

Treatment of Focal Segmental Glomerulosclerosis

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Overview

2

Pathogenesis

3

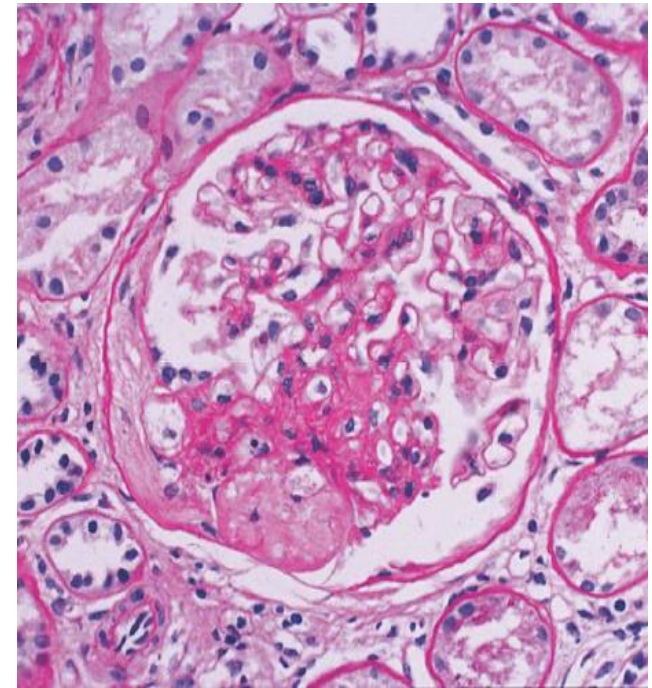
Diagnosis and Treatment





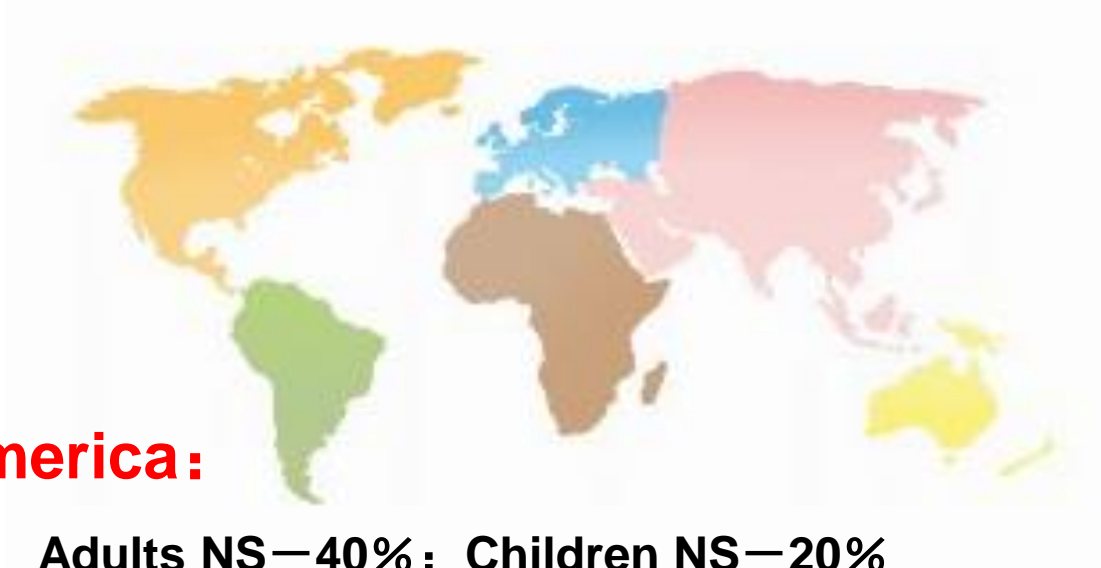
Focal Segmental Glomerulosclerosis

- First described in 1957 by Rich
- Accounting for 10%~ 30% of all renal biopsies
- Incidence has been increasing recently
- Typical manifestations:
 - ✓ Obvious proteinuria
 - ✓ Hypertension
 - ✓ worsening renal insufficiency
- One of the common causes of ESRD





The Epidemiology of FSGS (Global)



■ America:

- Adults NS—40%; Children NS—20%
- Incidence of FSGS-ESRD : **7 pmp**
- Renal biopsies:
12.2% (Haas M, 1995)-18.7% (D'Agati, 1994)

■ South Korea:

- Renal biopsies: 5.9% (1997-2001)





The Epidemiology of FSGS (China)



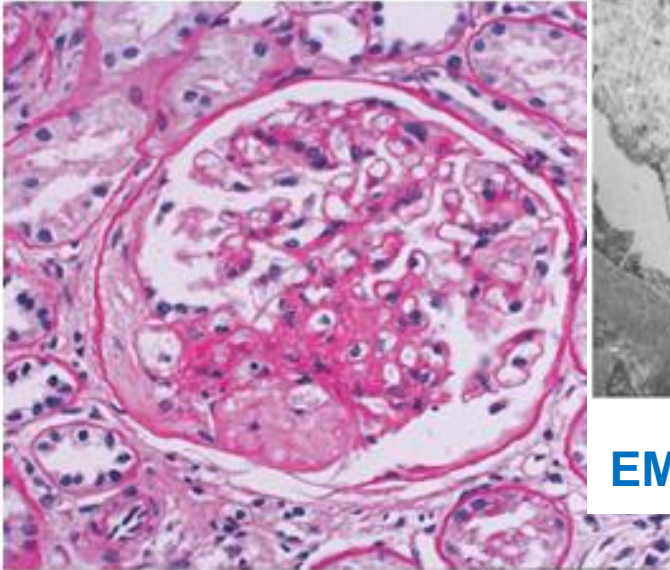
■ China:

- Peking University First Hospital : 3.3%(1998-2002), 3.8%(2003-2007)
- Nanjing General Hospital: 6.4%(1979-1989), 5.2%(1990-1999)
- Ruijin Hospital: 16.7%(2003-2005), 18.5%(2006-2008)

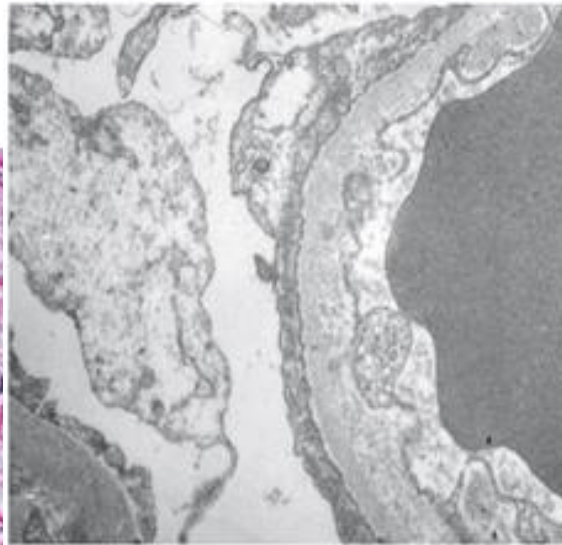




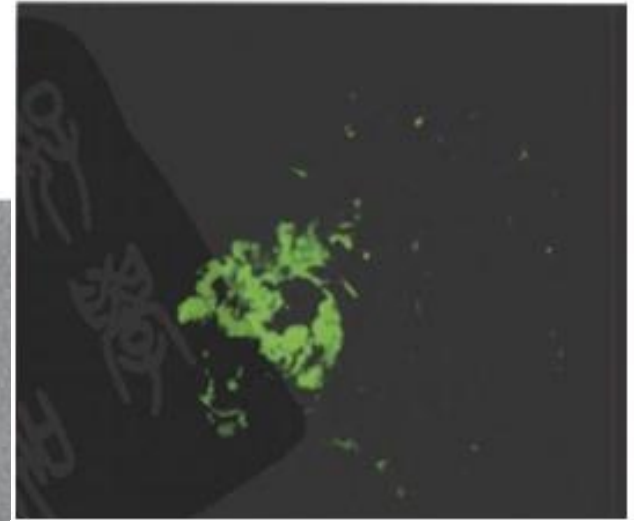
Pathological Feature



LM: Focal Segmental Glomerulosclerosis



EM: foot-process effacement

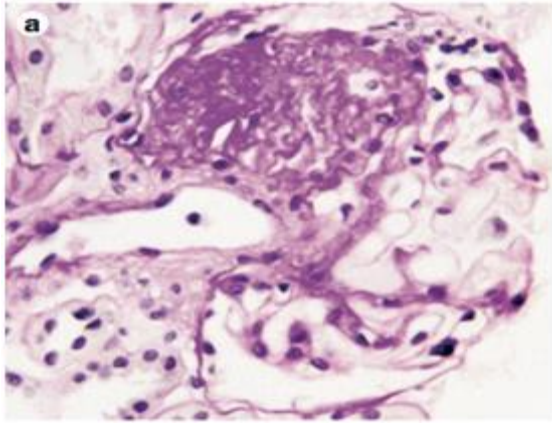


IF: deposits of IgM and C3

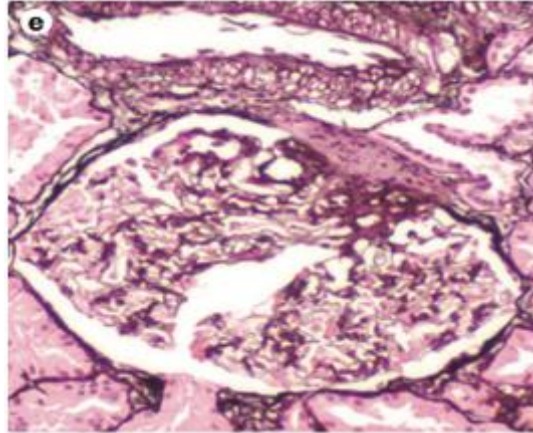




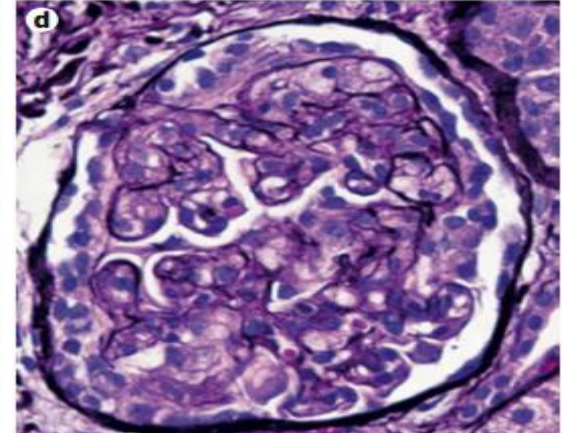
Pathologic Classification



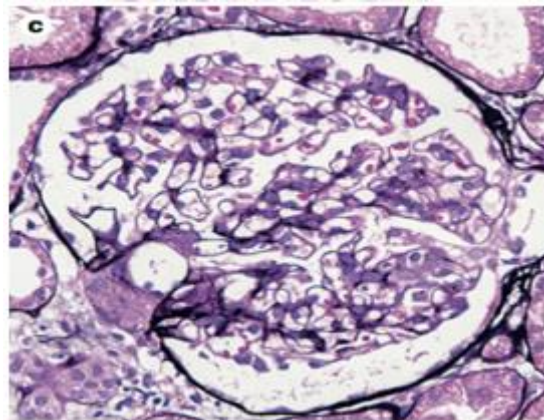
NOS



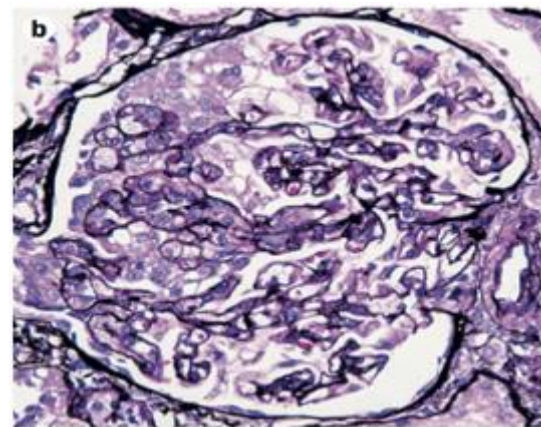
Hilar



Cellular



Tip



Collapsing

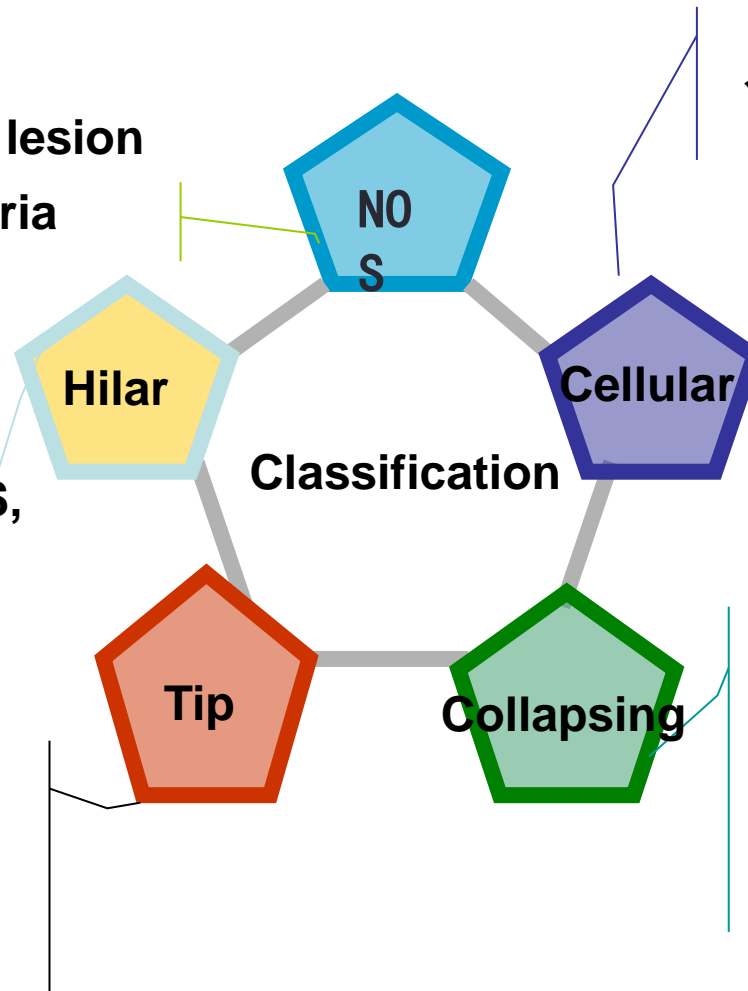




Characteristics & Outcomes

- ✓ Most common
- ✓ Tubulointerstitial lesion
- ✓ Massive proteinuria

- ✓ Accidental onset



- ✓ Also in secondary FSGS, DM, obesity etc.

- ✓ Poorest prognosis
- ✓ Frequent in African Americans
- ✓ Poor response to steroid

- ✓ Best prognosis
- ✓ Steroid-sensitive NS

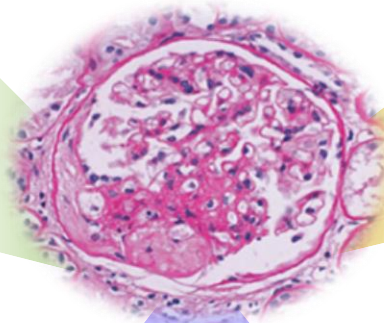




Etiology Classification

Primary FSGS

Unknown



Secondary FSGS

obesity, reflux
nephropathy,
drugs, infections
.....

Familial FSGS

Mutations of
podocyte related
molecules

Ren Fail. 2003; 25: 759-764

Kidney Int. 1996; 50: 1582-1590

Nat Med. 2011; 17: 952-960

Pediatr Nephrol. 2004; 19: 1075-1092





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Diagnosis and Treatment





Pathogenesis mechanism

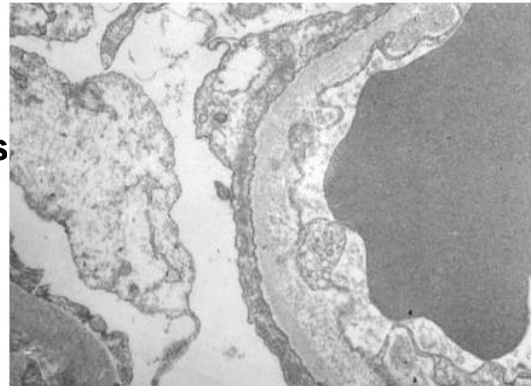
Permeability factors

Circulating factor

- suPAR
- IL-13

Podocyte-associated molecules

- Angptl4
- Integrin β 3



hemodynamics

- Decreased nephron mass
- Low birth weight
- Vessel dilation
- Hillar type is common



toxicity

- Viral infection, drugs, heavy metal, etc.
- Collapsing is common

heredity

- Podocyte genes mutations
- Other genes mutations





Familial FSGS genetic mechanism





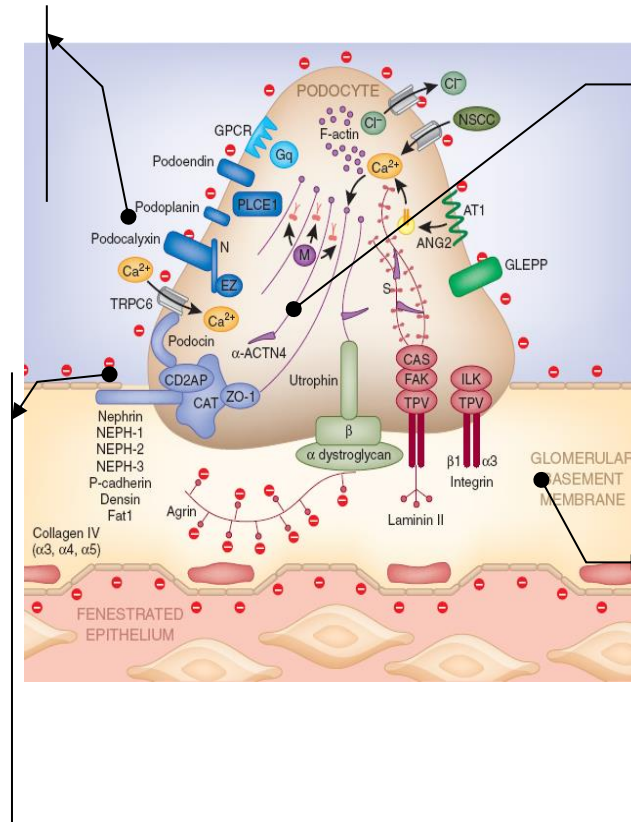
Pathogenic genes of FSGS

Membrane ionic channel

- **TRPC6 (adults)**

Slit Diaphragm

- **NPHS1 (children)**
- **NPHS2 (children, adults)**
- **CD2AP (children)**



Cytoskeleton

- **ACTN4 (adults)**
- **INF2 (adults)**

Transcription Factors

- **WT1 (children)**
- **PAX2 (children)**
- **Lmx1b (children)**

Kestila M *Molecular Cell*. 1998(1):575-582
Nat Genet. 2000 Apr;24(4):349-54.
Science. 2005 Jun 17;308(5729):1801-4

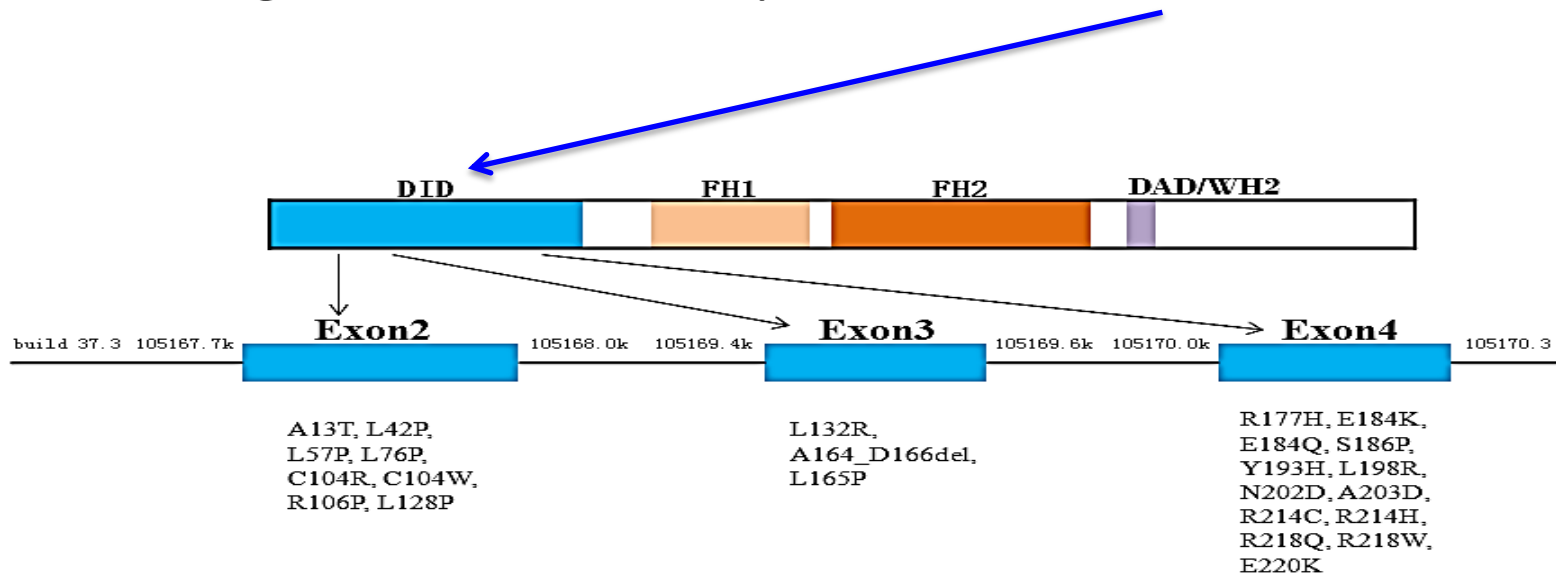
Putala H. *Hum Mol Genet* 10:1-8
Nat Genet. 2000 Mar;24(3):251-6
Nat Genet. 2010 Jan;42(1):72-6. Epub 2009





Podocyte cytoskeleton protein—INF2

- Location:14q32
- Encoding forming nucleation factor — actin-regulating proteins
- Mutations cause adult-onset FSGS (2010)
- Mutation rate : **9.8%-17%** (most common in Caucasian)
- Pathogenic mutations mainly locate in **DID domain**

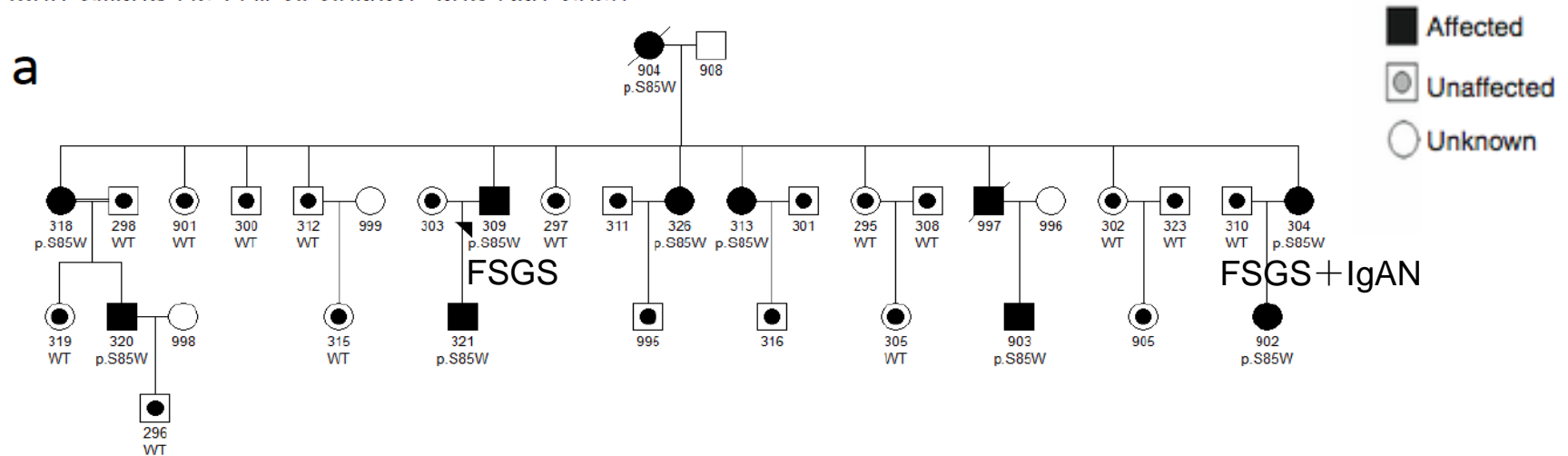




Family FSGS research 1 —linkage analysis and precise mapping

Novel mutations in the *inverted formin 2* gene of Chinese families contribute to focal segmental glomerulosclerosis

Jingyuan Xie^{1,6}, Xu Hao^{1,6}, Evren U. Azeloglu², Hong Ren¹, Zhaohui Wang¹, Jun Ma¹, Jian Liu¹, Xiaodan Ma³, Weiming Wang¹, Xiaoxia Pan¹, Wen Zhang¹, Fang Zhong¹, Yifu Li⁴, Guoyu Meng³, Krzysztof Kiryluk⁴, John Ciiiano He⁵, Ali G. Gharavi⁴ and Nan Chen¹

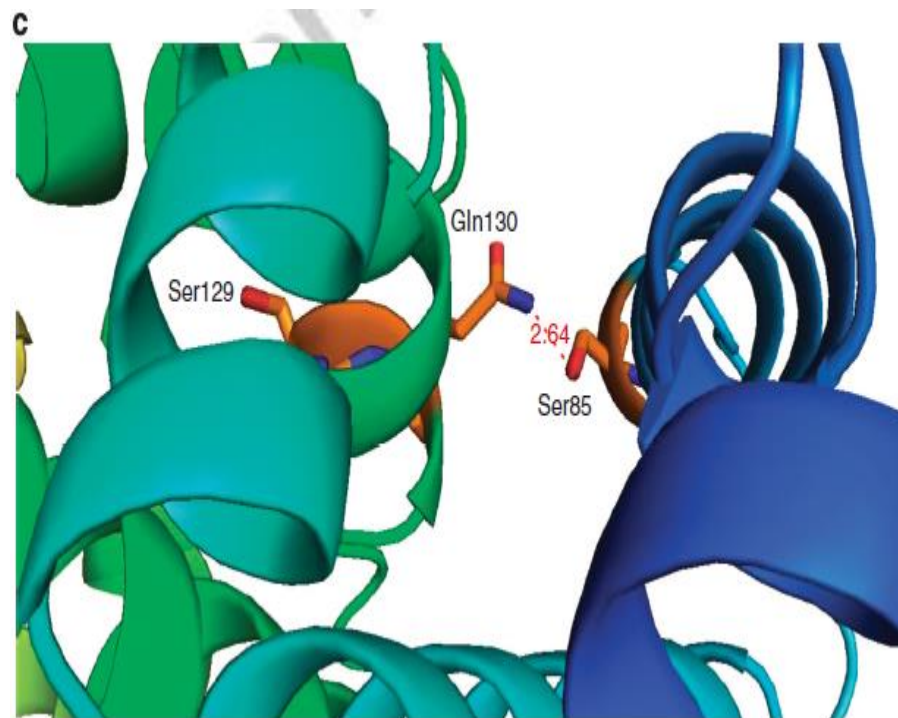
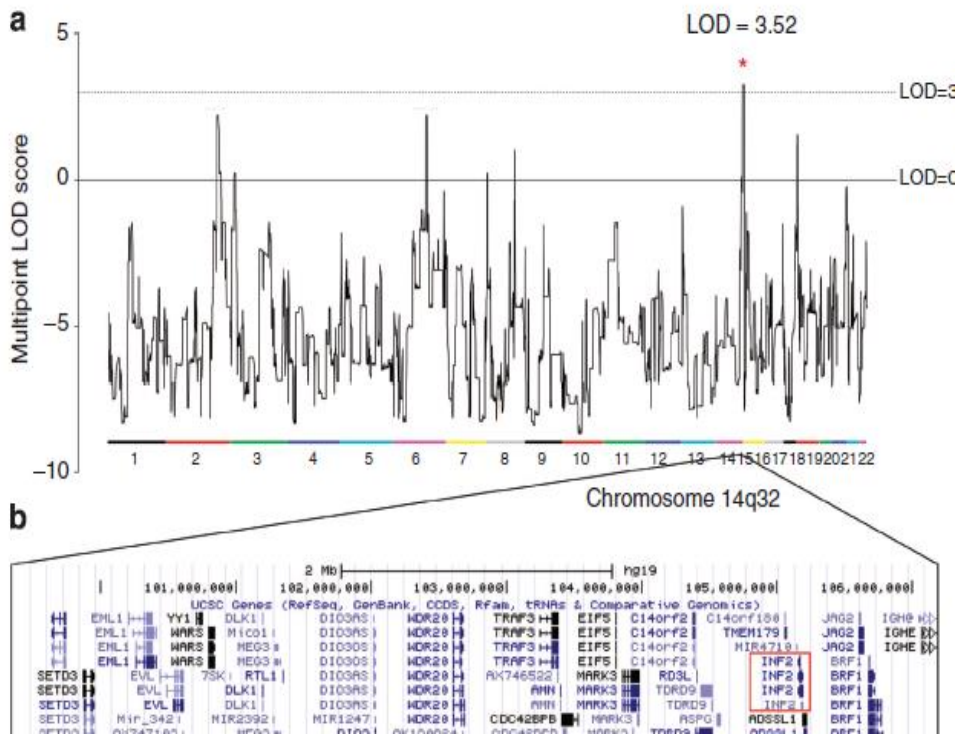


- **Autosomal Dominant**
- **55 adult-onset FSGS families, 34 IgAN families**
- **The largest pedigree including 100 family members, 11 affected individuals**
- **Onset age:20~60, massive proteinuria and ESRD**





Linkage analysis and structure prediction



14q32 LOD: 3.52

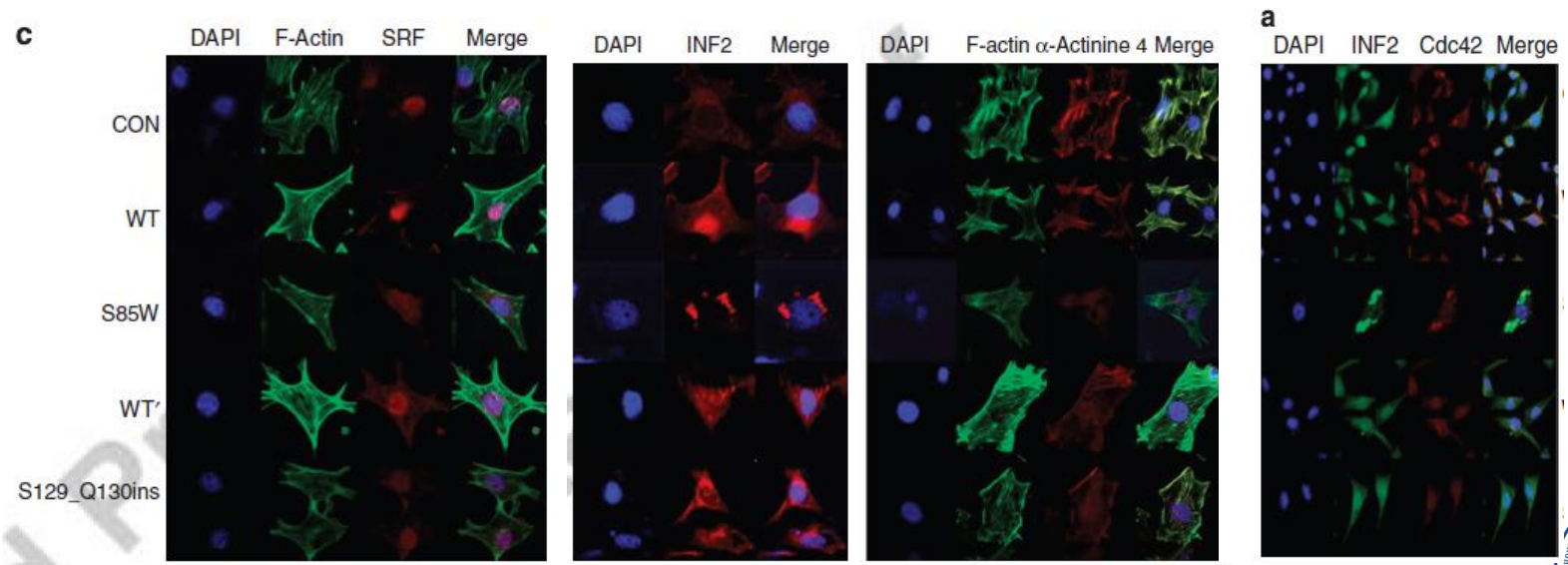
***p.S85W* abolishes the hydrogen bonding network in the inner core of the protein**





p.S85W affects the cytoskeleton

- Mutation rate in our study is 3.6%.
- *p.S85W* mutation affects the interaction between *INF2* and *Cdc42*, which may lead to decreased activation of SRF, leading to abnormal cytoskeletal and morphological changes of podocyte
- The mechanism of *p.S129_Q130insVRQLS* mutation is still unknown





Family FSGS research 2

— gene sequencing

498 | *Journal of Molecular Cell Biology* (2014), 6(6), 498–505

doi:10.1093/jmcb/mju040

Article

COL4A3 mutations cause focal segmental glomerulosclerosis

Jingyuan Xie^{1,†}, Xiaoxi Wu^{2,†}, Hong Ren¹, Weiming Wang¹, Zhaohui Wang¹, Xiaoxia Pan¹, Xu Hao¹, Jun Tong¹, Jun Ma¹, Zhibin Ye³, Guoyu Meng⁴, Yufei Zhu², Krzysztof Kiryluk⁵, Xiangyin Kong², Landian Hu^{2,*}, and Nan Chen^{1,*}

➤ 40 FFSGS families

50 sporadic FSGS

➤ Methods: exome sequencing and sanger sequencing

➤ Exclude patients with *ACTN4*, *TRPC6* or *INF2*

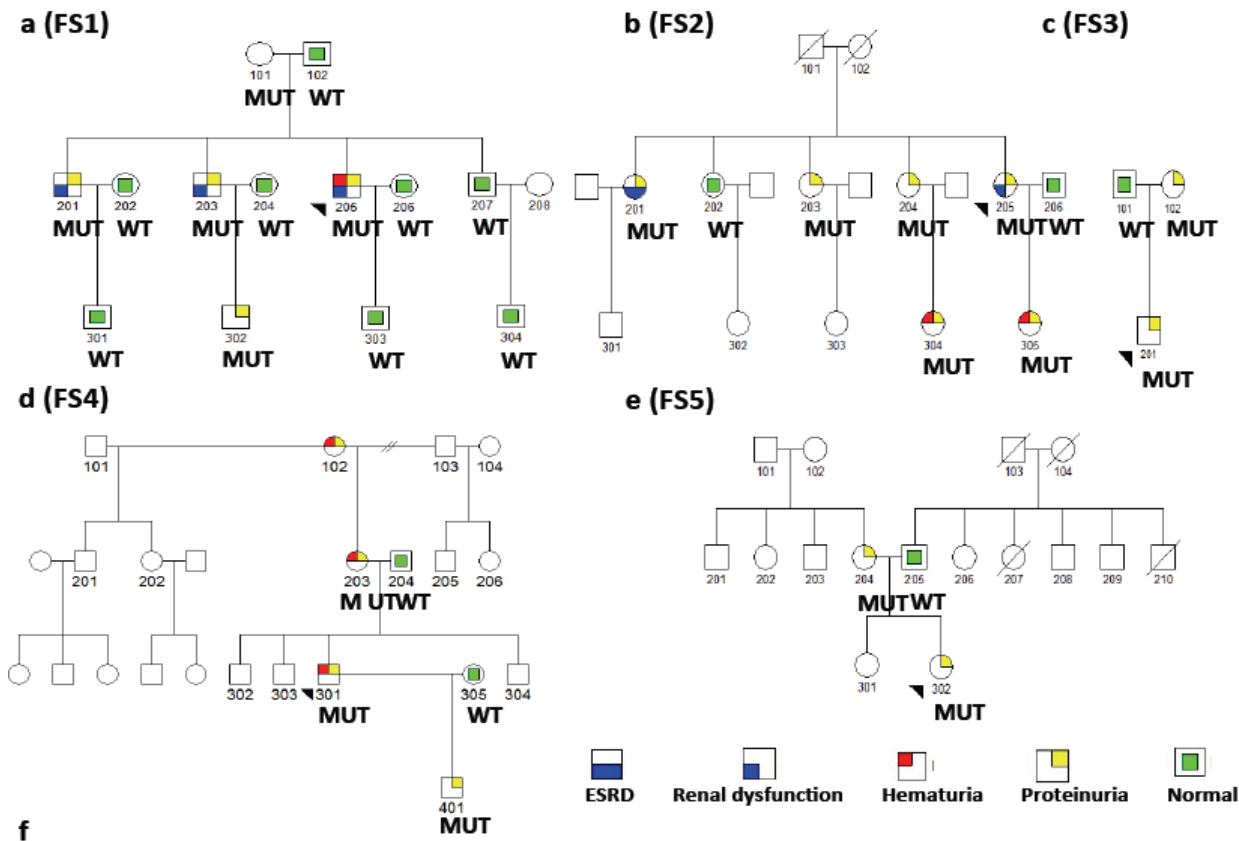
mutations

Xie J, Wu X, Ren H, Wang W, ..., Chen N
Journal of Molecular Cell Biology 2014





COL4A3 mutations cause FFSGS



✓ 5/40 (12.5%) family with *COL4A3* heterozygous mutation

✓ 1/50 sporadic patients with *COL4A3* heterozygous mutation



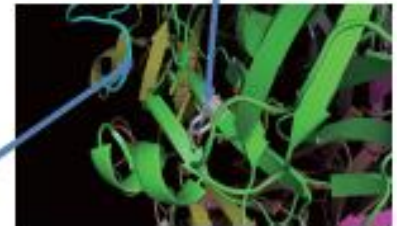
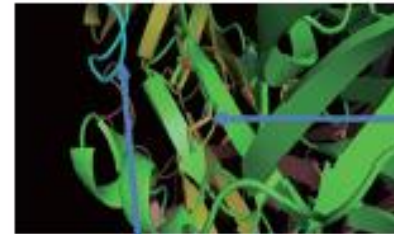


Located in highly conserved regions

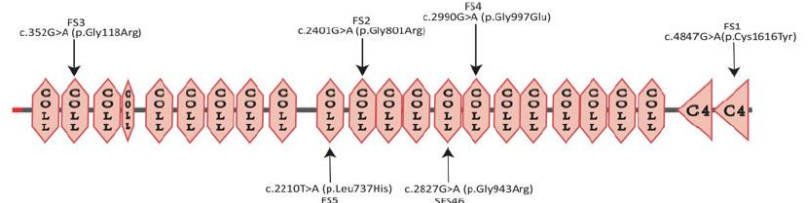
b

	F53 c.352G>A (p.Gly118Arg)	F55 c.2210T>A (p.Leu737His)	F52 c.2401G>A (p.Gly801Arg)
<i>Mus musculus</i>	PGLFGHPGFRRLAGLPGQNGS	GKFGRFGEPCIPGAKGEPVSG	FQDFGQFGSFGAKGSPGRICIP
<i>Rattus norvegicus</i>	PGIFGYPGFFLGLAGLPGQNGS	GRFGYFGEMGVPAKGEFVSG	FQDFGQSGFFPAKGFPPGRICIP
<i>Homo sapiens</i>	PGTFGNTGYPYVLGVVFGCSGS	GEFGLFGKPLGPAKGEPAVA	RGDFGQFGFFPEQGGPPGRCTE
<i>Pan troglodytes</i>	PGTFGNTGYPYVLGVVFGCSGS	GEFGLFGKPLGPAKGEPAVA	RGDFGQFGFFPEQGGPPGRCTE
<i>Macaca mulatta</i>	PGTFGNTGYPYVLGVVFGQNGS	GEFGLFGKPLGPAKGEPAVA	RGDFGQFGFFPEQGGPPGRCTE
<i>Oryctolagus cuniculus</i>	PGTFGNAGYPYVLGVVFGQNGS	GKFGIPGKPLGPAKGEFALA	RGDFGQFGFFPETGLPGRCTFQ
<i>Ailuropoda melanoleuca</i>	PFTFGHPGYFPAAGIFPGQNGS	GKFGSPGEPCLGGAKGEFGLA	QQDFGKFGFFPEKGGSPGRCMV
<i>Mustela putorius furo</i>	PGTFGHPGYFPAAGIFPGQNGS	GKFGSPGEPCLGPAKGEFGLA	QQDFGKFGFFPEKGGPPGRCTE
<i>Canis lupus familiaris</i>	PGTFGHPGYFPAAGIFPGQNGS	GKFGSPGEPCLGPAKGEFGLA	FQDFGKFGFFPEKGGPPGRCTE
<i>Felis catus</i>	PGTFGHPGYFPAAGIFPGQNGS	GKFGSPGEPCLGPAKGEFGLA	QQDFGKAGFFPEKGGPPGRCTE
<i>Equus caballus</i>	PGTFGHPGYVLGLAGLPGQNGS	GKFGSPGEPCLGPAKGEFGLA	RGDFGQSGFFPEKGLPGRCTE
<i>Bos taurus</i>	PGIFGHPGYVLGLAGLPGQNGS	GKFGSPGEPCLGPAKGEFGLA	RGDFGKFGFFPEKGGPPGRCTE
<i>Ovis aries</i>	PGIFGHPGYVLGLAGLPGQNGS	GKFGSPGEPCLGPAKGEFGLA	RGDFGKFGFFPEKGGPPGRCTE
<i>Dasybus novemcinctus</i>	PGIFGHPGYVLGLAGLPGQNGS	GKFGSPGEPCLGPAKGEFGLA	QQDFGQFGFFPEKGGPPGRCTE
<i>Monodelphis domestica</i>	PGTFGNTGYPYFPGIFPGQNGT	GKFGSPGEPCLGPAKGEFGLA	FQDFGKFGFFPEKGLPGLCIP
<i>Gallus gallus</i>	PGFPGIAGHPFRSGVFGQNGT	GEFGRAGNLQPEKGDGPII	RGDFGSSGVFSDPFGPKFAE

	SF546 c.2827G>A (p.Gly943Arg)	F54 c.2990G>A (p.Gly997Glu)	F51 c.4847G>A (p.Cys1616Tyr)
<i>Mus musculus</i>	KGERGEKGNFQFSQITLLKGD	PGLPGFFPRDITGSRGNPGR	EEFRASFFIECHGRGTCNYYS
<i>Rattus norvegicus</i>	KGERGEKGNFQFSQITLLKGD	PGLPGSGFPRDITGSSGDOPGR	EEFRASFFIECHGRGTCNYYS
<i>Homo sapiens</i>	KGEQDQGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Pan troglodytes</i>	KGEQDQGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Macaca mulatta</i>	KGEQDQGNFQFSQIISHVIGH	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Oryctolagus cuniculus</i>	EGERGDGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRANFFIECHGRGTCNYYS
<i>Ailuropoda melanoleuca</i>	KGEKDGKGNFQFSQIISNLLGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Mustela putorius furo</i>	KGEKDGKGNFQFSQIISNLLGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Canis lupus familiaris</i>	KGEKDGKGNFQFSQIISNLLGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Felis catus</i>	KGEKDGKGNFQFSQIISNLLGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Equus caballus</i>	KGEKDGKGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Bos taurus</i>	KGEKDGKGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Ovis aries</i>	KGEKDGKGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Dasybus novemcinctus</i>	KGEKDGKGNFQFSQIISHVIGD	PGFAGFFPRDGLSTGNPGE	EEFRASFFIECHGRGTCNYYS
<i>Monodelphis domestica</i>	KGERGNKGNFQFSQIISDDIGE	HGPPGFFPRDGLSTGNPGE	EDFRANFFIECHGRGTCNYYS
<i>Gallus gallus</i>	QGEFGKGNFQFSQITLVVIGN	EGVPGVGVKGMFLPGIPGG	EEFRALFFIECHGRGTCNYYS



Other COL4A3 subunit



➤ All six variants were located in a highly conserved region of *COL4A3*

➤ Structural predictions: The *p.Cys1616Tyr* variant disrupts the COL4A3 protein. **COL4A3 mutations cause FFSGS**





COL4 mutations

the most frequent mutations in adult FSGS

Study design:

- 81 adults from 76 families;
- 24 families had a history of renal diseases
- Target NGS panel: covering 39 genes

Results:

- Confirmed pathogenic mutations: 10/81(12%)
- Probably pathogenic mutations: 6
- Total mutation rate: 16/81 (20%)
 - Familial 22%, sporadic 10%
 - **COL4A3-5 mutation: 8/81 (9.8%)**

Gene	CHR	Exons	Size (bp)
ACSL4	X	15	2364
ACTN4	19	21	2736
ALG1	16	13	1395
APOE	19	3	954
APOL1	22	7	1289
ARHGAP24	4	12	2478
ARHGDIA	17	6	683
CD2AP	6	18	1920
CFH	1	23	3710
COLAA3	2	52	5013
COLAA4	2	47	5073
COLAA5	X	53	5383
COQ2	4	7	1266
COQ6	14	13	1495
INF2	14	23	3817
ITGB4	17	40	5628
LAMA5	20	80	11088
LAMB2	3	32	5397
LMNA	1	18	2452
LMX1B	9	10	1453
MYH9	22	40	5883
MYO1E	15	28	3327
NEIL1	15	11	1865
NPHP4	1	31	4505
NPHS1	19	29	3726
NPHS2	1	8	1152
NXF5	X	14	1098
PDSS2	6	8	1200
PLCE1	10	33	7300
PMM2	16	8	741
PODXL	7	9	1677
PTPRO	12	27	3655
SCARB2	4	12	1437
SMARCAL1	2	16	2865
SYNPO	5	5	7560
TRPC6	11	13	2796
WT1	11	12	1648
ZEB1	10	11	3445
ZMPSTE24	1	10	1428





Hereditary *COL4A3*/*COL4A4* variants mistaken for FSGS?

DUKE study: *COL4A3* mutation rate in FSGS is 10%

- ✓ Mutations in *COL4A3* and *COL4A4* are known to cause AS、TBMN
- ✓ Secondary FSGS is known to develop in classic AS at later stages
- ✓ In all 7 families, there were individuals with histologic features of FSGS by LM.
- ✓ In one family, EM showed thin GBM. other families had variable findings inconsistent with classical AS.





Redefining the spectrum of the disease?

- A subset of renal manifestations associated with *COL4A3* or *COL4A4* variants cannot be distinguished from FSGS by clinical data or histopathology
- Redefining the spectrum of Alport syndrome?



**Collegen IV
related
nephropathy**





Mutation rate of known genes in FSGS

	category	Number	Mutation rate	Reported rate
<i>NPHS2</i>	SRNS	44	0 (0)	0-18%(children)
	FSGS	77	1 (1.3%)	4-24%
<i>ACTN4</i>	FSGS	80	1 (1.25%)	3.5%
<i>TRPC6</i>	FSGS	80	2 (2.5%)	2.3~20%
<i>INF2</i>	FSGS	55	2 (3.6%)	16%
<i>COL4A3</i>	FSGS	40	5 (12.5%)	10%
Total			21.2%	35-80%

In over 70% pedigrees , no mutation is detected, indicating that lots of **potential pathogenic genes** need to be investigated.

Zhu B et al, *Mutation Res* 2009
Zhang Q et al, *Contr Nephrol* 2013
Clin J Am Soc Nephrol. 2011;6(5):1139-48





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Diagnosis and Treatment





Study of FFSGS from Ruijin hospital

- **Patients:**

- 2005~2012, 124 FFSGS patients (83 families)
- 124 primary FSGS patients

- **Diagnostic criterias of FFSGS:**

- ✓ More than one family member are biopsy-proven FSGS
- ✓ One has biopsy-proven FSGS, other have proteinuria, progressive renal dysfunction or ESRD
- ✓ AS, FD, TBMN or other hereditary kidney diseases were excluded





Study of FFSGS from Ruijin hospital

- **Basic information**
 - **Gender, age, blood pressure, first episode, disease course**
- **Laboratory examination**
 - **Blood/urine routine 、 24H urine protein, renal function**
 - **eGFR was calculated by MDRD**
- **Outcome**
 - **ESRD**
 - **Remission of proteinuria (at least 50% decline)**





Baseline characteristics

Ruijin Hospital

Table 1. Clinicopathological characteristics of FFSGS and SFSGS

	FFSGS	SFSGS	p value
Number	124	124	
Male/female	71/53	76/48	0.096
Follow-up, months	28.3±12.5	26.5±19.5	0.081
Age of onset	34 (10–62)	40 (12–80)	0.035
Hypertension, %	43.75	35.16	0.079
Serum creatinine, $\mu\text{mol/l}$	174.3±186.6	153.0±144.2	0.086
Uric acid	415.6±118.1	397.9±109.4	0.275
eGFR, ml/min per 1.73 m ²	69.49±38.27	81.76±50.47	0.069
Hematuria, %	62.9	22.9	0.0001
Proteinuria, g/day	1.1 (0.03–5.6)	1.5 (0.05–8.7)	0.003
Nephrotic syndrome, %	13.3	22.6	0.029
Focal glomerulosclerosis, %	13.23±11.83	15.71±13.49	0.141
Global glomerulosclerosis, %	25.56±21.4	16.52±20.96	0.0007
Tubulointerstitial lesion score	4.07±0.28	3.00±0.1	0.0004
Remission of proteinuria, %	23.08	48.39	0.042
End-stage renal disease, %	27.42	2.42	0.003

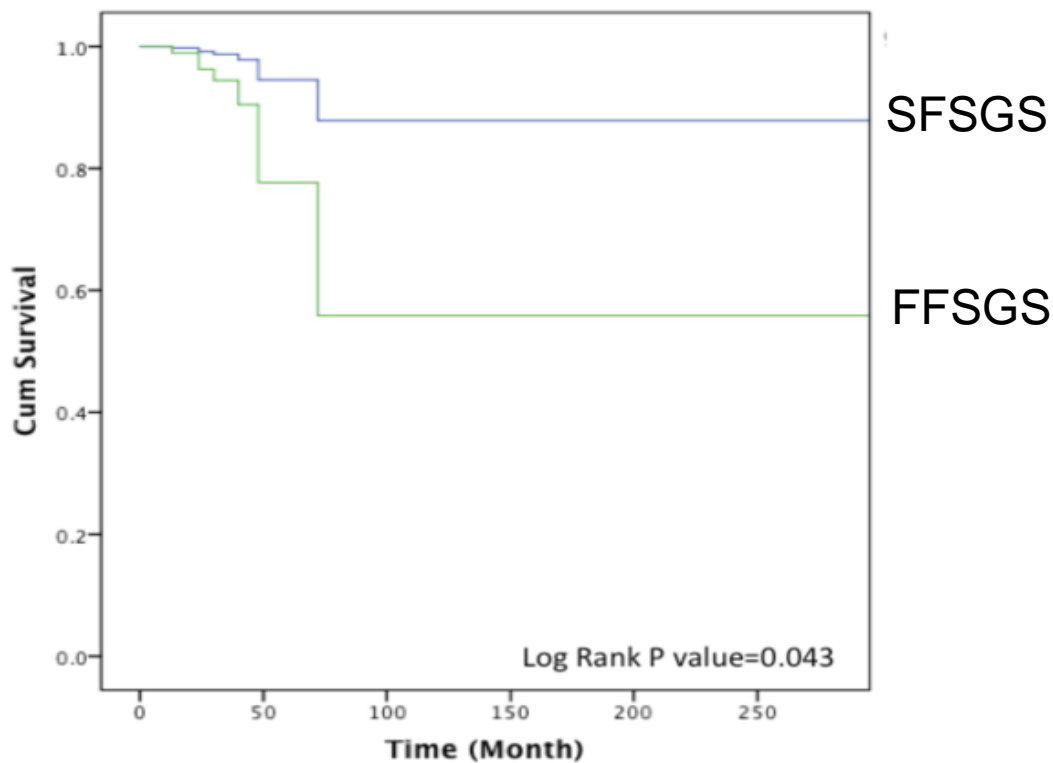
Remission of proteinuria was defined as a proteinuria decline of at least 50% of the baseline level. Hematuria was defined as more than 3 red blood cells in each high-power field.





Renal Outcome

	FFSGS	SFSGS	P
PR/CR	23.08%	48.39%	0.042
ESRD	27.42%	2.42%	0.003





Comparison between FFSGS and SFSGS

Ruijin Hospital



FFSGS

- More hematuria
- Lower proteinuria
- Less NS
- More severe pathological lesions
- Less CR/PR
- More ESRD



SFSGS

- Higher proteinuria
- More NS
- Less severe pathological lesions
- More CR/PR



Initial treatment

- Steroid and immunosuppressive therapy be considered **only in** idiopathic FSGS associated with NS. (1C)
- Prednisone: a daily single dose of 1 mg/kg (maximum 80 mg) or alternate-day dose of 2 mg/kg (maximum 120 mg). (2C)

Initial treatment

- **Initial high dose of steroids is given for a minimum of 4 weeks , up to a maximum of 16 weeks until complete remission (2D)**
- **Steroids be tapered slowly over a period of 6 months after achieving complete remission (2D)**
- **CNIs : first-line therapy ——patients with contraindications or intolerance to high-dose steroids (2D)**

Relapse treatment

- Same to MCD
- **CTX 2-2.5mg/kg/d, po. 8wks (2C)**
- For Recurrence patients after CTX treatment, CsA 3-5mg/kg/d or
FK506 0.05-0.1mg/kg/d, po (2C)



Question 1——

Corticosteroid benefit for moderate proteinuria?

Design (RCT study)

- ❖ **Participants:** primary FSGS patients with moderate proteinuria
- ❖ **ACEI/ARB group(n=30), corticosteroid group(n=30)**
- ❖ **Follow-up:12 months**
- ❖ **Results:** higher rate of proteinuria remission in corticosteroid group compared to ACEI/ARB group

	group	0 m	6 m	p value	12 m	p value
proteinuria (g/24h)	ACEI/ARB	1.96 ± 0.16 g	1.67 ± 0.33	0.432	1.71 ± 0.44	0.596
	corticosteroid	1.75 ± 0.19 g	0.69 ± 0.16	<0.001	0.48 ± 0.12	<0.001





Question 2—

Extend duration of initial steroid treatment ?

Treatment: full dose of steroid ,N=65

Steroid treatment: 8 vs 12wks

Duration	CR	PR	NR	Effective rate
8 wks	30	15	18	70.31
12 wks	30	16	15	71.87

P>0.05

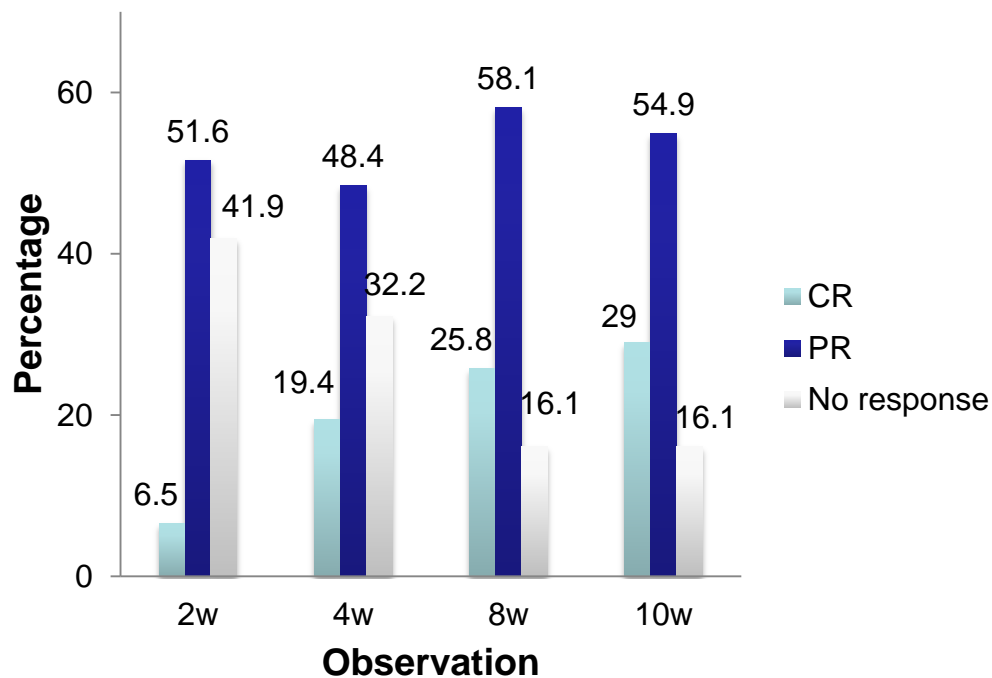
Extending steroid treatment to 12 wks resulted in a **similar** remission rate but **more side-effects**



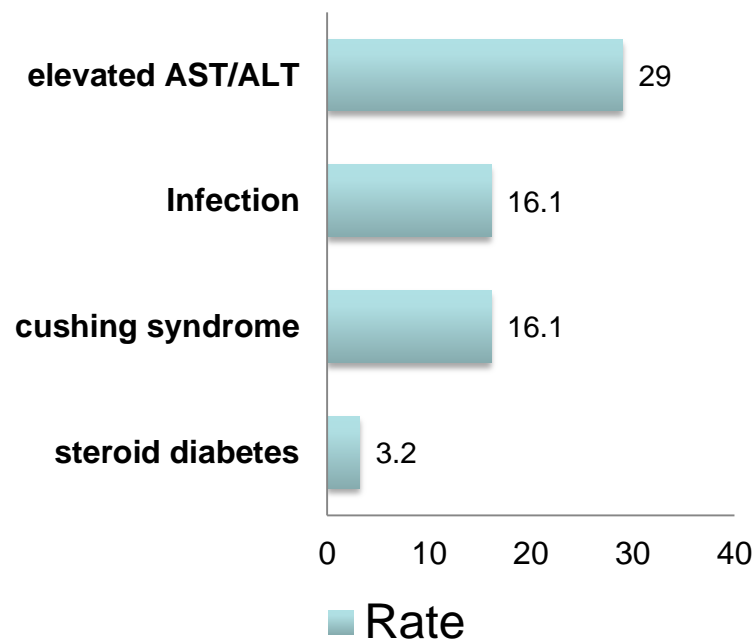


Question 2— Extend duration of initial steroid treatment ?

Extending treatment to **10wks** resulted in a same remission rate



Advent events in **week 10**





Question 3: Immunosuppressive Agents——CTX?

	CR	PR
steroid-sensitive+CTX	50%	25%
steroid-resistant+CTX	10%	10%

❖ CTX can be used as a second-line therapy?





Immunosuppressive Agents

FK-506

- ❖ **Participants:** steroid-resistant FSGS (NS) , N=44
- ❖ **Follow-up:** 24 months

Response to FK506

Parameter	Value	
Total remission	23 (52.2%)	
CR	17 (38.6%)	
Partial remission	6 (13.6%)	
Time to remit (weeks)	15.28 ± 6 (04–24)	
FSGS variants	TAC responsive	TAC resistant
FSGS-NOS (<i>n</i> = 33)	18 (54.5%)	15 (45.4%)
Cellular variant (<i>n</i> = 08)	3 (37.5%)	5 (62.5%)
Tip variant (<i>n</i> = 03)	2 (66.7%)	1 (33%)
Relapse during tapering	5 (21.7%)	
Relapse after completion of therapy	7 (30.4%)	
Complications	29 (65.9%)	

SR-FSGS, steroid-resistant focal segmental glomerular sclerosis; NOS, not otherwise specified; TAC, tacrolimus.





Immunosuppressive Agents CTX vs FK-506

- ❖ **Participants** : steroid-dependent or steroid-resistant FSGS
- ❖ **CTX group N=18, FK506 group N=15**
- ❖ **Follow-up:12 months**
- ❖ **Result: CTX and TAC had a **similar** efficacy**

Table 3. Comparison of remission rate between CTX and FK-506 groups, n (%)

Duration	CTX		FK-506		p		
	NR	remission		NR		remission	
		CR	PR			CR	PR
6 months	8 (44.4)	10 (55.6)	5 (33.3)	10 (66.7)	0.77		
		7 (38.9) 3 (16.7)		7 (46.7) 3 (20.0)			
12 months	6 (33.3)	12 (66.7)	4 (26.7)	11 (73.3)	0.97		
		9 (50.0) 3 (16.7)		6 (40.0) 5 (33.3)			

CTX is preferable for its price





immunosuppressive Agents

CTX vs FK-506

Adverse events	CTX (n = 18)	FK-506 (n = 15)	p
Infections*	9 (50.0%)	2 (13.3%)	<0.05
Sepsis	0	1	
Lung	2	1	
URTI	2	0	
UTI	4	0	
Herpes zoster	1	0	
Elevated glucose*	0	4 (26.7%)	<0.05
Elevated transaminase	1 (5.6%)	1 (6.7%)	NS
Deterioration of renal function	1 (5.6%)	0	NS
Menstruation disorders	1 (5.6%)	0	NS
Nausea/vomiting	2 (11.1%)	0	NS
Atopic dermatitis	1 (5.6%)	0	NS

NS = Non-significant difference; URTI = upper respiratory tract infection; UTI = urinary tract infection. * p < 0.05 indicates a significant difference between CTX and FK-506 groups.





immunosuppressive Agents

CsA vs MMF

- ❖ **Multicenter, RCT**
- ❖ **Participants:** **steroid-resistant FSGS, N=138**
(MMF group N= 66, CsA group N= 72)
- ❖ **Dose:** CsA 4.6 ± 1.7 mg/kg/d
MMF 26.2 ± 6.1 mg/kg/d
- ❖ **Result:**
 - Same PR and CR rate at 12 month (MMF 33% VS CsA 46%)
 - Same remission rate for extra 26 weeks.





Question 4- Rituximab

❖ **Single center, prospective**

❖ **FSGS-NS patients, N=18**

(Steroid-dependent/resistant, CNI/MMF-resistant)

Treatment: Rituximab

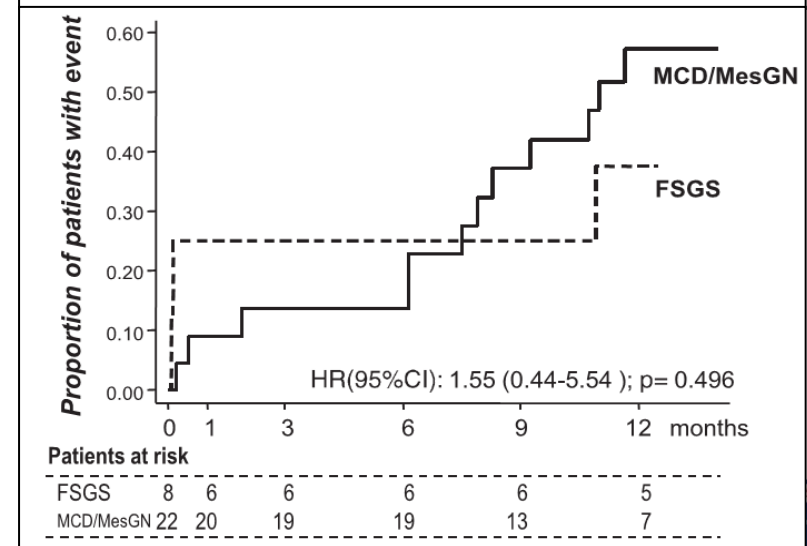
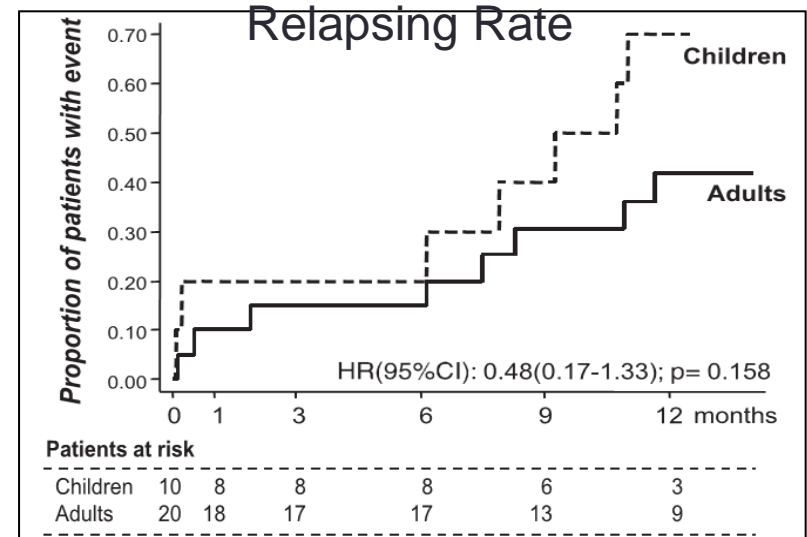
	Before treatment		8m later		12m later	
FSGS	PR	NR	PR	NR	PR	CR
	10	8	17	1	17	1





Therapy—Rituximab

- Multi-center, prospective
- N=30
 - ✓ Children (N=10)
 - ✓ Adult (N=20)
 - ✓ MCD/MesGN (N=22)
 - ✓ FSGS (N=8)
- SDNS OR FRNS
- 1-2 doses of RTX (375mg/m²)
- Follow-up: 1 yr
- Result:
 - ✓ 100% remission
 - ✓ 60% treatment-free
 - ✓ 50% never relapsed





Rituximab in treatment of refractory FSGS

- ❖ **Patients** : FSGS-NS, N=12 (SDNS, SRNS),
no response to CTX, MMF, CsA/Tac, average duration 2-10 years)
- ❖ **Treatment** :
 - ✓ 375mg/m²/w, 2-4 times
 - ✓ Tapering steroid and immunosuppressive agents within 3-6 months
 - ✓ Reapply Rituximab when CD19 count>1% or >15/ul
- ❖ **Follow-up** : 6-18 month
- ❖ **Efficacy** : All achieved CR , no relapse





Case Report

一般情况：男，17岁，学生

病史简介：

5年前无明显诱因下出现颜面、双下肢浮肿伴泡沫尿，2010年10月、2011年6月二次外院肾穿病理提示FSGS (NOS型)，期间予“激素+干细胞” (2010. 10-11. 6)、 “激素+环孢素” (2011. 6-2012/2)、 “激素+FK506” (2012. 2-2012. 8)、 “激素+FK506+MMF” (2012. 8-2012. 9) 治疗，治疗过程中NS一度缓解，但反复复发。2年前我院就诊，当时24蛋白定量16398-25889mg / 24h，Alb19g/l，Scr41umol / l，调整治疗方案为“激素+FK506+CTX” (2012. 9-2014. 10) 治疗，规范治疗15月后NS完全缓解，4月后因上感后NS再次复发。

查体：BP:115/62mmHg，神清、满月脸、双肺呼吸音偏低，心率80次 / 分，腹软，腹部皮肤紫纹明显，双下肢浮肿。





治疗经过

- 2014. 10考虑NS复发，在“激素40mg / d+FK506 4mg / d”基础上加用美罗华 375mg/m²治疗。
- 2次疗程后（2014. 10. 27、2014. 11. 3），NS完全缓解，予激素减至25mg / d。
- 2014. 12、2015. 2再次分别予美罗华治疗，治疗期间病情稳定，激素、FK506逐渐减药。
- RTX治疗期间无药物相关性不良反应。
- 随访期间无复发及不良反应反应。

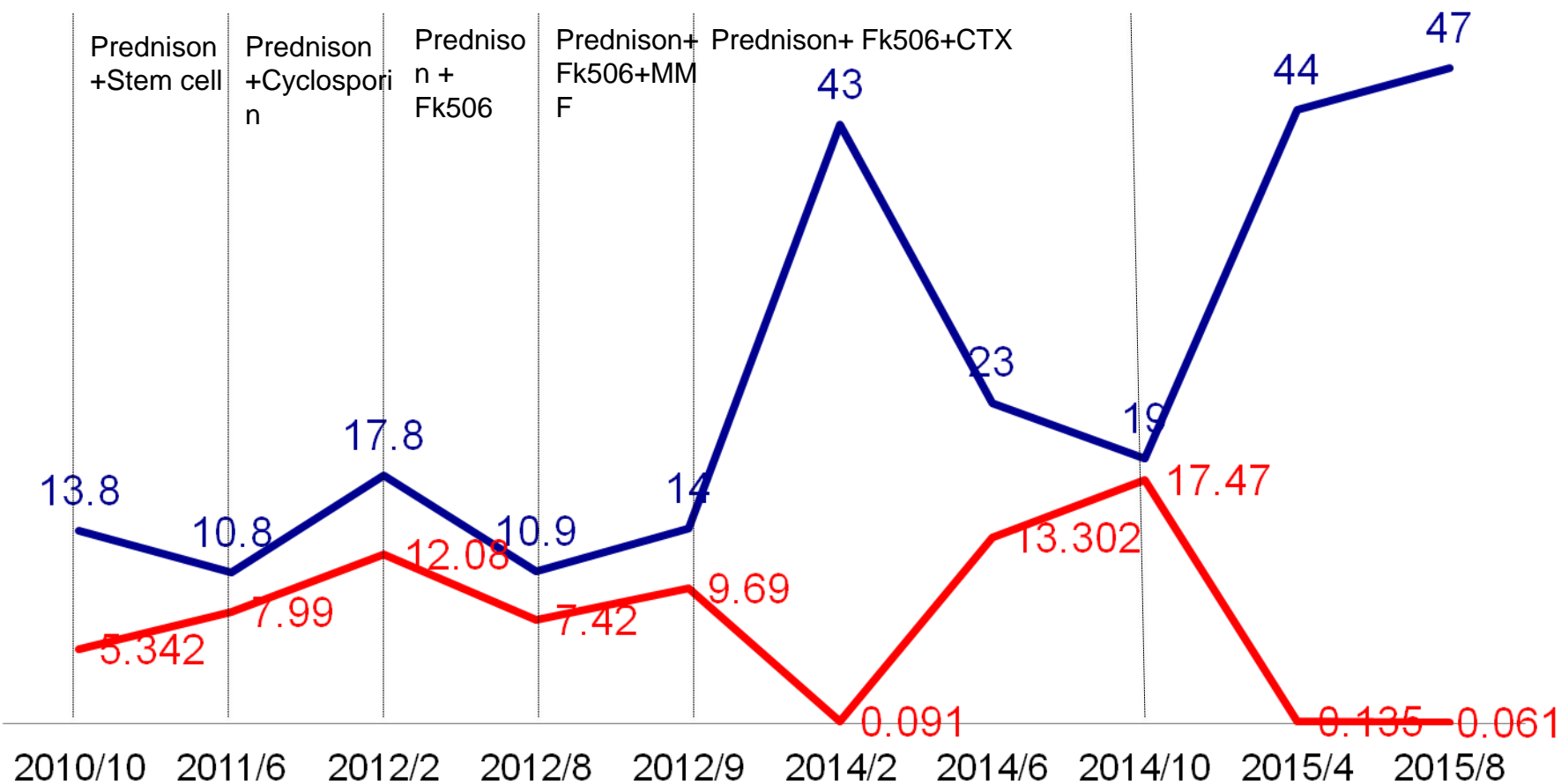




24h proteinuria and albumin before and after treatment

RTX + Prednison+FK506

— Alb (g/l) — 24h proteinuria(g/24h)



Ruijin Hospital unpublished data





B cell counts during the treatment

RTX Treatment	Duration (month)	Lymphocyte ($10^9/l$)	CD19+ (%)	CD20+ (%)	CD19+ ($\uparrow / \mu l$)
RTX ($375\text{mg}/\text{m}^2$) + Prednisolone (40mg) + FK506($4\text{mg}/\text{d}$)	1/4	4	4.2	4.4	168
	1	3.6	0.3	0.5	10.8
	2	3.5	0.1	0.5	3.5
	4	4.1	0.1	0.1	4.1
	6	4.1	0.1	0.1	4.1
	9	4.1	0.1	0.1	4.1

Ruijin Hospital unpublished data





Summary

- **Exact pathogenic genes only account for small part of FSGS**
- **Next-gen sequencing helps to explore novel genes**
- **Mutation rate of *INF2* in Chinese familial FSGS is 3.6%**
- ***COL4A3* mutations account for 12.5% of AD FSGS**
- **Reclassification of Alport syndrom is suggested, “Collagen IV related nephropathy” ?**





Summary

➤ Heterogeneous clinical manifestations of FSGS

❑ FFSGS less response to treatment and a worse renal prognosis

➤ Treatment:

-lack of high-quality RCT study especially in Chinese patients

-steroid therapy could be considered for **moderate proteinuria**

-extend steroid treatment may not be suggested

-rituximab is promising but need more data





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