

Idiopathic Hypogonadotropic Hypogonadism— An Update on the Aetiopathogenesis, Management of IHH in Both Males and Females—An Exhaustive Review

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How to cite this paper: Kulvinder, K.K., Allahbadia, G.N. and Singh, M. (2016) Idiopathic Hypogonadotropic Hypogonadism—An Update on the Aetiopathogenesis, Management of IHH in Both Males and Females—An Exhaustive Review. *Advances in Sexual Medicine*, 6, 50-78.

<http://dx.doi.org/10.4236/asm.2016.64007>

Received: July 13, 2016

Accepted: August 8, 2016

Published: August 11, 2016

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Abstract

Methods: Asystematic literature search was performed using PUBMED for all English articles up to April 2014. Although this review mainly focuses on published human studies, it also draws attention to where future research should be directed based on animal studies. **Results:** Besides the 9 known mutations widely quoted for KS namely KAL1, Fibroblast growth factor 8 (FGF8), fibroblast growth factor receptor 1 (FGFR1), prokineticin 2 (PROK2), PROK receptor 2 (PROKR2), WDR11, heparin sulfate-6-O-Transferase (HS6T1), chromodomain helicase DNA binding protein 7 (CHD7) and semaphorin 3A (SEMA 3A), additional mutations in “FGF8 synexpression” group e.g., FGF 17, ILRD, DUSP 6, SPRY4 and FLRT3 have been shown to be involved in CHH, mostly KS besides SEMA 7A. Although traditionally division has been based on anosmic/normosmic criteria, further genes found to cause so called nIHH like Gonadotropin releasing hormone receptor (GNRH). KISS1, TAC3, TACR3 have also been found to be associated with hyposmia on detailed testing on UPSIT and MRI for olfactory structures revealed absent OB. Further detailed examination of transcription factor genes have revealed involvement of HESX1, TSHZ1, AXL, SOX10 with a strong overlap of in transcription factors in development of septooptic dysplasia (SOD), combined pituitary hormone deficiency (CHPD) and KS. Treatment with rFSH/-hCG gives almost similar results to pulsatile GnRH therapy and should be based on cost factor, availability and in occasional cases specific treatment like kisspeptin therapy. **Conclusions:** Contrary to the traditional thinking, one should reconsider classifying cases of IHH simply on basis of anosmia/normosmia. Deafness calls for looking for mutations in Sox 10/CHD7/ILRD7 considering 38% associ-

ation of former. Therapy should be individualized based on availability of pulsatile GnRH, cost factor and in recalcitrant cases kp therapy may be of use with kp mutations and NKB mutations.

Keywords

Idiopathic Hypogonadotropic Hypogonadism, Kallmannsyndrome, Anosmia, Hyposmia, Normosmia, Gene Mutations

1. Introduction

Idiopathic hypogonadotropic hypogonadism (IHH) is characterized by the absence of spontaneous pubertal development in the face of low sex steroids and gonadotropin levels with otherwise normal pituitary function. When associated with anosmia this HH is termed **Kallmanns Syndrome (KS)** whereas isolated HH with a normal sense of smell is termed **normosmic IHH (nIHH)**. The incidence varies from 1:10,000 - 140,000 with male:female ratio 4:1 with males presenting every 1:8000 [1]. In most patients with IHH, physiological GnRH response restores normal levels of pituitary and Gonadal hormones, allowing for testicular growth and spermatogenesis in men and ovulation in women [2].

Studies on critical roles of mutated genes causing human GnRH deficiency in the fate specification, proliferation, developmental migration, secretory function and/or survival of GnRH neurons have formed the bases of much of our current understanding of GnRH biology [3].

Human studies in KS fetuses with olfactory bulb agenesis revealed that premature interruption of the olfactory, vomeronasal (VN), and terminal nerve fibers in the fronto nasal region disrupts the migration of GnRH-1 cells which normally migrate from the nose to the brain along these nerve fibers [4] [5].

2. Aetiopathogenesis of Ks

Developmental abnormalities in this migratory journey like those demonstrated with the deletion of the KAL1 gene results in KS [6] [7]. Besides that the fibro blast growth factor (FGF) signaling pathway genes (FGF8 and FGFR1) [8]-[10], Prokineticin 2 (PROK2) signaling pathway genes [11]-[16] and chromodomain helicase DNA-binding protein7 genes (CHD7) [17]-[19], and WDR11 [20], SEMA3A [21] [22] genetic pathways have been identified to have identical neurodevelopmental function and implicated in aetiopathogenesis of KS.

WDR11 is an intracellular protein that interacts with the transcription factor EMX1 and semaphorin 3A a secreted protein involved in axonal path finding acting through neuropilin (NRP)1 receptors [21]. Subsequently Cariboni *et al.* in mice studies suggested SEMA 3A signals redundantly through both the classical NRP1 and unconventional NRP2 while the usual NRP2 ligand SEMA 3F is dispensable for this process and

suggesting mice lacking it recapitulate the anatomical features of the single KS case analyzed [23]. Further Kansakoski *et al.* 2014 have further included SEMA 7A besides SEMA3A while screening 50 Finnish children (34KS, 16nIHH). They found 3 heterozygous gene variants c458A > G (p.Asn153Ser), c.1253A > G (p.Asn418Ser) and c.1303G > A (p.Val435Ile) in SEMA 3A in 3 KS patients who also had a mutation in FGFR1. Two rare heterogenous variants c.442 C > T (p.Arg148 Trp) and c.1421G > A (p.Arg474Gln) in SEMA 7A were found in one maleKS patient with a previously identified KISS 1R nonsense variant and one male patient with previously identified KAL1 mutation respectively. They thus concluded that SEMA 3A/SEMA 7A heterozygous missense variants may modify the phenotype of IHH but most probably are not sufficient to cause the disorder alone [24].

The identification of zinc finger homeodomain factor tea shirt finger family member 1 (TSHZ1), a key regulator of mammalian olfactory bulb (OB) development not only in mice but also in humans as revealed by Ragancokoya *et al.* has given some answers to a lot of unanswered questions regarding role of PROK2 ligands acting through their Gprotein coupled receptors PROKR2 regarding how their mutations cause both KS and IHH [25]-[28]. They investigated families with congenital aural atresia, heterozygous for loss of function mutations in TSHZ1, and found by microarray analysis, ISH, as well as CHIP that TSHZ1 bound to and regulated the expression of the gene encoding PROKR2, a GPCR essential for OB development and thus giving a clue to the link between PROK2 and KS despite GnRH neurons lacking PROK2 receptors [29]. The OB is one of the few areas in the mammalian brain that produce neurons throughout life. New interneurons originating from progenitors in the subventricular zone (SVZ) are continually added to the OB. While mRNA of both PROKR1/PROKR2 are expressed in the SVZ and OB, only PROK2 mRNA found here, which functions as a chemo attractant for these neuronal progenitors which follow a rostral migratory stream [28].

Some of the Prokineticin receptor 2 mutations are not routed to the cell surface, instead they are trapped in the cell secretory pathway. The cell permeant agonist/antagonists have been used to rescue some membrane receptors that are not targeted onto the cell membrane. Chen *et al.* chose three disease associated mutations W178S, G34D, P290S, which all resulted in retention of PROKR intracellularly. They showed that a small molecule, PROKR2antagonist A457 dramatically increased cell surface expression and rescued the function of P290S PROKR's but had no effect on W178S and G234DPROKR2, Interestingly one of PROKR2 mutations P290S is identified in more than 5% of Maghrebian KS patients who show PROKR mutations in 23.3% of their KS population as opposed to 5% in KS patients from general populations, which have a monogenicrecessive or digenic/oligogenic transmission [30]. Treatment with "chemical chaperones" like 10% glycerol significantly increased the cell surface expression and signaling of P290S and W178S PROKR2. They concluded that some KS-associated intracellularly retained mutant PROKR2 receptors can be functionally rescued suggesting a potential treatment strategy for the patients of KS bearing such mutations [31].

Further just like Axl/Tyr 3 null mice have delayed first estrus cycle and abnormal

cycles due to defect in GnRH neuronal migration and survival Salian-Mehta *et al.* sequenced coding regions for AXL in 104 unrelated KS and nIHH subjects and found 3 missense AXL mutations (p.L50F, p.S202C and p.Q361P) and one intro variant and 6 bp upstream from start exon 5 in two KS and two nIHH subjects and also tested in mice simultaneously and concluded that functional consequences of AXL sequence variants in patients with IHH support the importance of AXL and the Tyro3 family Axl, Mer (TAM) family in reproductive development [32].

Recently HESX2 homeobox genemutations have been reported in cases of KS and considered that they may cause KS. In a study of 217 cases of IHH/KS along with 192 controls 2 novel heterozygous missense mutations *i.e.* H424 and p.V75L and a previously identified heterozygous missense mutation p.Q6H were identified in 3 of 217 cases (1.4%). All were found in males with KS. This highlights the importance of HESX1 homeobox gene expressen in embryonic stem cells as another important gene in aetiopathogenesis of KS [33] [34]. See **Table 1**.

2.1. Validity of Classification of KS/Nihh on Basis of Smell/Genetic Testing

Historically KS and nIHH have been considered distinct clinical entities with KS subjects representing a neurodevelopmental phenotype with a primary defect in GnRH neuronal migration, whereas nIHH subjects represent a neuroendocrine defect in GnRH secretion Most genes identified in KS subjects have been shown to play a predominant GnRH migratory role (KAL1, NELF1, PROK2/PROKR2/FGF8/FGFR1) [35],

Table 1. KS Genes.

GENES	KAL1	FGF8&FGFR	PROK2 & PROK2R Autosomal recessive or oligogenic/digenic	WDR11	HSsST1	SEMA3A	Chd7
Modes of Transmission	Xlinked recessive	Autosomal Dominant, Incomplete penetrance, or oligogenic/digenic		?	Digenic/Oligogenic	Digenic or oligogenic	?
Strategy for gene identificaton	Cytogenetics	Cytogenetics and Mouse Models	Mouse Models	Cytogenetics	C.elegans MODEL	Mouse Model	Candidate genes
Prevalence of mutations in KS pts	8% of male patients	10% & <1%	7% & 3%	≤1%	<1%	6%	1% - 6%
References	[7] [8]	[9] [10]	[11] [12]	[20]	[34]	[21]	Ogata T <i>et al.</i> 2006 [17], Jongmans <i>et al.</i> 2009 [18], Kim <i>et al.</i> 2008 [19]

whereas genes identified in nIHH subjects have been shown to primarily affect neuroendocrine regulation of GnRH secretion/action (KISS1R, TAC3, TAC3R, GNRH1, GNRHR) [36]-[42]. The overlap seen in mutations of genes such as FGF8/FGFR1/PROK2/PROKR1 in both KS and nIHH subjects prompted Lewkowicz Shpuntoff *et al.* to study the detailed olfactory phenotypic spectrum in a large cohort of patients using MRI to determine olfactory structures and university of Pennsylvania Smell Identification Test (UPSIT), along with genetic testing of IHH n = 286 (201 male/85 female) with 2183 control (1011 male: 1172 females). They found 31.5% were anosmic, 33.6% hyposmic and 34.9% were normosmic. Most hyposmic (7/11) subjects with MRI data exhibited olfactory structure abnormalities. Of hyposmic subjects 39.5% harboured mutations in genes involved in neuronal migration (KAL1, PROK2, FGF) which was not unexpected but surprisingly also in genes associated with GnRH secretion like (GNRHR, KISS1, KISS1R and TAC3). The only explanation they could give was that either these patients harbor a oligogenic “second hit” as suggested by Sykiotis *et al.* that contributes to hyposmia or that there maybe a yet unrecognized neurodevelopmental role in addition to regulating GnRH secretion [43]. Besides that they suggested accurate olfactory phenotyping can inform regarding the pathophysiology and guide genetic testing [44].

Further Valletti *et al.* studied 36 patients (31 males) of IHH and on basis of smell test classified them to be normosmic (n = 21.58.3%) and anosmic/hyposmic (n = 15.41.6%) and found anosmia/hyposmia is significantly related to anatomical anomalies of the olfactory bulb/tracts, but the prevalence of other developmental anomalies, especially midline defects and neurosensorial hearing loss was high both in HH and KS independent of anosmia/hyposmia. Hence they postulated that from clinical standpoint KS and normosmic HH should be considered as the same complex, developmental disease. Since there are a lot of lacunae in this study, namely small numbers, not using UPSIT for smell detection and study being partially prospective, partially retrospective it is difficult to draw meaningful conclusions [45].

2.2. Paradox in ICA (Role of OB in GnRH Development)

However this has been further complicated by Moya-Plana reporting isolated congenital anosmia (ICA) and olfactory bulb agenesis without gonadotropin deficiency with three PROKR2 mutations previously described for KS along with one new PROK2 mutation, and incomplete penetrance on investigation of families, on screening for KAL1/FGF8/FGFR1/PROK2/PROKR2 mutations in 25 cases of ICA which suggests the considerable complexity of GnRH neuron development in humans [46].

While studying 13 most common IHH/KS genes in 48 patients the overall prevalence of digenic gene mutations was found to be 12.5% by Qaynori *et al.* In addition 30% of patients without a known mutation had mutation in a single gene. Similarly Sykiotis *et al.* as well found digenic disease in 11% of 89 patients who had known mutation in one gene and 10 (2.5%) of 397 among all patients which is the largest series published [43]. These findings suggest with current knowledge that most IHH/KS patients have monogenic etiology [47].

3. Role of FGF in GnRH Development

The discovery of mutations in FGF8-FGFR1 in CHH has demonstrated a previously unappreciated role of FGF8-FGFR1 signaling in GnRH ontogeny. Subsequently Chung *et al.* have established FGF8 as critical for both GnRH as well as olfactory system development [48]. Besides that ears, eyes, kidneys and limbs are also influenced by FGF8 [49]-[52], all of which can be affected in CHH [53]. Among the >15 genes implicated in CHH, mutations in FGF8-FGFR1 account for 12% of cases and importantly KAL1 and HS6ST1 (Heparan sulfate-6-O-sulfotransferase 1) [MIM 604846], two genes known to be mutated in CHH, also encode important components of FGF8-FGFR1 signaling. KAL1 encodes anosmin 1, which enhances FGFR1 signaling by direct physical interactions with the FGFR-FGF-heparansulfate proteoglycan (HSPG) complex on the cell surface. HS6ST1, which encodes a heparansulfotransferase enzyme was found mutated in CHH and that heparan6-O-sulfation was required for anosmin function *in vivo* [54]. Based on the fact that multiple genes from FGF family are mutated in CHH, Miraouli *et al.* studied 386 individuals from CHH group with 155 controls to study role of “FGF8 synexpression group” and found except for FGF18 (MIM 6063726) and SPRY2 (Sprouty homolog2) [MIM602466], all other genes were found to be mutated in CHH individuals FGF17 (n = 3) individuals, IL17RD (Interleukin 17 receptor D)[MIM6068807] (n = 8), DUSP6 (Dual Specificity phosphatase 6) [MIM602748] (n = 5), SPRY4 (Sprouty homolog 4) (Drosophila) [MIM607984] (n = 4), FLRT3 (Fibronectin leucine rich transmembrane protein 3) [MIM604808] (n = 3) while FLRT3 is an enhancer as compared to others which are inhibitors. They further concluded mutations in IL17RD were found only in KS individuals and were strongly linked to hearing loss individuals (6/8). Further mutations in genes encoding components of FGF pathway are associated with complex modes of CHH inheritance, and act primarily as contributors to an oligogenic genetic architecture underlying CHH [55]. Animal models indicate that genes involved in FGF8 signaling such as KAL1, FGFR1, FGF8 and CHD7 are crucial for both placodal development and cell specification as well as for NC formation, migration and cell survival [9] [56]-[60]. The identification of the neural crest (NC) origin for the olfactory ensheathing cells (OEC) which indicates that all ensheathing glial cells of the peripheral nervous system of head, cranial ganglia, olfactory and auditory system are of NC origin of the OEC and the identification of pluripotent NC derived progenitors within the placodally derived structures [61] [62], is an important point when considering the evolution of NC and placodal structures and more broadly the cellular and molecular readout of syndromic pathologies such as CHARGE specific clinical cases of KS [9] [63].

Using different knockout in mouse lines and cre-lox mediated line age tracing fgf8 expression and cell survival was analyzed in the developing nose in relation to the expression of Bone morphogenetic protein 4 (Bmp4) and its antagonist Noggin (Nog) by Forni *et al.* 2013. FGF8 is expressed by cells that acquire an epidermal/respiratory cell fate and not by stem cells that acquire neuronal olfactory/vomer nasal (VN) cell fate. Ectodermal and mesenchymal sources of Bmp4 control the expression of BMP/TGF β antagonist Nog whereas mesenchymal sources of Nog defines the neurogenic borders

of the olfactory pit(OP).fgf8 hypomorph mouse models displayed severe craniofacial defects, together with overlapping defects in the OP, including i) lack of neuronal formation ventrally where GnRH neurons normally form and ii) altered expression of Bmp4 and Nog with Nog ectopically expressed in the nasalmesenchyme and no longer defining the GnRH and VN neurogenic border and concluding i) FGF8 is not sufficient to induce ectodermal progenitors of the OP to acquire neural fate and ii) altered neurogenesis and lack of GnRH neuron specification after chronically reduced fgf8 expression reflected dysgenesis of the nasal region and loss of a specific neurogenic permissive milieu that was defined by mesenchymal signals [64].

Recently it was demonstrated that CHD7 is essential for the migration of multipotent migratory NC cells, which migrate from the neural tube to many regions in the embryo, where they differentiate into various tissues including craniofacial and heart structures. In view of this Schulz *et al.* undertook a genome-wide microarray expression analysis on wild type (WT) and CHD7 deficient (Chd7^{Whi+} and Chd7^{Whi/Whi}) mouse embryos at day 9.5 atimepoint of NC cell migration. They identified 98 differentially expressed genes between WT and Chd7^{Whi/Whi} embryos. Interestingly many misregulated genes are involved in NC and axon guidance; such as semaphorins and ephrin receptors. By performing knockout experinments for Chd7 in xaenopus laevis embryos they found abnormalities in the expression patterns of Sema3a,a protein involved in the pathogenesis of KS *in vivo*. In addition they found nonsynonymous SEMA3A variants in 3out of 45 CHD7 negative CHARGE patients..Thus concluding that CHD7 regulates genes involved in neural crest cell guidance demonstrating a new aspect in the pathogenesis of CHARGE syndrome. Sema3a, is conserved regulatory mechanism across different species, highlighting its significance during development [65].

4. Diagnosis and Differential Diagnosis

Since genetic testing is becoming complex and costly, Costa-Barbosa *et al.* suggested prioritizing genetic testing in patients with KS using clinical phenotypes. For example certain clinical feautres commonly associated with genetic causes are synkinesia (KAL1), dental agenesis (FGF8/FGFR1), digital bony abnormalities (FGF8/FGFR1) and hearing loss (CHD7) and these can be useful to prioritize genetic screening, although renal agenesis and cleft lip and palate did not emerge as statistically significant predictors [66]. This is in slight contrast with the report of Dode *et al.* where they report associations of renal agenesis with KAL1 and cleft lip/palate with FGF8/FGFR1 mutations which was not found in this study [8] [67].

However before coming to a diagnosis of CHH one must rule out the differential diagnosis of pituitary tumours, or pituitary infiltration by neuroimaging studies like MRI [68] [69], juvenile hemochromatosis by serum iron and serum ferritin levels [70], and a systemic disorder that by undermining nutritional status could affect gonadotropin secretion and pubertal development like anorexia nervosa, celiac disease [71]. Anterior pituitary function must be thoroughly evaluated to rule out hyperprolactinemia [69], primary hypothyroidism, GH, ACTH and investigate adrenal axis or somatotrope axis

specifically when pubertal delay accompanied with statural retardation and rule out multiple hormone deficiencies. Indeed diagnosis of any associated endocrinopathy of this type will reorient the etiologic diagnosis towards a specific lesional or genetic disorder [72]-[77], which will thus conclude that the HH is isolated. The most likely differential diagnosis before 18-year is constitutional delay of puberty. Since IHH may present as delayed puberty, it becomes essential to know how to distinguish constitutional delay of growth of puberty (CDGP) from isolated HH with definitive diagnosis of IHH awaiting lack of spontaneous puberty by 18 years. Although basal gonadotropins and GnRH Stimulation tests have limited diagnostic specificity, with overlap in gonadotropin levels between adolescents with CDGP and IHH, Stimulation tests using more potent GnRH agonists (especially leuprolide acetate) and/or human chorionic gonadotropin (hCG) may have better discriminatory value, but small study size, lack of replication of diagnostic thresholds, and prolonged protocols limit clinical application. Basal inhibin B may offer a simple discriminatory test, however in a recent metaanalysis Harrington J., 2012 didn't find any reliable diagnostic test and recommended this an important area for future investigation ([78] for review and **Figure 1**).

In paediatric endocrinology this differential diagnosis is far more difficult as CHH is rare whereas CDGP is infrequent [79]. Serum inhibin B levels in CHH males correlate with testicular volume and thus with clinical severity of gonadotropin deficiency [80]-[82], and with very broad and overlapping values this single marker is not dependent. In view of all difficulties classical clinical features distinguishing CHH from CDGP are still of practical value, especially observing testicular volume over time, in patients receiving exogenous testosterone. In male patient with pubertal delay and low gonadotropins, presence of micropenis and/or cryptorchidism practically rules it out since they are rarely seen in CDGP and favours CHH [79] [83]. Signs of a particular etiology are also useful like anosmia etc. **Figure 2** (see [84] for review).

On studying the Brain changes in 45 male patients with KS in a large imaging study Manara *et al.* found significant morphological and structural brain changes, likely driven by olfactory bulb hypoplasia selectively involving the basal forebrain cortex (which include corpus callosum changes, multiple sclerosis like white matter abnormalities, acoustic schwannoma in 1 patient). By specific analysis KS patients presentation with symmetric clusters of grey matter volume increase and decrease and white matter volume decrease close to the olfactory sulci, reduced sulcal depth of the olfactory sulci and deeper medial orbital-frontal sulci, lesser curvature of the olfactory sulcus and sharper curvature close to the medial-orbital-frontal sulci and increased cortical thickness within the olfactory sulcus [85].

KS, combined pituitary hormone deficiency (CPHD), and septo-optic dysplasia (SOD) all result from developmental defects of the anterior midline in the human forebrain. It has been recently shown that deficient migration of GnRH neurons is also a feature in forebrain formation defects [5]. Hence Raivio *et al.* studied 103 patients with either CPHD (n = 35), or SOD (n = 68) and investigated them for mutations in genes implicated in the etiology of KS (FGFR1, FGF8, PROKR2, PROK2 and KAL1). Muta-

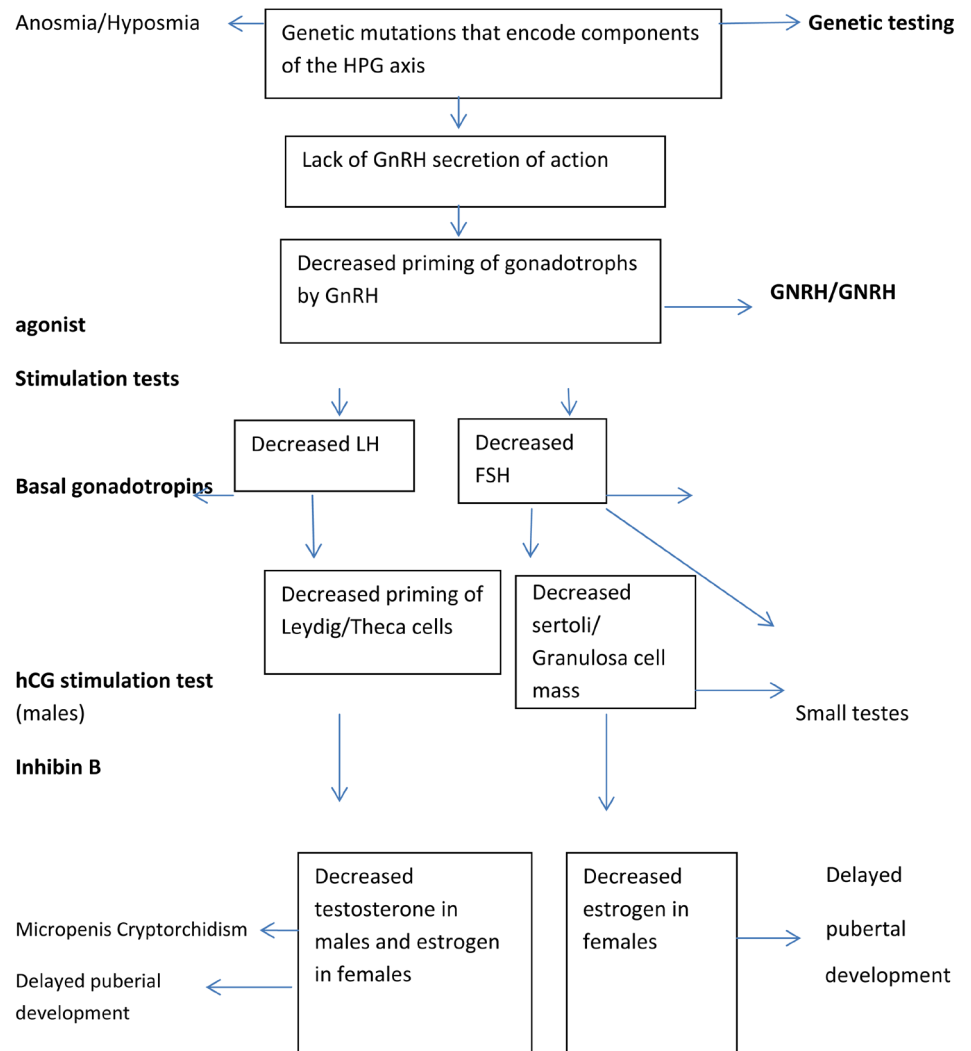


Figure 1. Courtesy ref [78]—In a subset of adolescents with IHH (and Kallmanns syndrome), mutations in genes that encode critical components of the HPG axis lead to either a lack of GnRH secretion, or action. The etiologies in the remaining cases are undetermined. The lack of GnRH action leads to a deficiency of both priming and hormonal secretion of the gonadotropins in the pituitary and of the leydig/theca cells of the gonads. These characteristics of the H-P-G axis form the physiological basis for the diagnostic tests (indicated in bold face) and typical characteristics (indicated in italic anosmia/hyposmia, small testes, micropenis, cryptorchidism) used to identify patients with a higher likelihood of IHH than CDGP.

tions in FGFR1/FGF8/PROKR2 contributed to 7.8% of their patients with CPHD/SOD which suggests a significant genetic overlap between conditions affecting the development of anterior midline in the human forebrain. Of the SOD 3 patients had heterozygous mutations in FGFR1, with these either shown to alter receptor signaling (p.S450F, p.P483S) or predicted to affect splicing (c.336C > T, p.T112T). One patient was found to have synonymous mutations in FGF8 (c.216G > A, p.T72T), that was shown to affect splicing and ligand signaling activity. Four patients with CPHD/SOD were found to harbor heterozygous rare loss of function variants in PROKR2 (p.R85G,

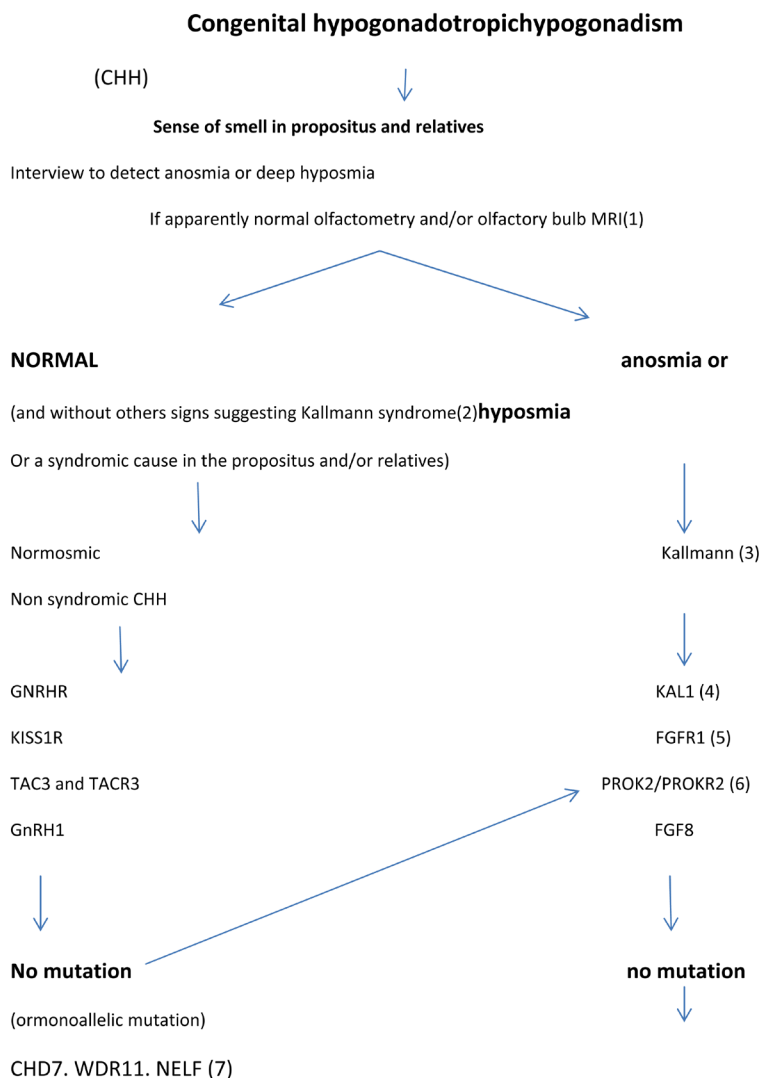


Figure 2. With permission from Dr Young courtesy ref [84]—Molecular studies performed in male patients of CHH categorized on the basis of smell 1) MRI, 2) Bimanual synkinesis, tooth agenesis, hearing impairment, renal agenesis, cleft lip/palate, high arched palate, pes cavus, ptosis, absent nasal cartilage, hand/foot skeletal abnormalities and iris coloboma 3) Step by step strategy based on familial history and putative mode of disease inheritance (pedigree), and the presence of additional clinical anomalies as mentioned above that may direct the geneticist towards a particular Kallmann gene 4 and 5) For instance 4) KAL 1 is analyzed especially in Kallmann men with mirror movements (bimanual synkinesis) and/or for kidney agenesis and/or when the pedigree suggests an X linked mode of inheritance, whereas 5) in subjects displaying cleft lip/palate FGFR1 mutations are searched in firstline whatever the apparent mode of inheritance 6) in subjects with monoallelic, PROK2 or PROKR2 mutations, search for mutations in other CHH genes to demonstrate a digenic or oligogenic mode of inheritance 7) Analysis of other large genes mentioned below performed in second line, given their lower or unknown prevalence among normosmic CHH and Kallmann men. Sizes of the genes currently sequenced in CHH patients: GNRH1; three exons, GNRHR; three exons, KISS1R; five exons, TAC3; six exons; TACR3; five exons; KAL1; fourteen exons, FGF8, six exons; FGFR1, 18 exons; PROKR2, two exons; PROK2, four exons; CHD7; 38 exons; WDR11; 29 exons; NSMF (NMDA receptor synaptonuclear signaling and neuronal migration factor—formerly known as NELF); 16 exons.

p.R85H, p.R268C) [87]. To further study the role of PROKR2/PROK2, McCabe *et al.* further studied 422 patients of congenital hypopituitarism (CH) and detected that variations in PROKR2 but not PROK2 are associated with CH and SOD. They detected 5PROKR2 variants in 11 patients with SOD/CH: novel p.G371R and previously reported p.A51T, p.R85L, p.L173R and p.R268C-the latter three being known as functionally deleterious variants [88]. Midline defects are encountered in all three KS [MIM; 147950], SOD [MIM; 182230] and holoprosencephaly [HPE; MIM 236100], a complex brain malformation that affects both the forebrain and face. In view of mutations identified in number of transcription factor genes in SOD cases such as SOX2, HESX1, SOX3, and OTX2, which are essential for normal forebrain development [89], and similarly in HPE mutations found in genes like SHH, SIX3, TGIF1, TDGF1, FOXH1, and GLI2 [90] [91], Vaaralahti *et al.* 2012 studied 19 subjects (18 males) with KS without known KS genes and screened them for mutations in SOX2, SHH, SIX3, TGIF1, TDGF1, FOXH1, GLI2 and GLI3. One male carried 2 heterozygous missense changes, one in SIX3 (c.428G > A, p.G143D) and the other in GLI2 (c.2509G > A, p.E837 K). Both of these genes have been implicated in etiology of HPE and none was present in 200 control subjects. Thus they concluded that KS and HPE may display a genetic overlap and in view of this the involvement of genes implicated in the etiology of midline defects in patients with KS warrants further studies [92].

Recently a novel syndrome has been defined known as **TUBB3E410 K Syndrome** where one of the eight missense mutations in TUBB3 gene, that encodes the neuronal specific protein β tubulin isotype 3, have congenital fibrosis of the extraocular muscles, facial weakness developmental delay, and possible peripheral neuropathy. This occurs due to c.1228G > A resulting in a TUBB3E410K amino acid substitution which directly alters a kinesin motor protein binding site. In detailed phenotype of eight unrelated individuals Chew *et al.* confirmed electrophysiology that a progressive sensorimotor polyneuropathy does indeed segregate with the mutation and expand the TUBB3E410K phenotype to include KS, stereotyped midface hypoplasia, intellectual disabilities and in some cases vocal cord paralysis, tracheomalacia and cyclic vomiting. Neuroimaging reveals a thin corpus callosum, and anterior commissure, hypoplastic to absent olfactory sulci, olfactory bulbs and oculomotor and facial nerves, which support underlying abnormalities in axon guidance and maintenance [86].

4.1. Management

Castaneyra-Perdomo *et al.* 2014 suggested that although normally treatment is started in KS patients just before puberty early treatment is advisable, as brain sexual maturation occurs long before puberty normally at perinatal age. As brain cells implicated in the development of the olfactory and reproductive system have a rostral and a caudal origin, and the rostral origin is affected by aplasia in KS and the caudal origin does not seem to be affected, the aim of early treatment of KS is to attain brain sexual maturation at the most appropriate age possible, which may prevent the eunuchoidal behavior and appearance observed in KS [93].

4.1.1. Role of Testosterone Therapy

The initial goal of treatment for adolescents and young men who present with CHH is to induce physical and behavioral development matching that of normal healthy subjects of same age. This includes development of secondary sex characters like pubic and axillary hair, increase in penis size, voice masculinization and development of muscle mass. Further one aims at correcting the delay in bone maturation and deficient bone mineralization, enhance libido and modify sex behavior. Mostly effective testosterone replacement therapy can lead to a spectacular improvement in quality of life, which demonstrates a causal relationship between testosterone deficiency and these patients symptoms.

Although it is more physiological to achieve such benefits with pulsatile GnRH administration or with combined gonadotropin therapy (human chorionic gonadotropin and FSH) [94] [95], with both therapies effectively inducing testicular growth and secretion of testosterone and estradiol [94]; we have to consider availability of GnRH infusion pumps, cost of treatment, patients requirements specially if patient presents as a partner of an infertile couple with spermatogenesis in view. Since longterm treatment is required even in west mostly testosterone therapy as (injectable esters) is generally preferred for convenience of infrequent injections and cost, DHT is not preferred as it can't be aromatized to estradiol, and hence can't serve the dual purpose of testosterone esters used for decades now as first line treatment. Tn enanthate is one of the cheaper preparations, used at a dose of 200-250 mg once every 2 or 3 weeks. Although these doses depend on age at diagnoses and local practices, Pediatric endocrinologists who see these patients at a younger age, initially prescribe lower doses, gradually increasing for fear of inducing abrupt virilization and bone maturation which could cause behavioral and relational problems. Endocrinologists see adult CHH patients at a later stage when main signs/symptoms are of severe hypogonadism, and usually require full dose. Although two approaches are not comparable patient should be counseled that he will need longterm androgen therapy. Once full virilization has been induced by exogenous testosterone, males whose testes have significantly increased in size (<5% cases), should be reevaluated off androgen replacement therapy to identify those with reversible forms [96] [97] who no longer require treatment.

In a detailed follow up study of 308 patients 44 underwent spontaneous reversal giving a lifetime incidence of 22%. 15 patients with reversal (30%) had KS of which 1 had undetectable OB on brain MRI. Maximum reversal occurred in neurokinin B (10% vs 3% in nonreversal) although reversal occurred across broad range of phenotypes and genotypes (including FGFR1, PROKR2, GNRHR). Sidhoum VF *et al.* 2014 suggested that despite the importance of NKB pathway for normal pubertal timing, its function is dispensable later in life and occurrence of reversal in patient with no OB demonstrates that this structure is not essential for normal reproductive function. Also patients with IHH require lifelong monitoring for reversal and if reversal occurs for subsequent relapse rates as evidenced in 5 patients who relapsed after reversal [98].

4.1.2. Role of Gonadotropin Therapy

Patients who wish to have an increase in testicular volume or fertility in developing

countries like ours where most centres don't have the facilities of infusion pump the approach of combination therapy with initial rFSH with the idea of stimulating proliferation of immature sertoli cells which are under control of FSH initial doses of 1.5 IU/Kg (180 - 450 u/week) \times 2 months - 2.8 yrs Puberty is then initiated with Hcg 500 - 4000 IU/week, 1 - 3 times/week sc and after onset of Hcg treatment if patient can't afford can shift to highly purified FSH. Raivio *et al.* found this induced prepubertal testes growth with increase in serum inhibin B levels, and 6/7 prepubertal boys displayed sperms despite extremely small initial testis primed with rFSH [95] [97] [99]. Similar approach in countries like china followed [100].

4.1.3. Role of Pulsatile GnRH Therapy

GnRH treatment is successful in inducing virilization and spermatogenesis in men with IHH, however a small subset of IHH men, fail to reach a normal testicular volume and produce sperm on this therapy [94]. Pitteloud in studying 76 IHH men undergoing GnRH therapy for 12 - 24 months to define predictors of outcome of longterm GnRH therapy concluded anosmia was not an independent predictor, however favourable predictors of achieving an adult testicular size and consequently optimizing spermatogenesis are prior history of sexual maturation, with a baseline inhibin (IB) > 60 pg/ml along with absence of cryptorchidism [101].

Further extending Sykiotis *et al.* studied 90 patients and classified patients into four groups according to the response obtained to longterm physiological pulsatile GnRH release. 67/90 subjects displayed normal expected response, with normal serum T (270 - 1100 ng/dl), LH (4.2 - 17 IU/L) and FSH (1.8 - 14 IU/L) and had sperm in their ejaculate and were labeled as **typical responders**. In rest 23 patients (26%) three distinct patterns were seen 1) 10 men remained hypogonadotropic and hypogonadal with low normal LH/FSH, serum T < 200 ng/dl and no sperm despite GnRH doses upto 800 ng/Kg-and thus labeled **Group 1 with triple defect with GnRH deficiency, pituitary resistance and testicular failure**. 2) 8 men achieved normal serum T and produced sperms but did so with high LH (>17 IU/L and FSH > 14 IU/L and thus labeled **Group 2 with dual defect, GnRH deficiency and testicular resistance**. 3) 5 men remained azoospermic after atleast 21 months despite achieving normal serum T, LH and FSH and thus labeled **Group 3 with GnRH deficiency with azoospermia**. Although typical responders showed mutations in all the IHH genes tested, atypical responders displayed mutations exclusively in KAL1 Gene [102]. Although Sinisi *et al.* reported a case with homozygous mutation in PROKR2 gene Val274 Asp which presented as reversible KS along with persistent oligozoospermia [103].

4.1.4. Role of rFSH Priming Followed by GnRH Therapy

Dwyer *et al.* 2013 conducted a randomized open label prospective trial to see if there is any benefit of giving recombinant FSH (rFSH) pretreatment for 4 months followed by pulsatile GnRH therapy vs GnRH therapy alone in a group of CHH patients with prepubertal testis (<4 ml), no cryptorchidism, and no prior gonadotropin therapy and found rFSH increased inhibin B levels into normal range and doubled testicular vo-

lume. Histological analysis showed proliferation of sertoli cells (SC) and spermatogonia, a decreased SC to germ cell ratio from 0.74 to 0.35 and SC cytoskeletal rearrangements. Although with pulsatile GnRH similar hormones and significant testicular growth was exhibited, all men receiving rFSH developed sperm in their ejaculate (7/7 vs 4/6 in GnRH only group) and showed trends towards higher maximal sperm counts and hence concluded that rFSH not only appears to maximize the SC population, but also induces morphologic changes, suggesting broader developmental roles [104]. This maybe in accordance with the study of Pitteloud *et al.* who found after studying 25 patients of IHH that. GnRH deficient men undergoing GnRH induced sexual maturation displayed an inverse responsiveness to GnRH and baseline testicular size and I(B) levels. This observation implied that increasing seminiferous tubule maturity represents the major constraint on FSH responsiveness to GnRH in early puberty. In contrast LH responsiveness to GnRH correlated directly with duration of GnRH exposure [105]. However in view of small numbers Dwyer *et al.* proposed that although data appears promising a multicenter trial is recommended to optimize the optimal strategy for treating the very severe CHH cases.

4.1.5. Insulin like Peptide 3 (INSL3)

A testicular hormone secreted during fetal life, neonatal period and after puberty was studied in 281 patients of CHH/KS for its diagnostic and investigative regulation. It was found that INSL3 is as sensitive a marker as T for the evaluation of altered leydig cell function in CHH/KS patients. INSL3 levels correlate with LH levels in CHH/KS men showing together with their rise in INSL3 levels during Hcg therapy and not treatment alone with T that INSL3 secretion seems not constitutively secreted during adulthood but is dependent on pituitary LH [106].

Many studies indicate that the profound FSH/LH deficit is already present during fetal life which explains the micropenis, cryptorchidism and marked testicular hypotrophy already present at birth. Besides that neonatal activation of gonadotropin secretion is compromised in severe CHH/KS, preventing the first phase of postnatal testicular activation. Finally CHH is characterized by the persistence of Gn deficiency in a vast majority of cases at the time of puberty and during adulthood. This prevents the normal pubertal testicular reactivation required for physiological sex steroid and testicular peptide production and for spermatogenesis. Trabados *et al.* thus proposed further that CHH/KS represents a pathological paradox that can help to unravel *in vivo*, the role of each Gn in human testicular exocrine and endocrine functions at different stages of development. Recombinant pure Gn's or pure FSH or LH activity have been used to stimulate leydig cells and sertoli cells respectively and thereby to clarify their paracrine interactions *in vivo*. The effects of these pharmacological probes can be assessed by measuring the changes they provoke in circulating testicular hormone concentrations. Thus Trabados *et al.* reviewed the impact of CG deficiency in endocrine function of interstitial compartment which contains T, E2, INSL3, secreting leydig cells, regulation of inhibin B and AMH secreted by seminiferous tubules. Further insights are provided by studies of human testicular stimulation with recombinant stimulation with recombi-

nant gonadotropins used individually or in combination [107].

4.2. IHH in Women

Shaw *et al.* retrospectively studied 248 female patients of IHH from 1980 to 2010 seen in Massachusetts general hospital. The clinical presentation varied from primary amenorrhea and absence of any sexual characteristics to spontaneous breast development and occasional menses. In this cohort rare sequence variants were present in all known genes associated with GnRH deficiency, including novel identification of GnRH deficient women with KAL 1 variants. They concluded that the pathogenic mechanism through which KAL 1 variant disrupts female reproductive development requires further investigation [108].

IHH in women (as in men) is treated with sex hormones. Initially low dose estradiol (1 mg) is begun in order to develop sexual characteristics and then it is gradually increased in doses [109]-[112]. From the second year of treatment estrogen supplementation with chlormadinone acetate is done [110]-[112]. Desire for pregnancy warrants pulsatile GnRH therapy to stimulate production of serum FSH and LH [2] [113]. Further the importance of GnRH administration should be intermittent and pulsatile to be able to restore activity of the reproductive axis in patients with HA and other disorders of GnRH deficiency was emphasized by Knobil *et al.* [114]. Tonic exposure of GnRH inhibited pituitary gonadotropin secretion paradoxically [115]. Although chances of success is good, where GnRH is not available, LH or FSH can be administered alternatively [116].

Just like in male subjects Abel *et al.* repeated the study in 37 IHH women treated with i/v pulsatile GnRH therapy (75 ng/kg/bolus) (retrospective study (1980-2012), all patients over 16yrs with 46%anosmic and tested for all 14 genes and found that during first cycle 60% (22/37) recreated normal cycles, 30% (12/37) demonstrated altered gonadotropin response indicating pituitary resistance and 10% (3/37) an exaggerated FSH response consistent with ovarian resistance. Mutations in CHD7, FGFR1, KAL1, TAC3, TACR3 were documented in IHH women with normal cycles, whereas mutations were identified in GNRHR, PROKR2 and FGFR1 in those with pituitary resistance. Women with ovarian resistance were mutation negative. Thus they concluded that although physiological replacement with GnRH recreates normal menstrual dynamics in most IHH ladies [117], Hypogonadotropic responses in first week of treatment identify a subset of women with pituitary dysfunction, only some of whom have GNRHR mutations. IHH women with hypergonadotropic responses to GnRH replacement, consistent with an additional ovarian defect did not have mutations in genes known to cause IHH similar to their findings in men with evidence of an additional testicular defect. Hence they hypothesized that identification of women with abnormal responses to physiological GnRH replacement would give greater insight into the pathogenesis of HH. Hyper/hypogonadotropic responses would implicate an ovarian/pituitary defect, respectively. Such findings would suggest that genes involved in gonadotropin or ovarian development and function that are also expressed in the hypothalamus should be given greater consideration in the search for new IHH genes.

Still controversy exists on the sexuality and intimate relations of men with severe CHH accompanied by cryptorchidism and micropenis [118]. Since there is a negative prognostic value of cryptorchidism and low testicular volume for the future fertility of patients with severe CHH, a trial of earlier gonadotropin therapy during the neonatal or normal pubertal period is warranted and just may prove beneficial, both in terms of testicular hypertrophy and in terms of future fertility [119]-[121].

5. Role of Estradiol (in Male CHH)

Trabados *et al.* studying 91 men of IHH, syndrome found male hypogonadism in CHH is associated with profound E2 insufficiency which can be overcome by aromatizable androgen (Tenanthate) or combined gonadotropin (FSH-HCG) therapy, but not dihydrotestosterone (DHT) contrary to Klinefelters syndrome [122]. This E2 deficiency is also associated with abnormal bone development, noteenage growth spurt and osteopenia or osteoporosis [123]-[125], reviewed in [126]. Further Rochira *et al.* reported 4 cases of tall stature without Growth hormone deficiency who had a impaired response of GH to GHRH-ARG as compared to normal subjects and who had significantly lower IGF1 levels as compared to normal subjects and both IGF1 peak and concentrations were not modified by estrogen therapy in men with aromatase deficiency and concluded insulin as the cause of tall stature rather than GH for the marked increase in height due to nonclosure of epiphyses [127]. Besides that for normal physiology, Pitte-loud *et al.* 2008 showed that for Inhibition of LH secretion by T in men aromatization is required for its pituitary effect but not its hypothalamic effect [128].

Further since Kp 10 is a potent stimulator of LH and increases pulse frequency in men and thereby LH and testosterone levels in normal men there may be a potential role of Kp agonists in HH due to KISS 1/KISS1R mutations ([129], reviewed in [130]). Although TAC3/TAC3R mutations also are associated with IHH, administration of NKB was not accompanied by increase in serum LH and testosterone levels, hence role of NKB doesn't appear to be useful in treating these patients presenting with TAC3/TACR3 mutations [131] however since KISS1 appears to be downstream of NKB signaling, pulsatile gonadotropin secretion is restored by kisspeptin administration to patients with HH secondary to NKB and or its receptor [132] [133].

6. Conclusions

The very concept of classifying IHH into anosmic/hyposmic KS and normosmic IHH gets challenged by the study of Lewkowitch Shpunoff although till now we have been tuned to working on the basis of KS/nIHH. This was further substantiated by the study by Valetti *et al.* where olfactory bulb was absent in patients with anosmic/hyposmic IHH, and otherwise there was no difference in the developmental anomalies in the 2 groups. Although this study was a not well conducted in terms of numbers, smell test and neither fully prospective but it does add to the results of Lewkowitch *et al.* in that there is no point in classifying as two separate disorders but both are part of same spectrum. Despite identification of so many genes like 9 of those which were earlier consi-

dered as for the KS group as they affect the migration or GnRH development eg KAL1, FGF8, FGFR1, HS-6TS1, CHD7, WDR11, PROK2/PROKR2/SEMA3A these account for not more than 30% - 32% cases of KS. Addition of “FGF Synexpression group” adds to the list besides the genes considered so far associated with GnRH function/secretion. Still a lot of controversial issues remain regarding role of PROK signaling as highlighted in ref. [27] regarding absence of PROK receptors on GnRH neurons, mode of inheritance-digenic/oligogenic [43] with presence of PROK2 mutations even in normosmic HH and thus extending the role beyond olfactory bulb development and GnRH neuronal migration and absence of any defect in homozygous mutations while most of human presentations being in heterozygous mutations. Although some answers are provided by the link between TSHZ1, which bound to and regulated the expression of gene encoding PROKR2, a GPCR essential for OB development. Further recently that there maybe an ethnic role is highlighted by greater presence of PROKR2 mutations in KS patients from Maghreb as compared to European origin patients (23.3% vs 5.1%) [30]. Further more work needs to be done on other transcription factors as has been shown for SOX10 *i.e.*, loss of function mutations in SOX 10 is associated with KS along with deafness though not in KS without deafness and highlighting importance of neural crest as well as part origin for GnRH neurons [134]. Similar work needs to be done for other transcription factors in view of genetic overlap of other midline fore-brain disorders like SOD and HPE with KS. Further genes involved in development of hypothalamus, pituitary and gonads in both males as well as females who fail to respond to pulsatile GnRH therapy should be targeted for understanding reason for 4 - 5 times greater preponderance in males—whether it is explained just by X linked transmission or we are missing some female cases due to underreporting in mild hypogonadism or primary amenorrhea not brought forward [135]. The role of SDF-GABA signaling in promoting linear axophilic migration of GnRH neurons through interaction of its receptors, chemokine receptor 4 and 7 (CXCR4/CXCR7), needs further investigation like other molecules, hepatocyte growth factor (HGF), deleted in colorectal cancer (DCC), Slit2 and Robo3 as suggested by Wray *et al.* and Cariboni respectively in animal models [136]-[139].

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