight-lifters suggesting that carrying this polymorphism is beneficial mainly for speed rather than strength sport events.

The effect of IGF-I polymorphism on muscle mass and fitness characteristics has been studied mainly among the elderly because of the age-related decrease in muscle mass and strength [22, 30], and the consequent reduction in functional abilities, increased risk of falling and fractures [21, 29]. Since the IGF-I gene regulates muscle hypertrophy, several studies have focused on the relationship between IGF-I polymorphism, IGF-I levels, muscle mass, and physical ability. The majority of these studies have demonstrated a significant relationship between the IGF 1245T SNP and both baseline fat free mass (FFM) and FFM response to training in the elderly population [8, 18, 19, 31].

Surprisingly, very few studies have examined variation in IGF-I polymorphism among professional athletes [20]. The frequency of the IGF-I promoter polymorphism was significantly greater among athletes (9.2%) compared to non-active controls (2.4%). Moreover, among the athletes, a substantially higher sequence frequency was found among strength (11%) compared to team-sport athletes (7.8%). We previously demonstrated [4] in a cohort of Israeli athletes a higher frequency of the IGF C1245T T/T polymorphism among athletes (4.8%), while none of the control participants carried that polymorphism. Interestingly, half of the T/T polymorphism carriers were endurance athletes and half were power athletes. However, despite the equal distribution among endurance and power athletes, one of the most striking finding of the study was that while the endurance athletes with the T/T polymorphism were of national level, the power athletes were elite-level athletes (international and Olympic). The results suggested that the IGF-I T/T polymorphism possible association with muscle mass increase is more important to power sport performance and excellence. The main contribution of the present study assessing the frequency of another polymorphism of the IGF-I gene (i.e. IGF-I rs7136446) is that carrying the GG genotype was significantly more frequent among sprinters compared to weight lifters. All together this may suggest that among power sports, IGF-I polymorphism is more important to speed rather than strength athletes. Previous studies have demonstrated other polymorphism frequency differences between speed oriented and strength oriented athletes [3].

It is very frequent in sports genetic research to combine athletes from different sports-types but with similar metabolic demands (e.g. endurance

athletes: long-distance running, swimming, cycling and triathlon; power athletes: sprinting, jumping and weight lifting). Interestingly, we did not find a higher frequency of the IGF-I rs7136446 GG polymorphism among swimmers. This is consistent with our previous reports demonstrating that in contrast to elite endurance and power track and field athletes, single nucleotide polymorphisms of IGF-I and the IGF-I receptor and myostatin were not frequent among elite Israeli short- and long-distance swimmers [2, 4-6]. These results may indicate rather uniquely that the myostatin-IGF-I-IGF-I receptor growth factor system is less important for competitive swimming and elite swimming performance than for runners. The mechanism that may explain the different role of myostatin-IGF-I-IGF-I receptor growth factor system polymorphism in swimming compared to running performance is currently unknown. One possible explanation is that excellence in swimming is largely affected by the swimming technique and by the swimmer's size (particularly limb length) [23] which may, at least partially, mask metabolic and muscle mass differences between swimmers, enabling technically-skilled and/or tall swimmers to excel at all swimming distances, and particularly in the sprints. All together these results indicate that different sports disciplines (e.g. swimming and track and field) may have different genetic polymorphisms, despite seemingly similar metabolic characteristics. Thus, it should be acknowledged that combining different disciplines for sports genetic research purposes should not be done, or at least be done with extreme caution.

In summary, although it is well recognized that a single polymorphism cannot determine an athlete's ability to succeed or fail in sports, the findings suggest a potential importance of IGF-I polymorphisms mainly to power sports and in particular to speed sport performance in athletes. Whether a multi-potent athlete who wants to develop a competitive career and carries a beneficial IGF-I polymorphism should prefer track and field over swimming is currently hypothetical and speculative. Moreover, one should acknowledge that while a favourable genetic predisposition is important, psychological and environmental aspects, including training equipment and facilities, nutrition, familial support, and motivational issues, are also crucial for elite-level sports accomplishment.

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