



Academisch Medisch Centrum
Universiteit van Amsterdam
LEIDEN UNIVERSITY MEDICAL CENTER

*A novel cause of congenital central hypothyroidism:
 From two cousins to
 a novel X-linked endocrine syndrome*

A.S.Paul van Trotsenburg & Jan-Maarten Wit

**Emma Children's Hospital AMC, Amsterdam, Netherlands
 Willem-Alexander Children's Hospital, LUMC, Leiden
 On behalf of the international IGSF1 consortium**






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1. How it started

- **Index case TE**, d.o.b. 29-10-1994
- BW +2.3 SDS, BL +1.1 SDS, BHC +3.5 SDS (40.5 w)
Jaundice for 2 weeks, hydrocephalus.
- Screening for congenital hypothyroidism (age 9 and 13 d):
 - T4 -2.4 and -3.3 SDS
 - TSH 5 mU/l

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LUMC *1. How it started* **amc**

- Hospital, AMC:
 - FT4 8.3 pmol/L, TSH 3.5 mU/L
 - TRH test: TSH max 10.4 mU/L (low)
 - GH, PRL, IGF-I, CRH and GnRH test nl.
 - Start L-T4 treatment.
 - TRHR and TSHB genes nl.
- MRI: external hydrocephalus. VP drain at 1.0 yr.
- At age 1.8 y: transfer of care to LUMC
- Height SDS → -1.3
- 11.0-15.2 y testes 4->16 mls, but low T until 15.2 yrs → T treatment; BMI +2.5 SDS. PRL borderline

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LUMC *His cousin* **amc**

- RH, dob 8-2-1991
- BW +2.2 SDS, BL +1.6 SDS (43 w)
- At 7.2 y referred to LUMC
 - growth deviation
 - height SDS -1.4,
 - BMI +2.2,
 - delayed bone age (3.1 y)

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LUMC *His cousin* **amc**

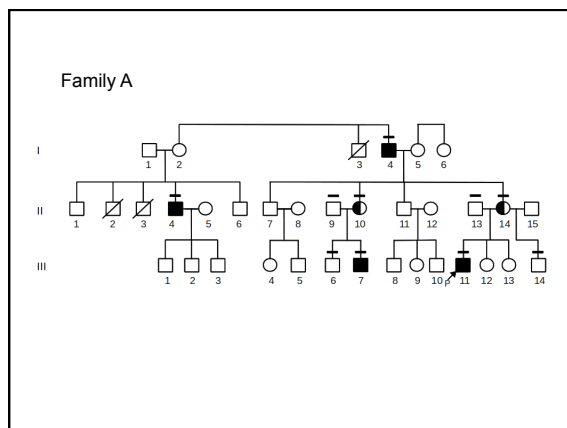
- Lab. investigation:
 - FT4 8.8 pmol/L, TSH 1.6 mU/L
 - TRH test: TSH max 4.3 mU/L, PRL low, IGF-I -1.3 SDS.
 - Start L-thyroxine
- Lab. investigation 2:
 - GH max 16.9 and 13.1 mU/L, IGF-I -1.0 SDS
- 8.8 y: start GH treatment: excellent catch-up, adult height 191.6 cm (+1.3 SDS) (Target height 0.6 SDS). BMI 24.4 (+1.5 SDS). Retesting: GH max normal.
- 9.9-14.2 y testes 4->16 mls, while T still low. GnRH test: LH 0.7-13.4, FSH 6.5-13.4 U/L. T↑ from 14.5 y

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LU MIC *Maternal grandfather A-I.4* **amc**

- Growth:**
 - normal stature, overweight (BMI 33), GH: normal
- Thyroid/prolactin:**
 - FT4 9 pmol/L, TSH 0.9 mU/L, PRL normal
- Adrenal:**
 - possibly central hypocortisolism. Low DHEAS
- Gonads:**
 - late puberty, large testis (removed after torsion), contralateral testis atrophic, postoperative primary hypogonadism
- CNS/behavior:**
 - normal (retired general practitioner)

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LU MIC *(Patho)physiology of thyroid hormone secretion* **amc**



The flowchart illustrates the (patho)physiology of thyroid hormone secretion. It starts with the Hypothalamus (paraventricular nucleus) secreting TRH (+), which stimulates the Pituitary (thyrotrophs) to secrete TSH (+). TSH (+) stimulates the Thyroid (thyrocytes) to produce T₄ and T₃. T₄ and T₃ are then transported to target tissues (nuclear receptors) and excreted. Liver conjugation is also shown. Feedback loops are indicated by (-) signs: T₄ and T₃ exert negative feedback on the Pituitary and Hypothalamus.

Causes of hypothyroidism
Hypothalamus/pituitary: TRHR and TSHB genes defect
 Central (sec./tert.) hypothyroidism

Thyroid: agenesis, hypoplasia, enzyme defects
 Primary hypothyroidism

CH screening in Netherlands: T4+TSH+TBG approach.
 If low T4 + normal TSH + low T4/TBG ratio: **central hypo?**

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Conclusion family A

Novel X-linked syndrome of:

- Pituitary TSH deficiency
- Variable prolactin deficiency
- Macroorchidism and delayed puberty
- Partial GH deficiency?
- Large birth size, overweight

Plan: exome sequencing of the X-chromosome

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AMC

5 ½ families with clinical picture of "X-linked" central hypothyroidism:

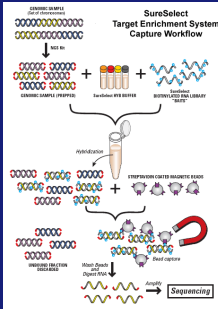
- TE (+RH) - nephews
- CS (1984) + YS (1989) - nephews
- AJ + MH (1990) + (1991) - nephews
- PK (1999)+ SK (2003) - nephews
- DL (1991) + SR (2002) - (half-)brothers
- JS (1995) + KS (2000) - brothers

2003/2004: linkage studies - not succesful
2010, plan: exome sequencing of the X-chromosome

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2. Discovery of the gene defect: Exome sequencing




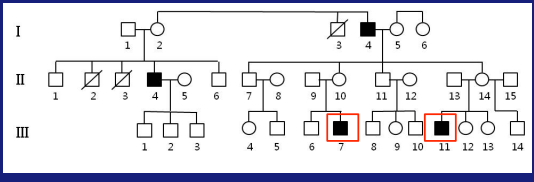
- Mutation detection in monogenic diseases
- Drawback
 - In principle only exonic regions
 - Variable capture efficiency
 - Difficulties in analysis (comparison)

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LU
MC

Exome sequencing Family A (Yu Sun)

- SNP array to detect candidate region (III.11 vs 14)
- Exome sequencing III-7 and III-11
- Sureselect Agilent, X-exome capture
- GAll Illumina, 51nt paired end run

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LU
MC

Standard variant calling and filtering

- BWA + samtools : SNVs and short indels
- Seattleseq Annotation
- 1000 Genomes Project
- Allele frequency Hapmap

	III-7	III-11	in common
#indels	735	770	510
#SNVs	2278	2366	1922
#SNVs not in 1000 Genomes Project			591
#SNVs hapmap freq <1%			471
#SNVs in region			129

SNV = single nucleotide variant

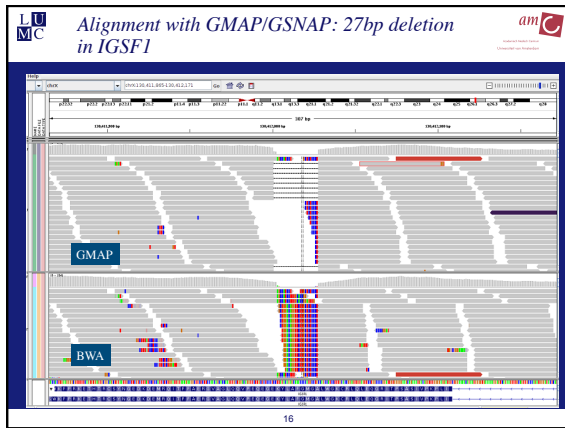
14

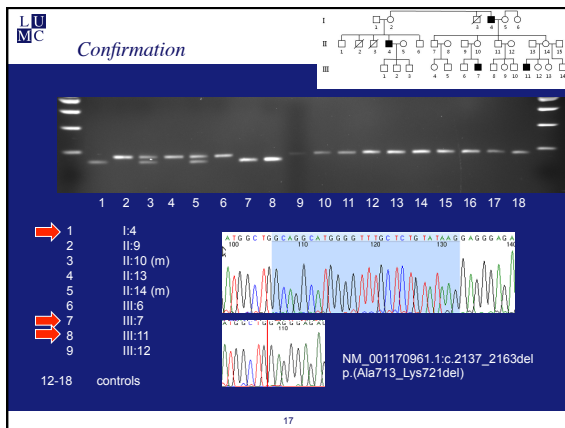
LU
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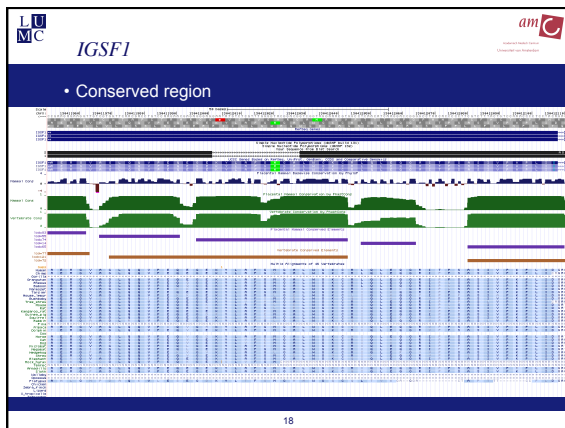
Searching for candidate gene



129 SNVs in region

	All	Not in dbSNP
nonsense	0	0
missense	10	2
silent	1	0
Splice-3	0	0
Splice-5	0	0
3-utr	11	3
5-utr	12	1
intron	63	20
intergenic	20	3
Near-gene-3	9	1
Near-gene-5	7	2









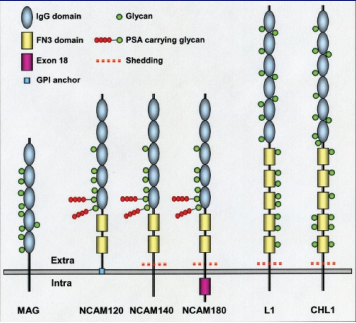
3. What was known of IGSF1

- Immunoglobulin superfamily member 1
- Membrane glycoprotein
- 3 isoforms (REFSEQ)
- Function
 - Unclear
 - Possibly a co-receptor in inhibin signaling, but not a high-affinity inhibin receptor.
 - Antagonizes activin A signaling in the presence or absence of inhibin B (by similarity).
 - Necessary to mediate a specific antagonistic effect of inhibin B on activin-stimulated transcription.

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Other members of the family



Structure of immunoglobulin cell adhesion molecules (IgCAMs)

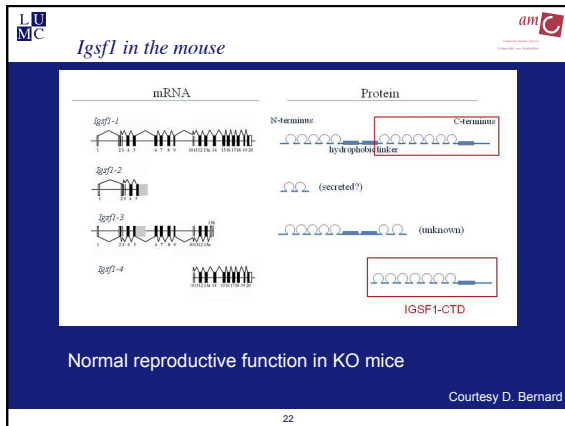
Irintchev, The Neuroscientist 2011 20

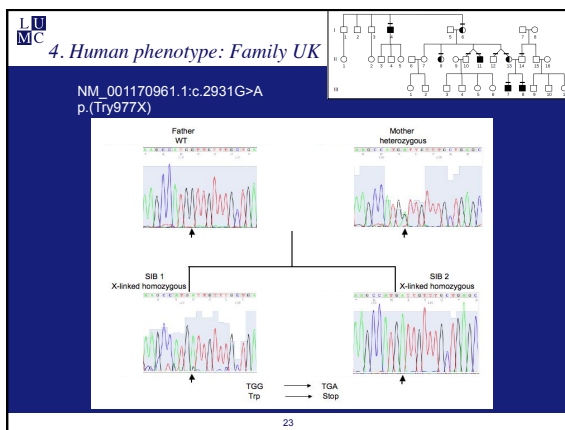



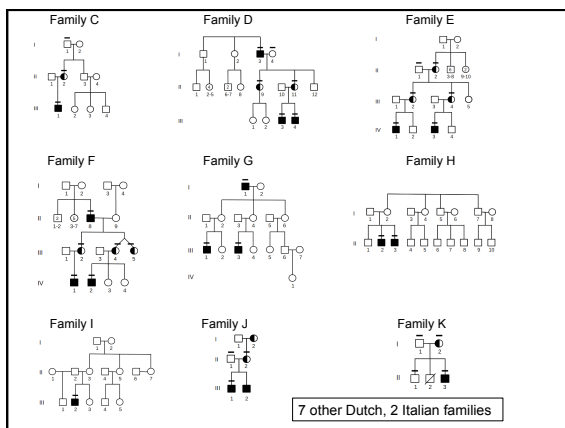
IGSF1

- Expression
 - Highly expressed in pituitary, hypothalamus, pancreas, testis and fetal liver.
 - Moderately expressed in heart, prostate and small intestine.
 - Expressed at very low levels in thymus, ovary, colon, fetal lung and fetal kidney.

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IGSF1 mutations predicted to result in loss of function

Whole gene deletions:
Family E 126kb deletion arr Xq26.1q26.2(130.386.267-130.512.002)x0 (hg19)
Family F 328kb deletion arr Xq26.1q26.2(130.310.905-130.639.353)x0 (hg19)

25 maandag 10 maart 14

The variants

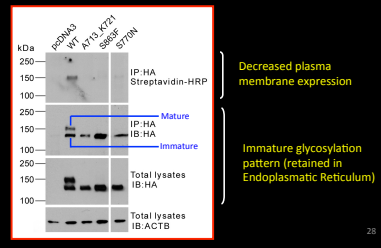
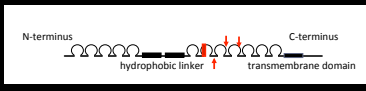
- Not present in database
 - Local inhouse database, dbSNP, 1000 Genomes Project, HGMD, LOVD
- Cosegregate with phenotype
- Might affect function of the protein

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IGSF1 truncation mutants are not trafficked to the plasma membrane

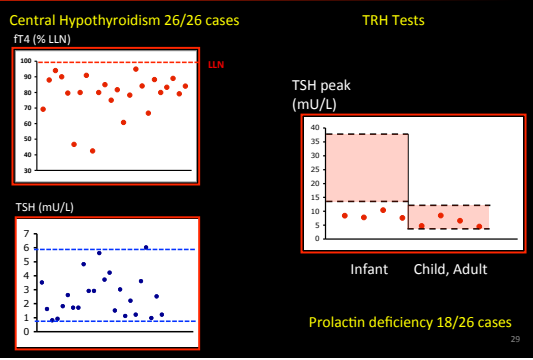
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IGSF1 in-frame mutants are inefficiently trafficked to the plasma membrane



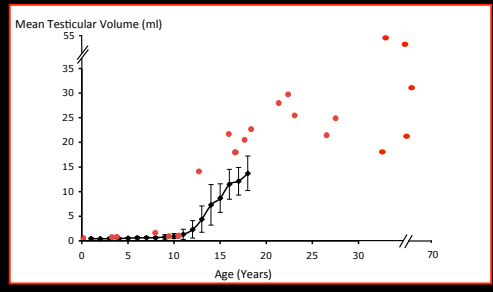
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IGSF1 Loss-of-Function Causes Central Hypothyroidism and Prolactin Deficiency



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IGSF1 Loss-of Function Causes Adult Macroorchidism

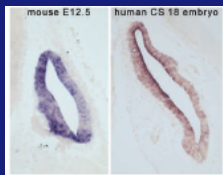


Pubertal development is disharmonious: testosterone low for testis size

Ultrasound references from Goede et al Horm Res Paediatr 2011; 76:56-64 30

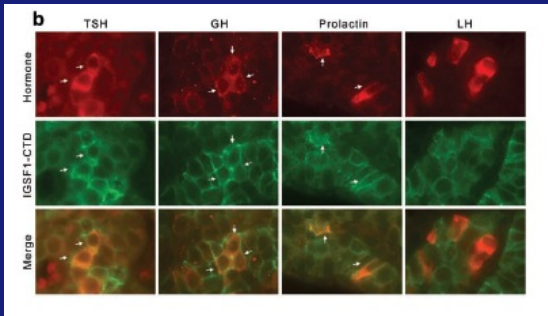
5. Mouse phenotype

- Expression of IGSF1/Igsf1 mRNA in murine embryonic day 12.5
- Also expressed in human embryo Carnegie stage 18 Rathke's pouch (in situ hybridization)



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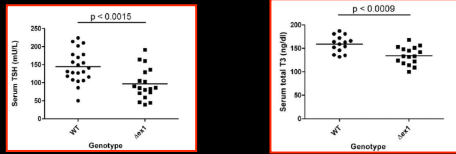
Igsf1 is expressed in TSH, GH and PRL secreting cells



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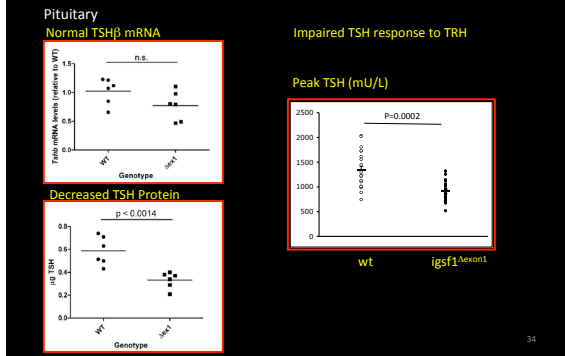
TSH and T3 levels are decreased in male Igsf1^{Δex1} mice

Decreased Serum TSH Decreased Serum T3

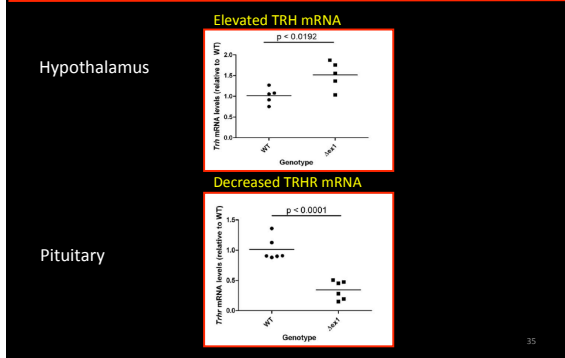


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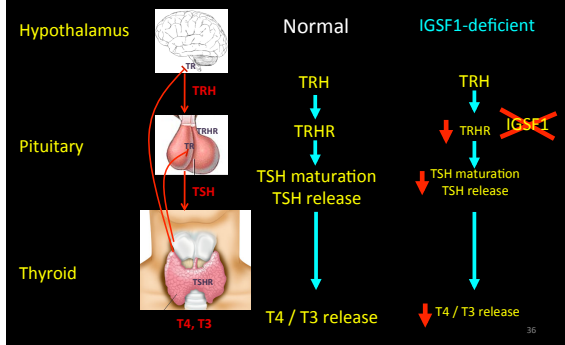
TRH signalling may be impaired in male *Igsf1^{Δex1}* mice





TRH signalling may be impaired in male *Igsf1^{Δex1}* mice



Central Hypothyroidism due to *IGSF1* Loss-of-Function



Conclusions

- *IGSF1* mutations cause a novel, X-linked syndrome of central hypothyroidism, testicular enlargement and variable prolactin deficiency
- Central hypothyroidism (and PRL deficiency?) caused by defects in TRHR expression and TRH signalling.
- croorchidism: mechanism unresolved
- Detection important because of theoretical reasons that hypothyroidism should be treated

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