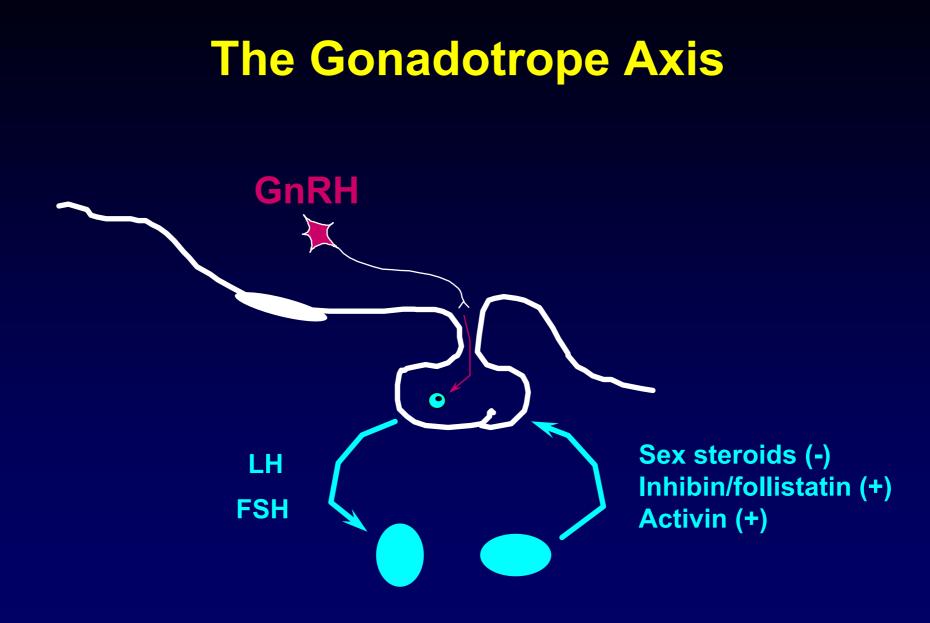
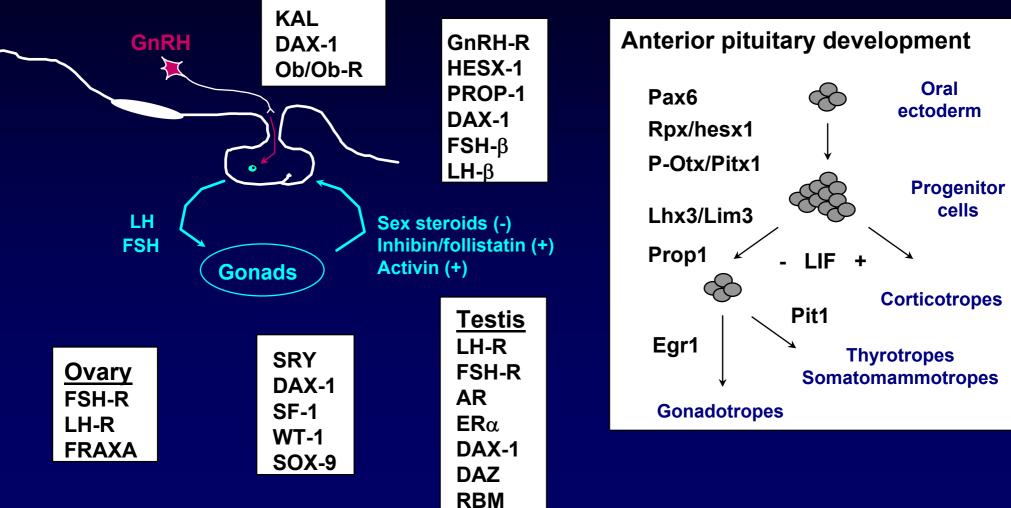
Genetics of Hypogonadism

François Pralong Division of Endocrinology Lausanne University Hospital



Single Gene Mutations and Hypogonadism



Achermann and Jameson, Mol Endo 13, 1999

- 1981 work up of hypogonadism, associated with bilateral cryptorchid testes
 - conserved sense of smell
 - family history negative for infertility
- LH 0.9 U/L FSH 0.4 U/L T<0.7 nmol/L
- Otherwise normal anterior pituitary function
- Normal CT of the hypothalamo-pituitary region

Isolated hypogonadotropic hypogonadism

HCG: 2000 IU 3x/week, replaced by HCG/HMG (1.82):

| Date | 3.8.81 | 7.8.81 | 1.9.81 | 6.11.81 | 7.12.81 | 13.1.82 | 2.4.82 |
|---------------|--------|--------|--------|---------|---------|---------|--------|
| T (nmol/L) | <0.7 | 1.4 | <2.0 | 1.9 | 23.2 | 36.3 | 19.4 |
| TV R (mL) | - | - | 3 | 4 | 4 | 4 | 4 |
| TV L (mL) | - | - | - | 3 | 3 | 3 | 3 |

1994 patient wants fertility

| Date | 10.10 | 23.10 | 9.11 | 20.11 | 1.12 | 8.12 |
|-----------------|-------|-------|------|-------|------|------|
| GnRH (ng/kg) | 60 | 250 | 250 | 250 | 250 | 250 |
| LH | <0.5 | <0.5 | <0.5 | <0.5 | <0.5 | <0.5 |
| FSH | <0.2 | <0.2 | <0.2 | <0.2 | <0.2 | <0.2 |
| Т | 3.7 | 2.7 | 1.9 | 1.8 | 1.4 | 0.9 |
| TV R | 8-9 | 8-9 | | | | |
| TV L | 5-6 | 5-6 | | | | |

Primary failure of pulsatile GnRH therapy

Rx HCG (500 IU 3x/week) and HMG (75 IU 3x/week)

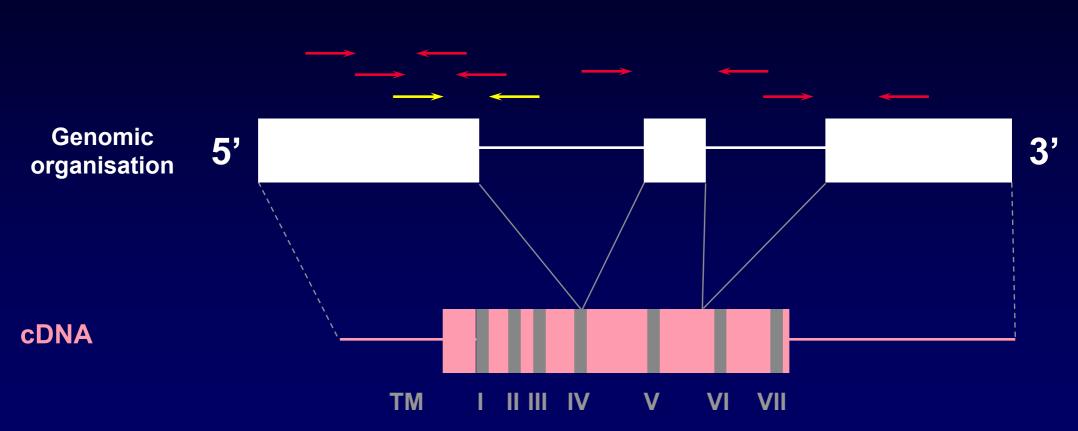
| Date | 13.1.96 | 23.2 | 19.7 | 28.10 | 6.3.97 |
|------|---------|-------|------|-------|--------|
| Т | 11.8 | 16.8 | 9 | 27 | 24 |
| TV R | 8-10 | 10-12 | 12 | 12 | 15 |
| TV L | 6-8 | 8-10 | 8-10 | 10 | 14-15 |

Spermogram:28.10.96: 1x103 sperm cells/mL20.02.97: 9x106 sperm cells/mL

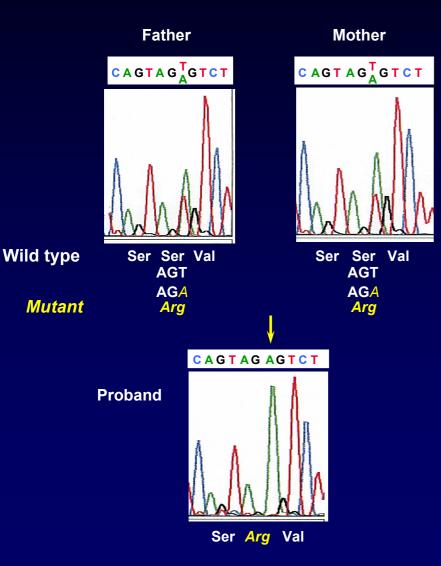
Hypothesis

- The origin of this patient's hypogonadism is at the pituitary level, rather than the hypothalamic level
- Presence of an inactivating mutation of the GnRH receptor gene

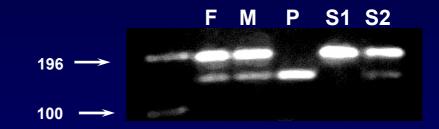
Sequencing Strategy

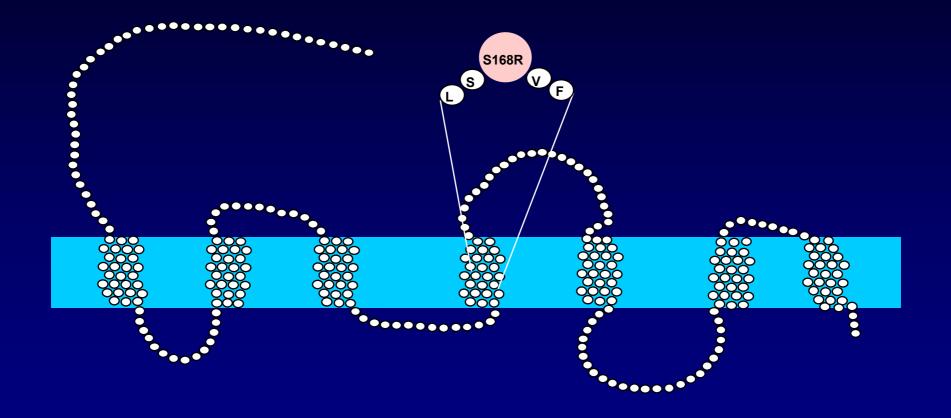


A T to A Point Mutation at Position 504 Results in a Change from Serine to Arginine at Residue 168 of the GnRH-R

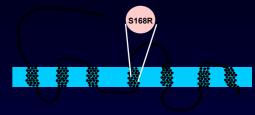


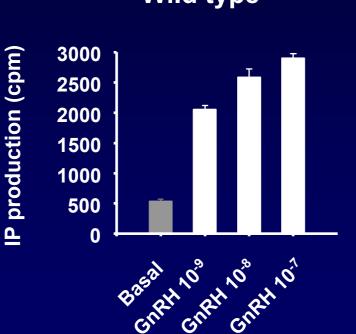
The T to A Mutation Introduces a Hinfl Restriction Enzyme Site





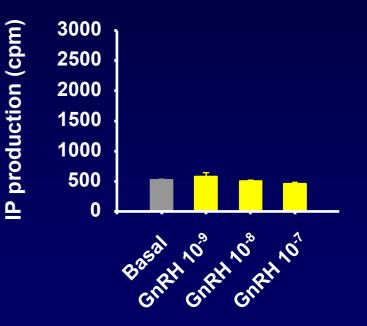
The S168R Mutation Is a Complete Loss-Of-Function Mutation



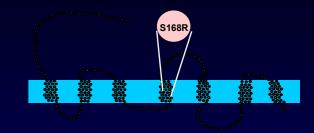


Wild type

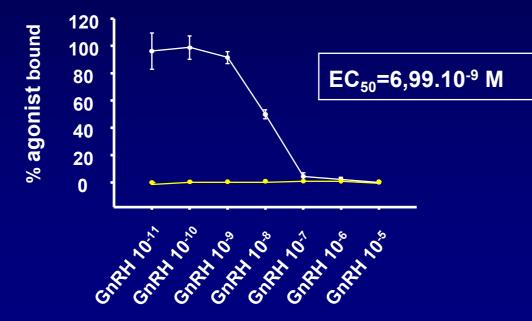
S168R mutant



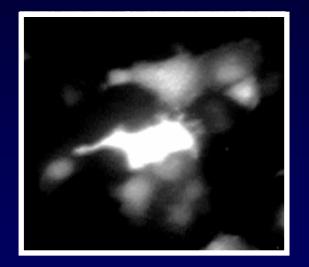
The S168R Mutation Abolishes Ligand Binding



Wild type receptor S168R mutated receptor



Intracellular Localization of the S168R Mutated GnRH-R





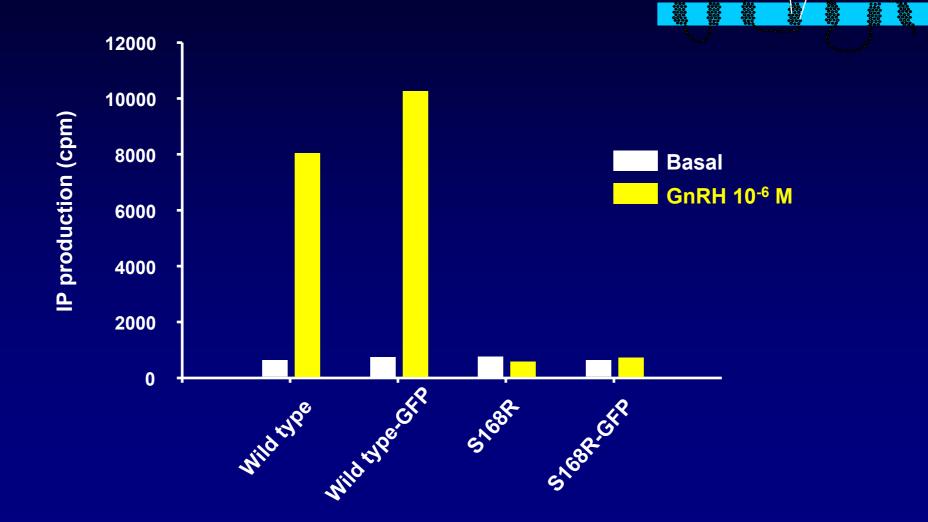


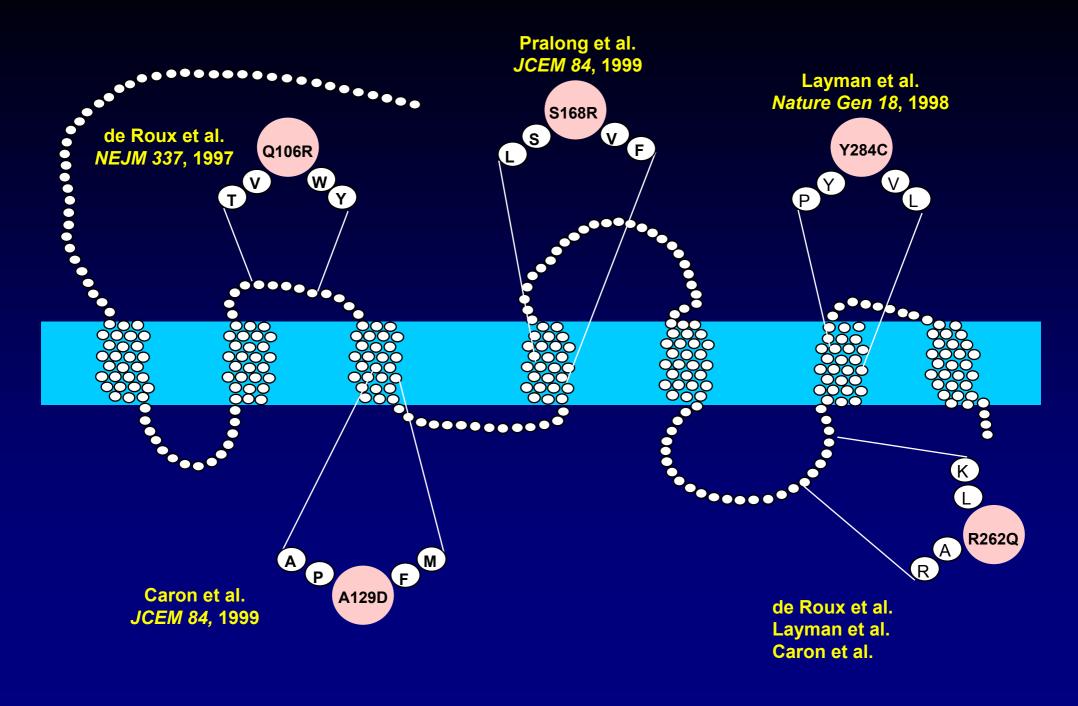
GFP

Wild type-GFP

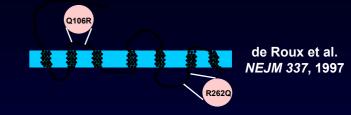
S168R-GFP

The GnRH Receptor-GFP Fusion Protein Is Functionally Active

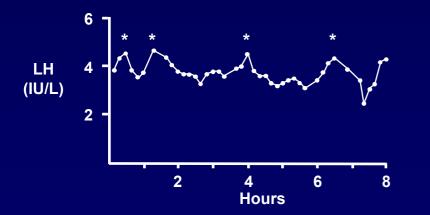




Phenotype of compound heterozygote patients



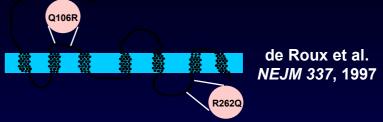
Male : scrotal testis (8 mL)
LH : 4.0 IU/L, FSH : 5.9 IU/L
Puberty at age 16 years
Basal LH secretion displays blunted pulsatility:

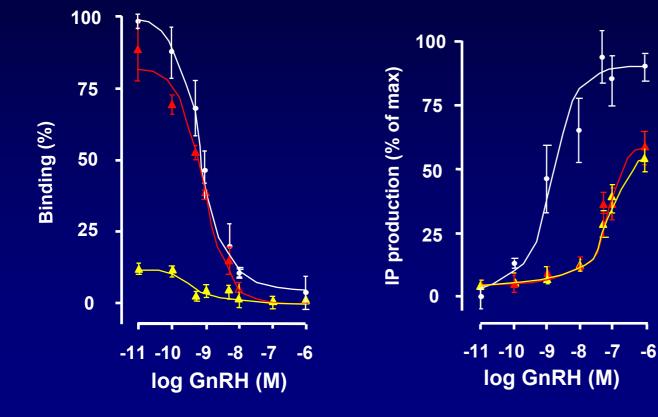


Normal response to GnRH (100 μg)
Female : primary amenorrhea and infertility
Telarche at age 14 years
LH : 5.0 IU/L, FSH : 5.2 IU/L, E2 128 pmol/L

Conclusion : incomplete hypogonadotropic hypogonadism

Functional characterization of compound heterozygote mutations





Q106R mutation R262Q mutation

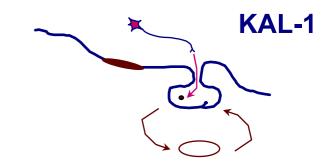
Conclusions

- There is a wide range of phenotypic expression of loss-of-function mutations of the GnRH-R, characterized by a variable degree of resistance to GnRH
- This phenotypic heterogeneity seems to correlate well with the degree of functional impairement of the receptor

Conclusions

Genotype/phenotype correlations performed on these patients provide invaluable information about structure-function relationships of the GnRH receptor

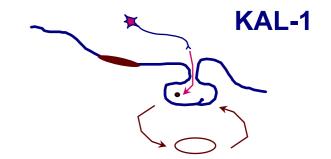




HH associated with anosmia

- X-linked syndrome
- due to mutations of KAL-1 (member of the superfamily of Neural Cell Adhesion Molecules)

Kalmann's Syndrome



The Candidate Gene for the X-Linked Kallmann Syndrome Encodes a Protein Related to Adhesion Molecules

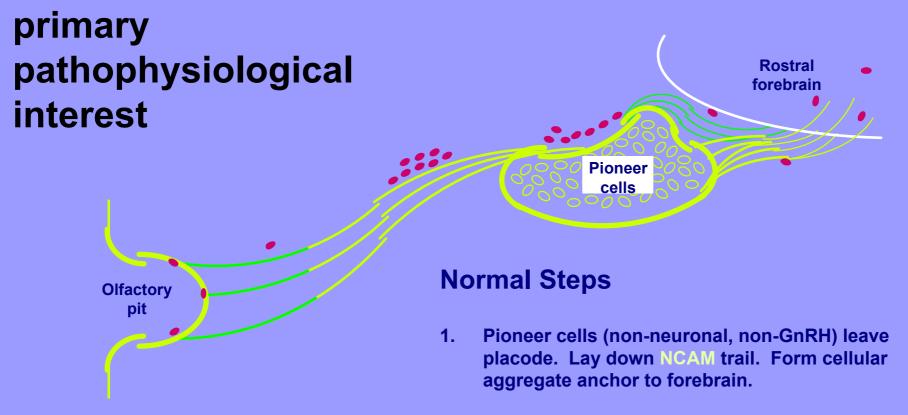
Legouis et al., Cell 67, 1991

A Gene Deleted in Kallmann's Syndrome Shares Homology with Neural Cell Adhesion and Axonal Path-Finding Molecules

Franco et al., Nature 353, 1991

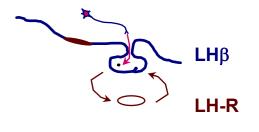
<u>KAL-1</u>

Schwanzel-Fukuda



- 2. Vomeronasal and terminal nerves follow trail.
- 3. **GnRH** cells follow nerves and trail.
- 4. Polysialated NCAM helps acceleration from placode and aggregate.

LH Deficiency - Males



LHβ

One single case described Bio-inactive LH

• Phenotype:

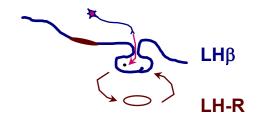
- normal male
- delayed puberty
- response to hCG: normal virilization, but not fertility
- Male heterozygotes: 3/4 infertile

Broad spectrum of phenotypic expression of inactivating mutations

LH-R

- pseudohermaphroditism and complete azoospermia
- micropenis, delayed puberty and arrest of spermatogenesis

LH Deficiency - Females



LH-R

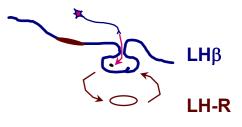
No LH-β mutation yet described in a female patient

- normal external genitalia
- normal pubertal development
- primary amenorrhea
- no pre-ovulatory follicles

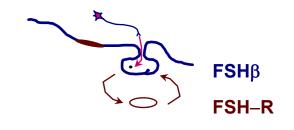
Role of the LH/LH-R System

- Important for normal male development
- LH-R plays a role in spermatogenesis as well as ovulation

LH-R is a candidate gene for male as well as female infertility



FSH Deficiency - Males



$FSH\beta$

Two cases described

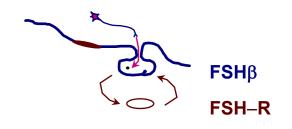
- Phenotype:
 - 1) delayed puberty, low testosterone and absent spermatogenesis
 - 2) normal puberty and virilization, spermatogenic arest



FSH-R

- normal virilization
- decreased testicular volume
- variable suppression of spermatogenesis

FSH Deficiency - Females



$FSH\beta$

Three cases described

- Phenotype:
 - delayed puberty
 - primary amenorrhea
 - normal response to FSH with achivement of fertility

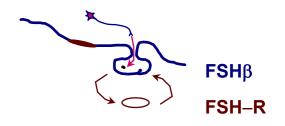
FSH-R

Finnish study

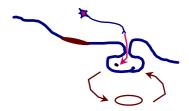
- Phenotype:
 - primary amenorrhea
 - ovarian dysgenesis with normal karyotype

Role of the FSH/FSH-R System

- Important for estrogen production, follicular maturation and fertility
- Role of FSH in spermatogenesis remains unclear:
 variable spermatogenesis in FSH-R mutations
 - absent spermatogenesis in FSH β mutations

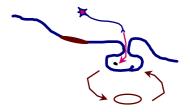


Conclusions



The study of inactivating mutations of several genes throughout the gonadotrope axis has provided invaluable insights into the physiology of reproduction in humans

Conclusions



These mutations offer a model of singlegene diseases, allowing genotype/phenotype correlations to be drawn and structure-function relationships to be inferred

Acknowledgments

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