

# 00029 Neuroendocrine phenotype, genetics and hormonal treatment outcome in idiopathic normosmic hypogonadism and Kallmann syndrome patients: a multicenter Belgian study

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**Aim:** To study the clinical phenotype, the genetics and therapeutic responses in a series of 35 consecutive patients with hypogonadotropic hypogonadism and normosmia (nHH) /hyposmia (KS).

**Methods:** The study of the genes FGFR1 and KAL1 (anosmin), is performed in our center since 2013. Recently, a panel of genes is available for analysis of the following genes: KAL1, FGFR1, PROKR2, PROK2, CHD7, FGF8, KISS1, KISS1R, APR3, TACR3, GNRHR, GNRH1, NELF, WDR11, HS6ST1, SEMA3A.

**Results:** the series includes 35 patients (32 H/3F, 18 ± 9 years) belonging to 31 families. We have identified by olfactometry 26 nHH and 9 KS. Brain MRI was performed in all patients: two patients had a malformation of Chiari I, two patients showed a partially empty sella, one patient had a cyst of the pouch of Rathke and another one had a cleft palate. Preliminary genetic analysis demonstrated a FGFR1 mutation in three patients and in a family. Identified mutations were: c.1663 + 1 G > A, c.1025T > A (p.Leu342\*) and c.937 - 1234C > T (new mutation: exon 8A of the isoform IIIb). An anosmin mutation was also identified in another patient: c.827\_856 + 49delins\_p.Ala276\_asp286delinsGlyAsn. A last patient had a new mutation TAC3 c.238 + 1 G > A. concerning fertility outcomes, an oligospermia was obtained in 6/12 men treated with hCG and FSH. Hormonal treatment allowed the development of secondary sexual characters in all patients. The patient with FGFR1:c.937 - 1234C > T showed a reversibility of hypogonadism, after 4 years of treatment.

**Conclusions:** Patients with nHH FGFR1 mutation may also present with neuro developmental anomalies, which they should be screened for. The association of normosmic IHH and Chiari malformation is intriguing: it was reported just once in the literature (Kulmar & al. Pituitary 2010). We demonstrated hypogonadism reversibility in a patient with one FGFR1 mutation. Finally, we report two novel TAC3 and FGFR1 mutations.