

Table S1 Genotype diagnosis by variant using literature review of each gene and variant that have been previously reported in

IHH/KS, self-limited DP and unaffected individuals. Het: heterozygous, Hom: homozygous, KS: Kallmann syndrome, ID: Intellectual disability, SLDP: self-limited DP, SNHL: sensorineural hearing loss

Genes	Associated phenotype	IHH	SLDP	Normal	Domain disease causing	Variants	Protein variants	Zygoty	Variant lies in specific-domain	Variant specific for phenotype	Genotype Dx
<i>CHD7</i> (1-4)	CHARGE features	Het, oligogenic het	Het	Het	Mutations were found along the protein in both specific and non-specific domains	c.3738G>A	p.M124I	Het	Non-specific	No	Inconclusive
<i>DCC</i> (5, 6)	Facial asymmetry, MR, obesity, SNHL	Het, oligogenic het	No	Het	5 mutations have been reported which are located in specific domains including Ig-like C2 domain type 2 and fibronectin type III domain 1, 3, 5	c.1933C>T	p.P645S	Het	Fibronectin type III domain 3	p.P645S (het) has KS	IHH

<i>DMXL2</i> (7, 8)	none-immune DM, demyelinating polyneuropathy, MR	Hom	No	Het	3 cases with homozygous deletion of 5 amino acid in exon 24 (c.5824_5838del, p.1942_1946del), which is in non-specific domain, have IHH - Parents and a sister who carry heterozygous of the above mutation have normal pubertal timing	c.2540C>T	p.T847I	Het	non-specific domain	No	Inconclusive
<i>GNRHR</i> (9-14)	-	Hom, compound het	Het	Het	R262Q (hom) has been reported in CDGP	c.317A>G	p.Q106R	Hom	Extracellular loop domain2	-Patients with compound het of p.Q106R and other mutation in GnRH gene have been found to cause IHH - Parent of proband who carries p.Q106R (het) has normal puberty	IHH
						c.436C>T	p.P146S	Het	Intracellular loop domain2	- 2 sisters carry p.P146S (het) have IHH - Mother of 2 sisters carries same mutation	Inconclusive
<i>IGSF10</i> (15)	-	No	Het, digenic het	Het	2 het mutations in leucine-rich repeats (LRR) domains 1-7 have been reported to cause self-limited DP	c.7124A>G	p.N2375S	Het	Ig domain	No	SLDP

<i>KISS1R</i> (16-19)	-	Het, hom, compound het	Het	Het	More than 30 mutations have been reported to cause DP. The zygoty includes heterozygous, homozygous or compound heterozygous mutation. - Some variants in heterozygous state such as p.A287E, p.P476R do not cause IHH.	c.-249G>A	-	Het	96 base pairs before transcription start	No	Inconclusive
<i>OTUD4</i> (20, 21)	Progressive dementia & ataxia	Digenic hom	No	Het	- Digenic hom in RNF216 and OTUD4 cause IHH - Individuals with het mutation are unaffected	c.458_460delCTG	p.A153del	Het	Within catalytic domain	No	Inconclusive
<i>PROKR2</i> (22-27)	-	Hom, digenic het & het	No	no	Most mutations cause IHH/KS	c.809G>A	p.R270H	Hom	3rd intracellular loop domain	-This variant (het) with <i>SEMA3A</i> p.R617Q (het) has been found with IHH phenotype	IHH
<i>SEMA3A</i> (28-30)	-	Het, oligogenic het	No	No	Heterozygous mutation of <i>SEMA3A</i> located along the gene have been found to cause IHH/KS.	c.1849C>T	p.R617Ter	Het	Ig domain	p.R617Q mutation combined with heterozygous mutation of <i>CCDC141</i> and <i>PROKR2</i> was found in IHH.	IHH
<i>SEMA3E</i> (31, 32)	-	Oligogenic het	No	No	Oligogenic het mutation in SEMA and Ig domain have been found in IHH/KS.	c.398G>T	p.C133F	Het	SEMA domain	Mutation in the same domain, p.M102T, has been found in KS when combine with heterozygous mutations of	Inconclusive

										<i>PLNXA1</i> and <i>CCDC141</i> .	
TAC3 (33-38)	-	Hom, digenic het	Het	No	Patient with IHH has homozygous or digenic heterozygous mutation of this gene. - This gene is also found in family members of probands with IHH. The reported mutations are missense and mutation in splice site of TAC3.	c.209-1G>C		Hom	-	Heterozygous splice site mutation (TAC3 g.18595G>T) has been reported in individuals with delayed puberty; homozygous carriage of this variant has been reported in individuals with IHH with reversal	IHH
						c.*2-1G>T		Het	-		SLDP
TACR3 (33, 35-38)	-	Hom, compound het, het	Het	Het	- Mutation in this gene have been found in both IHH and self-limited DP. Heterozygous mutation of this gene can cause both conditions.	c.1090C>T	p.R364Ter	Het	Intracellular region	No	IHH

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