



KEY MESSAGES

- Hormone levels were reassuring
- > Oligo-/azoospermia:
 - 30% of AGA cases with complex hypospadias
 - 60% of SGA cases
- pathogenic monogenic variants or > No oligogenic variant combinations found

NTRODUCTION

Background: Hypospadias affects approximately 1/200 newborn males and is sometimes included in the DSD spectrum. Few data currently support this hypothesis.

Aims: To assess endocrine and reproductive outcome of young men born with hypospadias and explore genetic mechanisms underlying the development of hypospadias and eventual testicular dysfunction.

METHODOLOGY

Study design: Cross-sectional study (Ghent and Vienna University Hospitals)

Participants: Young adult men (16-21 years) born with non-syndromic hypospadias (n=193) and healthy male controls (n=50)

Assessments: Physical exam, hormonal status (gonadotropins, androgens, insulin-like factor 3 (INSL3) & inhibin B), semen analysis and molecular genetic analysis (NGS-based panel of 474 genes & Oligogenic Resource for Variant AnaLysis (ORVAL))

ΓA	\mathbf{K}	PA	S

	Appropriate for gestational age (AGA)	Small for gestational age (SGA)	
Distal hypospadias	122/167 (73.1%)	10/26 (38.5%)	
Midshaft hypospadias	33/167 (19.8%)	5/26 (19.2%)	
Proximal hypospadias	12/167 (7.2%)	11/26 (42.3%)	
Complex hypospadias	16/167 (9.6%)	7/26 (26.9%)	
Controls	n=50		

ADOLESCENTS & YOUNG ADULT MEN WITH HYPOSPADIAS:

TESTICULAR FUNCTION AND IN-DEPTH GENETIC SCREENING

L Tack, AF Spinoit, P Hoebeke, S Riedl, A Springer, U Tonnhofer, M Hiess, J Weninger, A Mahmoud, K Tilleman, E Van Laecke, A Juul, J Albrethsen, E De Baere, H Verdin and M Cools

Physical exam & hormone assays

Physical exam:

- Smaller stretched penile length in SGA cases
- Regardless of number of penile surgeries and height
- All had a normal timing and pace of puberty

Gonadotropin levels:

- Gonadotropin & INSL3 levels were higher in cases compared to controls • Few above laboratory thresholds
- Few cases had and rogen levels below the laboratory thresholds
 - All with isolated hypospadias
 - And in absence of elevated gonadotropins

Genetics

- No (likely) pathogenic monogenic variants (according to ACMG criteria) - No digenic variant combinations (predicted ≥95%-likely disease-causing) were withheld

AGA cases									
	Phenotype	Gene	Segregation analysis	HGVS transcript	HGVS Protein	rsID	REVEL score	GnomAD VAF	GnomAD HMZ
se 1	M, Cb	BNC2	De novo	c.1022A>G	p.(Asn341Ser)	-	0,156	-	-
se 2	Μ	CHD7	Paternal/de novo ⁺	c.3730A>G	p.(Thr1244Ala)	-	0,722	-	-
		NR5A1	Paternal/de novo ⁺	c.629C>T	p.(Pro210Leu)	rs900214501	0,18	0.000006569	0
se 3	D	ESR2	Maternal	c.1123G>C	p.(Glu375Gln)	-	0,866	-	-
se 4	D	ZFPM2	NA	c.89A>G	p.(Glu30Gly)	rs121908601	0,329	0,002970	2
se 5	D, Mic, B	DGKK	NA	c.255_290dl	p.(Ser91_Ala102del)	-	-	0,0009738	26*
se 6	Μ	GATA4	NA	c.1037C>T	p.(Ala346Val)	rs115372595	0,592	0,001432	0
se 7	Р	LHCGR	NA	c.1046C>T	p.(Ala349Val)	rs758729322	0,596	0.000006572	0
se 8	Р	ERBB4	NA	c.2444T>C	p.(lle815Thr)	rs1264168721	0,756	0.00001314	0
SGA cases									
	Phenotype	Gene	Segregation analysis	HGVS transcript	HGVS Protein	rsID	REVEL score	GnomAD VAF	GnomAD HMZ
se 9	Μ	BNC2	Maternal	c.2618C>T	p.(Pro873Leu)	-	0,494	-	-
se 10	Р	ESR2	NA	c.64A>G	p.(lle22Val)	rs76299711	0,269	0.0006192	0
se 11	Р, В	LHCGR	NA	c.1435C>T	p.(Arg479Ter)	rs757225917	-	0.000003981	0
se 12	P, Cb	NR5A1	NA	c.374C>T	p.(Pro125Leu)	rs780952265	0,24	0.00002207	0
se 13	D	ZNRF3	NA	c.925T>A	p.(Ser309Thr)	rs769697204	0,09	0.00001227	1
se 14	Р	LHCGR	NA	c.1847C>T	p.(Ser616Phe)	-	0,866	-	-
se 15	D	EP300	NA	c.5869C>T	p.(Pro1957Ser)	rs1301322622	0,217	0.00001415	0
		ESR2	NA	c.661A>G	p.(Arg221Gly)	rs78851986	0.794	0.002047	0

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number of homozygotes or hemizygotes (*); +: No DNA was available of the deceased father (who had hypospadias at birth). GnomAD v3.1.1.

CONCLUSION

 \geq All participants had a normal timing and pace of puberty > Mainly subclinical hormone abnormalities > Semen quality is a major concern in hypospadias, especially in SGA and complex hypospadias cases > No recurrent genetic causes were identified, no arguments for oligogenic effects > Non-coding variants? Epigenetics? Placental insufficiency? Multifactorial?

RESULTS







SGA

Hormone levels as predictors of oligo- / azoospermia:

AGA complex

AGA isolated

FSH: sensitivity 9.4%, specificity 100% Inhibin B: sensitivity 21.9%, specificity 97.9% ROC curve analysis: criterion FSH 4.11U/L ; Inhibin B 196.4ng/L Higher INSL3 levels (p=0.042)



normone levels within laboratory reference values

