

Hyperplasie macronodulaire des surrénales : stratégies de recherche de nouveaux gènes de prédisposition

Pr Guillaume Assié

Service d'Endocrinologie, Cochin
Institut Cochin, INSERM, CNRS, Université Paris Descartes



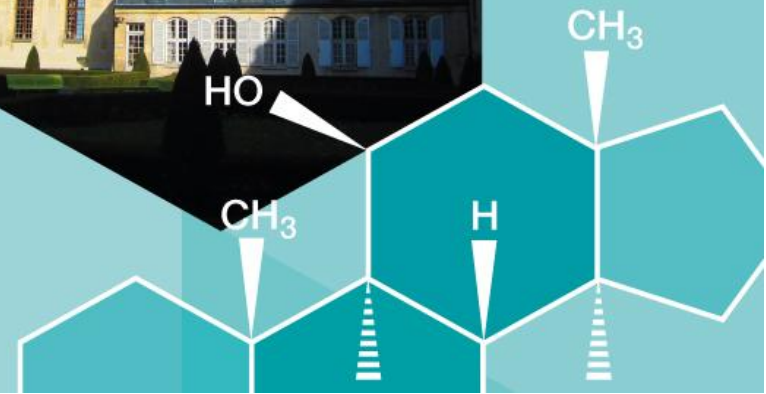
JOURNÉE
ANNUELLE

4^{ème} ÉDITION

FIRENDO

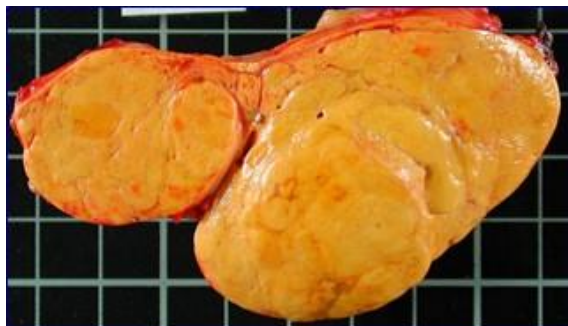
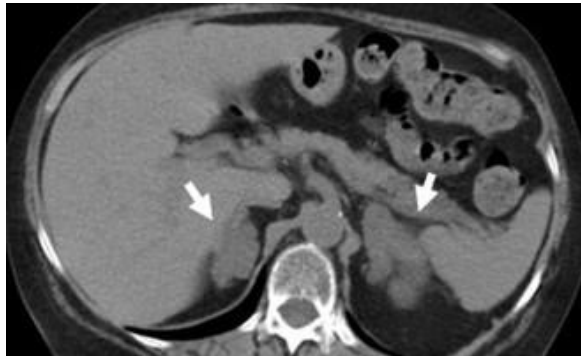
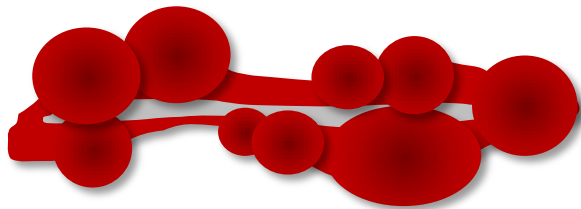
FILIÈRE MALADIES RARES ENDOCRINIENNES

Mercredi
13 décembre
2017



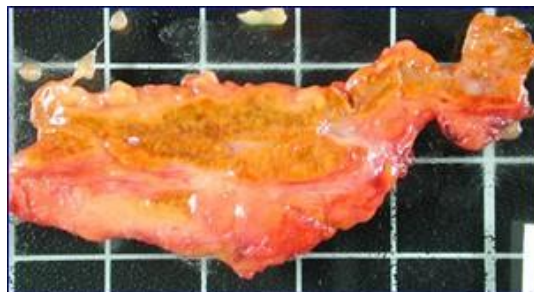
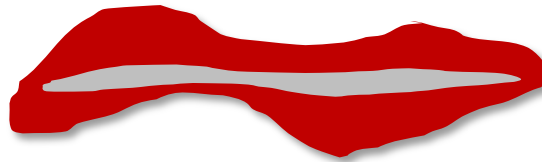
Bilateral enlargement of both adrenals

Macronodular Hyperplasia



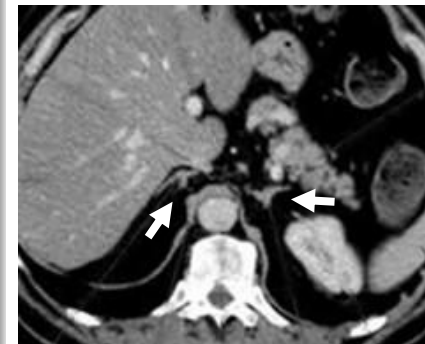
F Tissier

Diffuse Hyperplasia

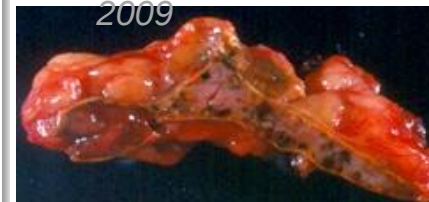


F Tissier

Primary Pigmented Nodular Adrenal Dysplasia



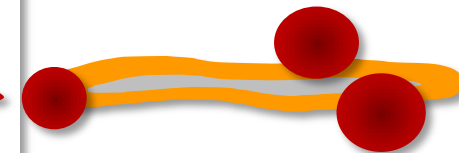
Hsiao, JCEM 2009



Meador et al, JCEM 1967

Carney et al, Medicine 1985

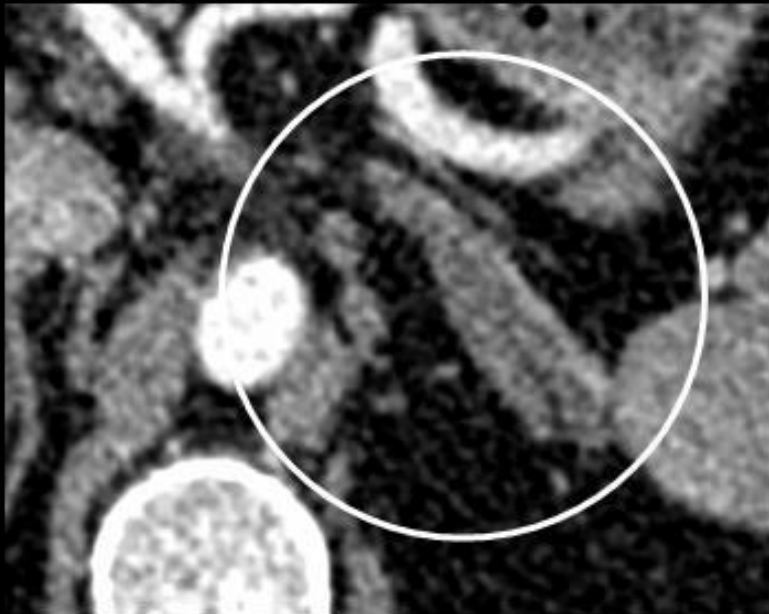
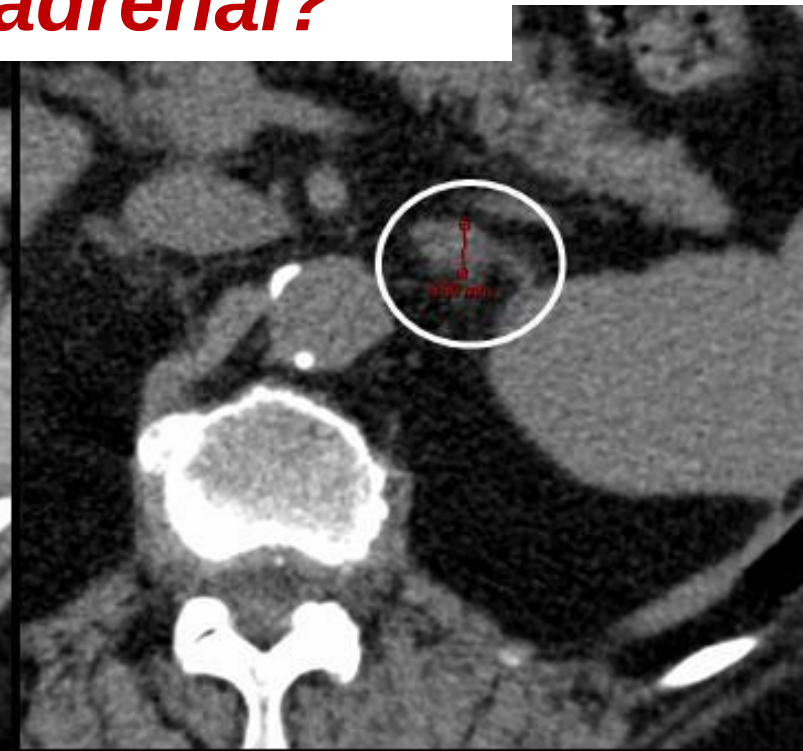
Multiple adenomas



Differential diagnoses

- CAH
- Hemorrhagia
- Metastases, lymphoma
- Pheo
- ACTH-dpt hypercortisolism

What is a normal adrenal?



Variable Hypercortisolism

Cushing syndrome

Non-
secreting

Subclinical
hypercortisolism

Overt
hypercortisolism

Hypertension
Diabetes
Osteoporosis
Infarct, Stroke
Thrombo-embolic
disease



Prevalence?

- 2% of overt hypercortisolism
- A « significant proportion » of incidentalomas (incidentalomas: 1-5% of population)

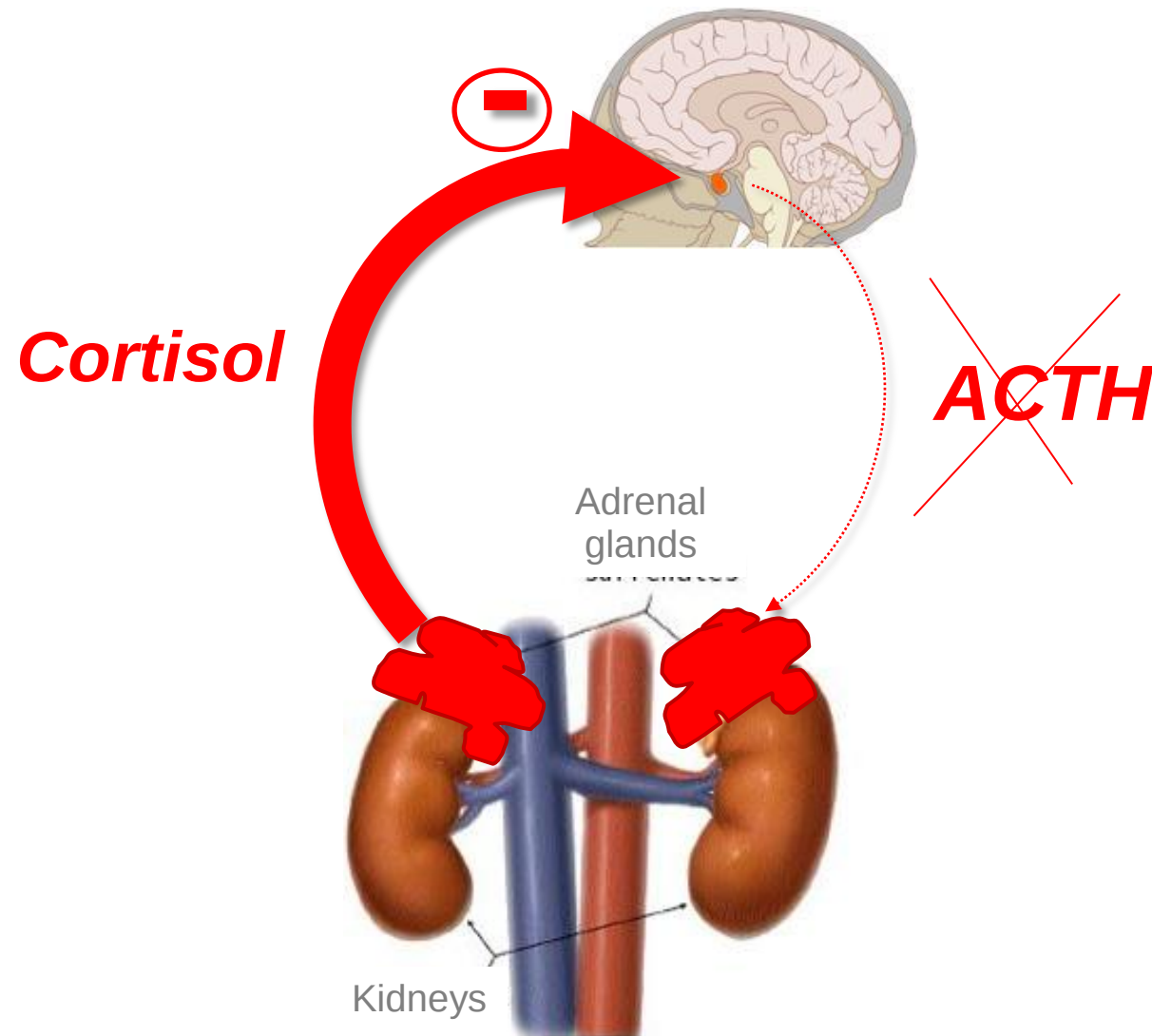
The adrenal networks....



CCTN

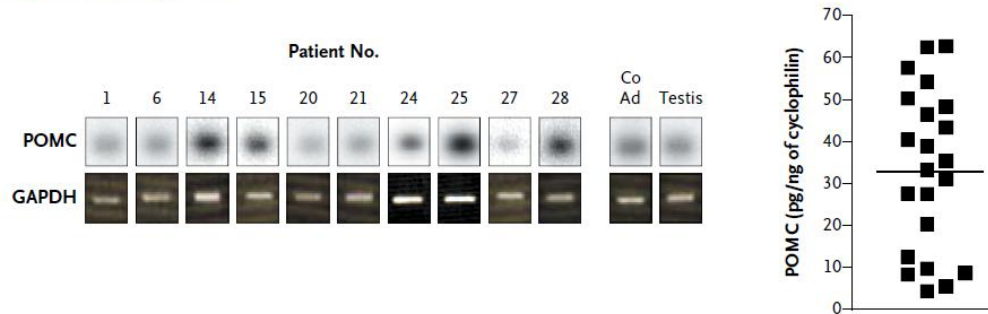


« ACTH-independant Adrenal Hyperplasia... »

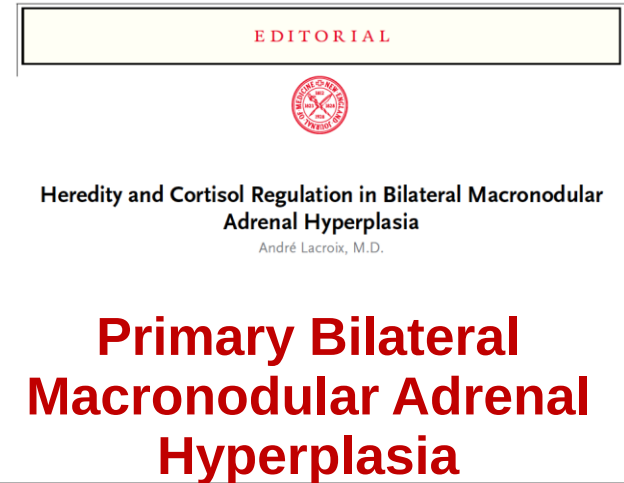
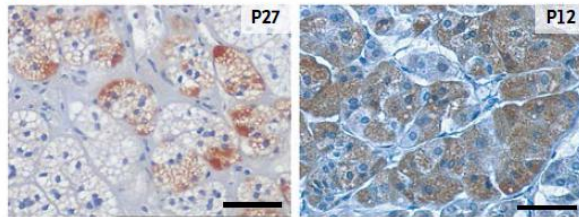


...But is it really « ACTH-independent? »

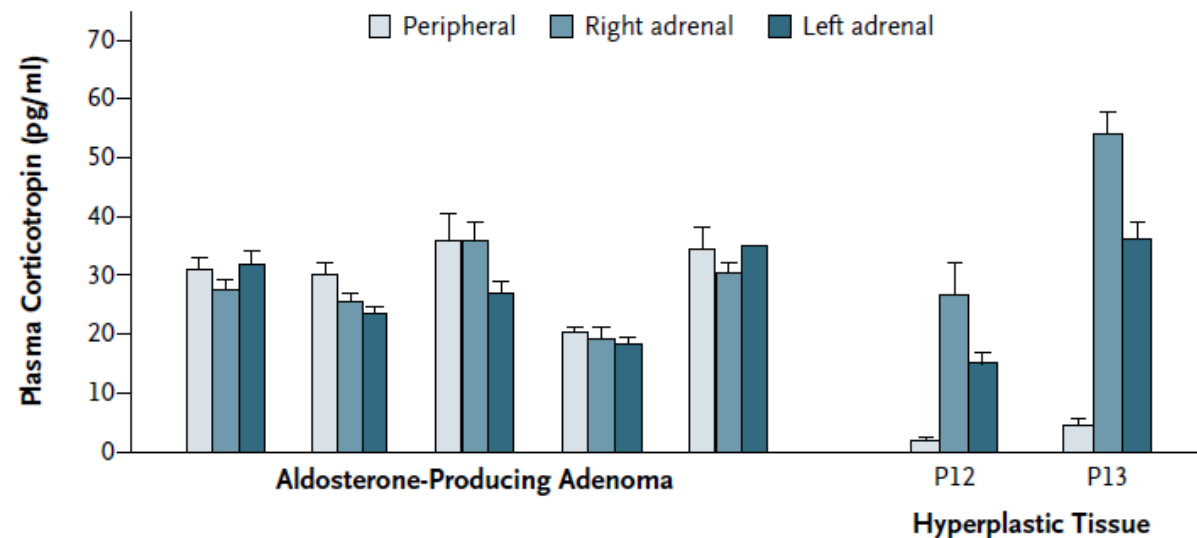
A POMC mRNA Expression



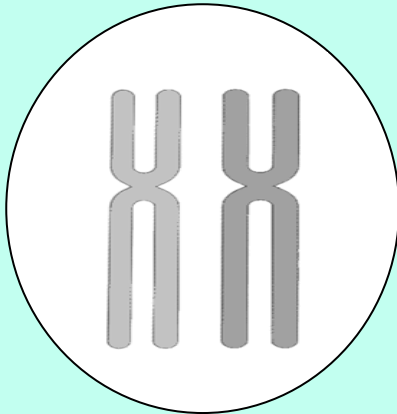
B Immunohistochemical Staining for Corticotropin



Louiset et al., NEJM 2013

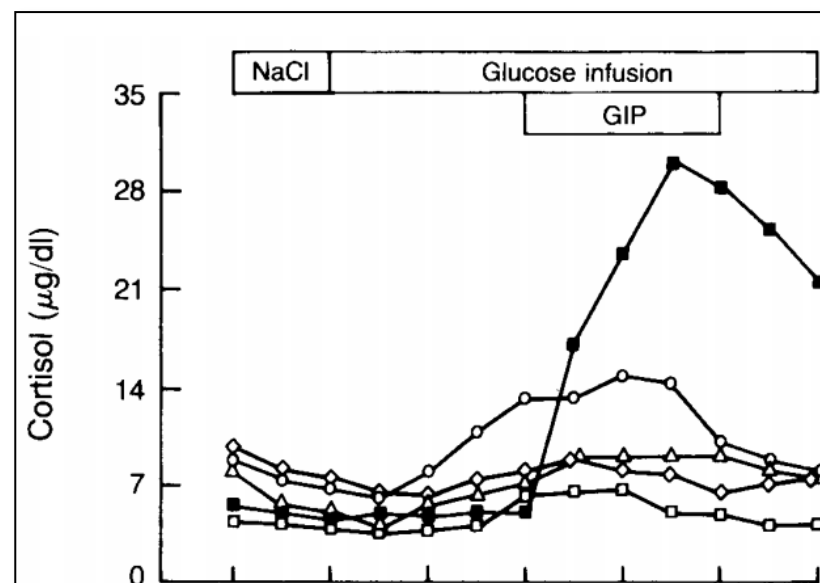
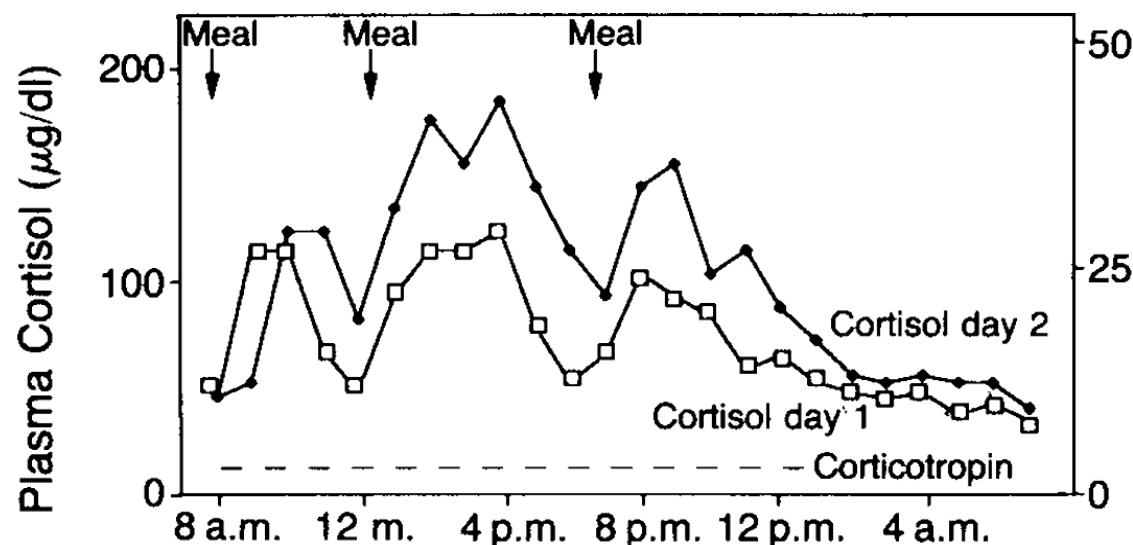


ACTH



FOOD-DEPENDENT CUSHING'S SYNDROME MEDIATED BY ABERRANT ADRENAL SENSITIVITY TO GASTRIC INHIBITORY POLYPEPTIDE

YVES REZNIK, M.D., VERONIQUE ALLALI-ZERAH, M.D., JEAN A. CHAYVIALLE, M.D.,
ROBERT LEROYER, PH.D., PIERRE LEYMARIE, M.D., GEORGES TRAVERT, PH.D.,
MARIE-CHRISTINE LEBRETHON, M.D., ILSE BUDI, M.D.,
ANNE-MARIE BALLIERE, M.D., AND JACQUES MAHOUDEAU, M.D.



GASTRIC INHIBITORY POLYPEPTIDE-DEPENDENT CORTISOL HYPERSECRETION — A NEW CAUSE OF CUSHING'S SYNDROME

ANDRÉ LACROIX, M.D., EDOUARD BOLTÉ, M.D., JOHANNE TREMBLAY, PH.D., JOHN DUPRÉ, M.D.,
PIERRE POITRAS, M.D., HÉLÈNE FOURNIER, M.D., JEAN GARON, M.D., DOMINIQUE GARREL, M.D.,
FRANCIS BAYARD, M.D., PH.D., RAYMOND TAILLEFER, M.D., RICHARD J. FLANAGAN, PH.D.,
AND PAVEL HAMET, M.D., PH.D.

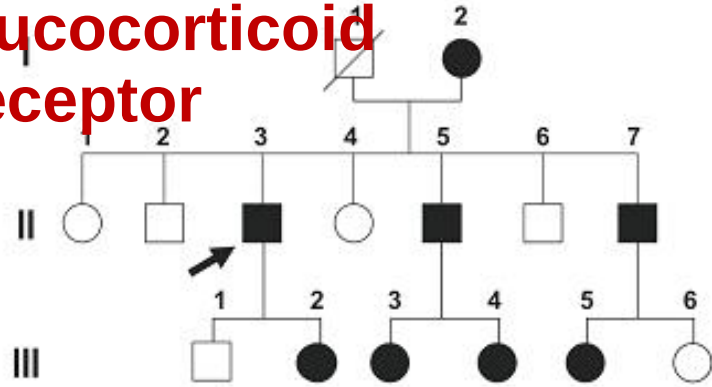
List of ectopic receptors

<i>Ligand</i>	<i>Receptor</i>	<i>In vivo response</i>
GIP	GIPR	Food
Vasopressin	V1a	Postural Test
Epinephrin	ADRB	Postural Test
LH	LH/HCG R	Pregnancy, menopause
Serotonin	HTR4	Metoclopramide

Duplication of GIPR

LeCoq et al, JCI Insight 2017

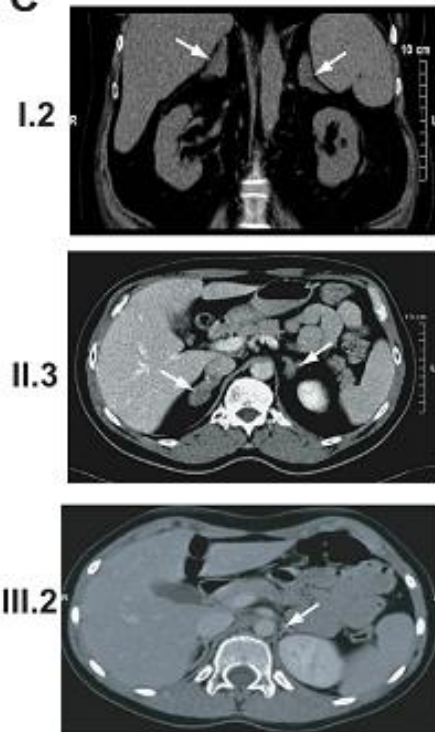
Glucocorticoid Receptor



B



C



Bouligand et al, PlosOne 2010

APC (Familial Adenomatous Polyposis)

Hsiao et al, 2009
Ymakita et al, 1997
Marchesa et al, 1997

MEN 1 (Multiple Endocrine Neoplasia Type 1)

Skogseid et al, 1992
Burgess et al, 1996
Hsiao et al, 2009

« Common » PMAH? A genetic disease?

- Bilateral affection
- Rares familial reports

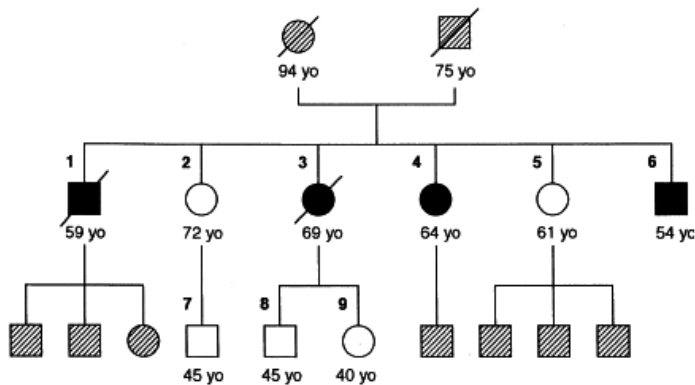
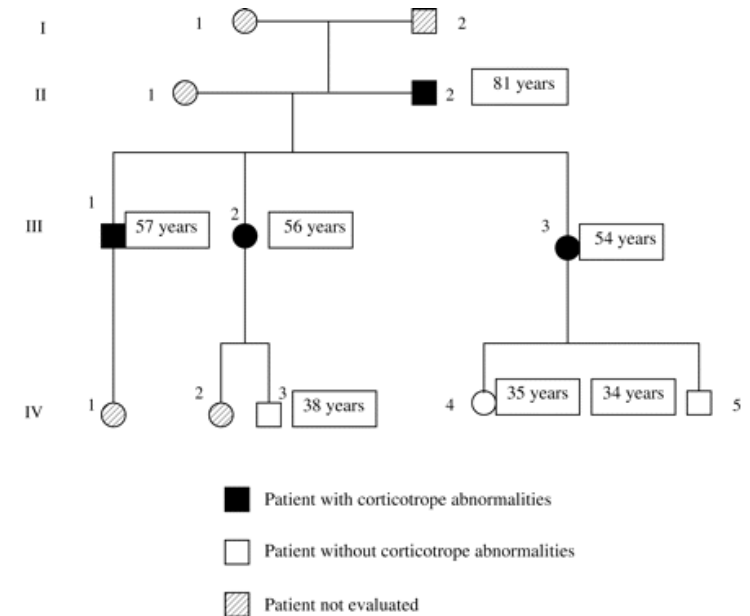


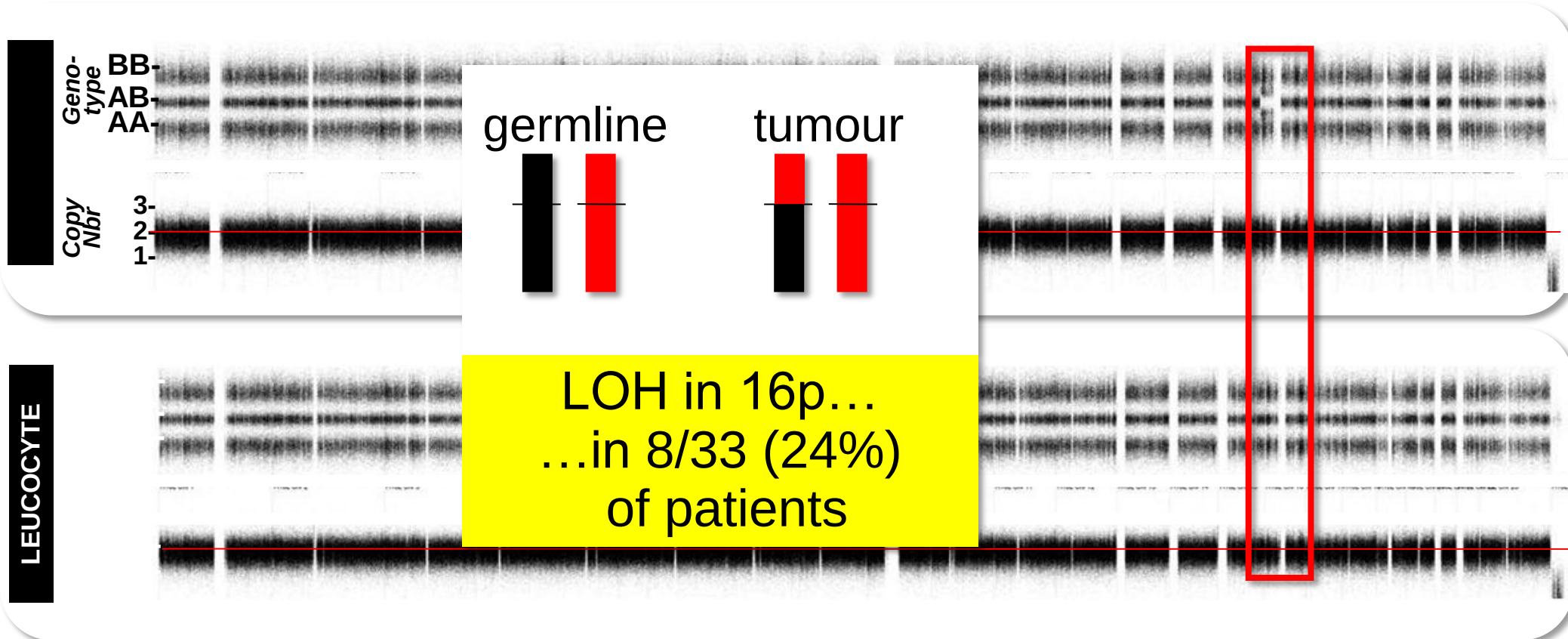
Fig. 1 Pedigree of the family. □, ▨, ■, Male family members; and ○, ●, ●, female family members. ■, ●, Affected members; □, ○, unaffected members; ▨, ●, not examined; /, deceased members. The examined members are numbered.

Nies et al. 1996



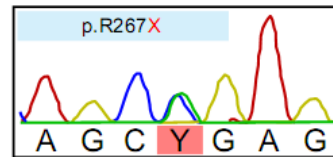
Vezzosi et al, EJE 2007

SNP Tumor Profiles in PMAH

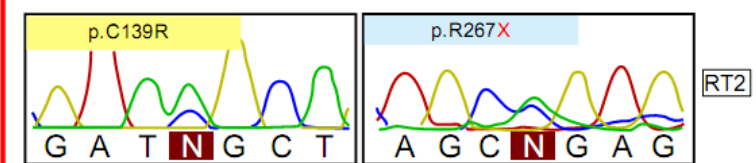
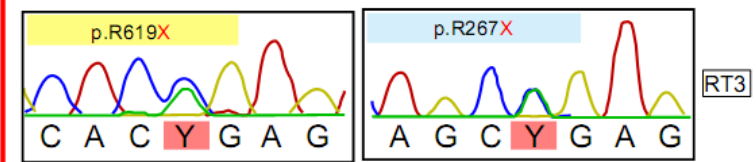
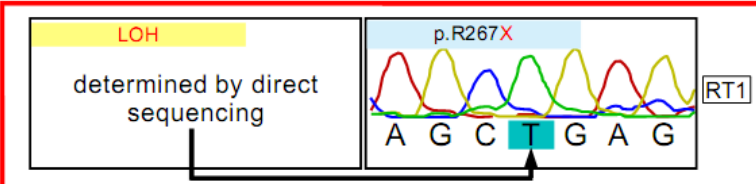


ARMC5: a tumor suppressor gene

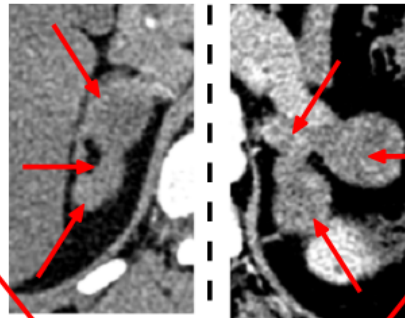
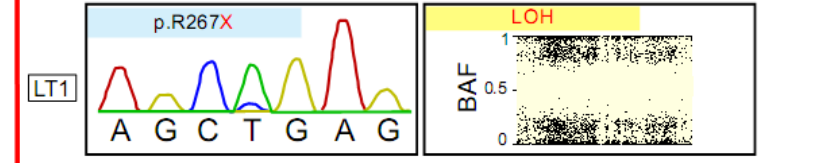
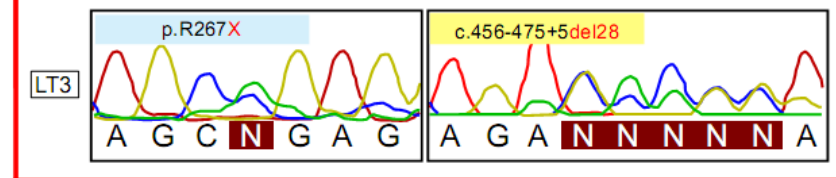
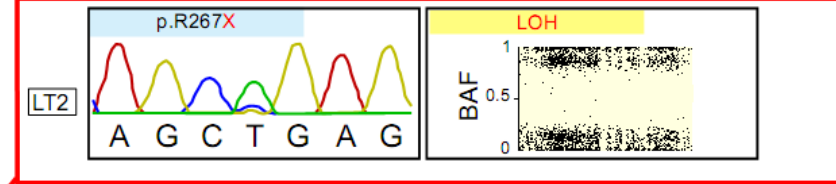
Germline DNA



Tumor DNA



Tumor DNA



Right adrenal

Left adrenal

2 hits on *ARMC5*:

- One germline
- One nodule-specific

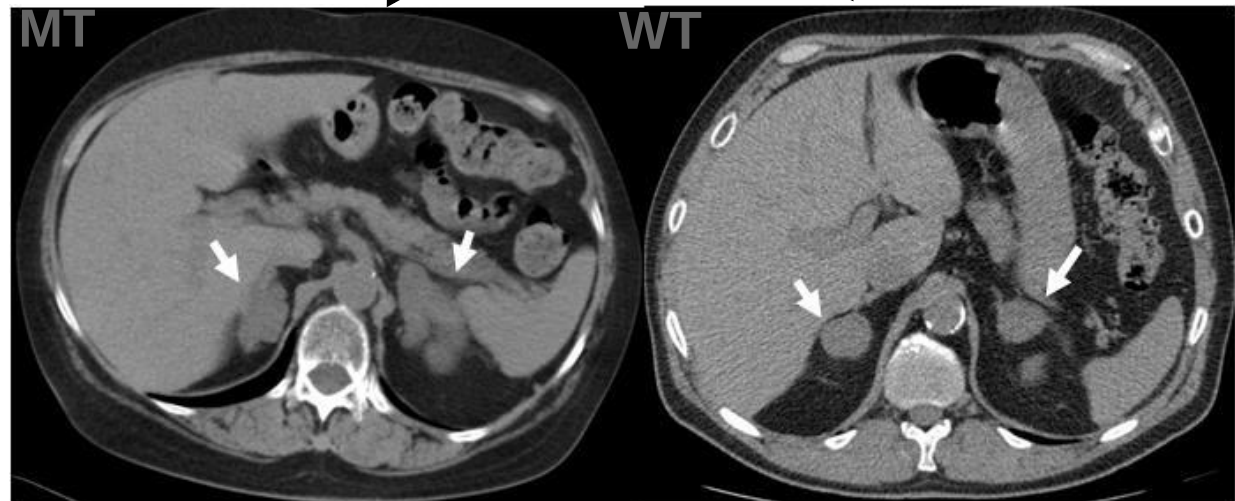
Correlation genotype-phenotype



Characteristics	Mutated (n=24) mean [95%CI]	Wild-type (n=68) mean [95%CI]	P-value
Surgery	20Y/4N	23Y/45N	<0,001
Cortisol after Dexamethasone suppression (µg/dl)	18 [0-38]	6 [0-19,7]	<0.001
Total weight (gr)	107 [28-186]	50 [0-128]	0.001
Number of nodules			<0,001
2	3	42	
3	1	9	
4	1	3	
5 or more	14	7	

→ Better
classification?

Espiard et al, JCEM 2015
Gagliardi et al, JCEM 2014
Alencar et al, JCEM 2014
Faucz et al, JCEM 2014



Prevalance of ARMC5 mutations in PMAH —

- 50% Operated

Assié et al, NEJM 2013

- 25% “Overt” PBMAH

*Albiger et al, Endocrine
2017*

Espiard et al, JCEM 2015

Gagliardi et al, JCEM 2014

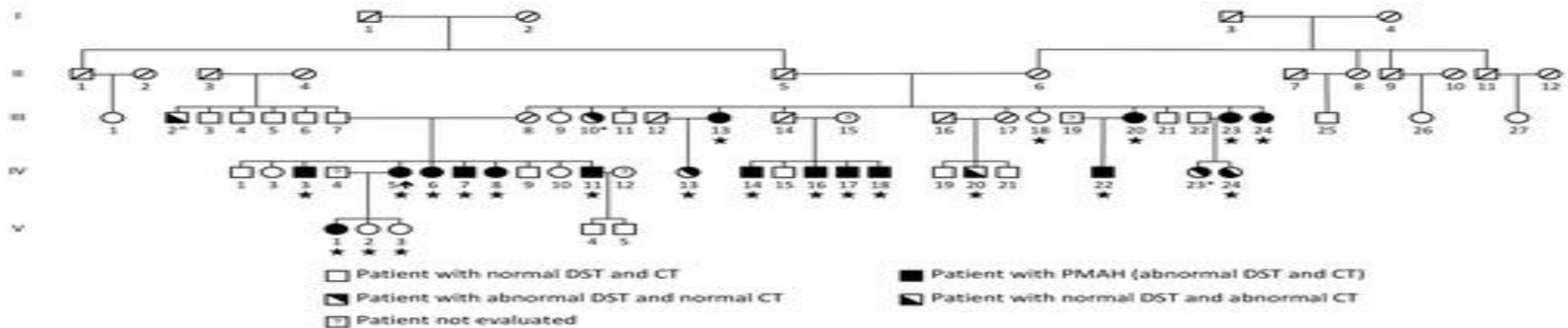
Alencar et al, JCEM 2014

Faucz et al, JCEM 2014

- <5% Bilateral Incidentalomas

Ems et al, Endocrine 2016

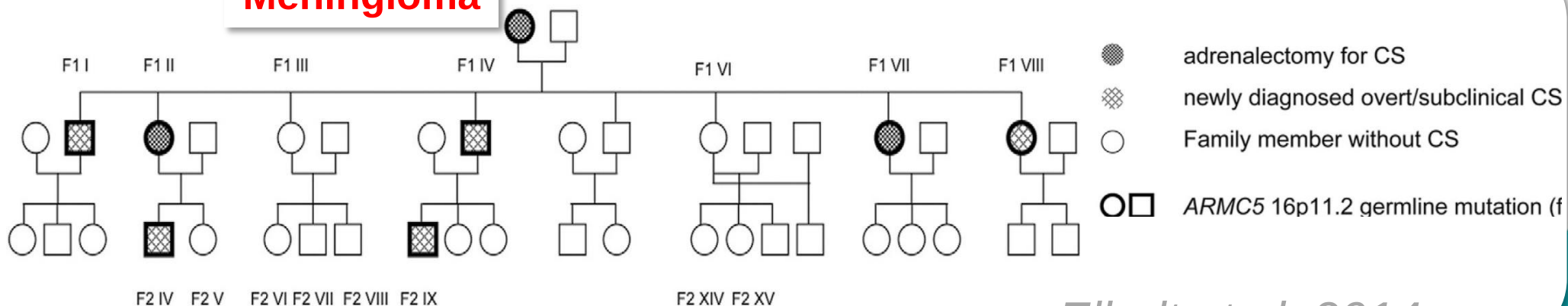
ARMC5-related PMAH is not sporadic, but familial...



Alencar et al, JCEM 2014

... and potentially syndromic

Meningioma



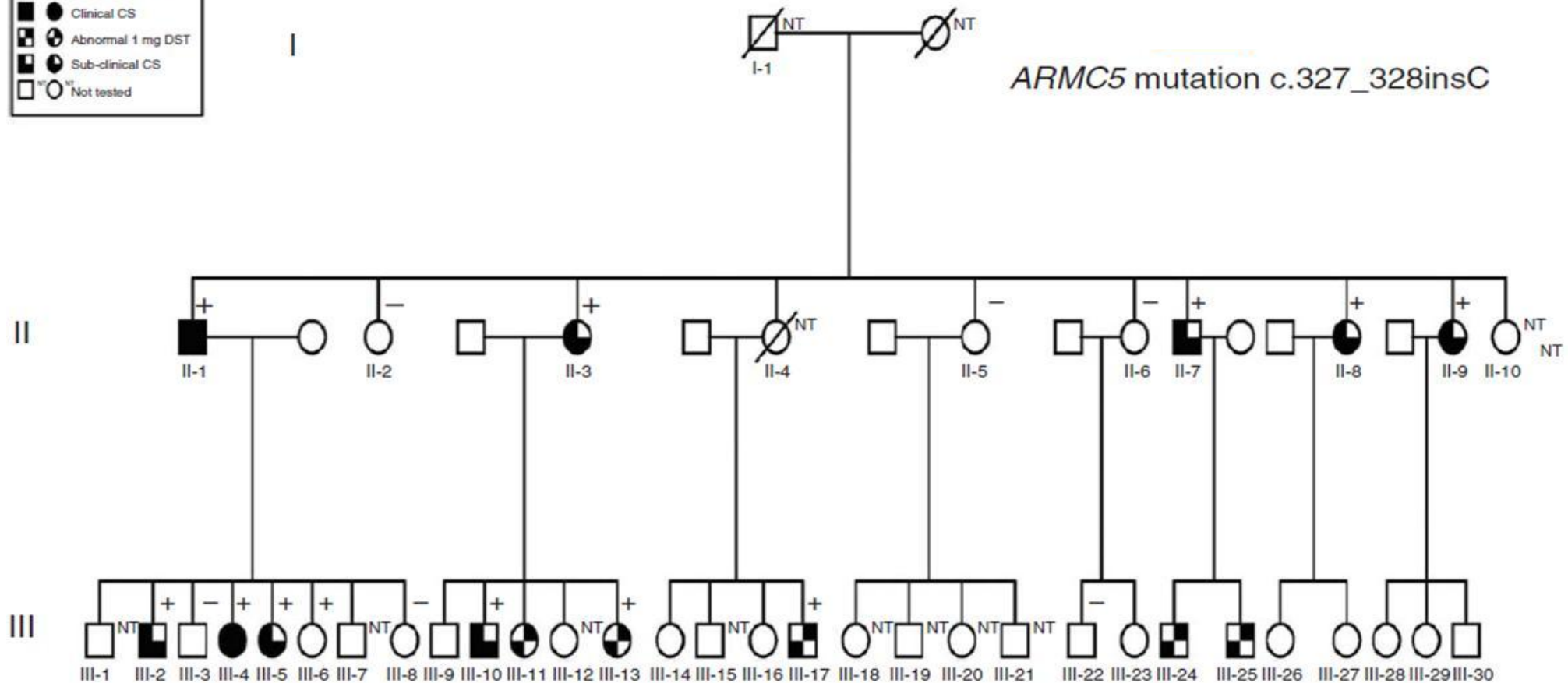
Elbelt et al, 2014

ARMC5-related PMAH is not sporadic, but familial...

Aberrant β -adrenergic and vasopressin response

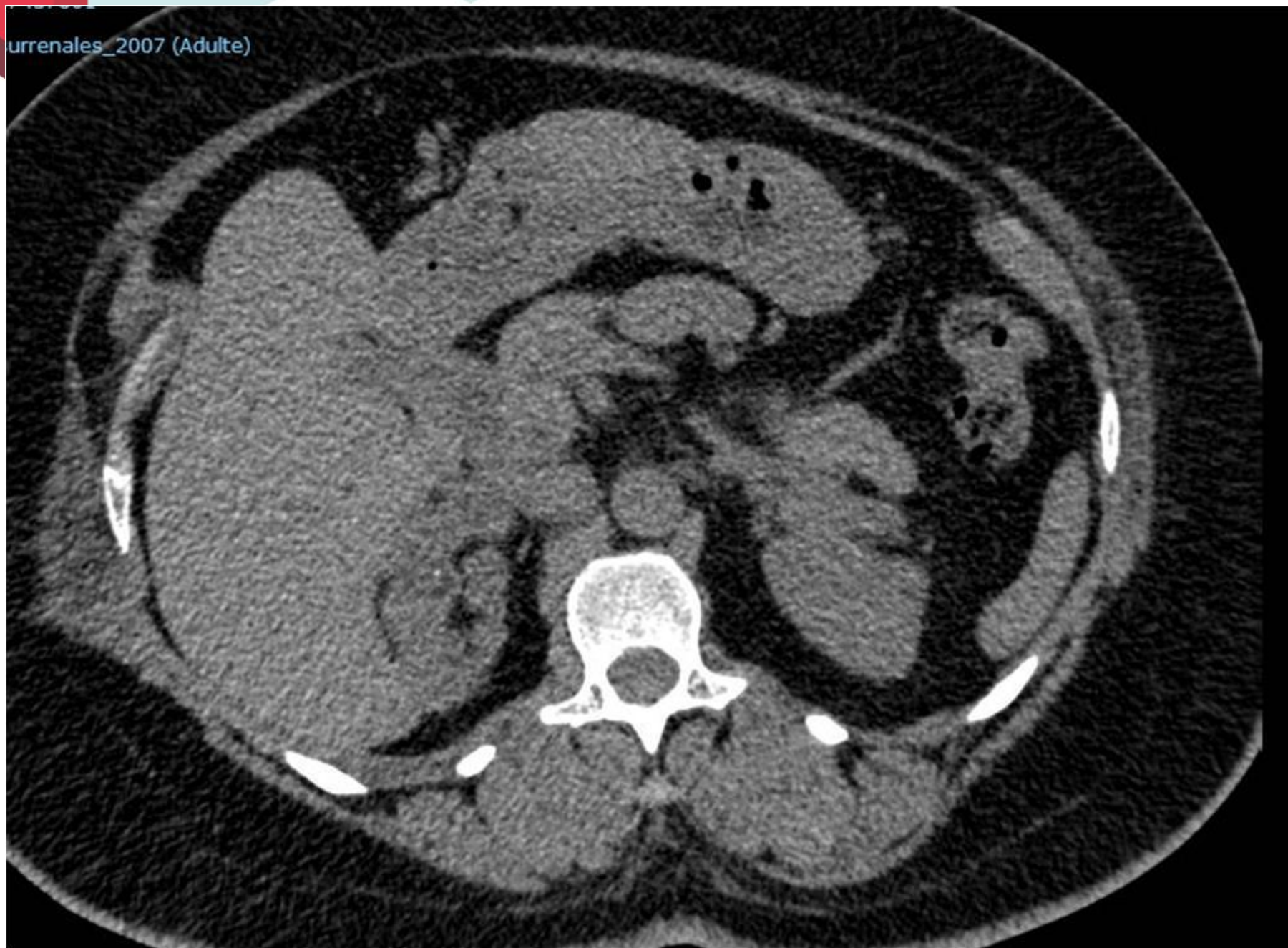
Symbol definitions

□	○	Normal 1 mg DST
■	●	Clinical CS
◼	◐	Abnormal 1 mg DST
◼	◑	Sub-clinical CS
□	○	NT
NT	NT	Not tested





urinaires_2007 (Adulte)



Adrenal hyperplasia / dysplasia gene hunt:

What are we looking for?

Different sources of DNA

- Somatic « mutations » (tumor suppressors only)
- Germline « mutations » in families
- Germline « mutations » in index cases

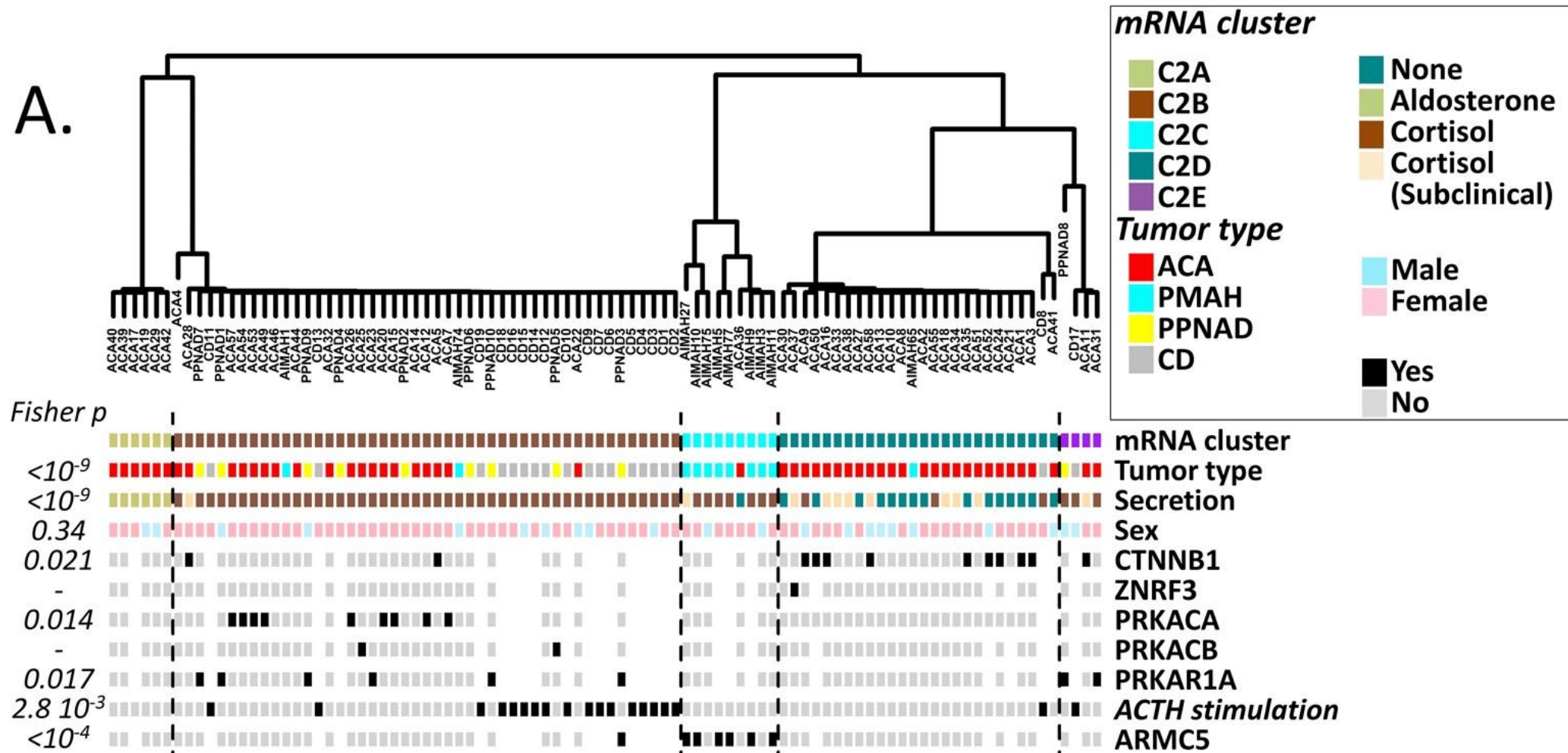
Different types of « mutations »

- DNA sequence variations *exome, WGS*
- DNA copy number variations *SNP arrays*
- Chromosome rearrangements neutral for CN and allelic ratios (inversion, translocations, mobile elements insertions) *WGS? Long range seq?*
- Epimutations *WGS?*

Transcriptome

4 main subtypes of benign tumors

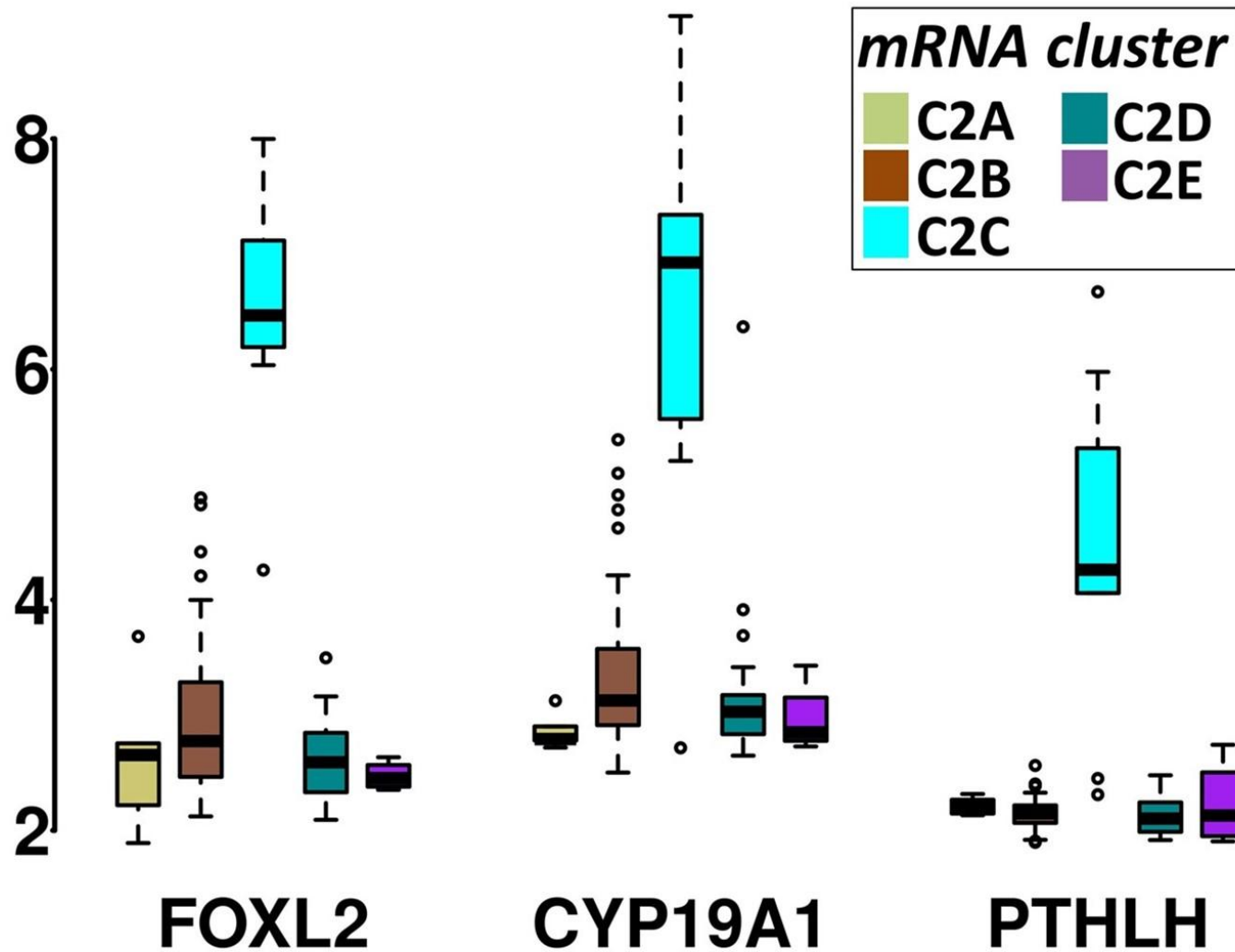
A.



Faillot et al, unpublished

ARMC5-mutated PMAH

A gonadal signature



Genes expressed in ovaries
Faillot et al, unpublished

Thank you !



guillaume.assie@aphp.fr

COMETE

Hervé Lefebvre
Antoine Martinez
Olivier Chabre
Eric Baudin
Anne Paule Gimenez-Roqueplo
Antoine Tabarin
Pierre François Plouin
Laurence Amar

**Cochin
Hospital**

Xavier Bertagna
Laurence Guignat
Olivia Barreau
Léopoldine Bricaire
Eric Clauser
Najiba Lalhou
Jean Guibourdenche
Mathilde Sibony
Bertrand Dousset
Paul Legmann
F Tenenbaum

Dpt of Genetics of Cochin

Michel Vidaud
Juliette Netoux
Kareen Leroy



ENS@T

Wiebke Arlt
Felix Beuschlein
Martin Fassnacht
Bruno Allolio
Marcus Quinkler
Massimmo Mannelli
Massimo Terzolo
Franco Mantero
Thomas Papatomas
Ronald De Krijger

Bioinformatics P.5 Univ.

Patrick Nietschke

Genomics Cochin Institute

Franck Letourneur team



Bioinformatics CIT

Aurélien de Reyniès
Eric Letouzé

GENOM'IC

