Association of Follicle Stimulating Hormone Receptor Single Nucleotide Polymorphisms with Fertility in Greek men.

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OBJECTIVES

epidemiological Although several studies have been conducted, the of Follicle Stimulating impact (FSHR) Receptor Hormone polymorphisms on male infertility remains unclear. The aim of this study was to investigate prevalence of specific FSHR Single Nucleotide Polymorphisms (SNPs) in the Greek population and associate the latter with the clinical phenotype.

METHODS

Our cross-sectional, single-center study prospectively enrolled 96 subjects. Men with idiopathic non-obstructive azoospermia (n=78) were compared with a control group with fertile men (n=18) for SNPs in FSHR in the promoter at position -29 (G-29A, rs1394205), in the coding region at position 566 (566 C/T, Ala189Val, rs121909658), position 919 (919A/G, Thr307Ala, rs6165), position 2039 (2039A/G, Asn680Ser, rs6166).

The SNP in position 566 was assessed by polymerase chain reaction restriction fragment length polymorphism (PCR-RFLP) and the other three SNPs' (-29, 919, 2039) with Single Strand Conformation Polymorphism (SSCP); All of them were validated with DNA sequence.

Graphs and tables

Association of hormonal and semen parameters with 680 and 566 SNPs.

Association of normo	onal and semen	parameters with	680 and 566 SNPs.
	2039 SNP 2	2039 SNP 3	566 SNP 2
Testis (R)	0.039	0.008	0.585
Testis (L)	0.192	0.588	0.581
FSH	0.660	0.380	0.327
LH	0.479	0.363	0.456
Prolactin	0.869	0.379	0.374
T	0.455	0.517	0.674
Inhibin B	0.322	0.054	1.000
AMH	0.908	0.419	0.569
Semen volume	0.450	0.343	0.794
Sperm count	0.822	0.149	0.839
Sperm motility	0.900	0.707	0.119
Sperm morphology	0.960	0.316	0.227
2039 SNP 1 vs. 2039 FNA score	9 SNP 2, 3, in P 0.550	value: 566 SNP 0.481	1 vs. 566 SNP 2, in P v 0.106

RESULTS

No SNPs were detected in positions -29 and 919.

The heterozygous SNP (AG) at position 2039 (680) was associated with different size of the right testis (p = 0.080).

There was no association between the 566 SNPs polymorphism and hormonal or semen parameters.

The combination SNP 2039 AA with 566 CT revealed significant association with FSH and LH concentrations.

2039 and 566 SNP distribution.

	Fertile	Infertile
AA ¹	5 (28%)	21 (27%)
AG ¹	4 (22%)	39 (50%)
GG ¹	9 (50%)	18 (23%)
566C ²	4 (22%)	19 (24%)

¹ Chi-square, AA vs. A56677d²GG, p = 0.1941(;7²89%) squa569, 6766% ys. 566T, p = 0.848).

CONCLUSIONS

The prevalence of 566CT polymorphism of the FSHR gene was high in the Greek population, yet no association was found with male infertility. The present study, showed no association between FSHR SNPs and male infertility, in the Greek population.

In conclusion, our study indicates that despite the ethnic differences in FSHR, SNPs do not appear to play any specific role in male infertility. Further studies with larger populations may be required to confirm our findings in a more definitive fashion.

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