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Unilateral Renal Agenesis as an early marker for genetic screening in Kallmann Syndrome.

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Abstract

Isolated GnRH Deficiency (IGD) that is either displayed as Kallmann Syndrome (KS) or normosmic idiopathic hypogonadotropic hypogonadism (nIHH) is a rare Mendelian disorder with wide clinical and genetic variability. Apart from the arrest of pubertal development, IGD is also characterized by a variety of non- reproductive features including unilateral renal agenesis (URA), midline defects, dental & ocular defects and many more. In this analysis we explored the role of unilateral renal agenesis, as a screening tool for detection of genetic changes associated with IGD. We performed detailed genetic screening with Sanger sequencing in 14 genes associated with Isolated GnRH Deficiency as well as screening of intragenic deletions in the gene of anosmin 1-ANOS1 with MLPA. No genetic variation was detected, suggestive of an alternative genetic cause for URA in these normosmic patients, highlighting the necessity of further genetic screening.

Keywords

GnRH; Unilateral renal agenesis; genetic screening

Declaration of interest: no completing interests

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Dear Editor,

Isolated Gonadotropin-releasing hormone (GnRH) Deficiency (IGD), is a rare Mendelian disorder, characterized by genetic and phenotypic heterogeneity. Apart from the failure of sexual maturation and pubertal development (i.e. the normosmic form of idiopathic hypogonadotropic hypogonadism (nIHH) or the anosmic form of Kalmann Syndrome (KS)), the disorder is also characterized by various **non-reproductive features** including unilateral renal agenesis (URA). Interestingly, family members of the affected patients usually display milder forms of IGD or its non-reproductive features, such as isolated anosmia and URA highlighting the variable expressivity of the disorder^{1,2}

We have previously reported the prepubertal diagnosis of KS in a male patient with URA, who carried a mutation in the gene of anosmin 1- *ANOS1*, previously known as kallmann-1 (*KALI*) ³, whereas other non-reproductive IGD features have been utilized for an early KS diagnosis ⁴. With this letter we would like to share our experience in examining the role of URA as an early prepubertal marker for IGD.

Nine prepubertal normosmic male patients were clinically evaluated and screened for mutations in 14 IGD genes. Genomic DNA was obtained from peripheral blood samples by standard phenol-chloroform extraction. Exonic and proximal intronic (15 bp from splice sites) DNA sequences of the following genes was amplified by PCR and determined by direct sequencing: *ANOS1* (OMIM 308700), *GNRH1*(OMIM 152760), *GNRHR* (OMIM 138850), *KISSIR*(OMIM 604161), *KISS1*(OMIM 603286), *CHD7*(OMIM 608892), *NMSF*(OMIM 608137), *FGF8*(OMIM 600483), *FGFR1*(OMIM 136350), *PROK2*(OMIM 607002), *PROKR2* (OMIM 607212), *HS6ST1* (OMIM 604846), *TAC3*(OMIM 162332), and *TACR3*(OMIM 162332). To detect exonic deletions/duplications across the entire coding region of *ANOS1*, gene dosage analysis was performed using the SALSA MLPA kit P132 Kallmann-1 (MRC Holland). Informed consents were obtained.

A summary of the clinical characteristics is shown in Table 1. Importantly, none of the children evaluated was anosmic, whereas signs of GnRH development disruption, such as cryptorchidism, as well as other non-reproductive features were seen in family members. A rare sequence variant (RSV) was defined as a) a variant affecting splice junctions within lObp of coding sequence; or a protein-altering/protein-truncating non-synonymous variant; and b) present in <1% minor allele frequency (MAF) in Exome Variant Server, NHLBI GO Exome Sequencing Project (ESP), Seattle, WA (URL: http://evs.gs.washington.edu/EVS./), [February 2016], 1000Genome project [http://www.1000genomes.org/home] ⁵ and the non – Finish European population of the Exome Aggregation Consortium (ExAC), Cambridge, MA (URL: http://exac.broadinstitute.org) [February 2016].

Neither RSVs in any of the 14 genes nor intergenic dosage alterations in *KAL1* were detected. The lack of a genetic association in this prepubertal normosmic cohort could be attributed to (i) the clinical presentation of our patients; even though nIHH cases with URA have been previously reported⁶, the majority of studies have linked URA to KS and *ANOS1* genomic alterations, (ii) a missed genetic alternation in the rest of the 37 known IGD genes not screened here, or (iii) an independent genetic cause for URA; since nephrogenesis is

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controlled by genes enhancing or inhibiting precursor cell growth⁷, a different genetic cause for isolated URA cannot be excluded in these patients. In conclusion, the genetic screening for the 14 IGD genes in a small cohort of normosmic children with isolated URA has not revealed any genomic alterations. However, further genetic screening may reveal a genetic association, important for genetic counseling and clinical follow-up. Such genetic screening is especially recommended in patients with a strong family history and/or additional nonreproductive features of IGD.

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Table 1:

Clinical Characteristics of 9 prepubertal male patients with URA.

Fx	Anosmia in both grandfathers	Cleft lip and palate in maternal aunt, Psychiatric disorder in two grandparents	Depression paternal great uncle	T1D father, Nephrolithiasis paternal grandfather, Hypothyroidism paternal grandmother, Hyperthyroidism paternal great uncle, Epilepsy maternal uncle	Anosmia paternal great grandmother, Kidney malformation in maternal great uncle, Verterbal collum lipoma maternal aunt	T1D maternal aunt, Gastroschisis paternal grate grandfather	Hyposmia paternal great aunt, Skoliosis father, Skoliosis, osteoporosis and Parkinson's disease in paternal grandfather, Parkinson's diseaseptaternal great uncle, Psoriasis paternal grandmother	Strabismus mother, Hypothyroidism- Thyroidectomy-T2D maternal grandmother	Strabismus mother, Hypothyroidism- Thyroidectomy-T2D maternal grandmother
Fx of renal agenesis	n/a	n/a	ри	п	n/a	n/a	ъ́л	brother with URA	brother with URA
Ass. Abn	ou	ou	оц	verbal disorder, speech therapy	ou	ou	hydrocele L, balanoposthetical adhensions	ou	stravismus
UT/ nephro defects	no	ureteropel vic junstion stenosis R, hydronephrosis R, chronic renal failure	оп	оп	0	ои	01	ou	hydroureter R, nephrocalycosis R
Birth order	second	third	first twin birth	second	third	third	third	second	first
prematurity	ou	ОП	ou	ou	ou	ou	ou	ou	ло
anosmia	ou	no	ou	ou	ou	ou	ou	ou	no
cryptorchidism	ou	ou	ou	ou	оп	cryptorchidiam R- orcheopexy at 18 m	ou	ou	ou
Age at Dx	2.5 m	7 month of embryonic development	2nd trimeter	2nd trimester	2nd trimester	2nd trimester	at birth	2nd trimester	9 m
Age	6 yo	15 yo	6 yo	8 yo	5 yo	6 yo	11 yo	6 yo	10 yo

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presence of urinary tract or renal defects, associated abnormalities, family history of unilateral renal agenesis or other diseases and disorders, including reproductive or non-reproductive features of IGD. Dx: Table 1 shows the clinical characteristics of the 9 male prepubertal patients analyzed including their age, age at diagnosis, the presence of cryptorchidism, anosmia or prematurity, the birth order, the Diagnosis, UT: Urinary tract, Ass. Abn: Associated Abnormalities, Fx: Family history, N/a: not available, nl: normal, T1D: type 1 diabetes, T2D: type 2 diabetes.