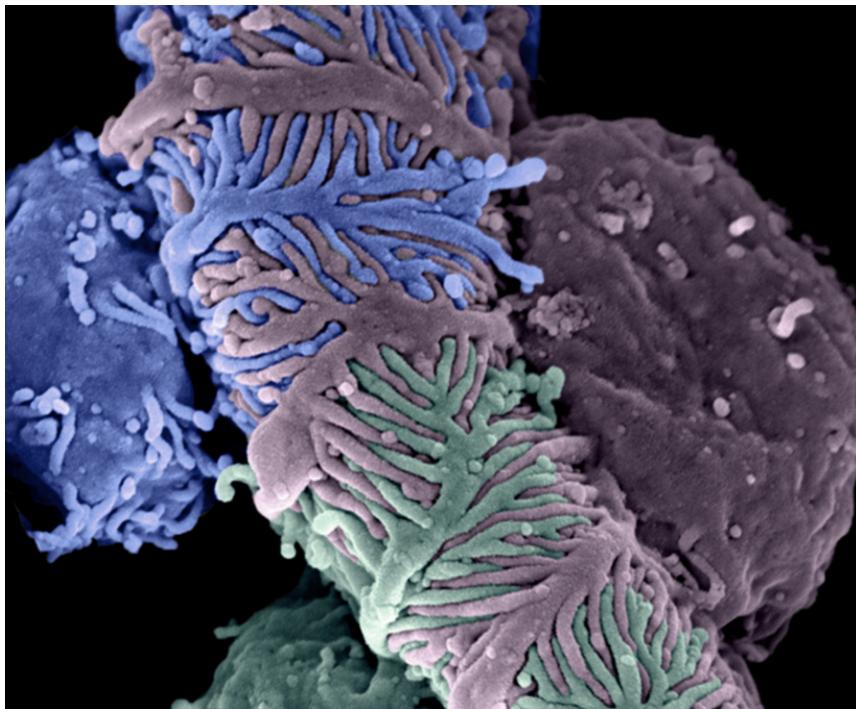


Monogenic podocytopathies in adults



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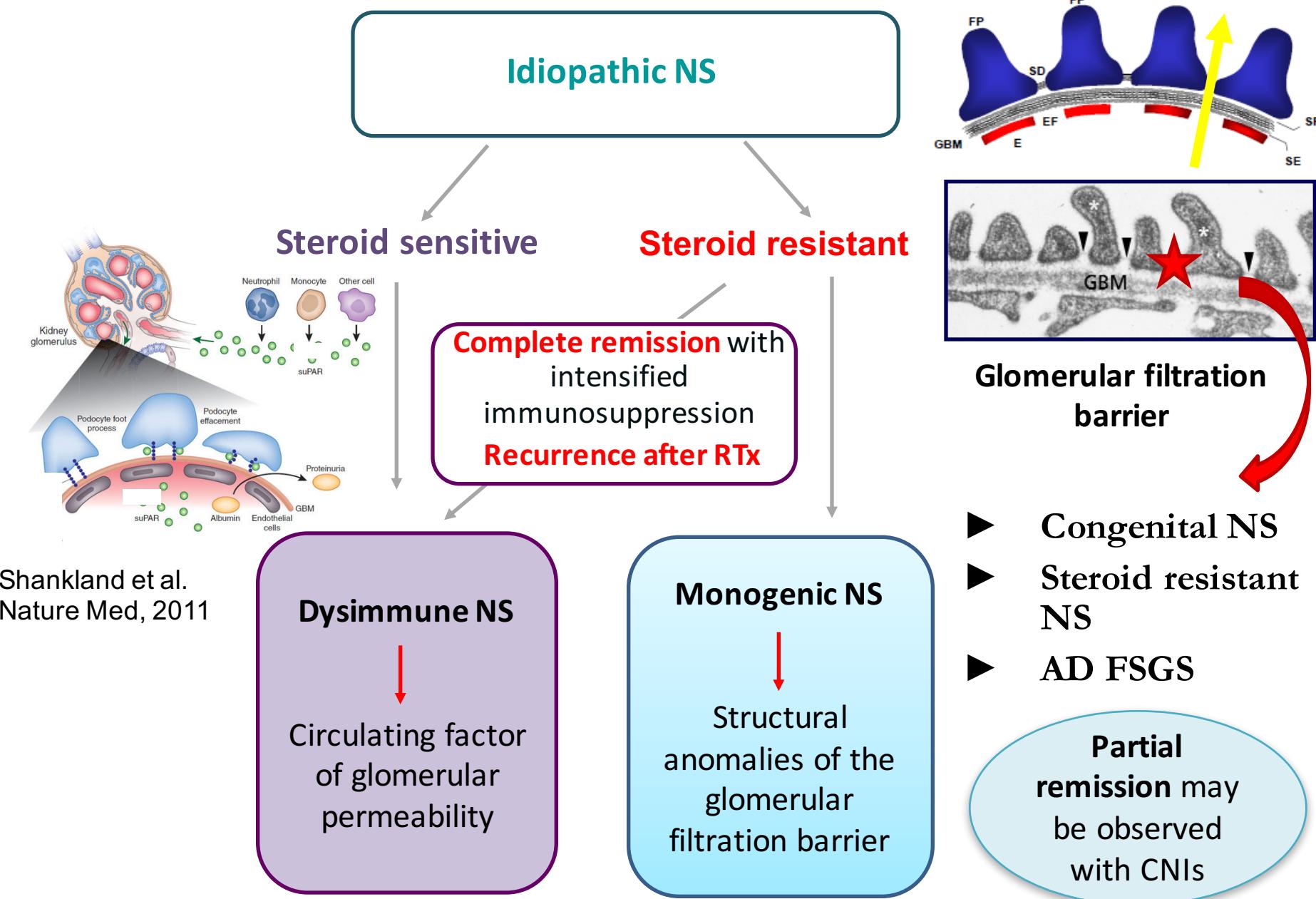


ORKiD FILIÈRE ORPHAN KIDNEY DISEASES

imagine
INSTITUT DES MALADIES GÉNÉTIQUES

UNIVERSITÉ
PARIS
DESCARTES

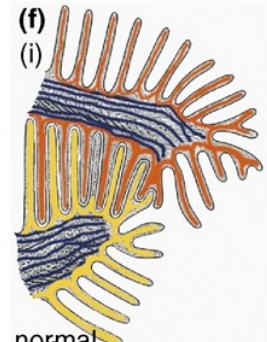
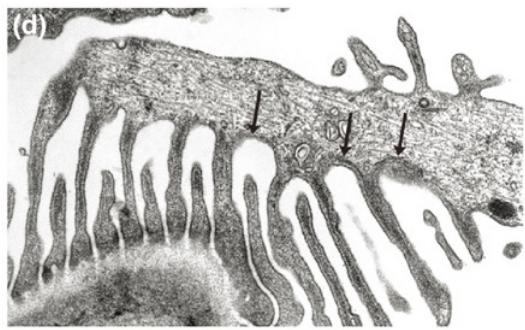
Different etiologies of nephrotic syndrome (NS)



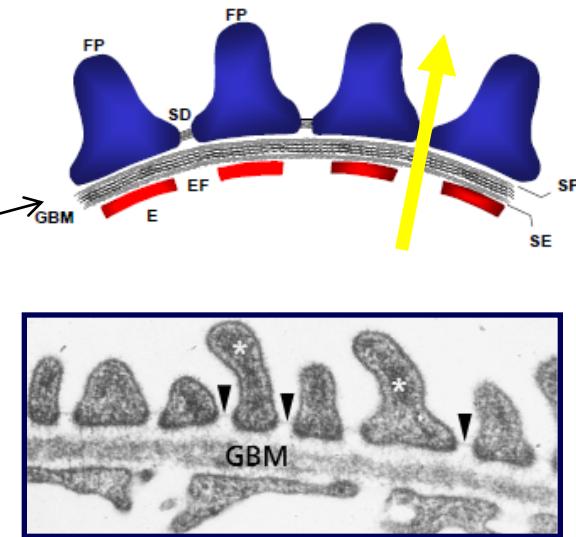
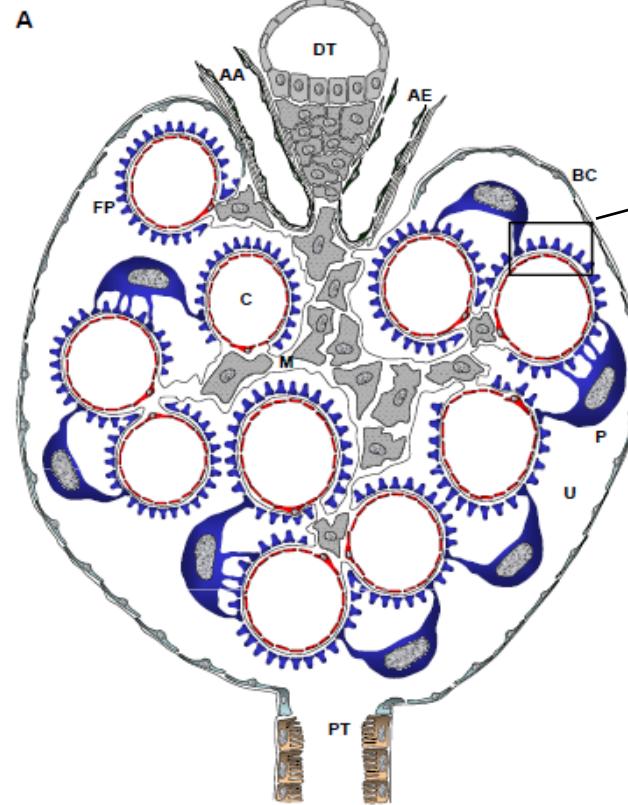
Podocyte is a fascinating cell



Octopus-like shape



Podocyte cytoskeleton



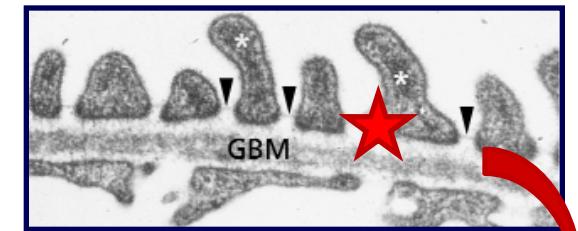
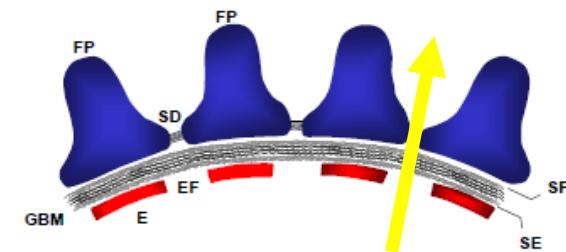
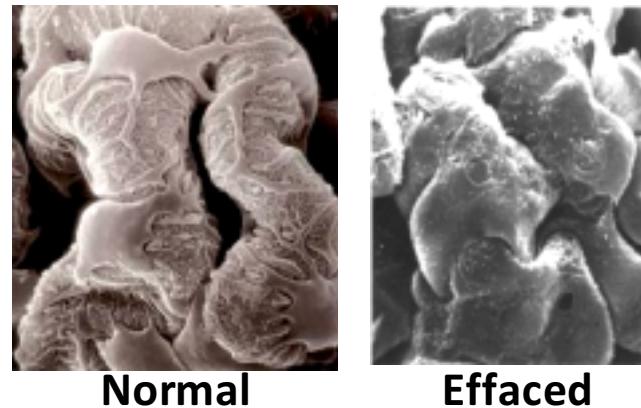
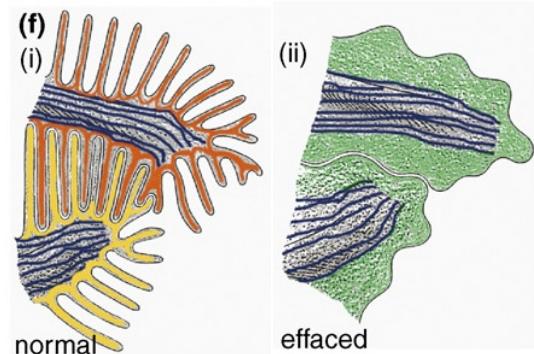
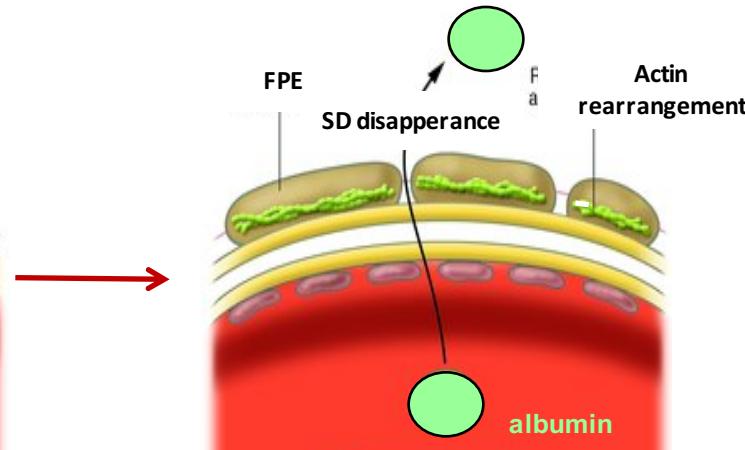
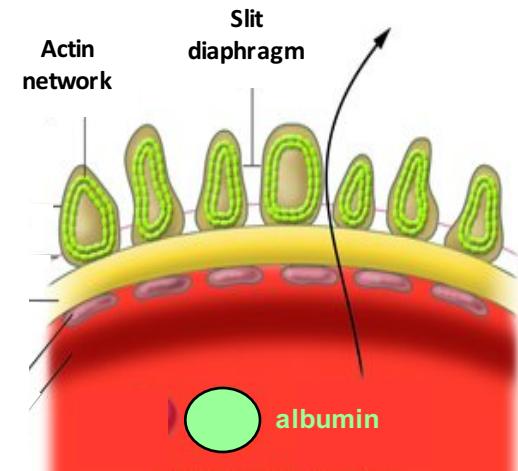
Glomerular Filtration Barrier

Hereditary podocytopathies

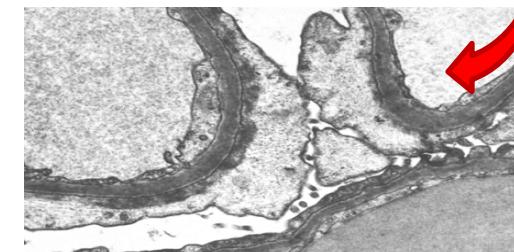
- Congenital NS
- Steroid-resistant NS
- Idiopathic FSGS

Machucca et al. HMG 2009
Faul, Trends Cell Biol 2007
Welsh, Nat Rev Nephrol 2012

Podocyte gene mutations result in actin rearrangement and foot process effacement



Glomerular Filtration Barrier



Foot Process Effacement

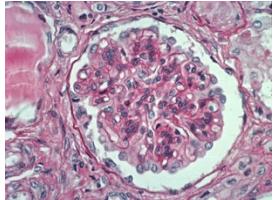
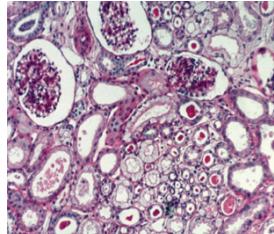
Machucca et al. HMG 2009
Faul, Trends Cell Biol 2007
Welsh, Nat Rev Nephrol 2012

Clinical forms of hereditary podocytopathies

Congenital & infantile NS

Antenatal onset – 1 yr
Enlarged hyperechoic kidneys
Increased α -FP
Enlarged placenta
Massive Pu, anasarca
→ ESKD

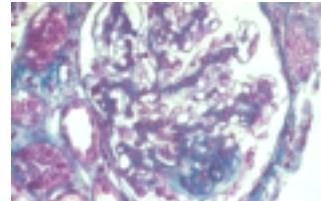
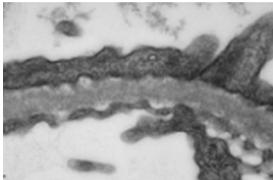
Tubular dilation (Finnish NS) or DMS or FSGS



Steroid-resistant NS

Childhood onset
Pu > 50 mg/kg/d
Albuminemia < 25-30 g/d
No remission with steroids and other IS (cyclosporine, MMF, ...)
→ **50% ESKD at 10 years**

MCD or FSGS

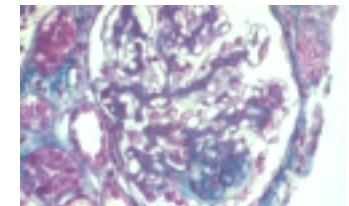


Glomerular Pu ± NS with FSGS

Late-onset (adolescent, adult)

Fortuitous Dx ++ MCD or FSGS

IS rarely used
Treatment with ACEi/ARBs



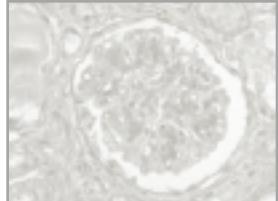
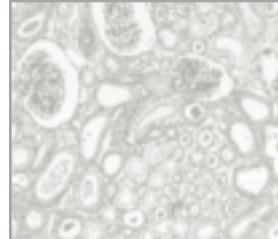
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→ ESKD

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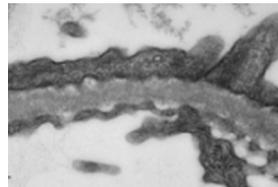


Steroid-resistant NS

Childhood onset
 $Pu > 50 \text{ mg/kg/d}$
Albuminemia $< 25\text{-}30 \text{ g/d}$

No remission with steroids and other IS (cyclosporine, MMF, ...)
→ 50% ESKD at 10 years

MCD or FSGS

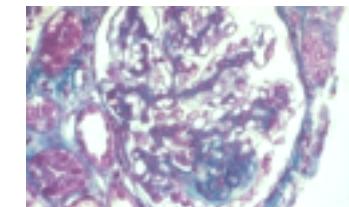


Glomerular Pu ± NS with FSGS

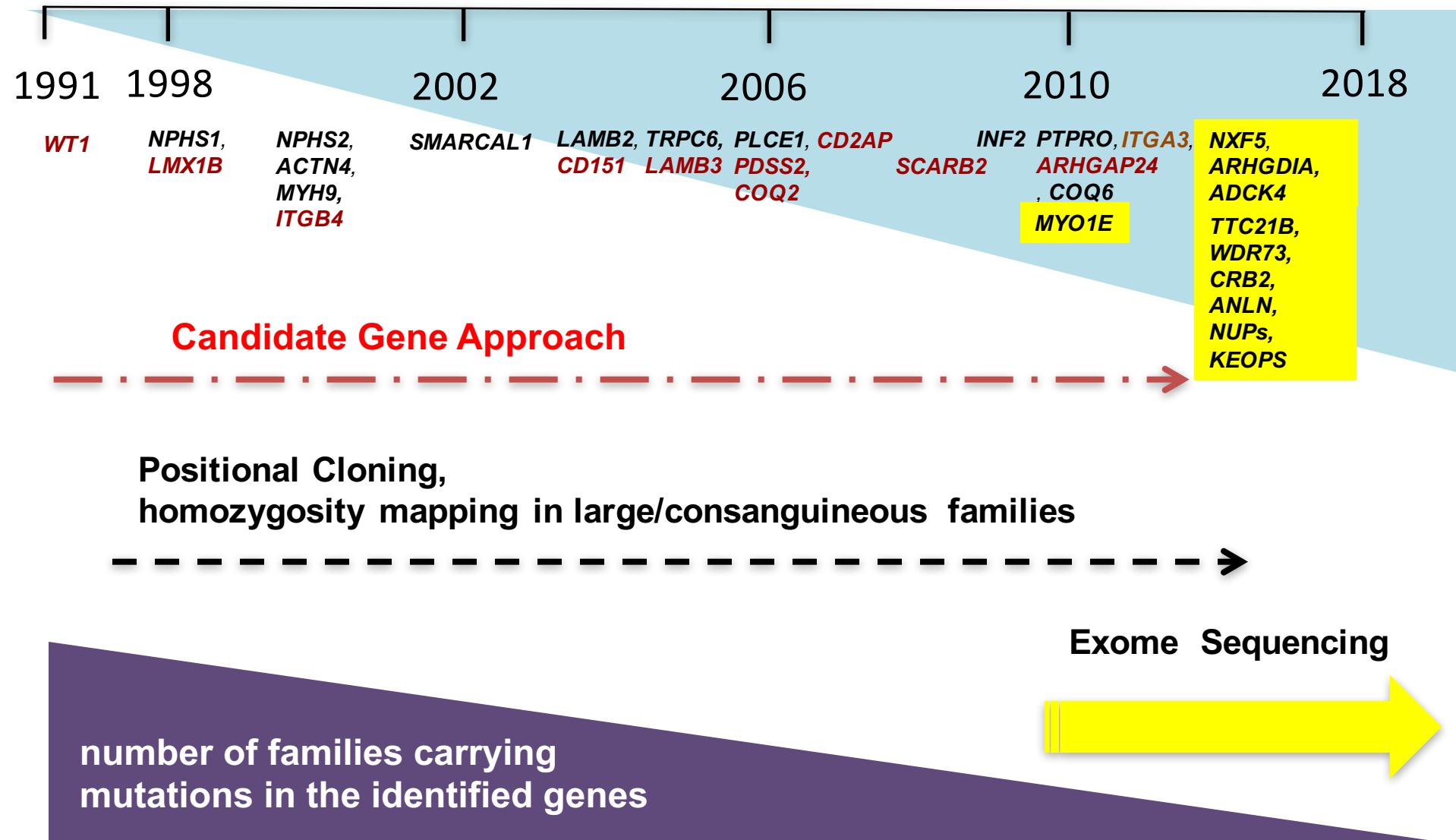
Late-onset (adolescent, adult)

Fortuitous Dx ++
K Bx: MCD or FSGS

IS rarely used
Treatment with ACEi/ARBs



Growing number of genes involved in podocytopathies



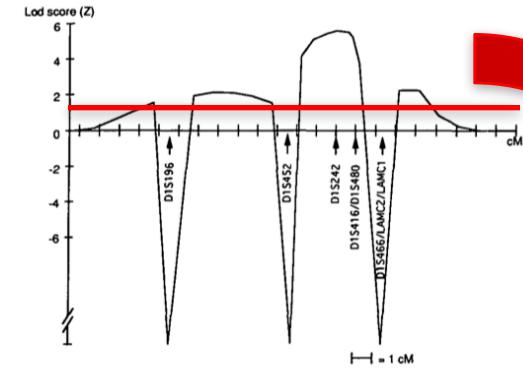
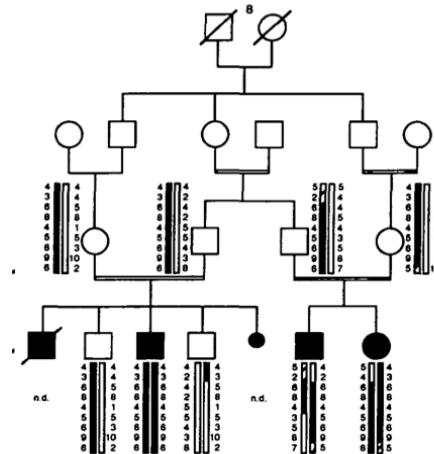
Input from positional cloning

Analyses the cotransmission of a morbide trait (NS/Pu) with microsatellite markers spread along the whole genome (linkage analyses, homozygosity mapping) → location in the genome

Sequencing of candidate genes at the locus
→ mutation

Direct identification of genes involved in hereditary diseases, which pathophysiology is unknown

Requires several informative families or large consanguineous families
Limited by incomplete penetrance



NPHS2 mutations (podocin) are the 1st cause of autosomal recessive steroid-resistant nephrotic syndrome

AR-SRNS

NS \approx 4 years, ESKD \approx 10 years
> 110 mutations
- 42% familial cases
- 10-30% sporadic cases

R138Q (32%): retained in the endoplasmic reticulum

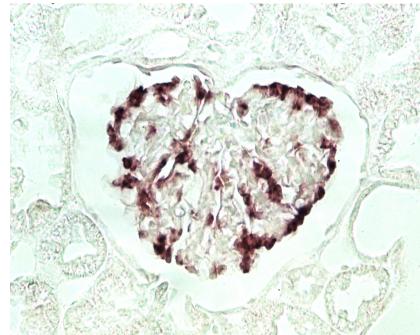
Age at Dx 2 years < 10 years for mutants located at the plasma membrane

Congenital NS

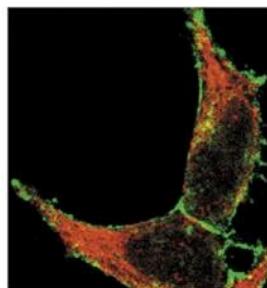
15-40 % of cases (ESKD \approx 6 years)

Late onset NS/Pu

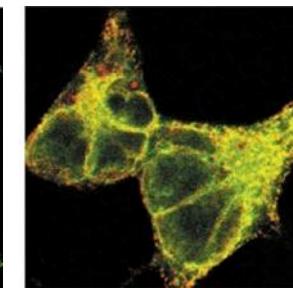
R229Q + 1 pathogenic mutation
19% of adult NS (Pu \approx 25 years,
ESKD \approx 32 years)



Expression in podocytes



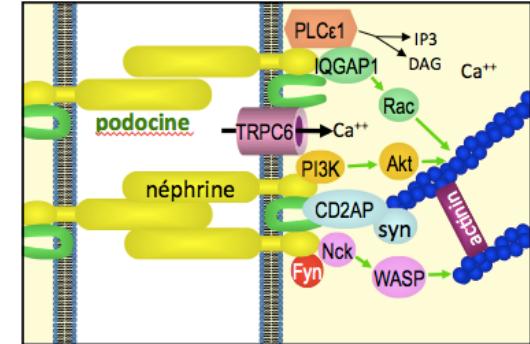
WT



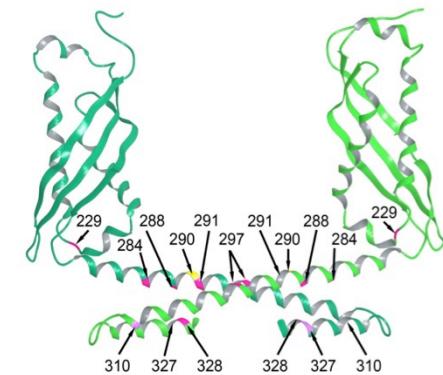
R138Q

 Podocine

 Calnexin (RE)



Slit diaphragm protein

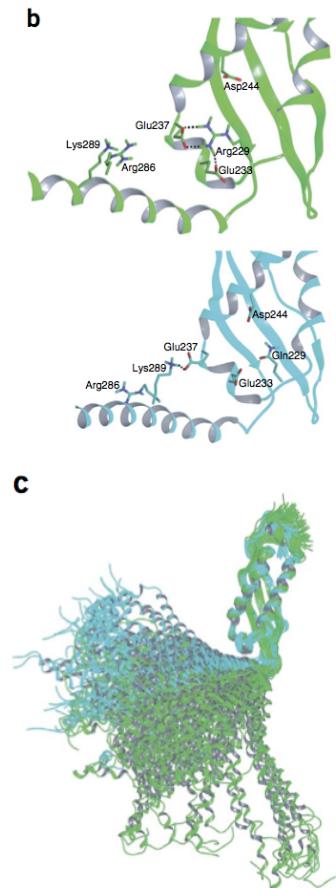


Homodimer

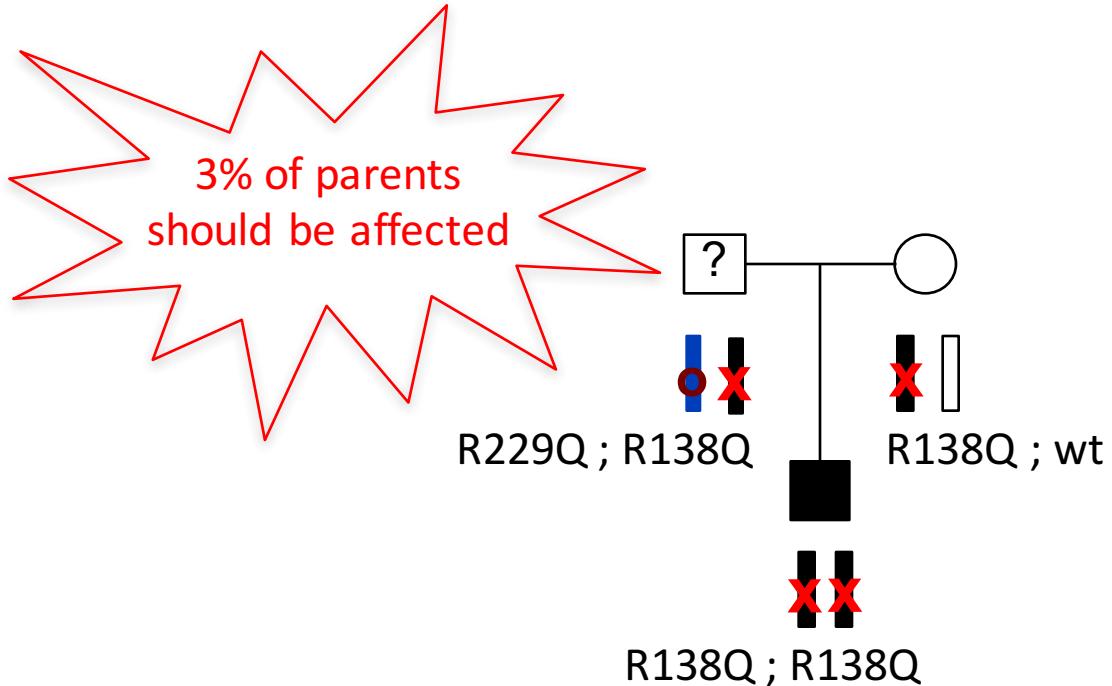
Adult patients carry the *NPHS2* polymorphism p.R229Q + 1 pathogenic mutation

Series of 105 cases / 96 families

- No adult patient with 2 pathogenic mutations
- 18 patients with 1 pathogenic mutation + p.R229Q
- P.R229Q: non neutral polymorphism (~ 3%) → reduced interaction with nephrin
- Age at Dx: 24.9 years (18.5 – 39 years)
- Age at ESKD: 32 years (27.7 - 38.5 years)



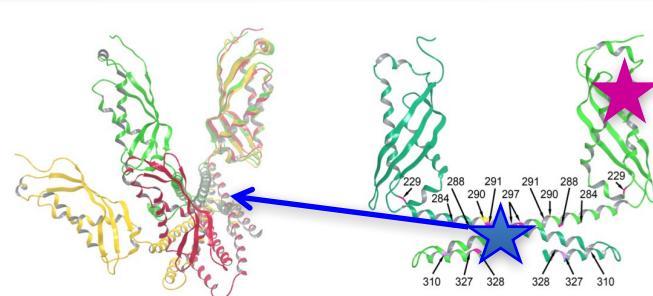
NPHS2 mutations and incomplete penetrance



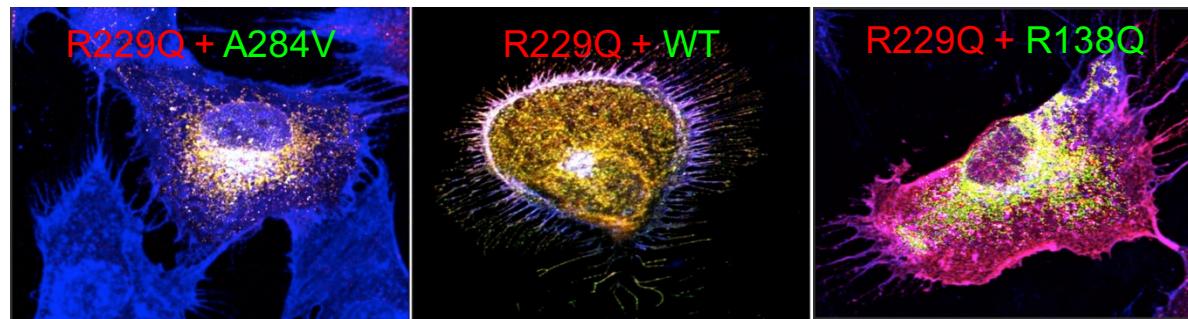
→ 6/129 parents of affected children are asymptomatic carriers of the association [p.R229Q];[mutation]

The effect of the p.R229Q *NPHS2* mutation depends on the 2nd mutation

Pathogenic dimers
[p.R229Q];[C-terminal mutation]



Non pathogenic dimers
[p.R229Q];[N-terminal mutation]



Podocin retained in the cell

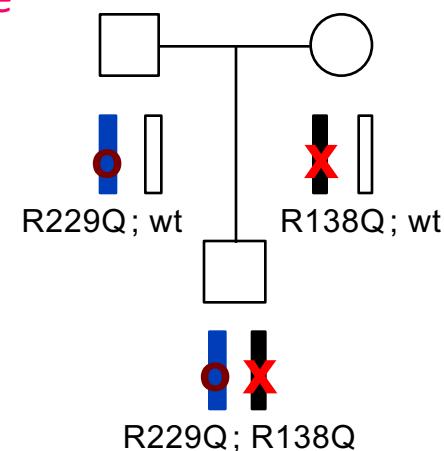
FSGS

Podocin at the plasma membrane

unaffected

→ [p.R229Q];[N-terminal mutation] DO NOT EXPLAIN NS => immune NS?

→ **Genetic counselling:** A couple carrying an *NPHS2* mutation in exons 1–6 in one member and R229Q in the other is not at risk of having an affected child



TRPC6 and *ACTN4* mutations are responsible for \approx 5 and 4% of AD FSGS

TRPC6

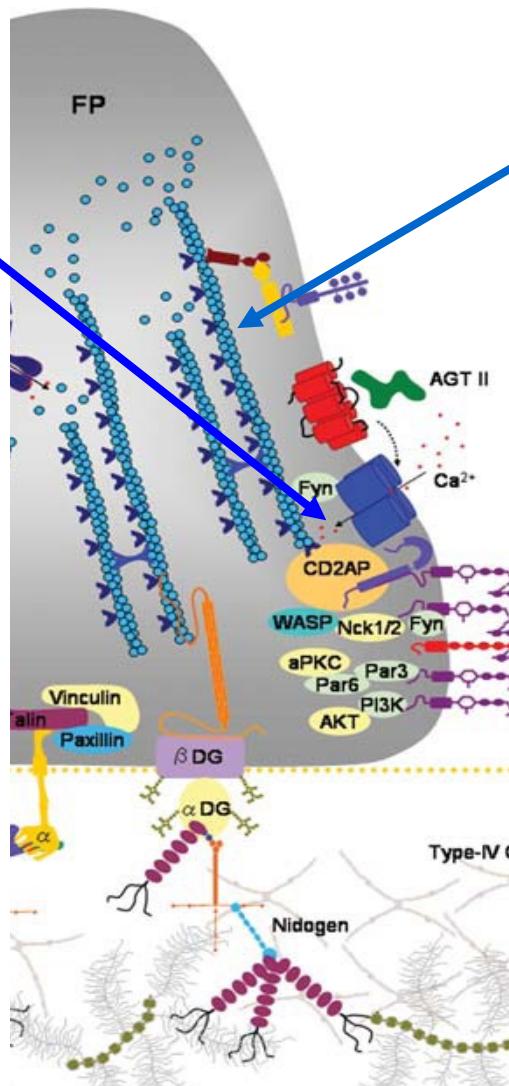
Calcium channel

Identified in a large family from New Zealand with late onset Pu and ESKD in 60% of patients

Mutations in **9/179 families (5%)**

↑ Intracellular Ca influx

Incomplete penetrance



ACTN4 (α -actinin 4)

Actin-bundling protein

Missense mutations in 3 large pedigrees with **AD FSGS**

5/141 familial FSGS (4%)

Proteinuria with FSGS (2nd decade), progressive CKD late-onset ESKD in some patients (~50 yrs)

Incomplete penetrance

Winn et al., Kidney Int 1999

Winn et al., Science 2005

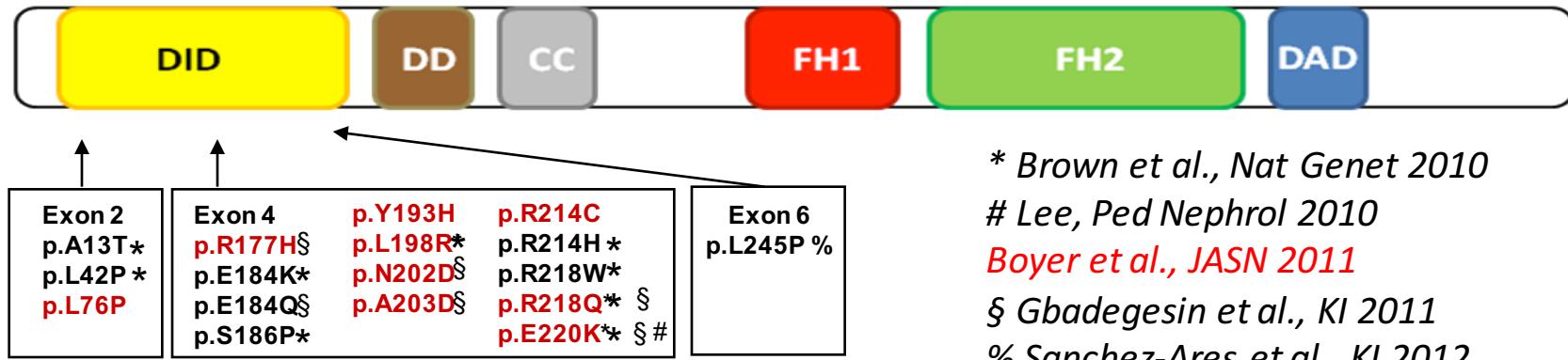
Reiser et al., 2005

Kaplan et al., Nat Genet 2000

Weins et al., JASN 2005

Weins et al., PNAS 2007

Mutations in *INF2* are the main causes of AD FSGS



* Brown et al., Nat Genet 2010

Lee, Ped Nephrol 2010

Boyer et al., JASN 2011

§ Gbadegesin et al., KI 2011

% Sanchez-Ares et al., KI 2012

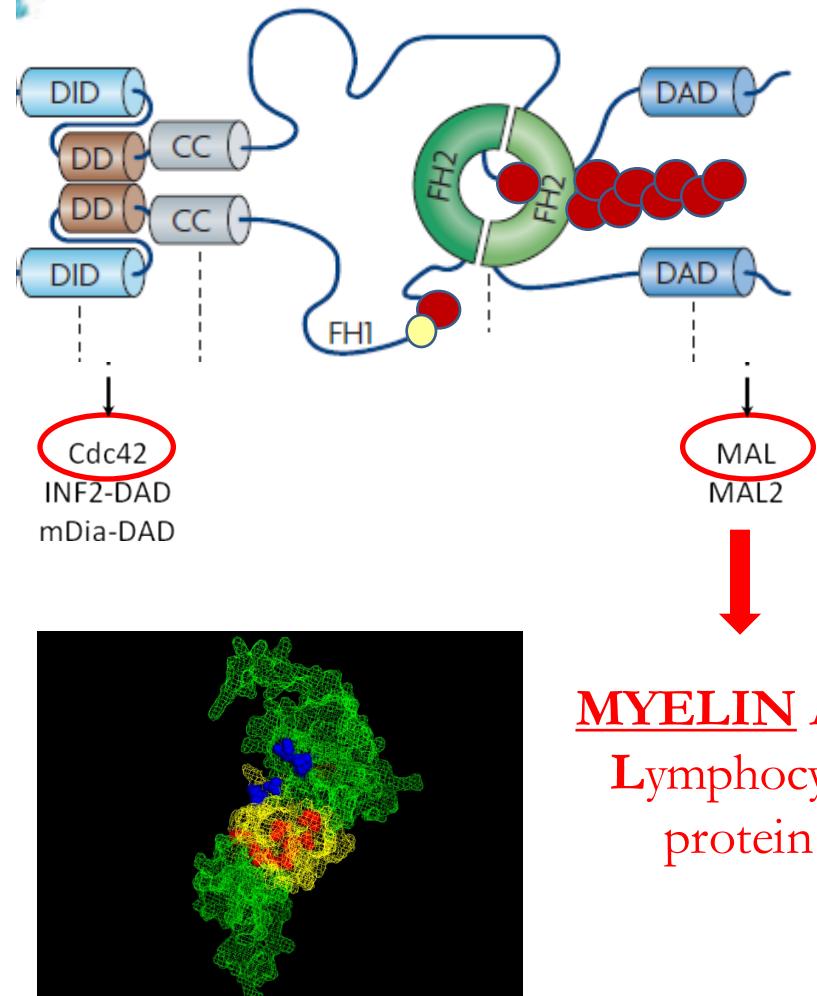
Reference	# tested families	# mutated families	% mutations	Age at onset Pu	Age at ESKD
Brown 2010	215	20	9%	11-72 yrs	13-67 yrs
Barua 2012					
Lee 2010	9	1	11%	7-30 yrs	14 yrs
Boyer 2011	54	9	17%	5-44 yrs	20-70 yrs
Gbadegesin 2011	49	8	16%	14-46 yrs	33-45.5 yrs
TOTAL	327	38	12%	5-72 yrs	13-70 yrs

Interindividual variability & incomplete penetrance

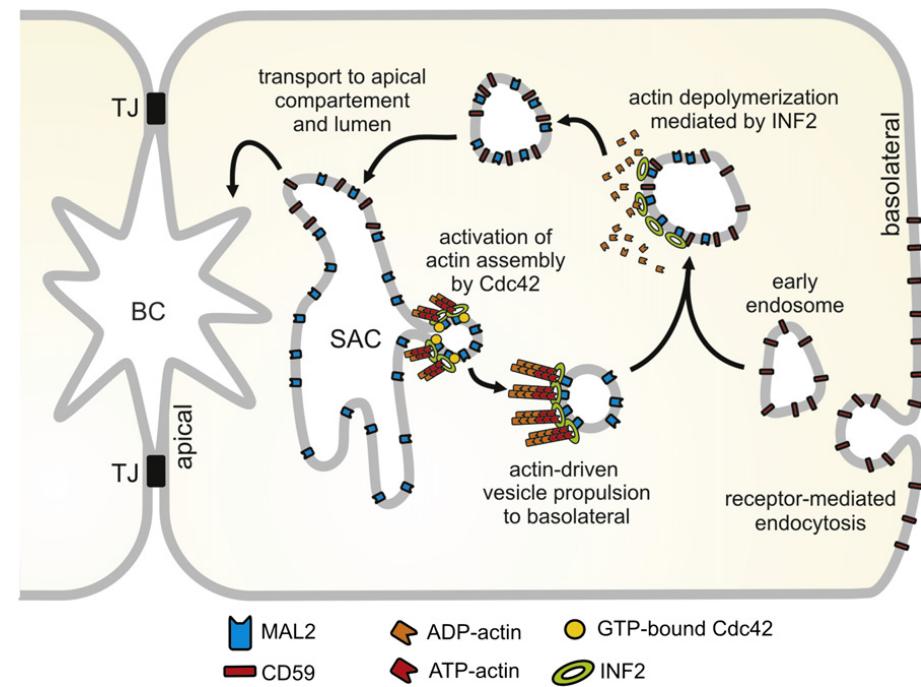
3/405 = 0.8% of sporadic cases

Brown E ... Pollak M. Nat Genet 10
Boyer O, ... Antignac C. JASN 11

Mutations in the INF2 DID alter the localisation of the protein and actin polymerization



- Enhances actin polymerization
- Promotes actin depolymerization and filament sievering
- **Role in intracellular protein transport in association with Cdc42 and MAL**



Madrid, Dev Cell 2010

Andres-Delgado, Blood 2010

Bown E ... Pollak M. Nat Genet 10

Input from candidate gene approaches



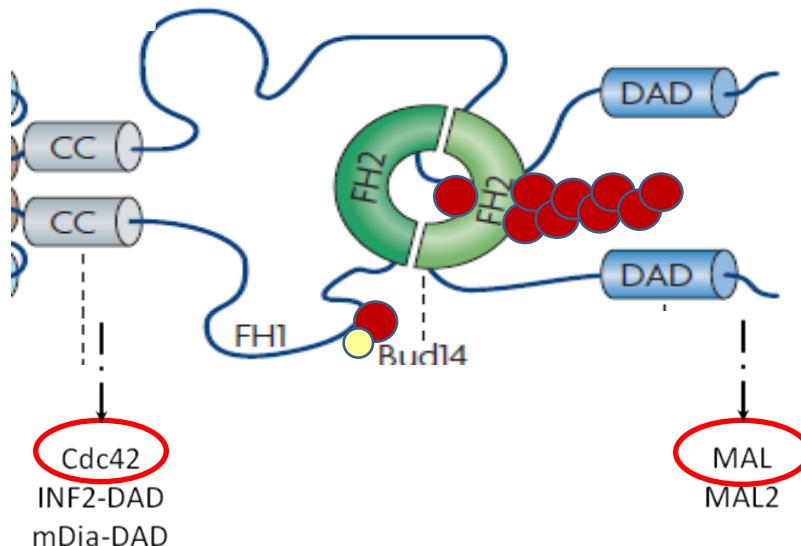
Amyotrophy
Steppage gait



Prevalence: 1/2500



Reduced deep tendon reflexes



CMT + FSGS

13 reported cases
Age Pu 1-28 yrs
Nephrotic syndrome 3/11
ESKD after 0.5-15 yrs



Common genetics?



INF2 partners Cdc42 and MAL are crucial for myelination

INF2 = good candidate?

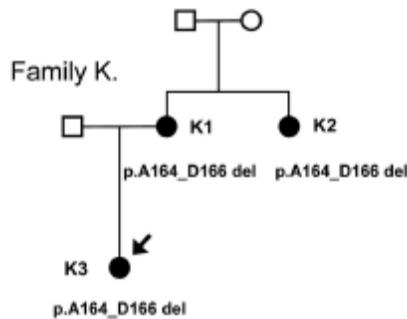
Lemieux et al., Can Med Assoc J 1967

Mutations in *INF2* are the main causes of FSGS associated with Charcot Marie Tooth disease

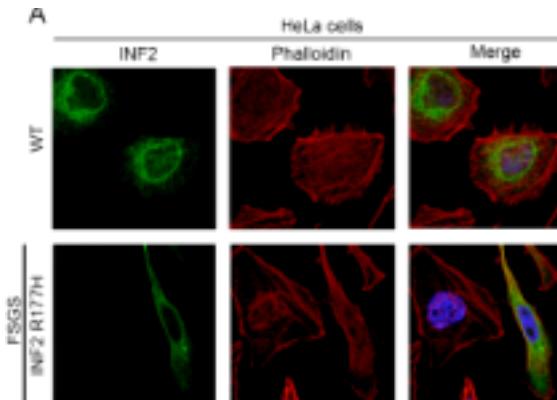
Cohort of 16 unrelated families with FSGS-CMT :

→ mutations in 75% cases

No mutation in 150 patients with CMT and no renal involvement

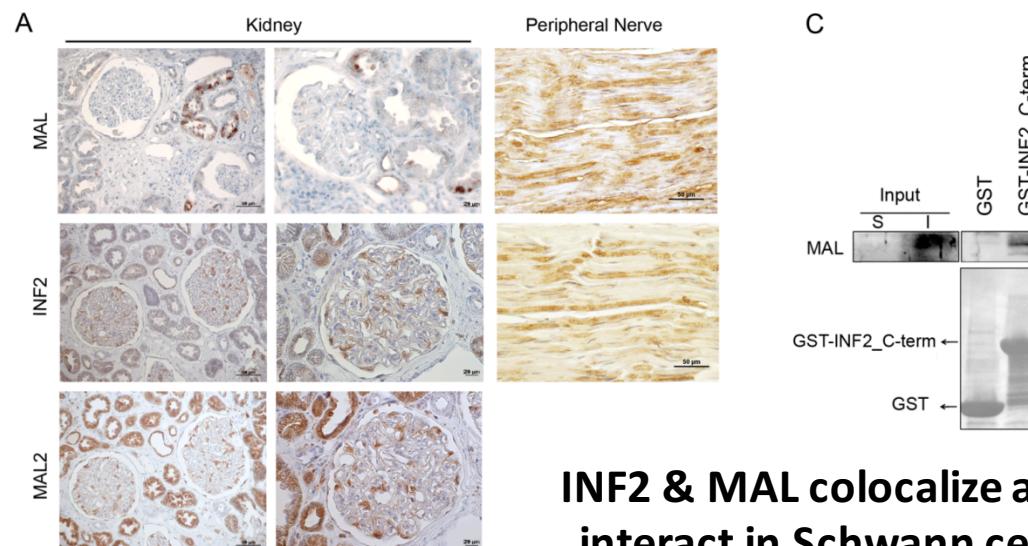


Dominant inheritance



Mutants alter the INF2-mediated polymerization & depolymerization

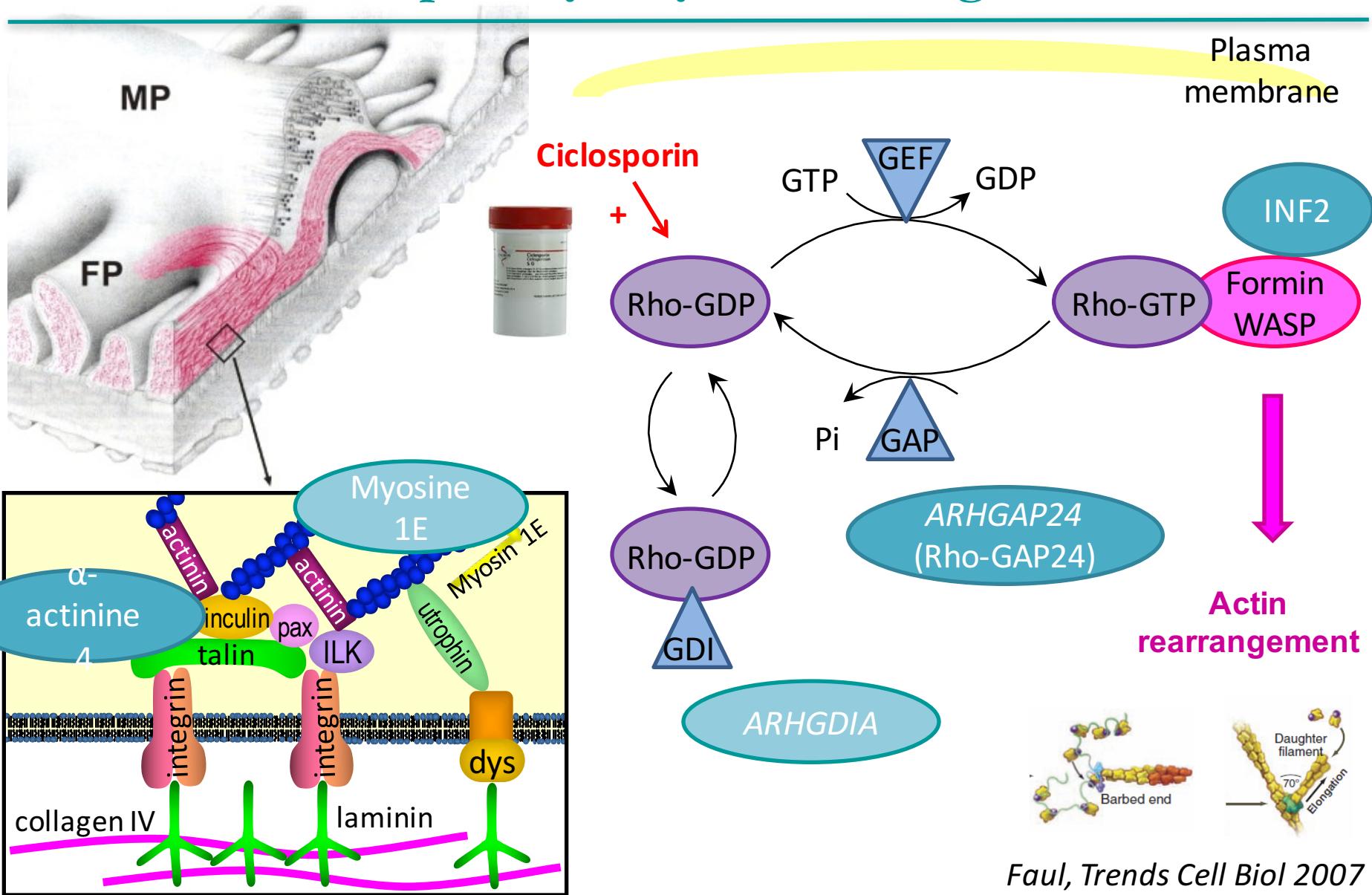
CMT : 13 years (5-28 years), 4 with deafness
Proteinuria : 18 years (10-21 years)
ESKD: 21 years (12-47 years)



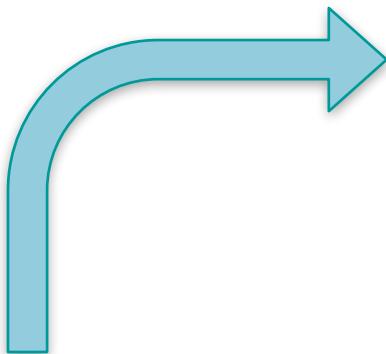
INF2 & MAL colocalize and interact in Schwann cells

Input from candidate gene approaches

Other podocyte cytoskeleton genes



Input from high throughput sequencing



XXXXX
variants



Filters



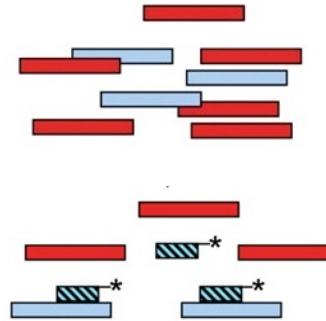
Candidate genes
Segregation analysis
Functional studies



Disease-causing
variant

Exome sequencing

Genomic DNA

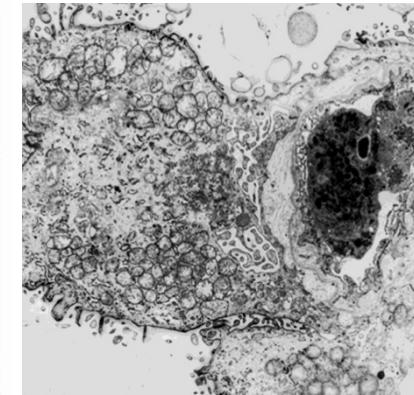
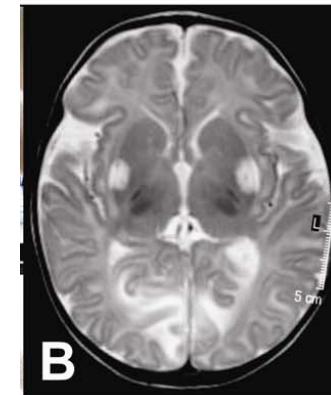
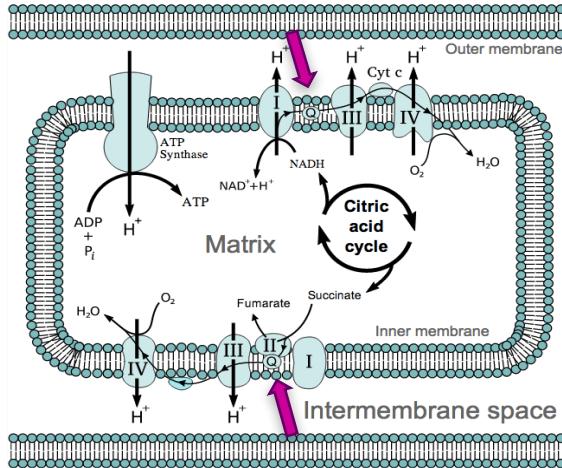


Exome
capture

Sequencing



1) Identification of new genes: *ADCK4* involved in coenzyme Q₁₀ (CoQ₁₀) biosynthesis

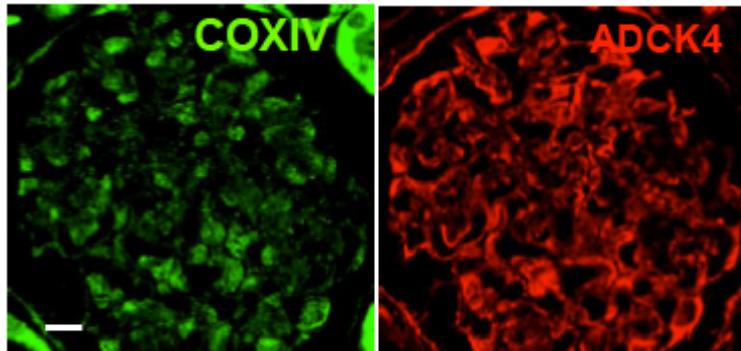


Gene	Kidney phenotype	Extra-renal features	COQ10 effects
<i>COQ2</i>	CNS, SRNS (FSGS)	Encephalopathy, MOF	?
<i>PDSS2</i>	Congenital/infantile NS	Deafness, MR, Leigh	?
<i>COQ6</i>	CNS, SRNS (FSGS/DMS)	± Deafness, seizures, ataxia	↓ Pu ± deafness
<i>ADCK4</i>	SN infantile, SNCR (HSF)	1 heart disease, 1 MR/15	↓ Pu

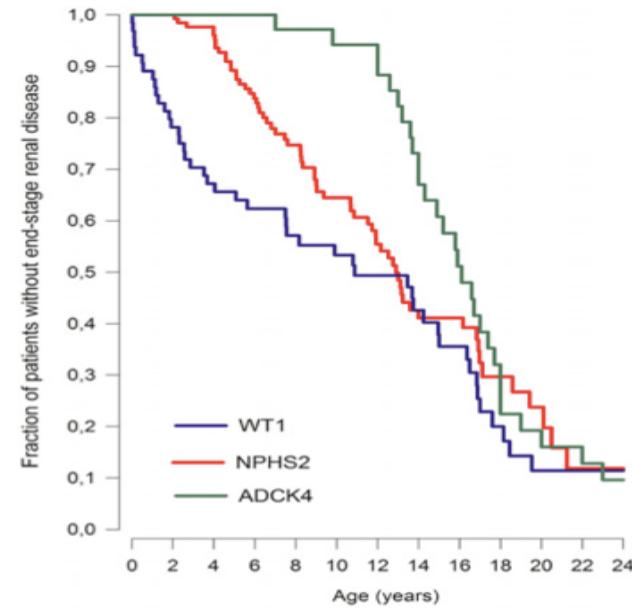


1) Identification of new genes:

***ADCK4 (COQ8B)* involved in coenzyme Q₁₀ (CoQ₁₀) biosynthesis**

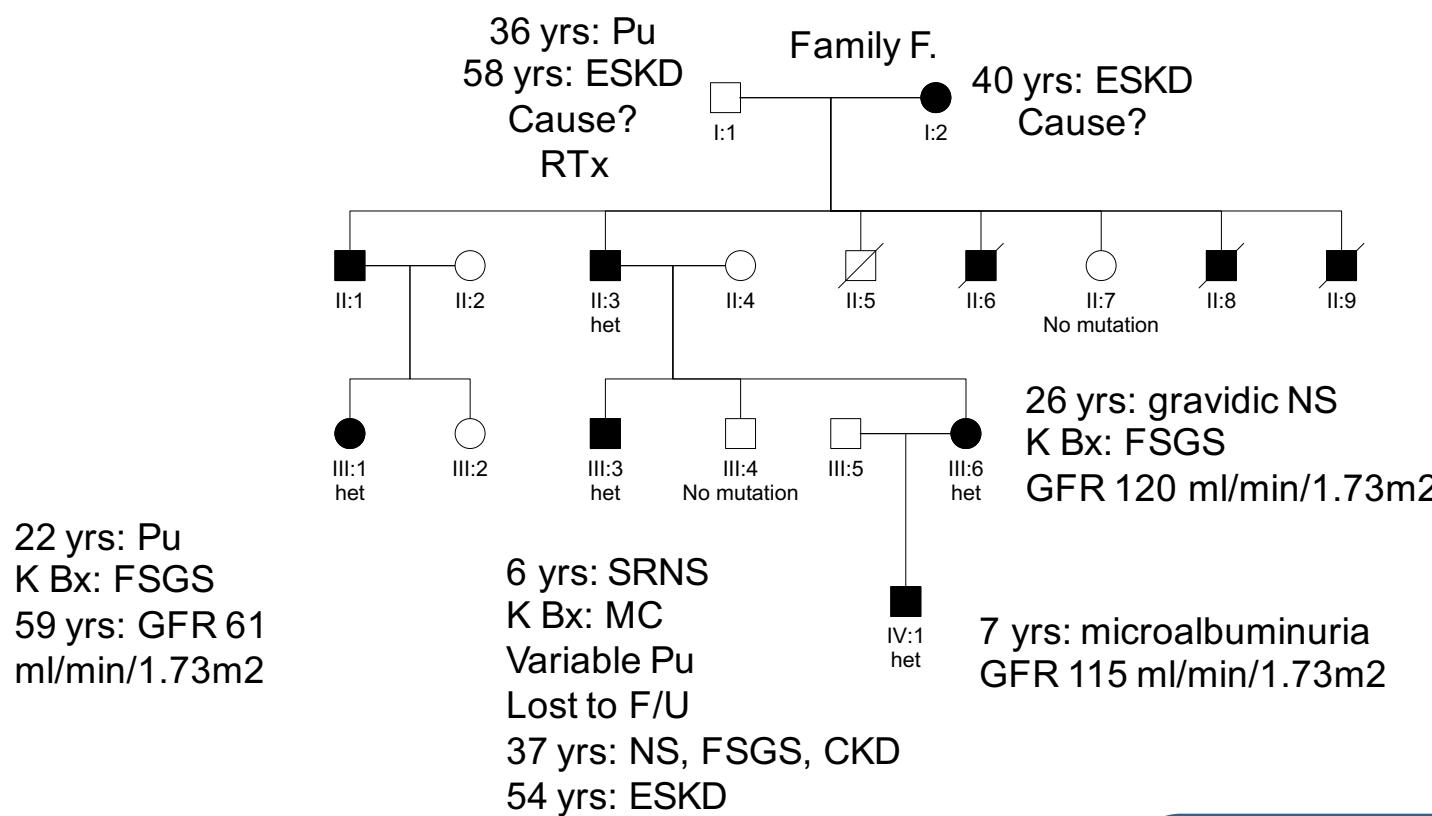


- Identification of *ADCK4* mutations in 15 patients (8 families) with SRNS by linkage analyses and exome sequencing
- Expression in podocytes (mitochondria and foot processes)
- ↓ level of CoQ₁₀ in lymphocytes or fibroblasts
- ↓ cell migration by *ADCK4* knockdown, reversed by CoQ₁₀
- ↓ proteinuria in 1 treated patient

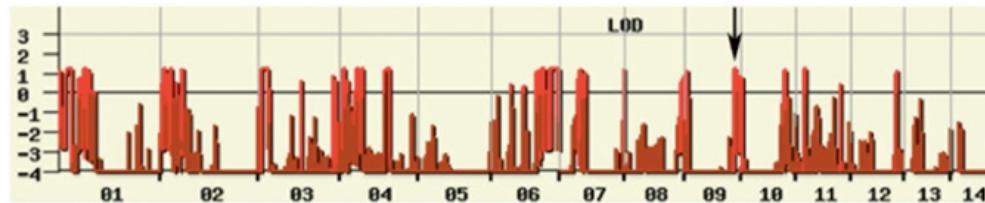


- 26 patients from 12 families
- median age, 14.1 years (11-17)
- nephrotic-range proteinuria in 44%
- advanced CKD in 46% of patients at time of diagnosis
- Age at start RRT 14-18 years
- Mostly isolated FSGS

2) Mutations in “syndromic NS genes” in patients with isolated glomerular diseases



**10 affected
4 generations
5 with DNA
No extra-renal
involvement**

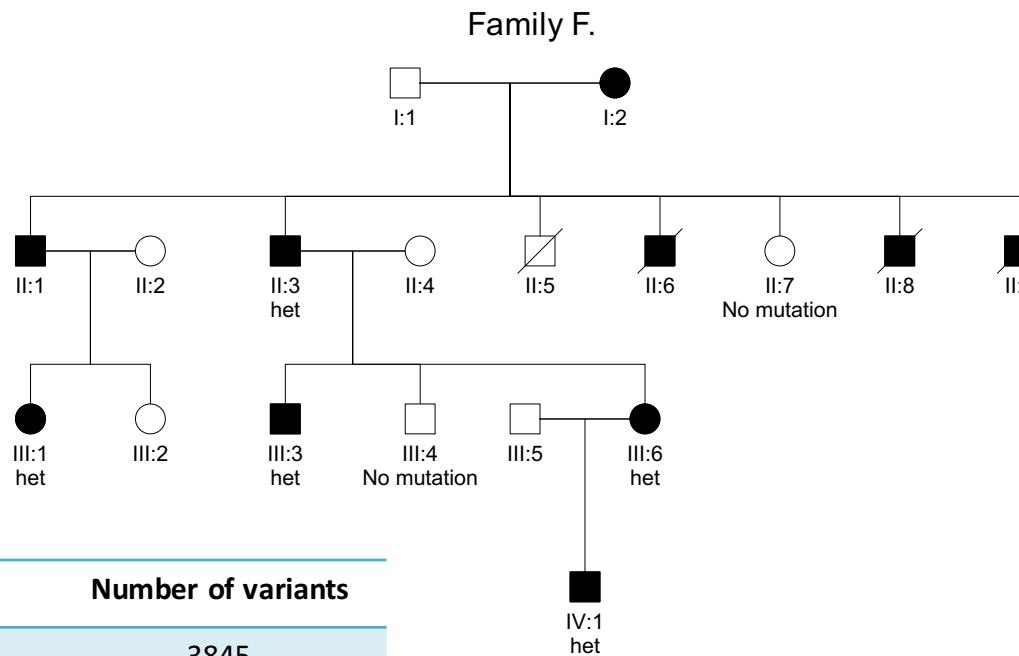


Linkage analyses:

Large region of interest
118 Mb on chromosomes
1,2,3,4,6,9,11,15,17,19
1000 genes

2) Mutations in “syndromic NS genes” in patients with isolated glomerular diseases

3845 variants



Needle in a haystack

***LMX1B* Mutation**

p.R246Q segregating in the family

p.R246Q and p.R246P in 2 others

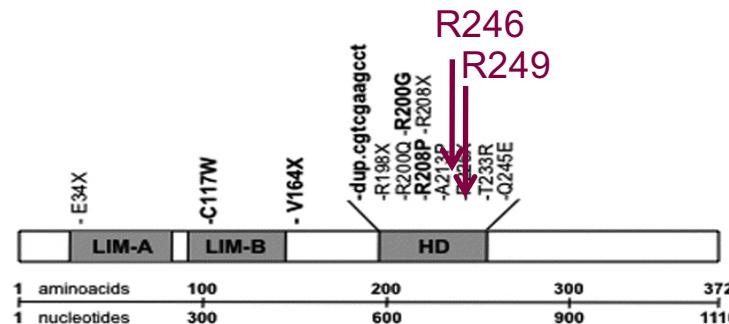


2) Mutations in “syndromic NS genes” in patients with isolated glomerular diseases: *LMX1B* mutations without Nail Patella

Nail-Patella syndrome

Developmental defects of dorsal limb structures, kidney and eye, with **nail dysplasia**, **patellar abnormalities**, elbow dysplasia, **iliac horns**, nephropathy (2-65%) and glaucoma

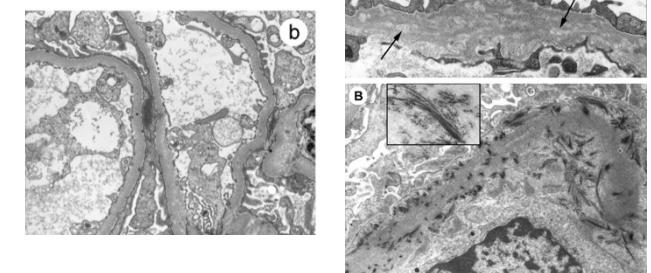
- **No extra-renal disorder** recorded in the patients
- **No typical EM lesions of the glomerular basement membrane** in the biopsy sample



Patients
Nail Patella

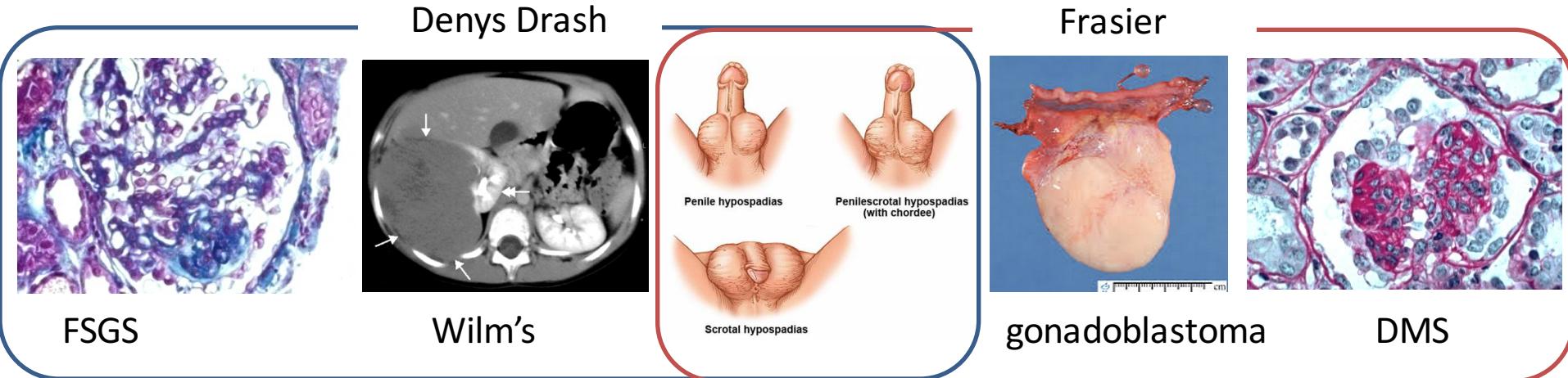


Patients

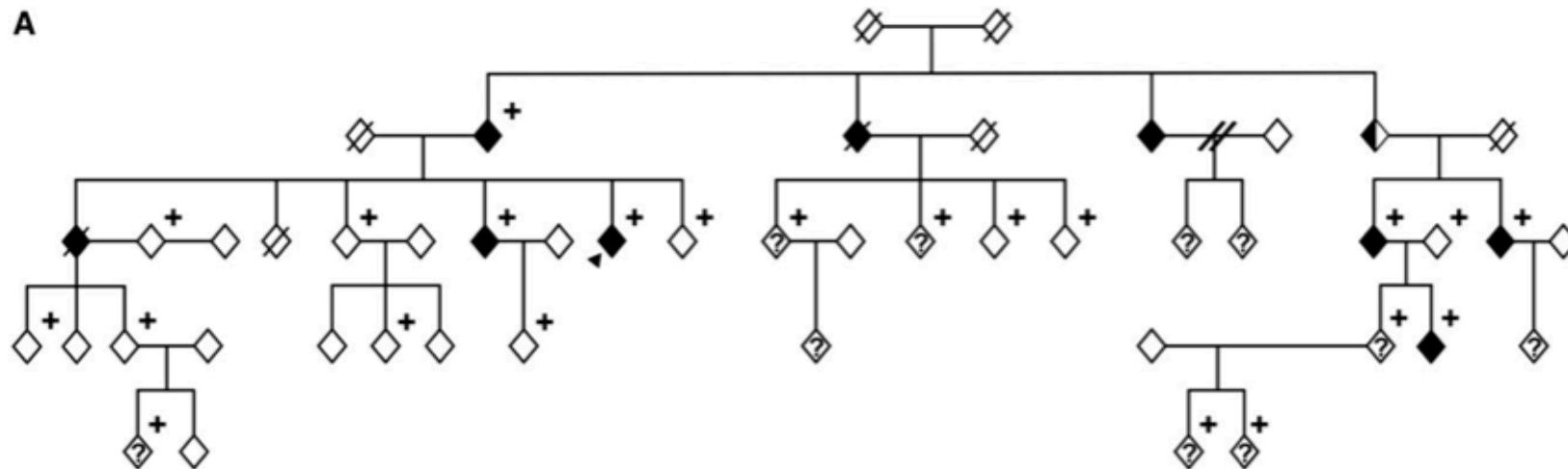


Patients
Nail Patella

2) Mutations in “syndromic NS genes” in patients with isolated FSGS: *WT1* mutations without Denys Drash or Frasier Sd



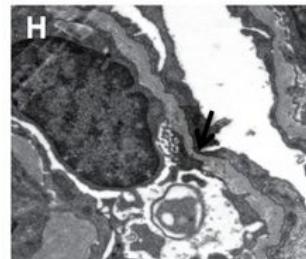
Identification of a *WT1* mutation segregating with the disease in 2 large families (with at least 5 males) with FSGS (diagnosed between 16-30 yrs)



3) Mutations in genes involved in other kidney diseases: *COL4A3-5* and *PAX2* gene

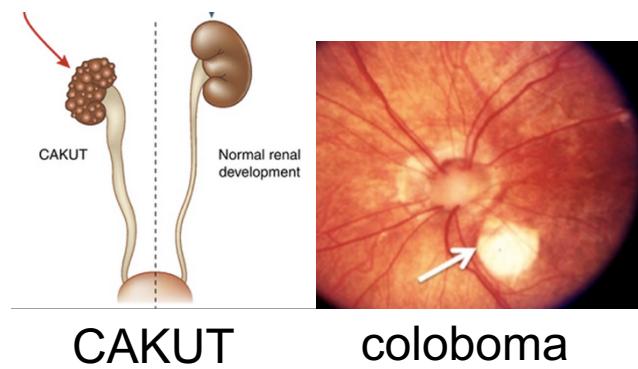
COL4A3-5 mutations

- ⇒ *COL4A3* and *COL4A4* mutations: AD and AR
Alport Sd and Familial Benign hematuria
- ⇒ *COL4A3* heterozygous mutations in **~10% families with AD FSGS – no deafness before molecular testing**
 - Some missing clinical data (Hu? EM?)
 - Some patients clearly have isolated proteinuria without hematuria or any GBM changes



PAX2 mutations

- ⇒ CAKUT and papillo-renal syndrome
- ⇒ Identified in 4% of adult-onset familial FSGS and no extra-renal features



Malone et al., *KI* 2014
Xie et al., *J Mol Cell Biol* 2014

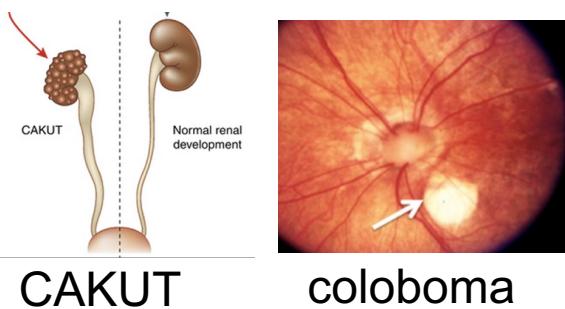
Barua et al, *JASN* 2014



Include *COL4A3-5* & *PAX2* genes in FSGS genetic testing

PAX2 (CAKUT)

- ⇒ CAKUT and papillo-renal syndrome
- ⇒ in 4% of adult-onset familial FSGS and no extra-renal features



Barua et al, JASN 2014

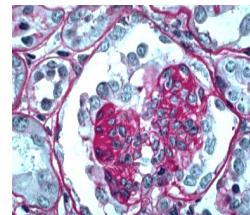
Tubulopathies

- CBN (cubilin) (Megaloblastic anemia)
- OCRL et CLCN5 (Dent)
- => Chronic Pu and FSGS
- => ↓ albumin reabsorption in the DCT
- ⇒ Secondary FSGS? Intrinsic role ?

Ovunc et al, JASN 2011

WT1 (syndromic NS)

Denys Drash

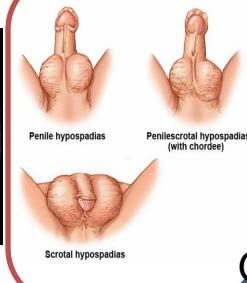


DMS



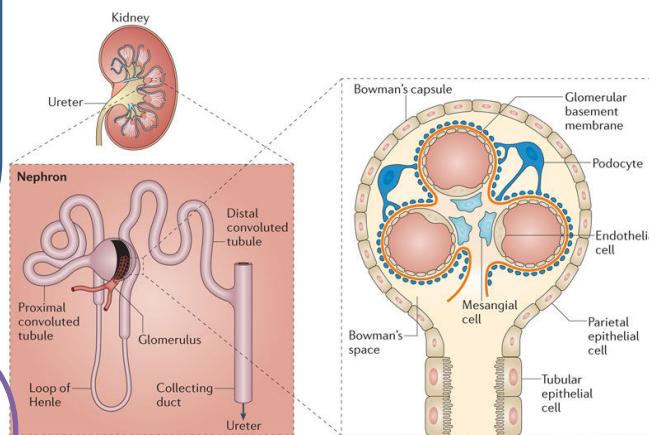
Wilms

Frasier



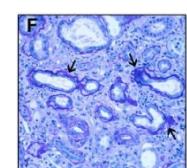
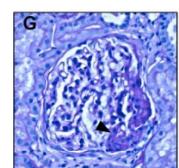
Gonadoblastoma FSGS

Hall G et al., JASN 2015

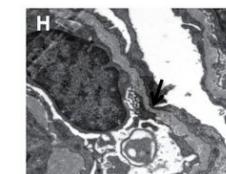


TTC21B (ciliopathy)

- Pu (1-7 g/l) et HBP ++
- ESKD 22 yrs
- ⇒ FSGS and nephronophthisis
- Magrheb and Portugal



Huynh Cong E et al., JASN 2014



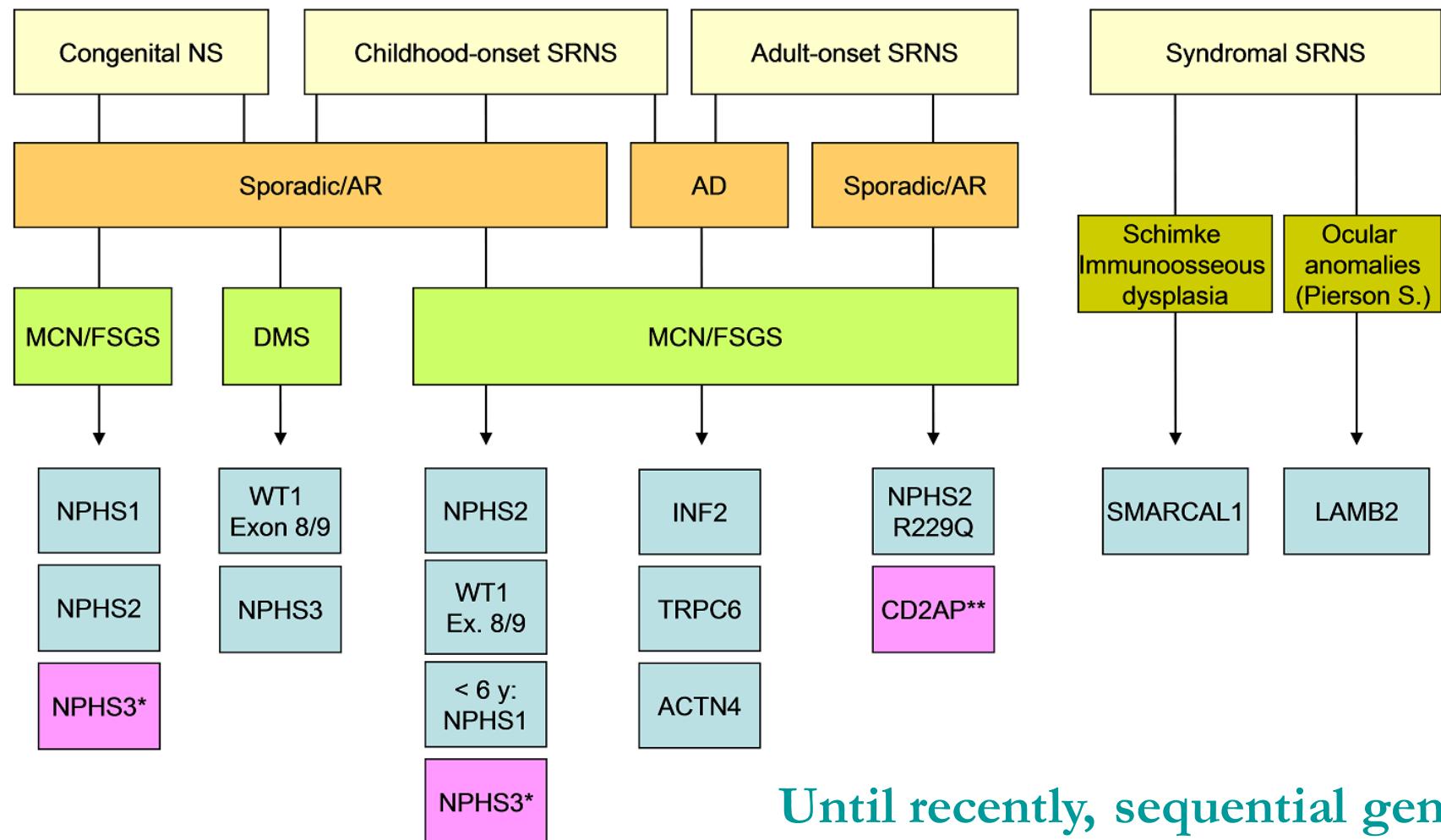
Malone et al., KI 2014

COL4A3-5 (Alport)

- ⇒ Heterozygous COL4A3 mutation in ~10% AD-FSGS families
- No deafness before genetic testing

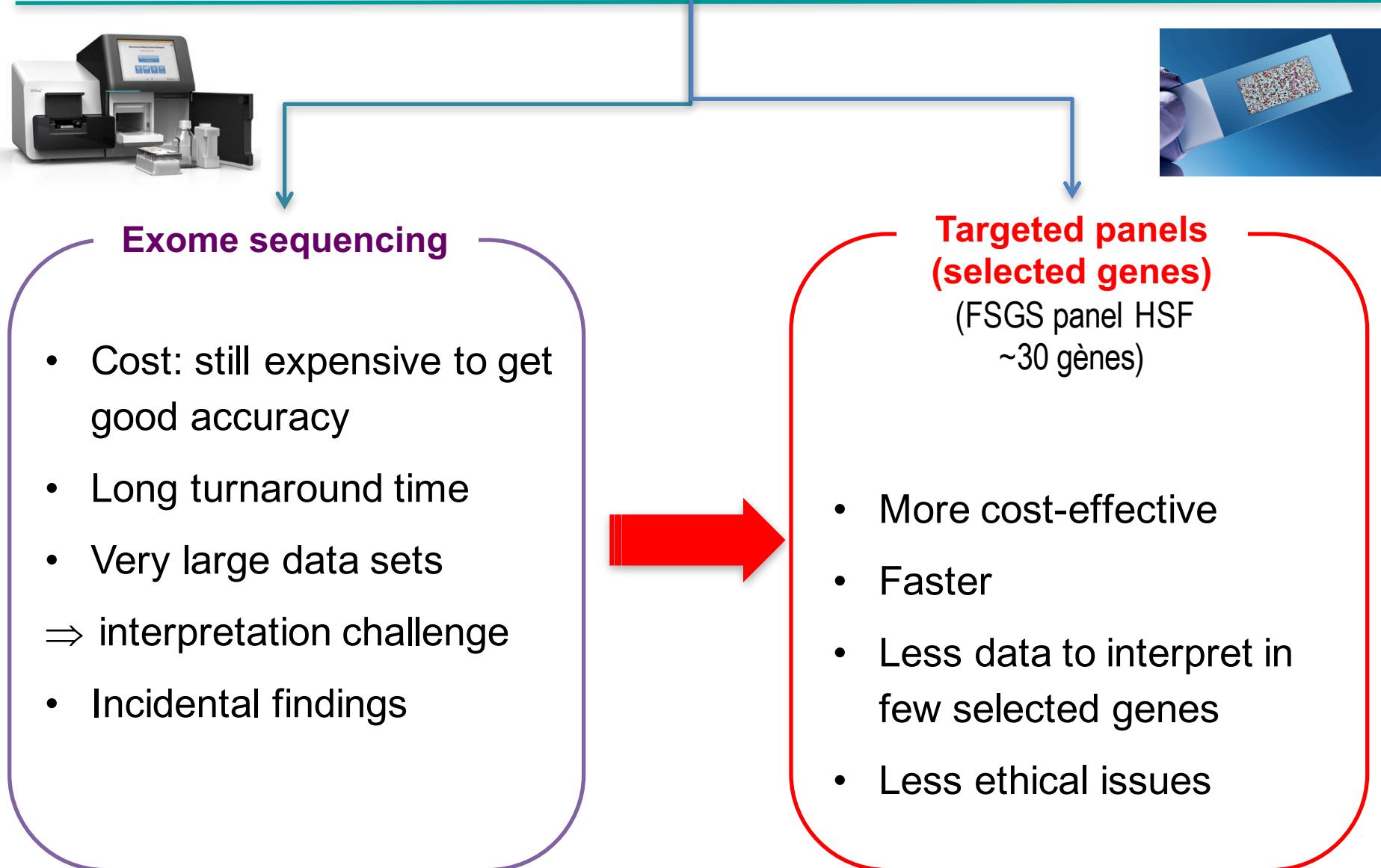
4) Strategies for molecular diagnosis of SRNS/FSGS ?

Suggested Strategy of Genetic Mutation Screening in SRNS



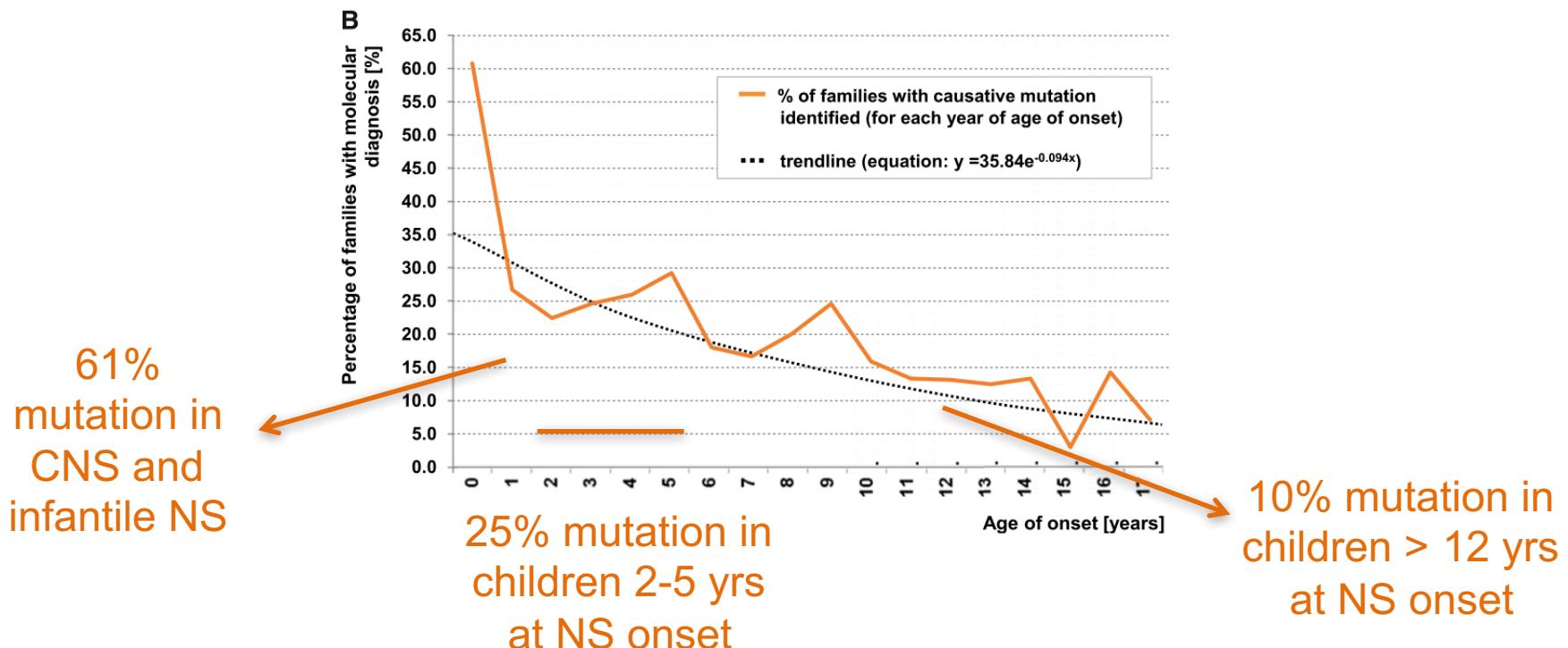
Until recently, sequential gene screening by Sanger sequencing

4) Strategies for molecular diagnosis in FSGS/SRNS



Targeted sequencing – Next Generation Sequencing (NGS) panel sequencing in SRNS-FSGS

- 1783 families with SRNS < 25 yrs : Sanger and targeted NGS of 27 genes
- 29.5% disease causing mutations in a single gene
- 49.5% in consanguineous families and only 25% in non consanguineous families



Adult forms (Necker's cohort)

- 135 adult patients
- Sporadic cases
- Mean age at Pu : 30 (18-84) years
- 58% edema at diagnosis
- 50% ESKD at a median age of 37 (20-72) years
- **16/135 (12%) patients with identified mutation**
 - Mean age at Pu 25 yrs (18-46)
 - All diagnosed < 25 yrs except 2 *COL4* gene mutations



A. Servais



C. Antignac
Necker, Paris

imagine
INSTITUT DES MALADIES GÉNÉTIQUES

Autosomal Dominant (31%)

INF2 2
WT1 1 (Frasier)

PAX2 1

LMX1B 1

TRPC6 0

ACTN4 0

Collagen genes (44%)

COL4A3 4
COL4A4 1
COL4A5 2

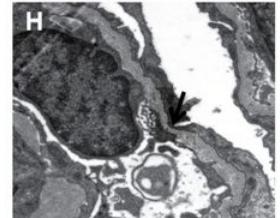
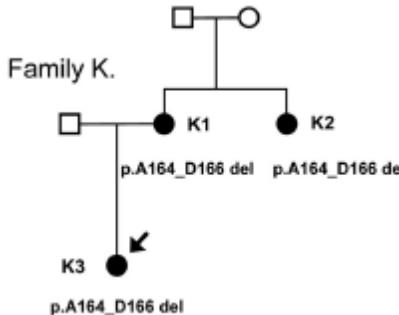
Autosomal Recessive (31%)

NPHS2 2
MYO1E 1
CD2AP 1

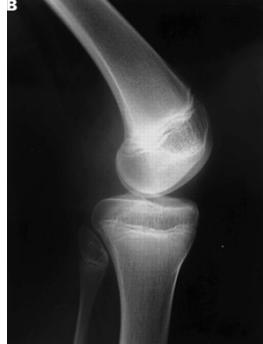
Genetic testing in nephrotic syndrome: who, when, how?

Search for hereditary forms

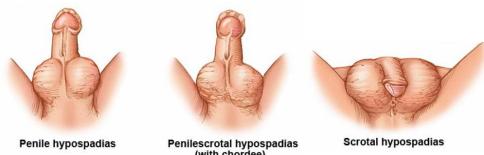
- * Tests the parents: Pu, Hu
- * search extraR features



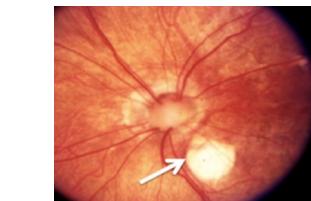
Alport Sd



Nail Patella



DDS and Frasier Sd



Papillo-renal



Mitochondrial disease



CMT

Genetic testing in nephrotic syndrome: who, when, how?

Who?

- All childhood cases with SRNS: no or partial remission under immunosuppressive therapy
 - ➡ Not in sporadic immune forms (negative Pu or post-Tx recurrence)
- Familial adult cases : Yes
- Sporadic adult cases ? < 25 years ?

When?

At the end of the first failed corticosteroid course.

How?

Gene panels Until whole exome/genome sequencing for all...

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