

肾脏疾病诊断和疗效预测的生物标志物



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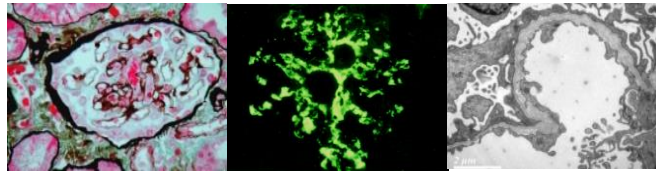
国家肾脏疾病临床研究中心

全军肾脏病研究所

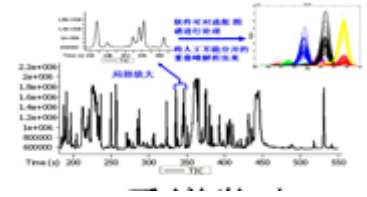
肾脏疾病的诊断和治疗



临床综合症诊断



组织形态学诊断

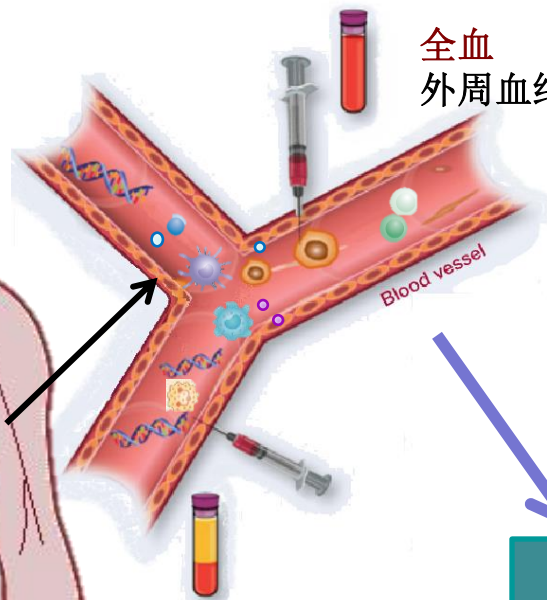
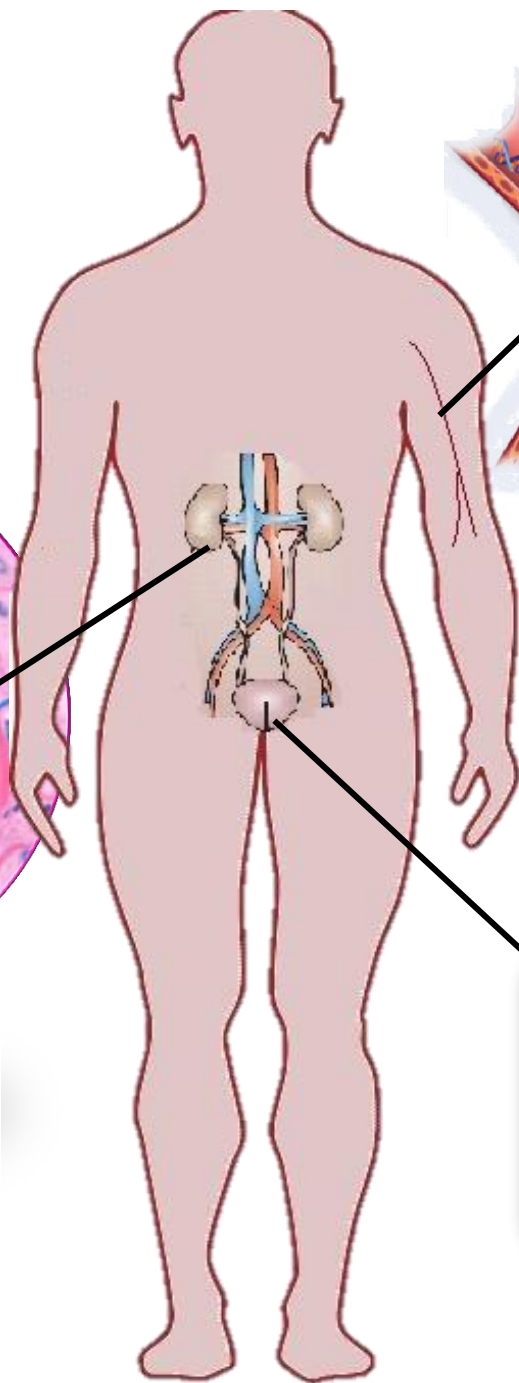
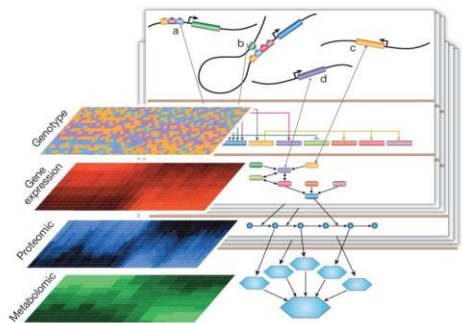
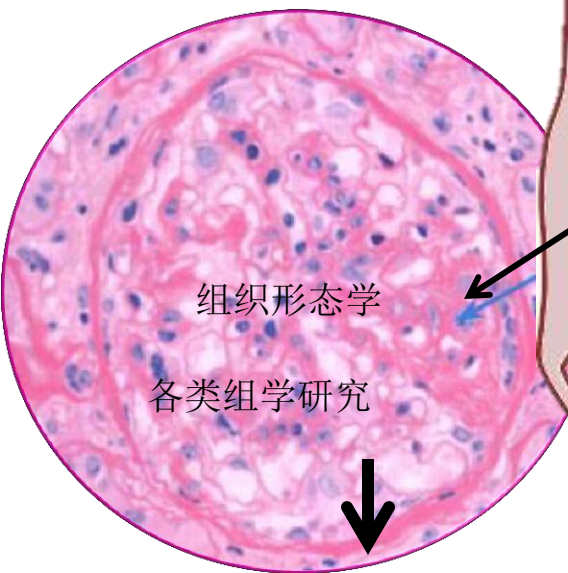


分子表型诊断

病因诊断，分子分型，基于发病机制的靶向治疗

精准医学
(Precision Medicine)

肾活检 Renal Biopsy

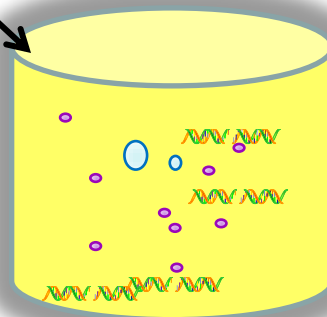


全血
外周血细胞成分分析

血浆
循环miRNA/外泌体等

体液活检 Biofluid Biopsy

非创伤性
连续动态监测

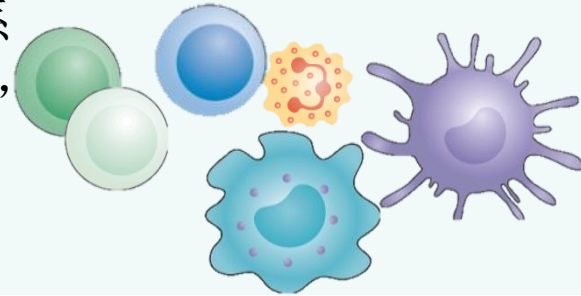


尿液
miRNA/蛋白质/外泌体等

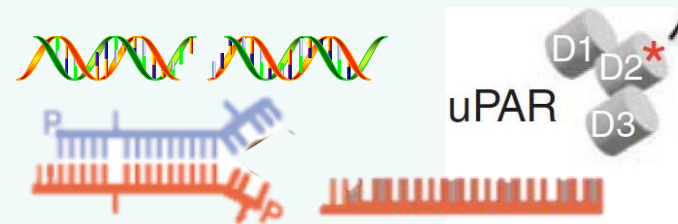
体液中的生物标志物

Biofluid-based Biopsies

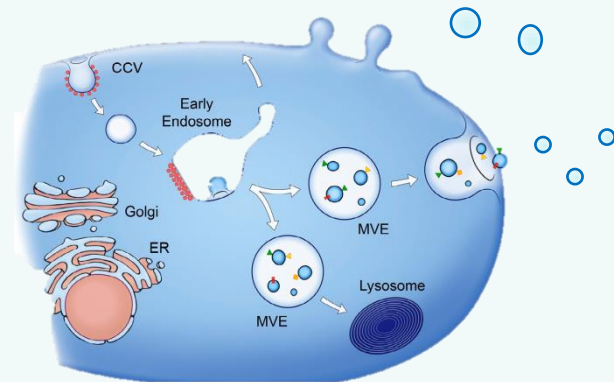
细胞：例如循环肿瘤细胞，髓系来源抑制性细胞，调节性T细胞，T细胞，B细胞

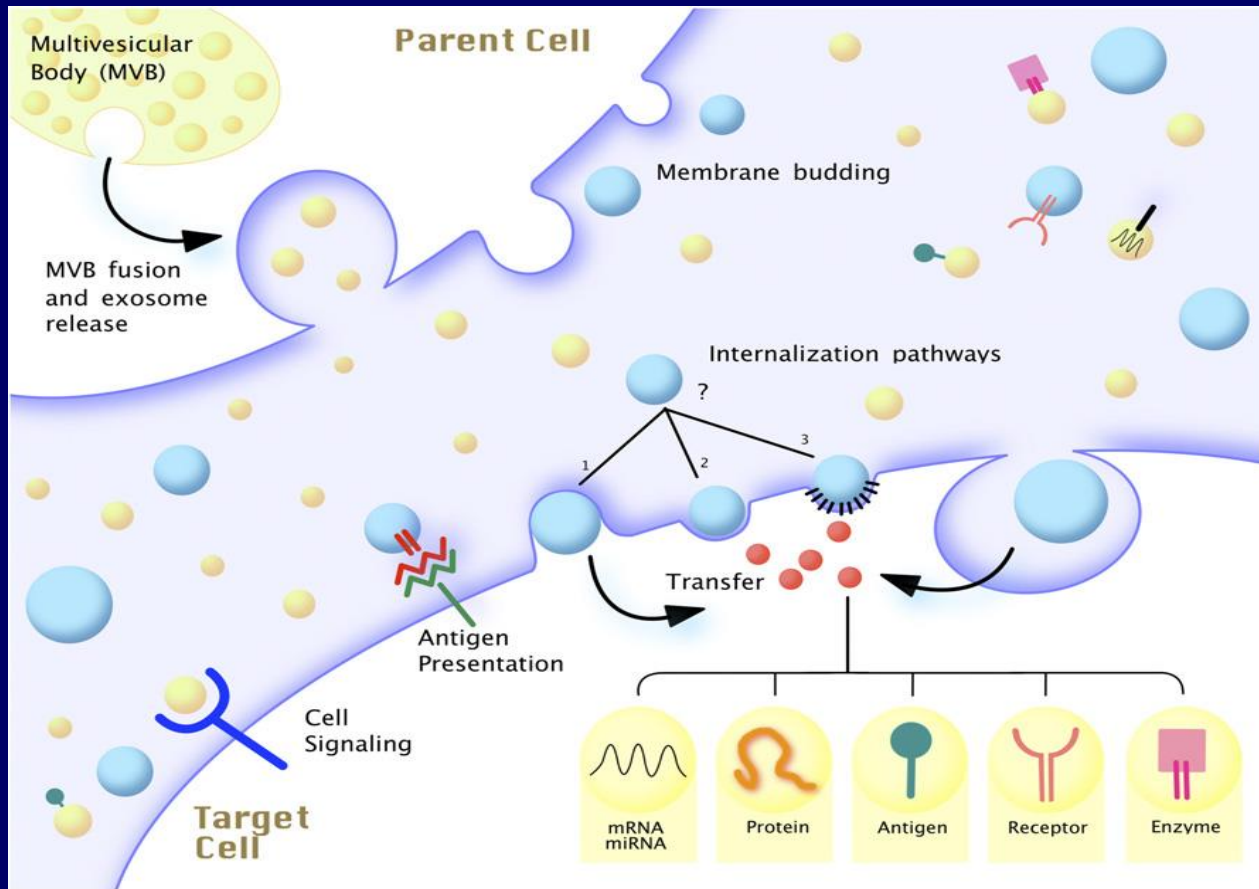


循环中的核苷酸，microRNA
蛋白质，酶，膜受体等



外泌体(Exosome): 装
载miRNA、DNA膜受
体、酶等





miRNAs对维持肾脏的生理和病理过程至关重要。循环和尿液miRNAs可能作为一种无创的生物标志物帮助诊断和监测疾病。

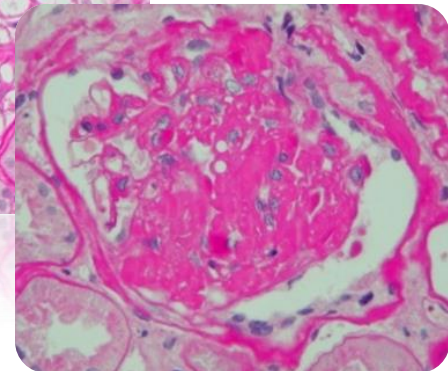
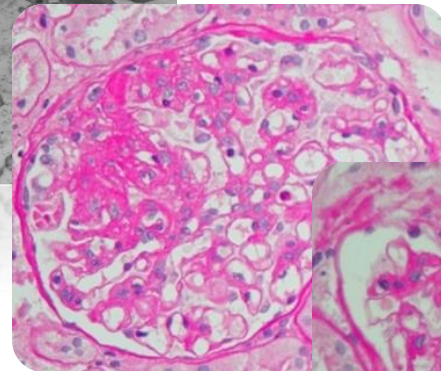
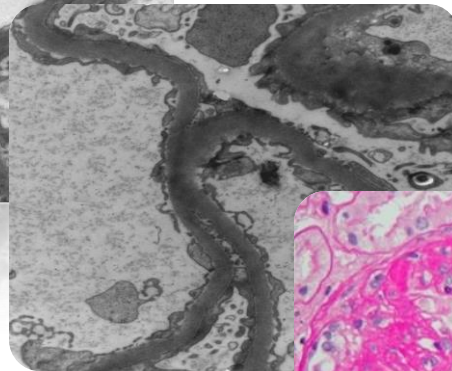
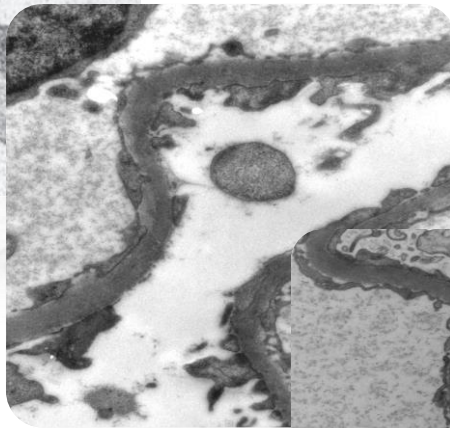
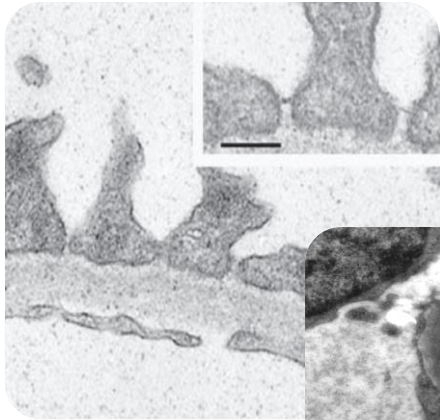
原发性肾小球疾病

(18813 cases, 2000-2010, 南京)

	n	%
IgAN	8580	45.61
FSGS	2400	12.87
MN	2422	12.76
IgMN	240	1.28
MPGN	242	1.29
EnPGN	213	1.13

FSGS accounts for 20%–25% of adult patients undergoing biopsy for evaluation of idiopathic GN in western countries.

足细胞损伤：FSGS的中心环节



早期肾小球硬化

FSGS 病因分类

Primary alterations of glomerular epithelial cell

- Primary (idiopathic) FSGS

- Viral diseases (HIV-associated nephropathy, parvovirus B19, hepatitis C)

- Drugs (heroin, pamidronate, lithium, anabolic steroids)

- Genetic disorders (podocin, α -actinin 4, transient receptor potential action channel 6)

 - Familial

 - Sporadic

Secondary to reduced nephron mass/glomerular adaptations

- Reflux nephropathy

- Renal dysplasia

- Oligomeganephronia

- Obesity-related glomerulopathy

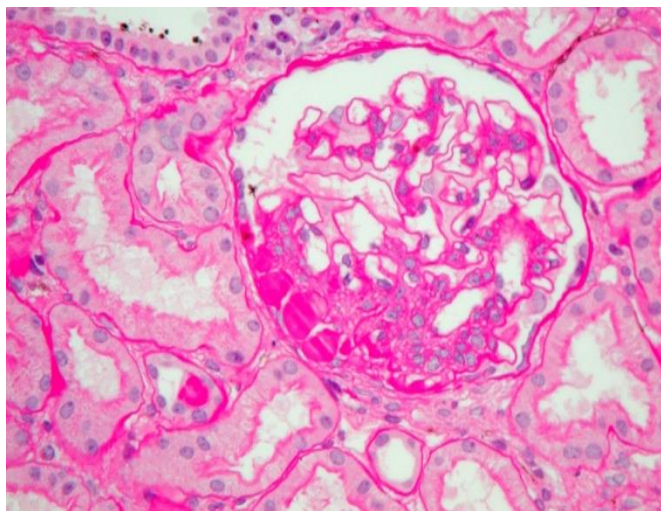
- Sickle cell disease

- Primary glomerular diseases

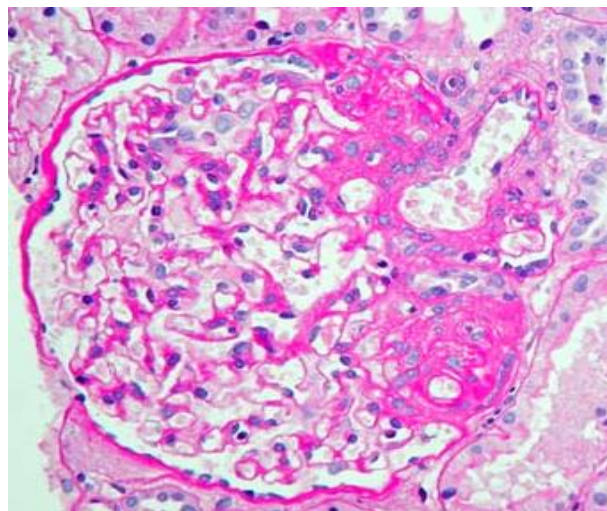
Secondary to focal proliferative GN

Secondary to hereditary nephropathies (Alport syndrome)

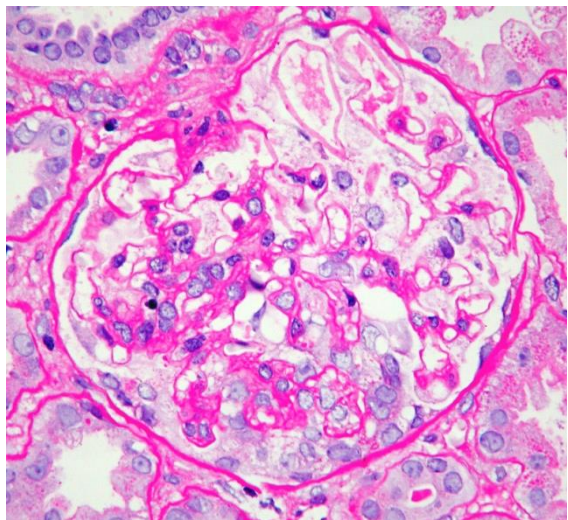
经典型



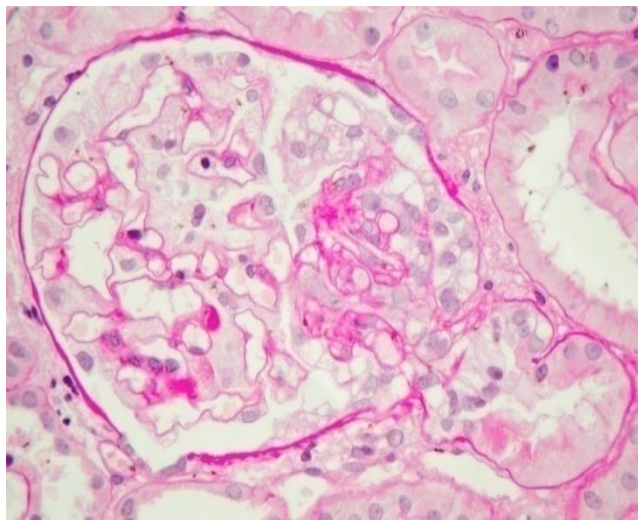
门周型



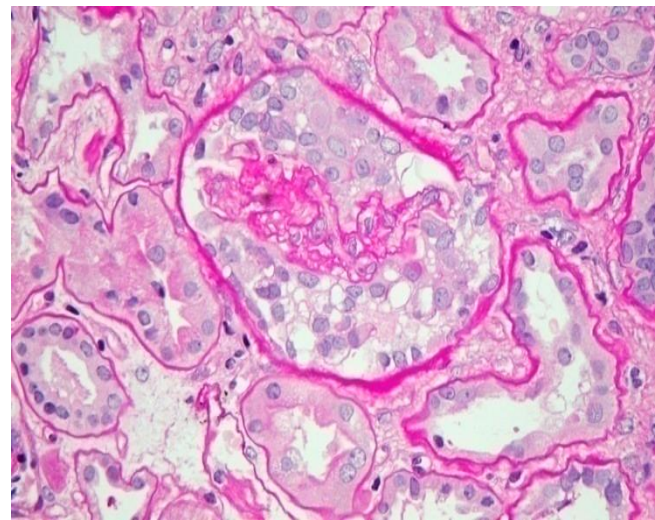
FSGS的病理分型



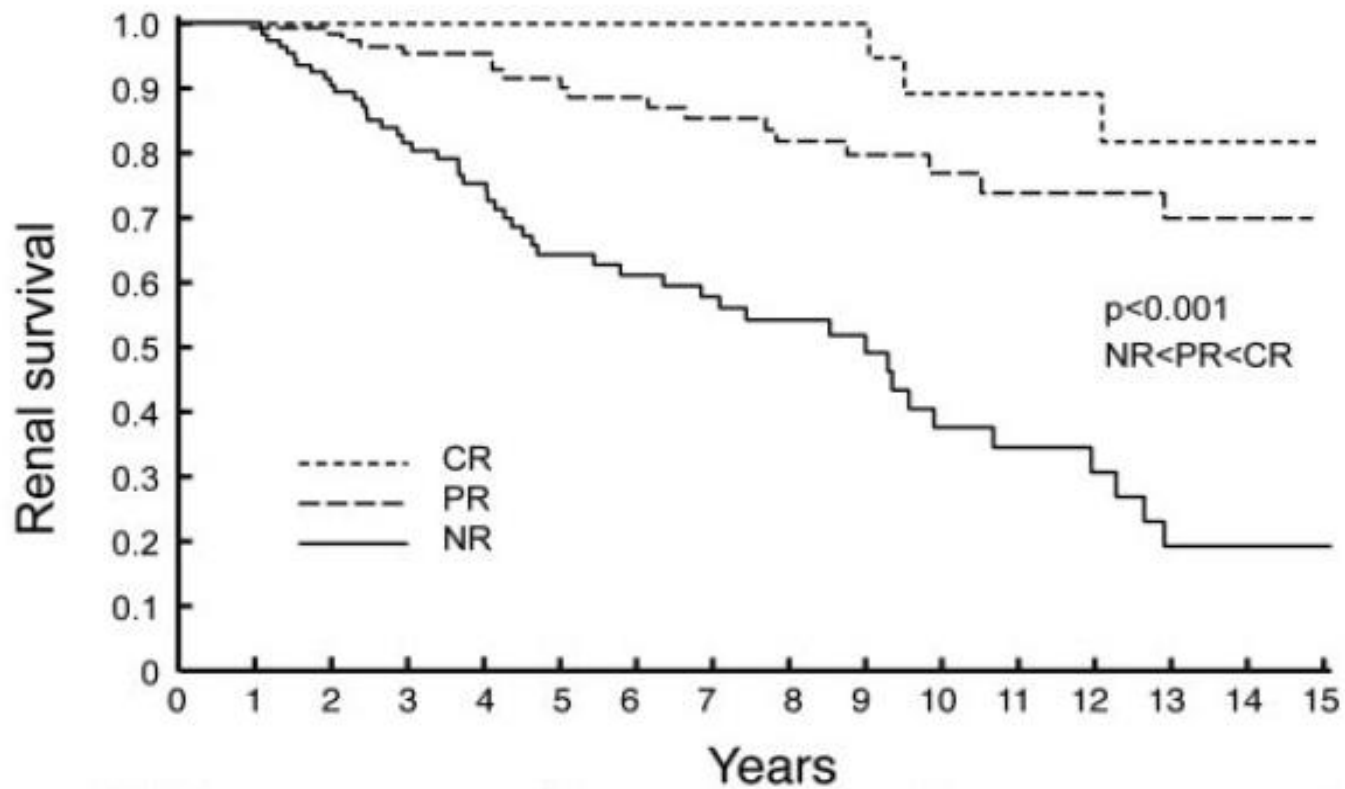
细胞型



顶部型



塌陷型



治疗后完全缓解的患者预后明显优于无缓解的患者，即使部分缓解者也呈现较好的预后。

生物标记物 (Biomarkers)

Detection of disease

Monitor of the disease in

Severity

Progression

Regression

Predict response to therapy

生物标记物 (Biomarkers)

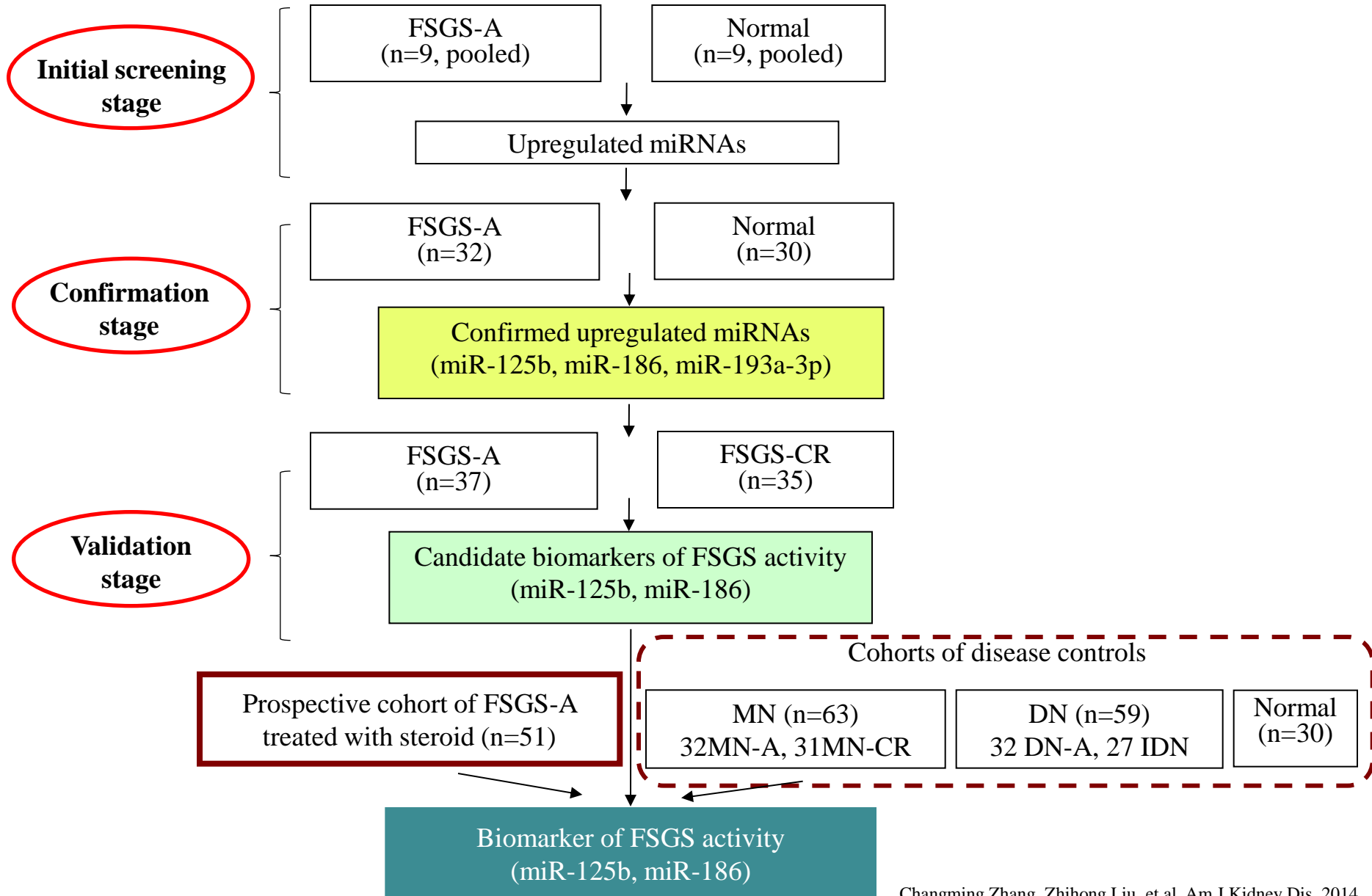
从哪里来? (Donor)

到哪里去? (Recipient)

从血浆 miRNAs 中寻找 FSGS 标志物



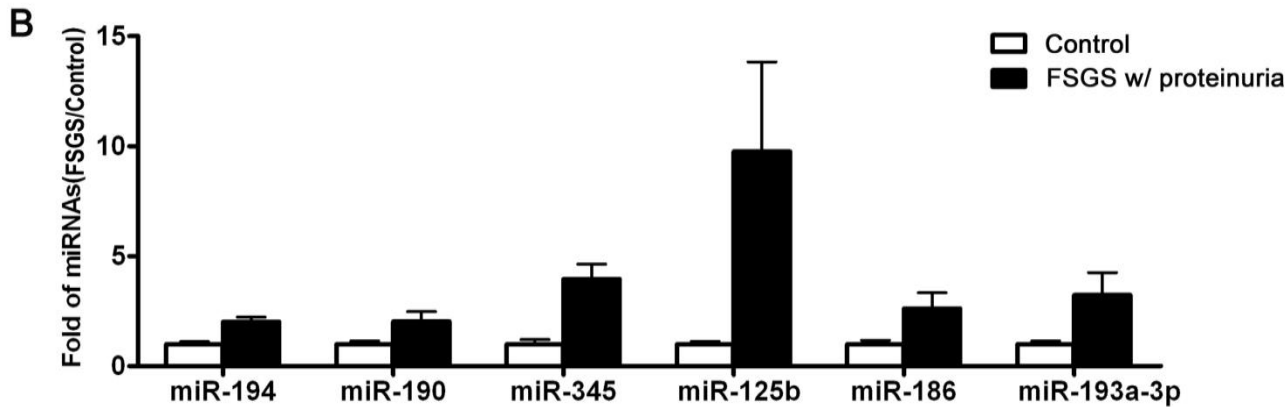
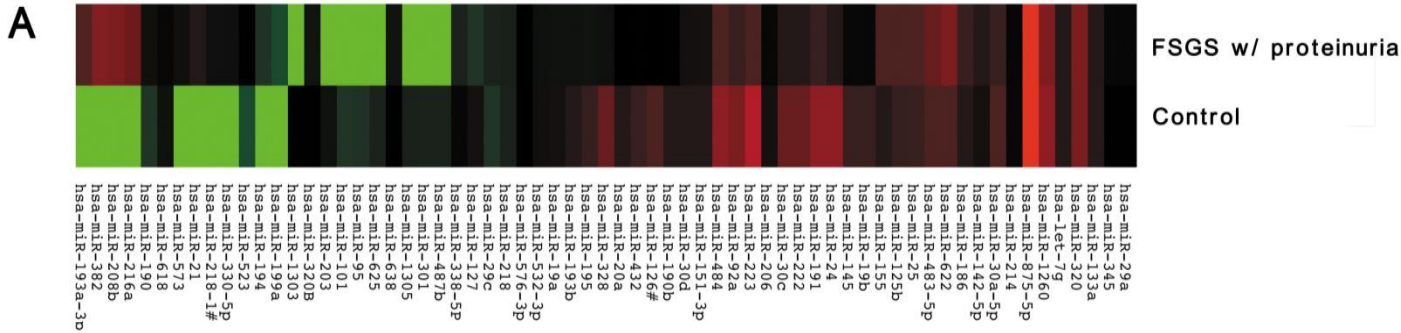
研究技术路线



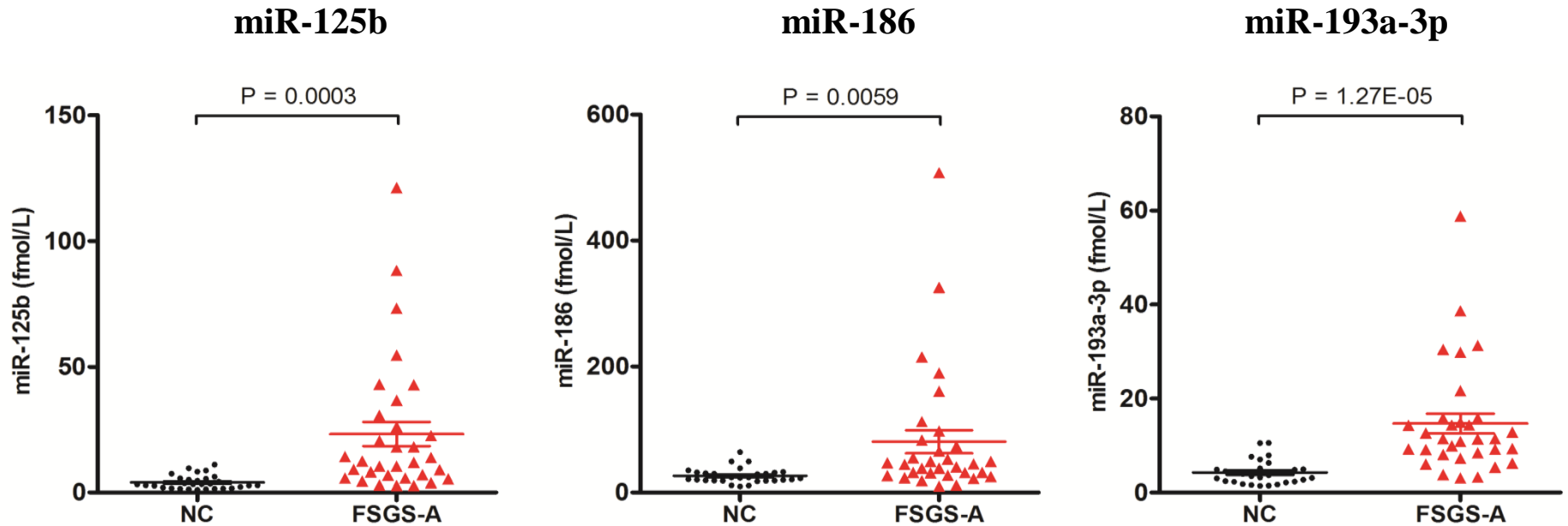
寻找并验证在活动性FSGS患者中升高的血浆miRNAs

The FSGS with nephrotic proteinuria had a plasma miRNA profile that was different from controls (miRNAs 45)

Individual qRT-PCR analyses confirmed the upregulation of 6 miRNAs in FSGS.

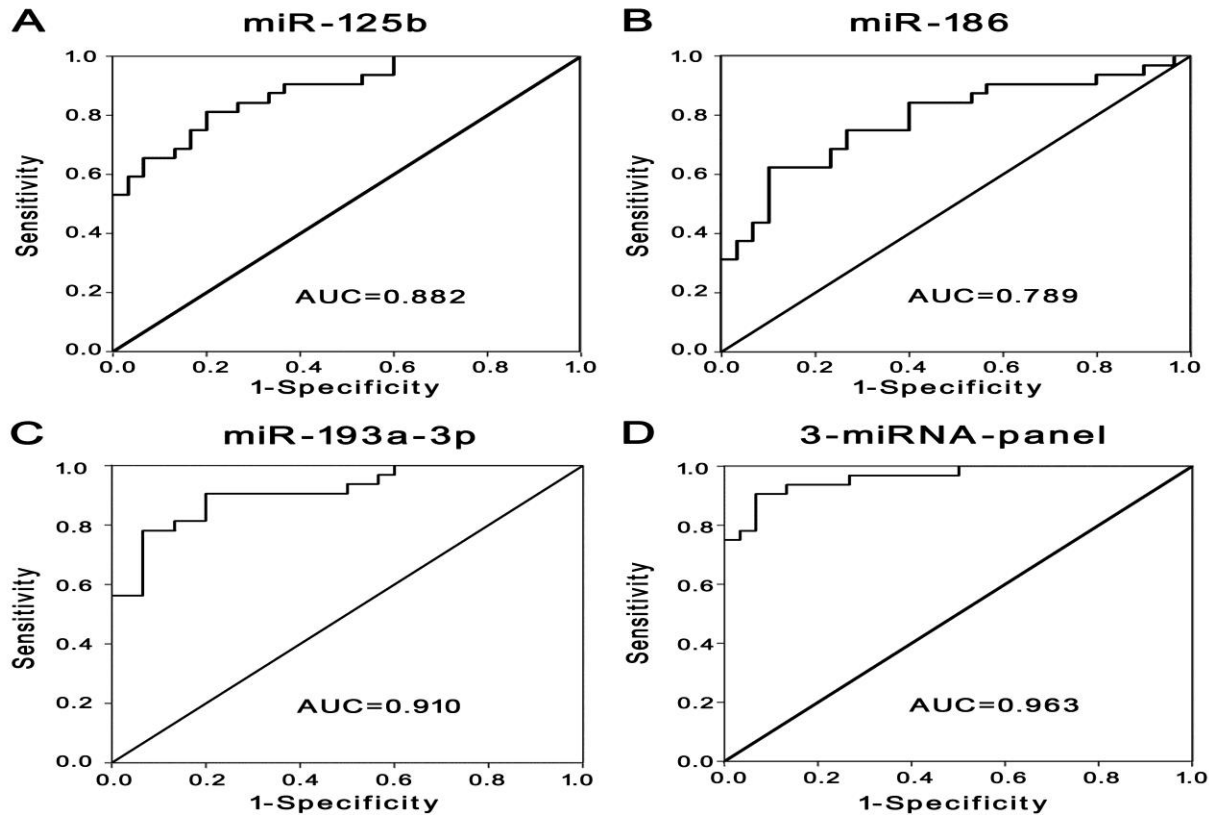


独立队列样本验证中血浆miRNAs的表达



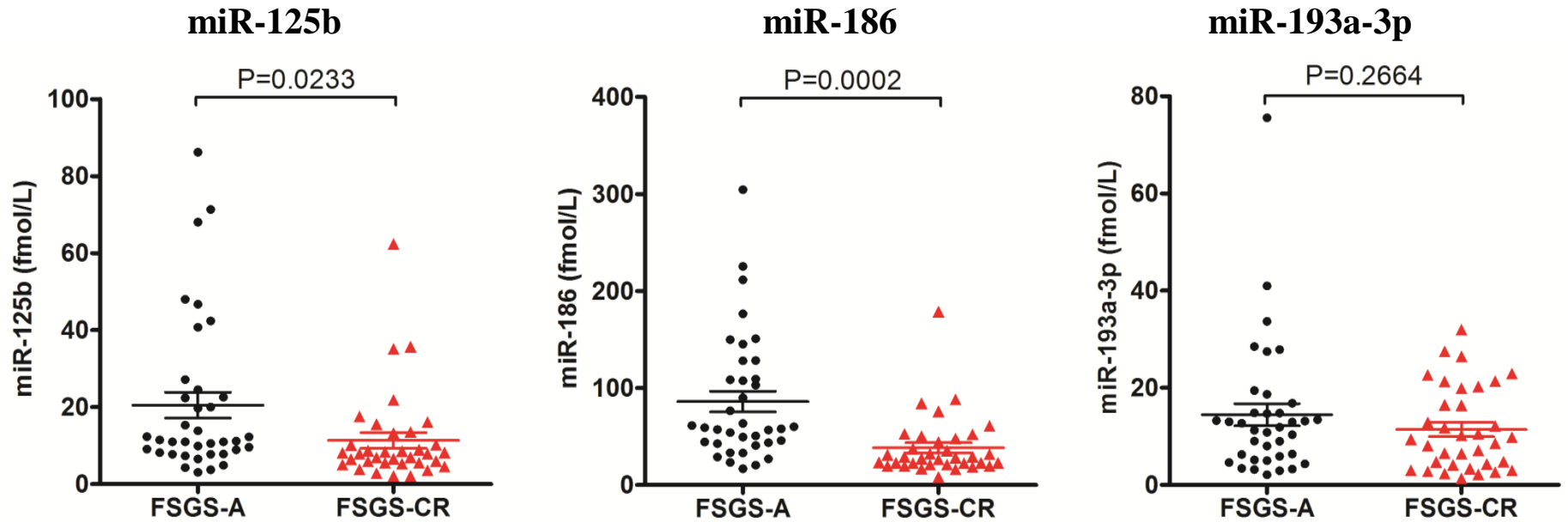
活动性FSGS患者血浆miR-125b, miR-186和miR-193a-3p较正常对照明显升高

血浆miR-125b, miR-186和miR-193a-3p区分活动性FSGS和正常对照的价值



三种miRNAs均可用于区分活动性FSGS和正常对照，三者联合则显示更好的预测效力。

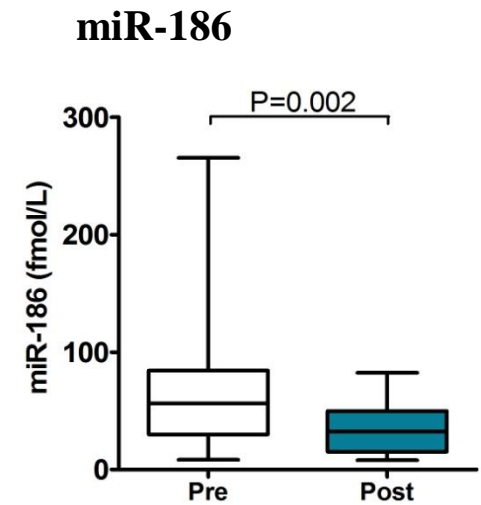
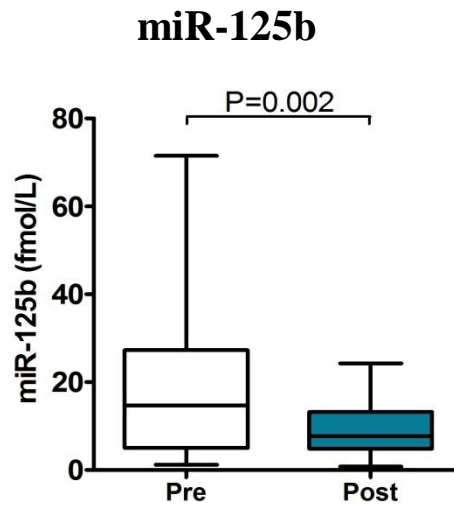
不同疾病状态下FSGS患者血浆miR-125b, miR-186和miR-193a-3p表达



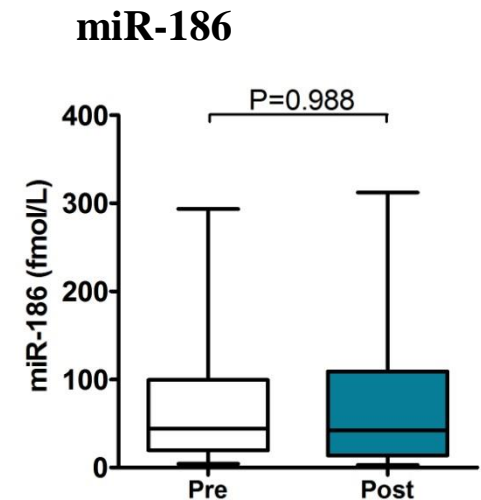
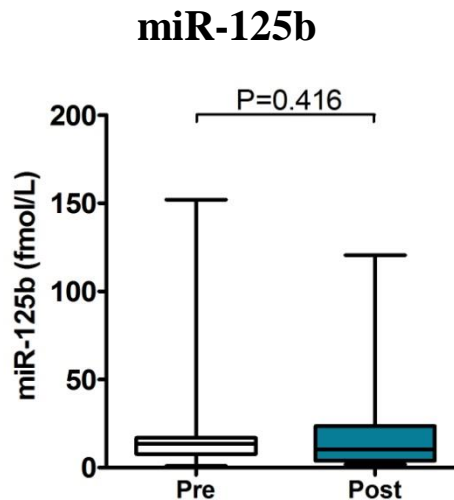
活动性FSGS患者血浆miR-186, miR-125b水平显著高于完全缓解FSGS

FSGS患者接受激素治疗前后血浆miR-125b, miR-186水平的变化

**Steroid-sensitive
(n=29)**

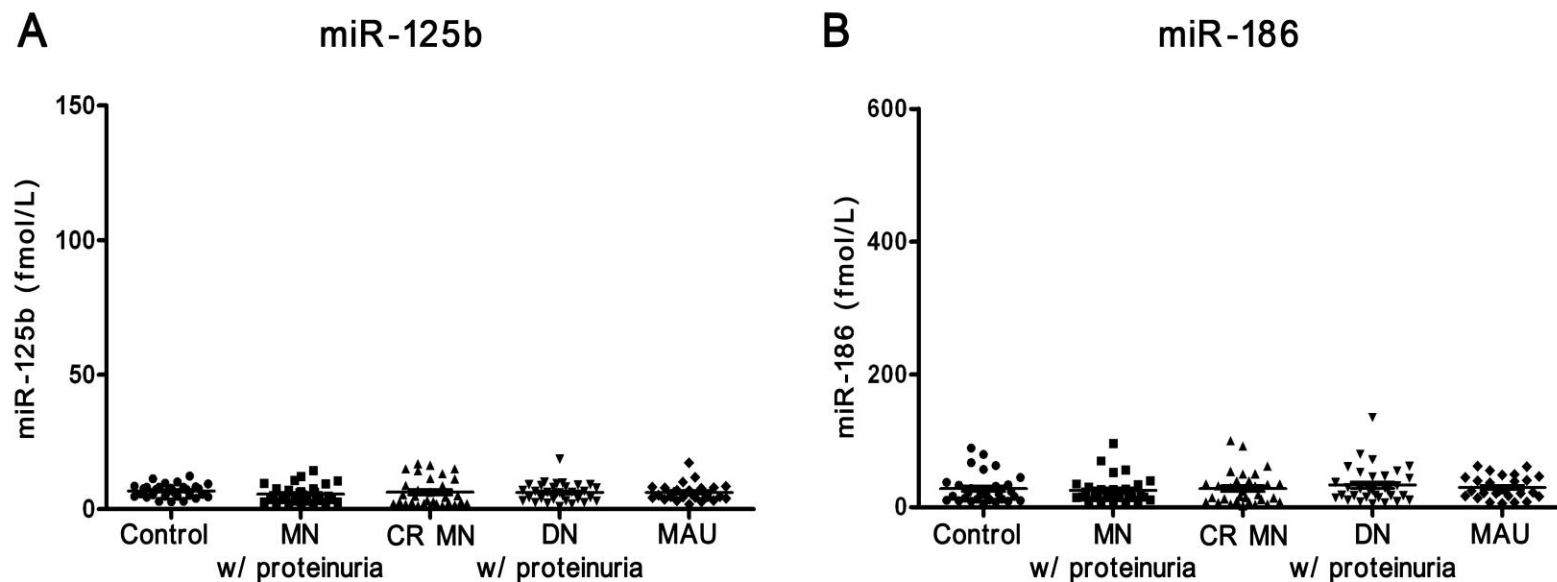


**Steroid-resistant
(n=21)**



激素敏感的FSGS患者血浆miR-125b和miR-186水平在治疗后显著下降

膜性肾病和糖尿病肾病患者血浆miR-125b和miR-186表达

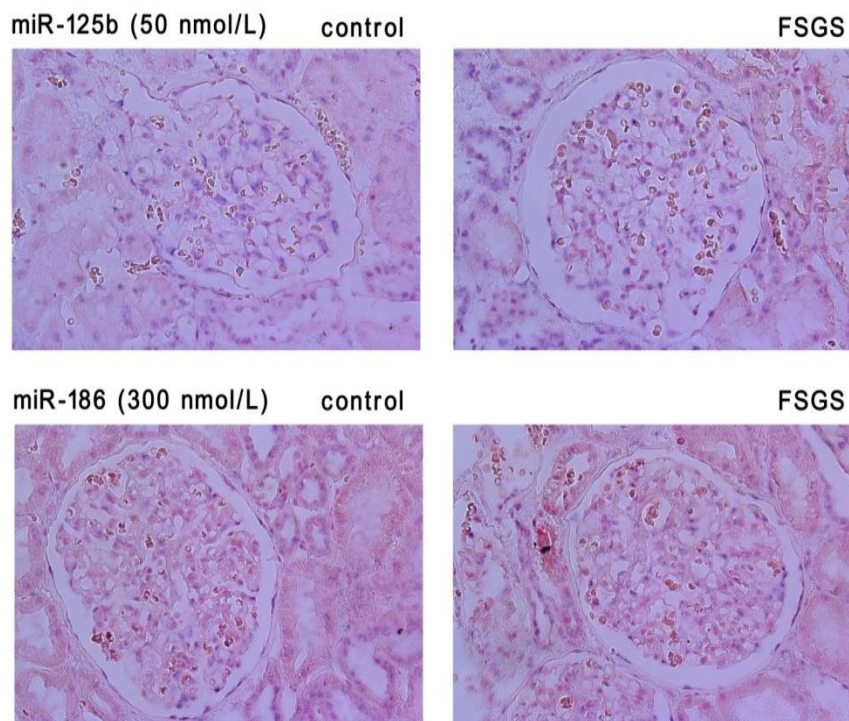
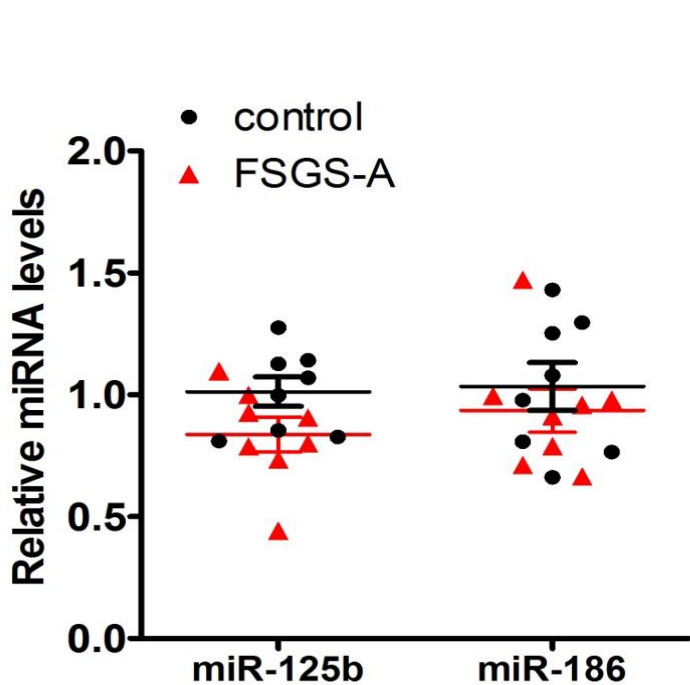


不论蛋白尿水平如何，MN和DN患者血浆miR-125b和miR-186水平与正常对照相比均无明显变化

血浆miR-186水平与FSGS疾病活动密切相关。激素敏感患者，血浆miR-125b和miR-186水平在治疗后较治疗前显著下降，激素抵抗患者则不然，表明二者可能作为监测FSGS疾病活动和预测激素疗效的生物标志物。

FSGS患者血浆miR-125b和miR-186变化的意义

FSGS患者肾组织miR-125b和miR-186表达



qRT-PCR及原位杂交结果均显示：与正常对照相比，FSGS患者肾组织miR-125b和miR-186表达均无上调。

PATHOGENESIS OF LIPOID NEPHROSIS: A DISORDER OF T-CELL FUNCTION

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Washington D.C. 20422, U.S.A.*

Summary

Clinical observations suggest that lipoid nephrosis is produced by a systemic abnormality of T-cell function resulting in the secretion of a circulating chemical mediator toxic to an immunologically innocent glomerular basement membrane. The lack of evidence of a humoral antibody response, remission induced by measles which modifies cell-mediated immunity, the therapeutic benefits of steroids and cyclophosphamide which also abate cell-mediated responses, and the occurrence of this syndrome in Hodgkin's disease support this hypothesis. The susceptibility of untreated patients to pneumococcal infections may be of primary or secondary pathogenetic importance. Taken together, the data suggest that this syndrome is a clinical expression of a self-limited primary immune-deficiency disease. ^a

The Lancet

September 7, 1974

Role of Myeloid-Derived Suppressor Cells in Glucocorticoid-Mediated Amelioration of FSGS

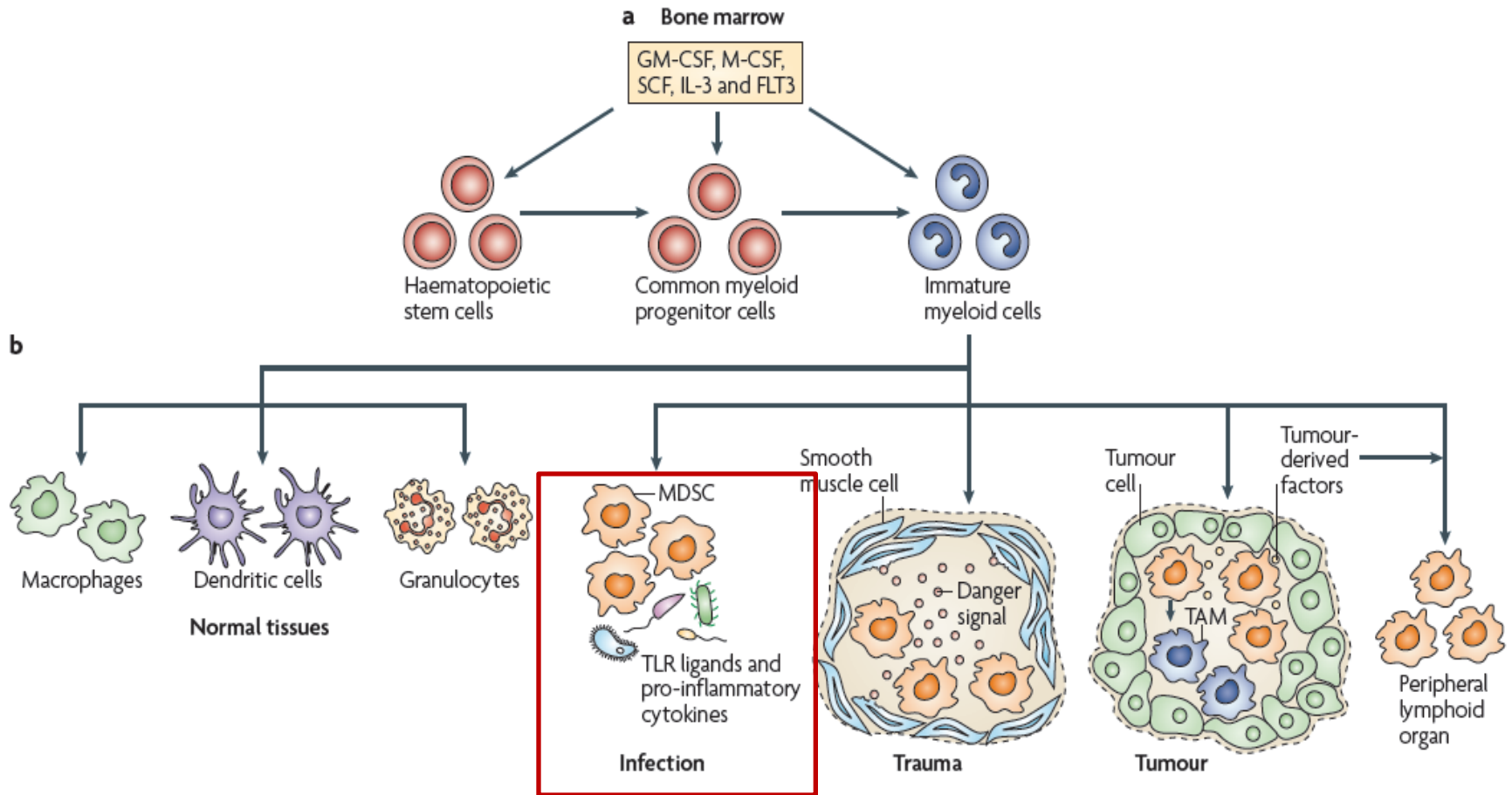
Limin Li,* Tao Zhang,[†] Wenli Diao,* Fangfang Jin,* Lei Shi,*[‡] Jiao Meng,[†] Huan Liu,*
Jing Zhang,* Cai-Hong Zeng,[†] Ming-Chao Zhang,[†] Shaoshan Liang,[†] Yuan Liu,[‡]
Chen- Yu Zhang,* Zhihong Liu,[†] and Ke Zen*[†]

*State Key Laboratory of Pharmaceutical Biotechnology, Jiangsu Engineering Research Center for MicroRNA Biology, Nanjing, China; [†]National Clinical Research Center of Kidney Diseases, Jinling Hospital, Nanjing University School of Medicine, Nanjing, China; and [‡]Center for Inflammation, Immunity and Infection, Department of Biology, Georgia State University, Atlanta, Georgia

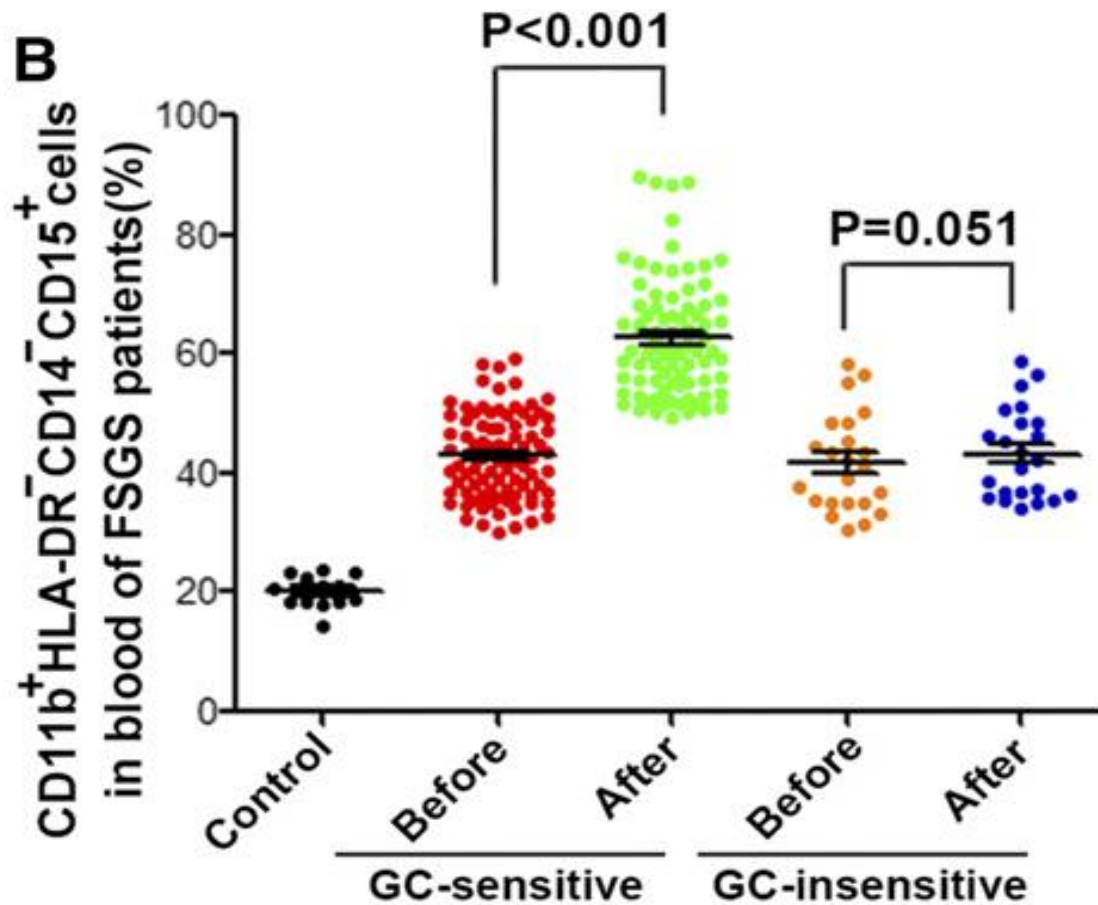
ABSTRACT

These rapidly induced MDSCs after steroid treatment would suppress the proliferation and activation of T cells and thus attenuate the inflammation and renal injury in steroid sensitive patients.

Myeloid-derived suppressor cells (MDSCs)

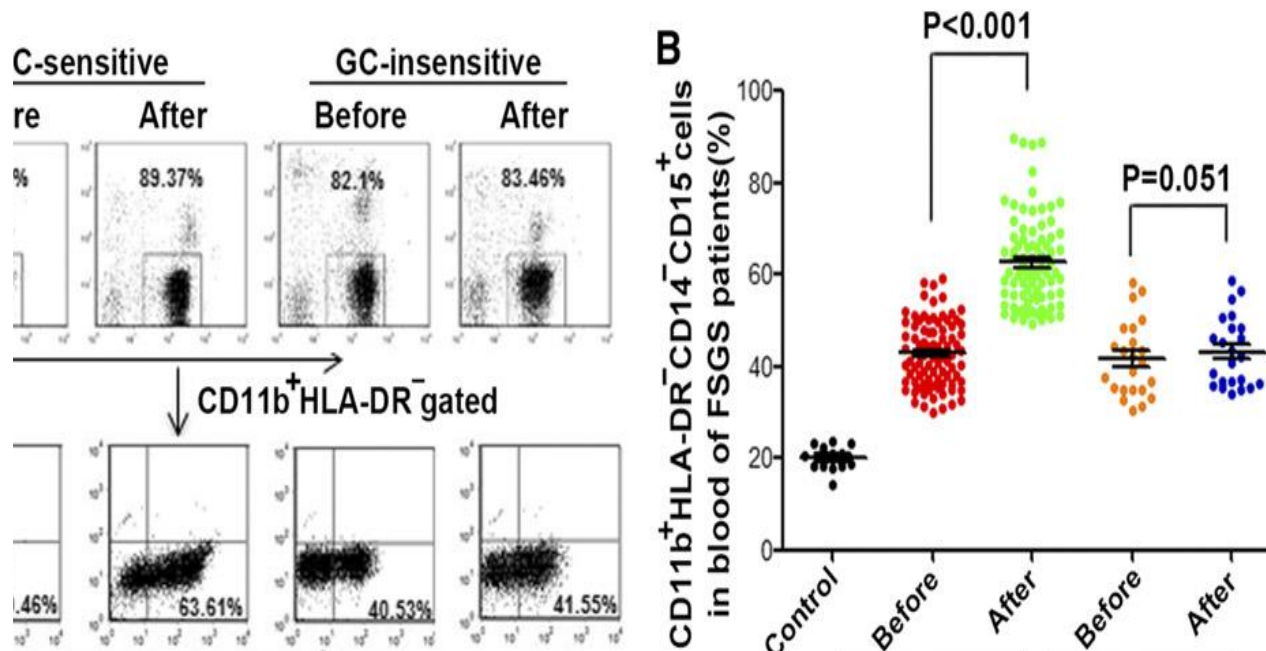


➤ MDSC, a heterogeneous population of immature myeloid cells, which have an ability to suppress T cell responses.

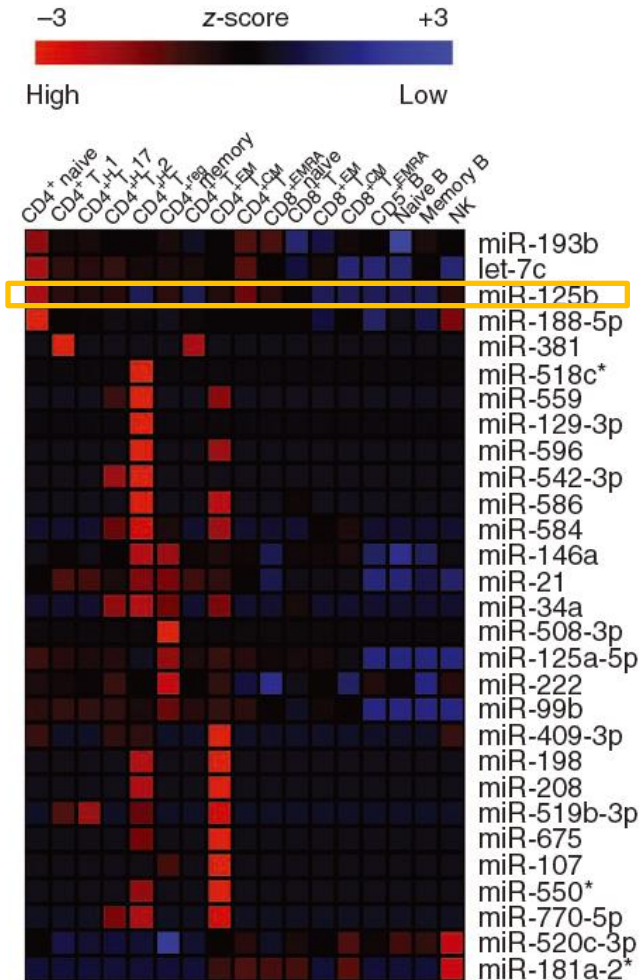


激素敏感FSGS患者治疗后MDSC增多，激素抵抗患者则不然。

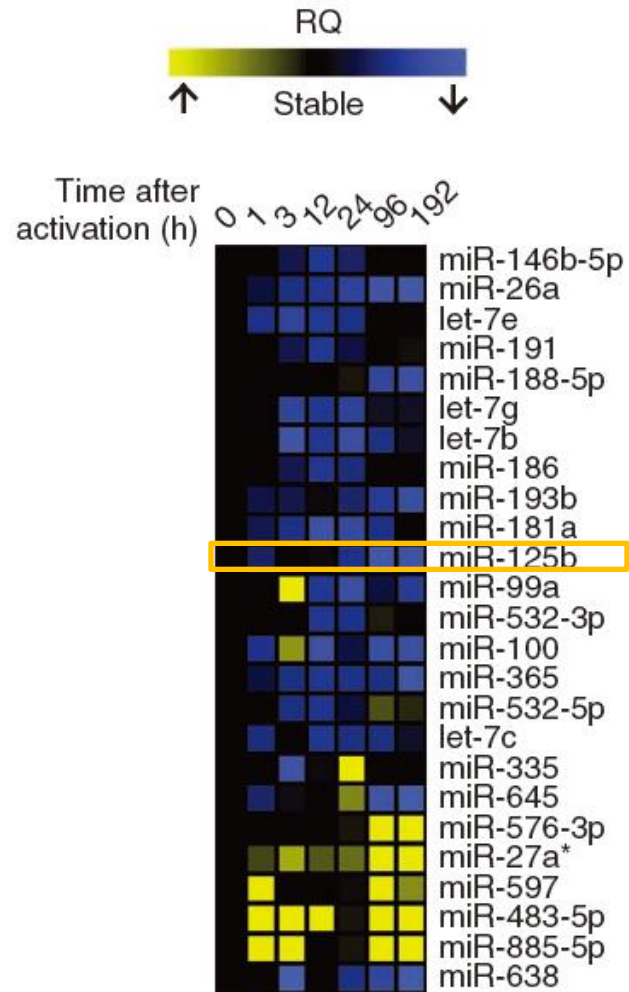
激素敏感FSGS患者MDSC能显著抑制T细胞增殖



MiR-125b特异性的表达于静息CD4⁺ T细胞

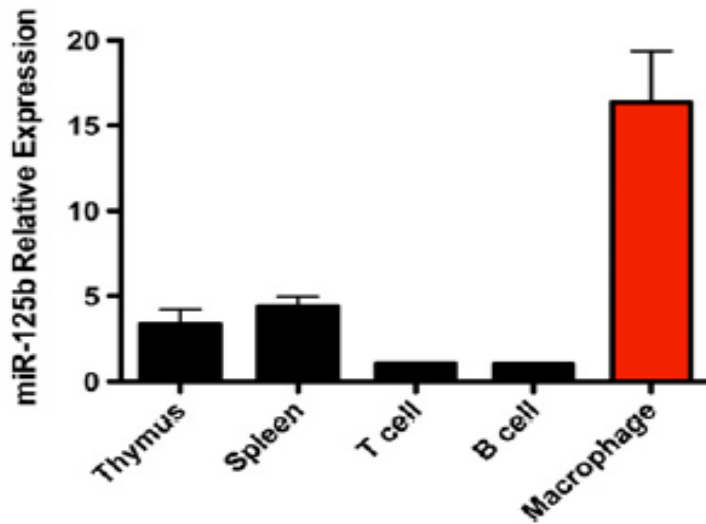


miR-125b as signature of human lymphocyte subsets , selectively expressed in naive CD4⁺ T cells



miR-125b with high expression specifically in resting naive CD4⁺ T cells were downregulated after activation with CD3 and CD28

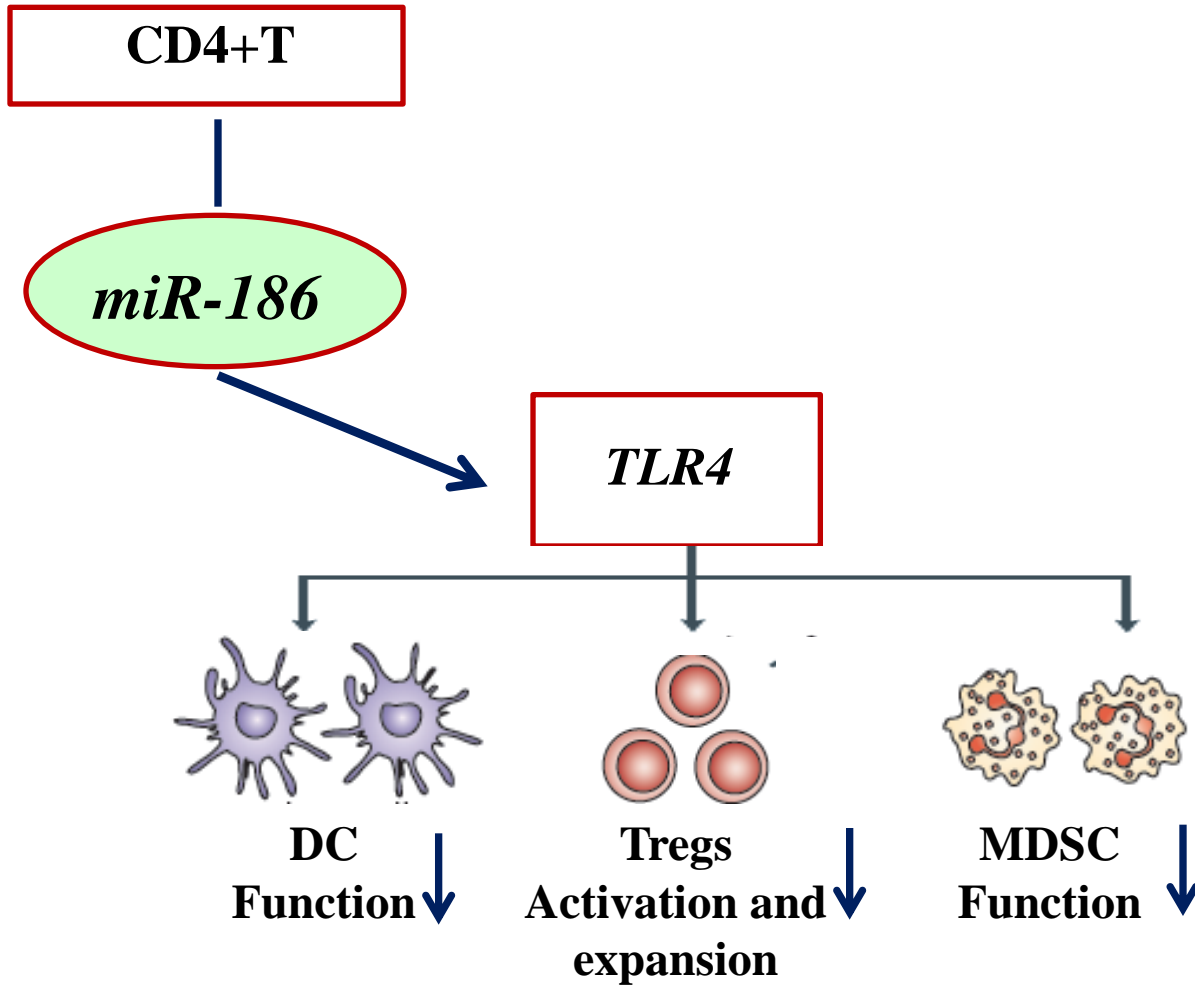
巨噬细胞富含MiR-125b



miR-125b level in total splenocytes, thymocytes, splenic T cells, splenic B cells, and peritoneal macrophages from C57BL/6 mice.

miR-125b is enriched in macrophages compared with lymphoid cells and whole immune tissues. Enforced expression of miR-125b induces increased costimulatory factor expression and elevated responsiveness to IFN-g

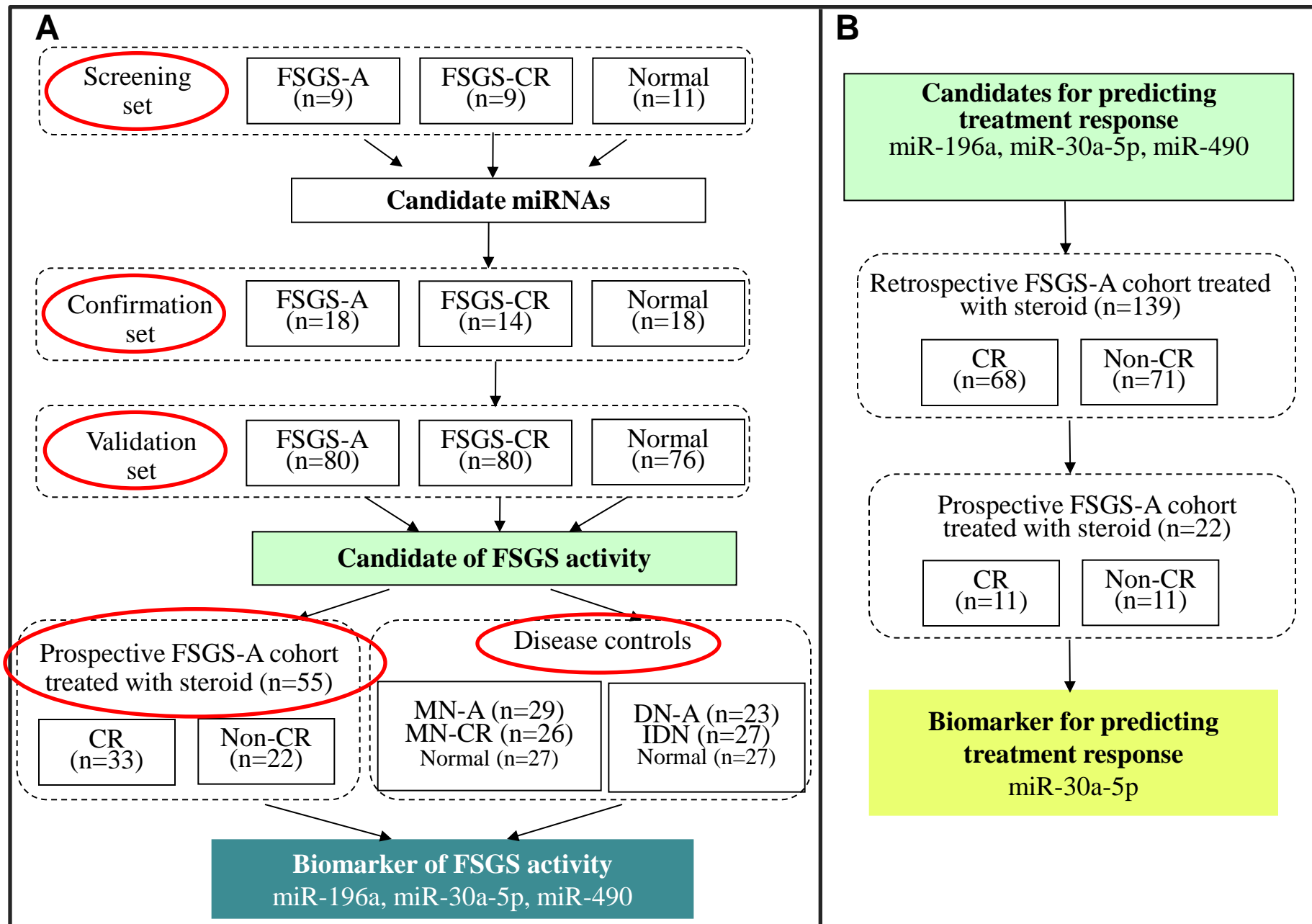
miR-125b能促进巨噬细胞的抗原提呈和诱导T细胞活化，增强巨噬细胞介导的获得性免疫。活化的T细胞又能分泌**IFN-g**，进一步增强巨噬细胞的活化。



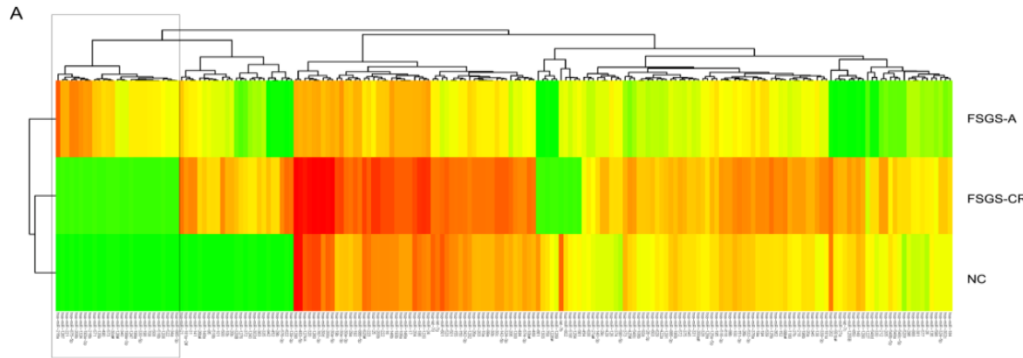
从尿液 miRNAs 中寻找 FSGS 标志物



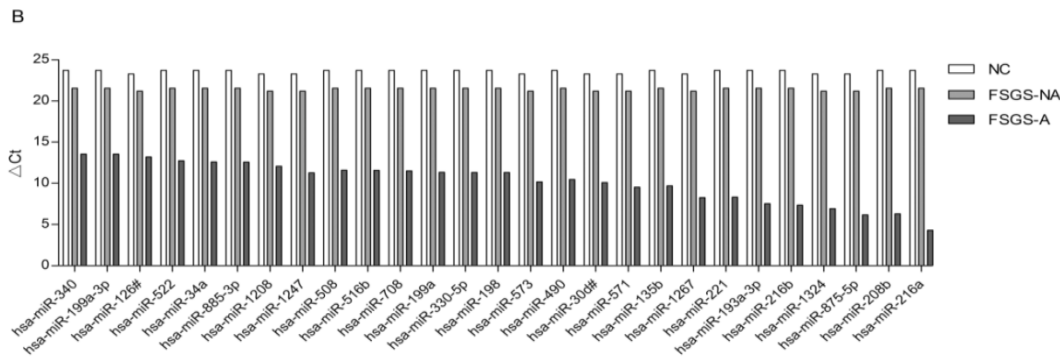
研究技术路线



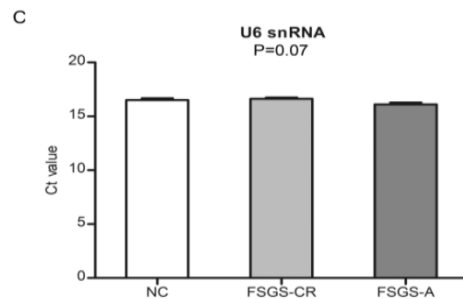
筛查候选 miRNAs



A distinct miRNA profiles among patients with FSGS-A, FSGS- CR, and normal controls



Individual qRT-PCR analyses confirmed the upregulation of 27 miRNAs in FSGS-A than that from patients with FSGS-CR and controls



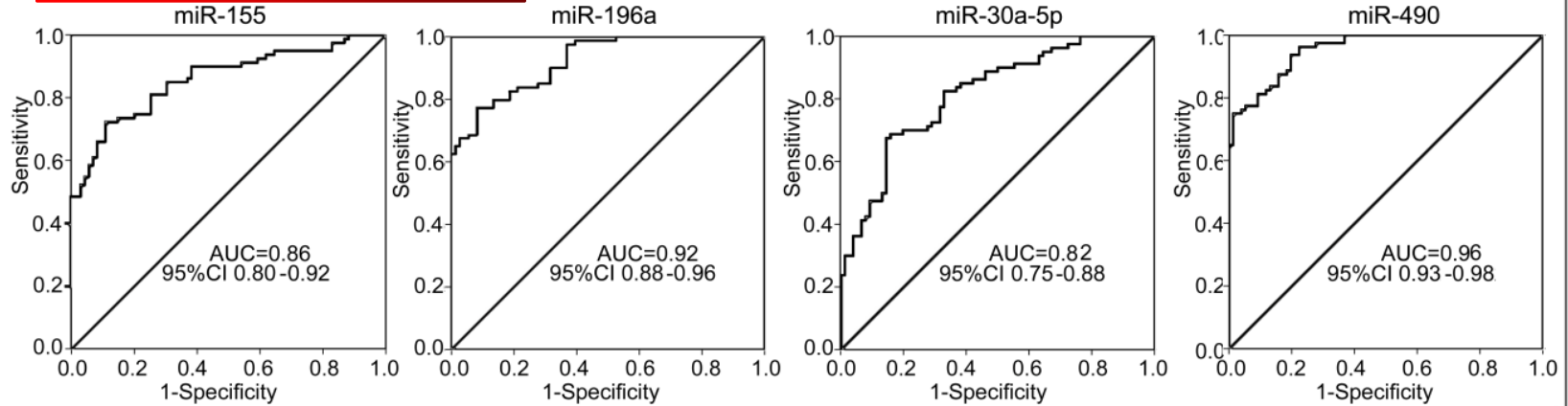
独立队列验证FSGS患者尿液miRNAs表达

microRNAs	Concentration of Urinary miRNAs (fmol/ L)				Fold Change for FSGS-A versus Controls		Fold Change for FSGS-A versus FSGS-CR		
	Controls	FSGS-CR	FSGS-A	P Value	Mean	P Value	Mean	P Value	
Confirmation set									
hsa-miR-135b	46.56 45.1	48.86 28.4	1066 85.6	0.01	2.30	0.004	2.22	0.01	
hsa-miR-155	25906 1970	69906 5750	13,3006 11,100	0.001	5.14	, 0.001	1.91	0.04	
hsa-miR-196a	2086 99.3	1906 80.6	13506 1040	0.01	6.51	0.004	7.11	0.01	
hsa-miR-30a-5p	2816 196	3976 164	8556 673	0.001	3.05	, 0.001	2.15	0.01	
hsa-miR-320	2386 123	7636 615	15906 1060	, 0.001	6.70	, 0.001	2.09	0.004	
hsa-miR-490	33.16 28.3	27.56 12.5	1076 96.0	0.03	3.24	0.03	3.89	0.03	
Validation set									
hsa-miR-135b	1056 159	1176 152	1406 165	0.70		0.43		0.58	
hsa-miR-155	23806 2080	49206 4770	12,3006 10,900	, 0.001	3.33	, 0.001	2.49	, 0.001	
hsa-miR-196a	1656 140	2056 195	12506 1090	, 0.001	6.74	, 0.001	6.08	, 0.001	
hsa-miR-30a-5p	4896 326	3906 328	12306 1110	, 0.001	2.81	, 0.001	3.15	, 0.001	
hsa-miR-320	19806 1890	19606 1950	20606 2280	0.27		0.11		0.17	
hsa-miR-490	42.96 33.0	69.36 56.0	2626 219	, 0.001	4.68	, 0.001	3.80	, 0.001	

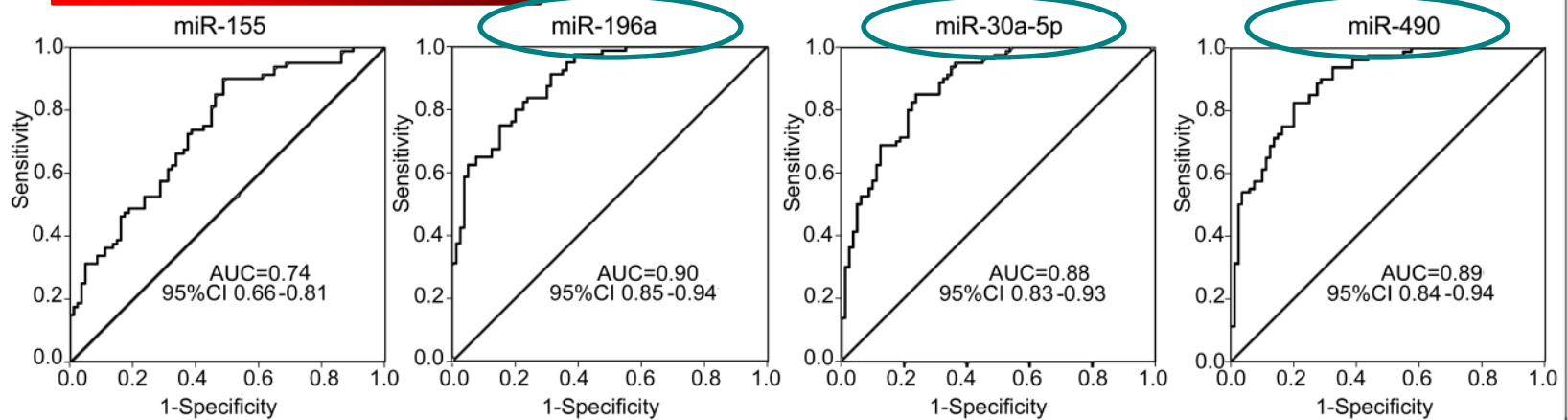
与正常对照和完全缓解FSGS患者相比，活动性FSGS患者尿液miR-30a-5p, miR-196a, miR-155和miR-490水平显著升高。

尿液miR-196a, miR-30a-5p和miR-490与FSGS疾病活动

A Discriminating FSGS-A from NC

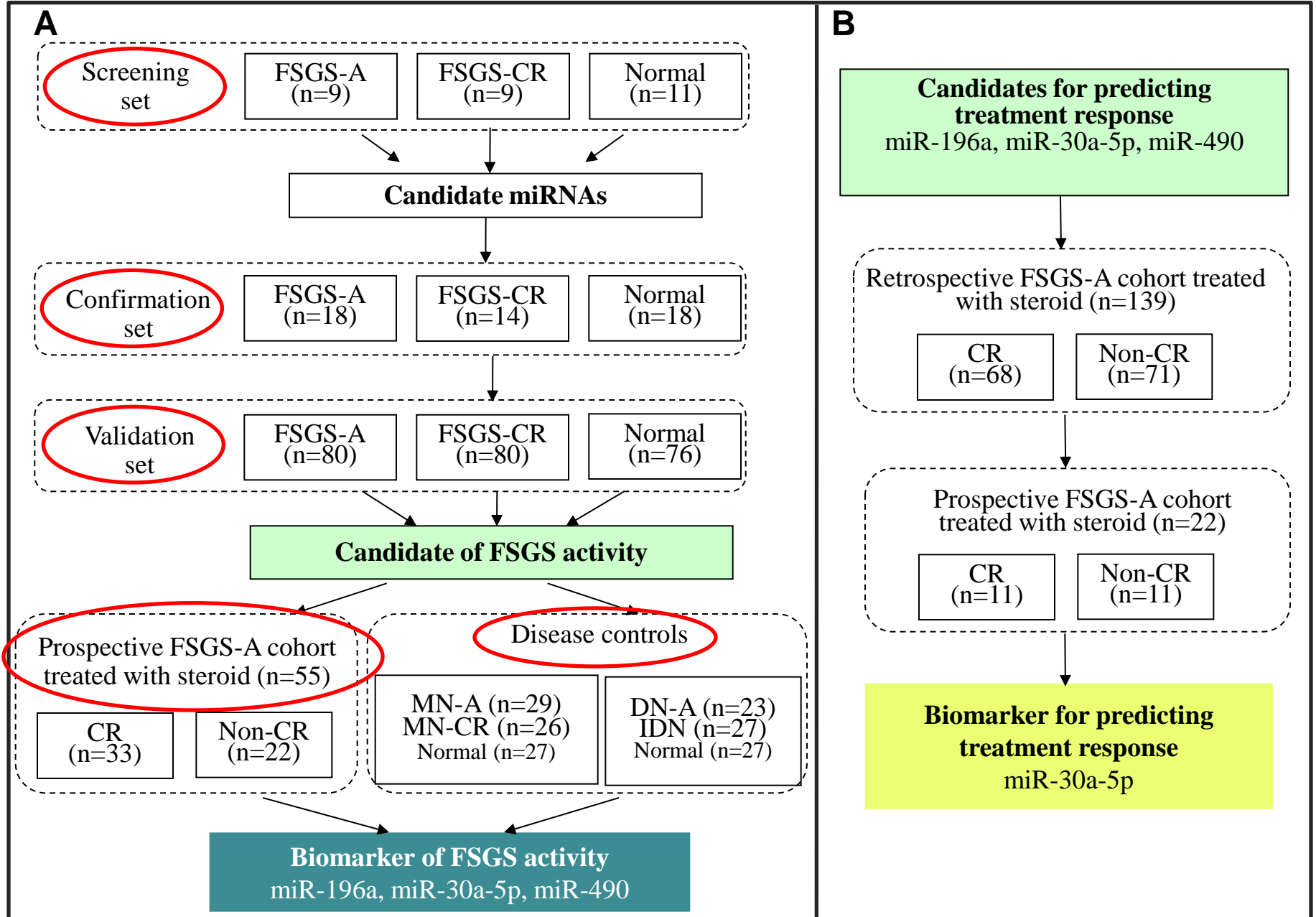


B Discriminating FSGS-A from FSGS-CR

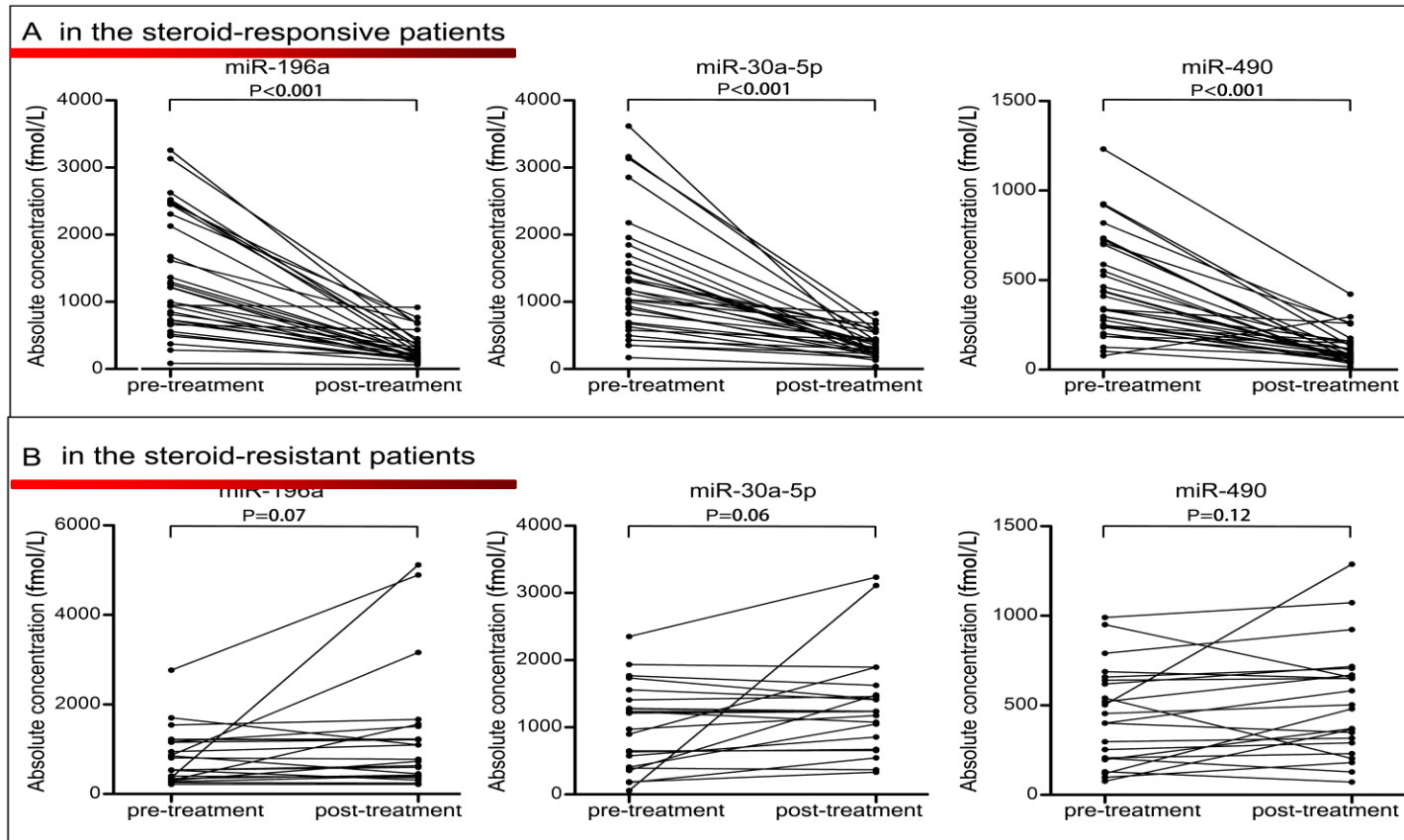


尿液miR-196a, miR-30a-5p and miR-490能很好的区分活动性FSGS和完全缓解FSGS

研究技术路线



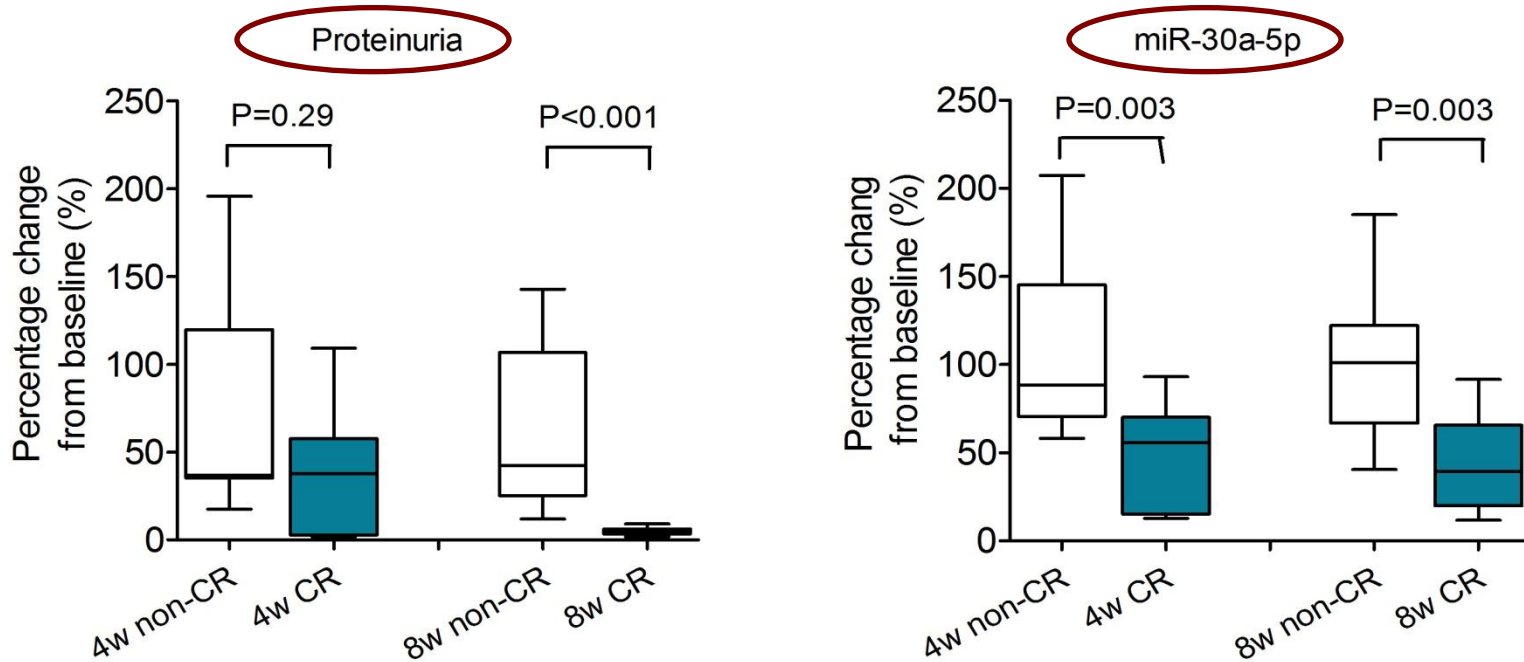
激素治疗前后尿液miR-196a, miR-30a-5p和miR-490表达变化



激素治疗后，激素敏感患者尿miR-196a, miR-30a-5p和miR-490显著下降。

尿液miR-30a-5p预测FSGS患者对激素治疗的反应

In a prospective study (n=22)



激素敏感患者，激素治疗4周后尿miR-30a-5p水平即显著下降，早于尿蛋白的缓解。

尿液miR-30a-5p, miR-196a和miR-490可能作为FSGS疾病活动的潜在标志物, 并且尿液miR-30a-5p的变化可能预测FSGS患者对激素治疗的反应。

FSGS患者尿液miR-30变化的意义

肾组织中miR-30的表达

各组织中miR-196a和miR-30a拷贝数

miRNA	liver	brain	spleen	kidney	esophagus	intestine	lung	uterus	heart	colon
miR-196a	0.32	0.19	26.78	2061.8	12.38	55.35	1.9	605.13	0.64	2.55
miR-196b	1.1	0.38	28.66	4706.24	19.99	4.61	8.54	1569.33	2.97	75.09
miR-30a	14532.42	10437.57	1561.69	35190.85	1526.49	1822.76	23887.77	2124.95	8607.81	76.13

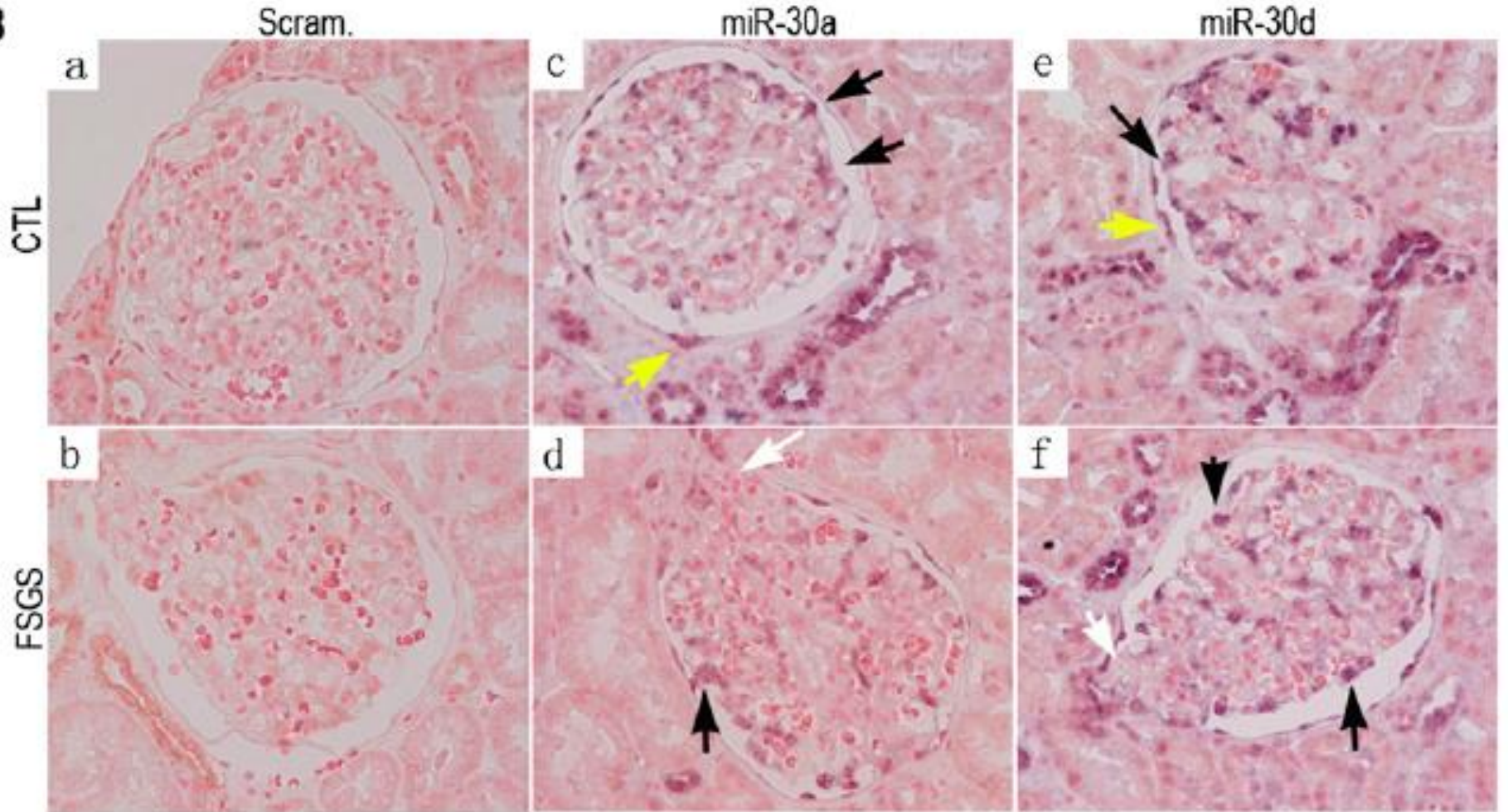
Downregulation of MicroRNA-30 Facilitates Podocyte Injury and Is Prevented by Glucocorticoids

Junnan Wu, Chunxia Zheng, Yun Fan, Caihong Zeng, Zhaohong Chen, Weisong Qin, Changming Zhang, Wanfen Zhang, Xiao Wang, Xiaodong Zhu, Mingchao Zhang, Ke Zen, and Zhihong Liu

Research Institute of Nephrology, Jinling Hospital, Nanjing University School of Medicine, Nanjing, Jiangsu, China

ABSTRACT

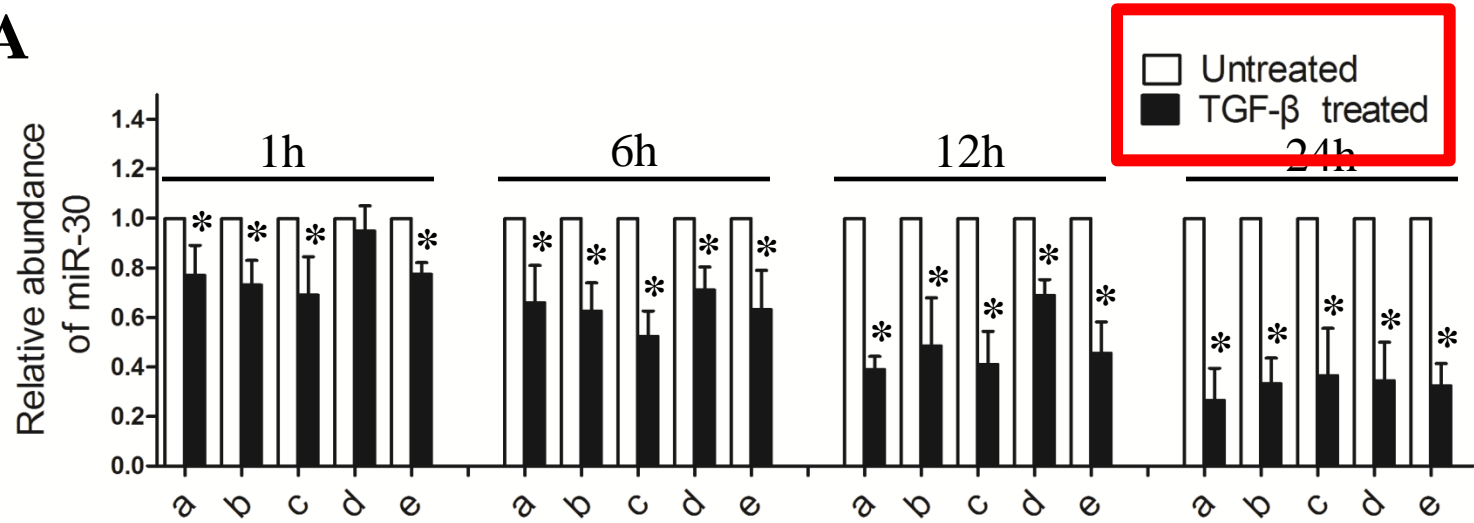
Downregulation of miR-30s in podocyte can induce proteinuria and podocyte injury. We found that miR-30s protect podocyte from injury by targeting Notch1 and p53. Glucocorticoid treatment sustains miR-30 expression in cultured podocytes treated with TGF- β or PAN and in the podocytes of PAN-treated rats.

B

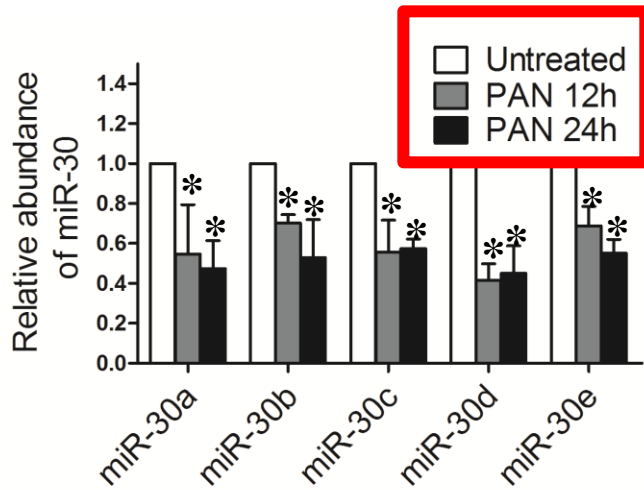
miR-30s表达于人足细胞，并且在FSGS患者肾组织中表达下调

损伤因素刺激后足细胞miR-30s表达下降

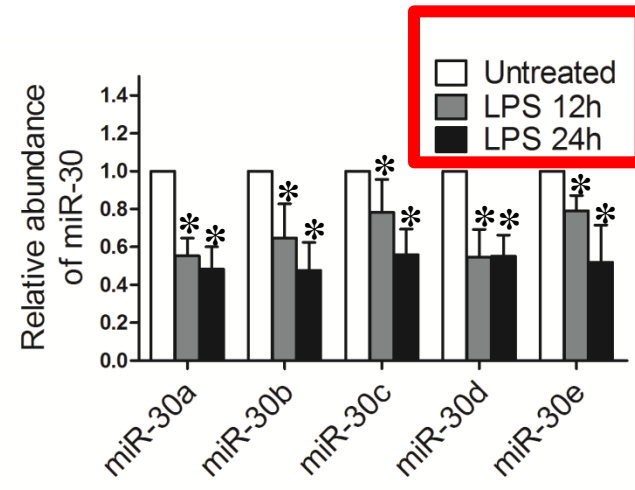
A



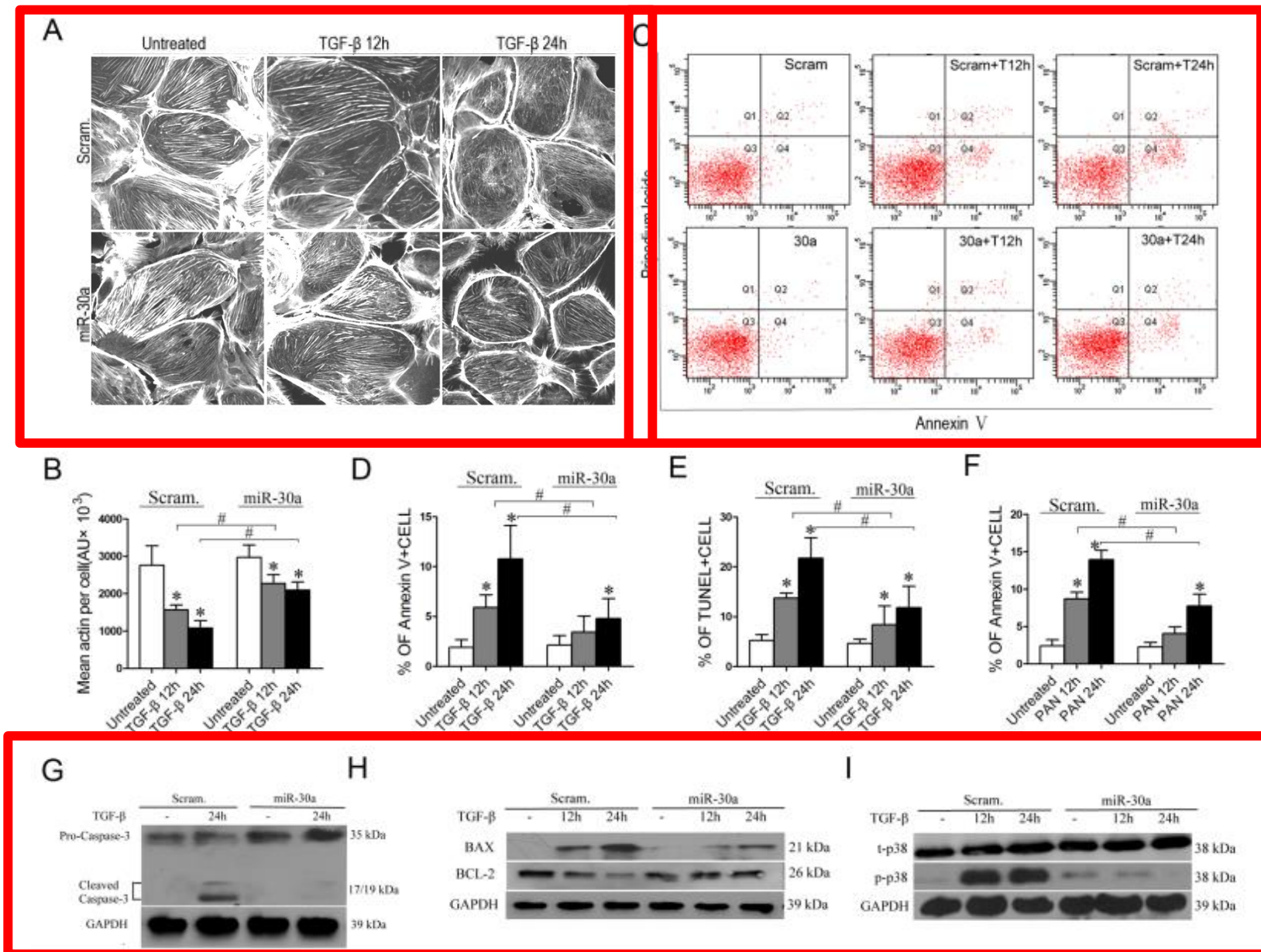
B



C

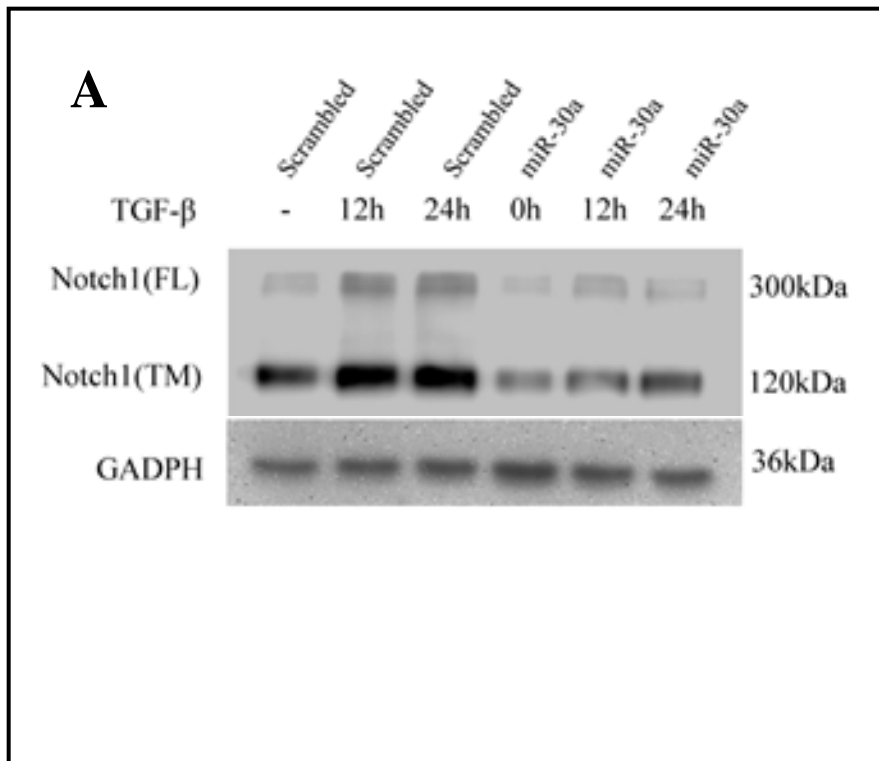


miR-30s表达下降参与足细胞损伤



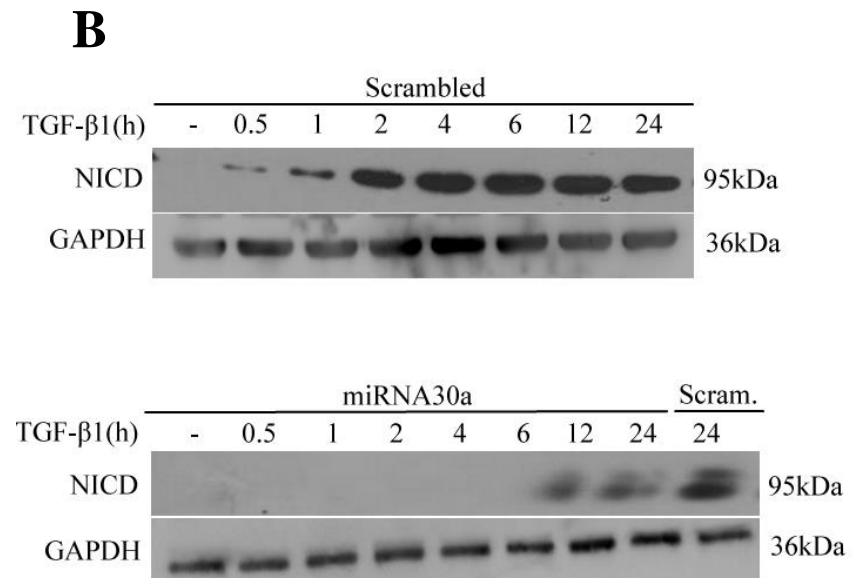
足细胞转染miR-30a质粒通过抑制Notch1受体表达 从而抑制Notch1信号通路活化

Notch1 production



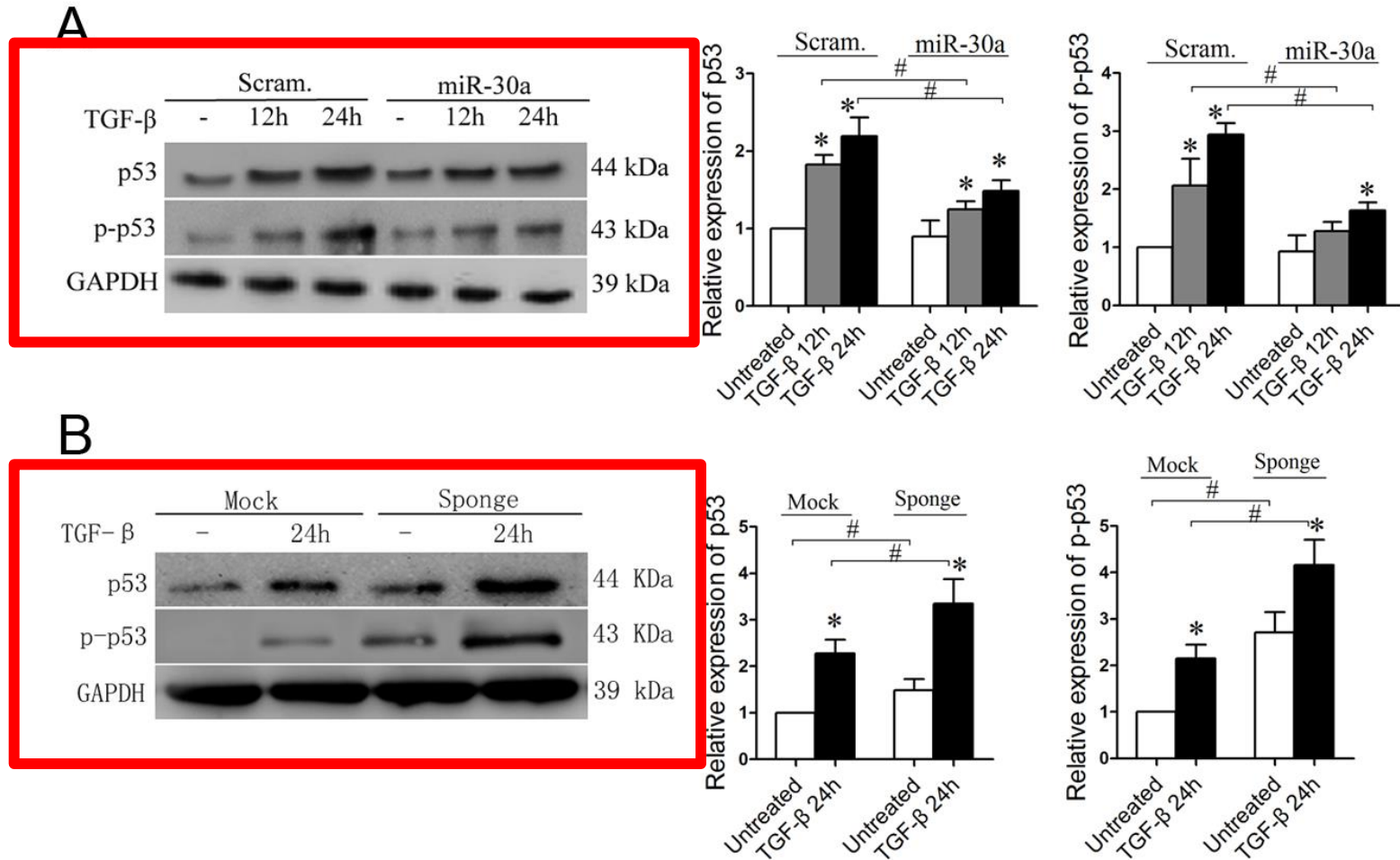
Western blot: 足细胞Notch1受体检测 (图A)

Notch1 activation

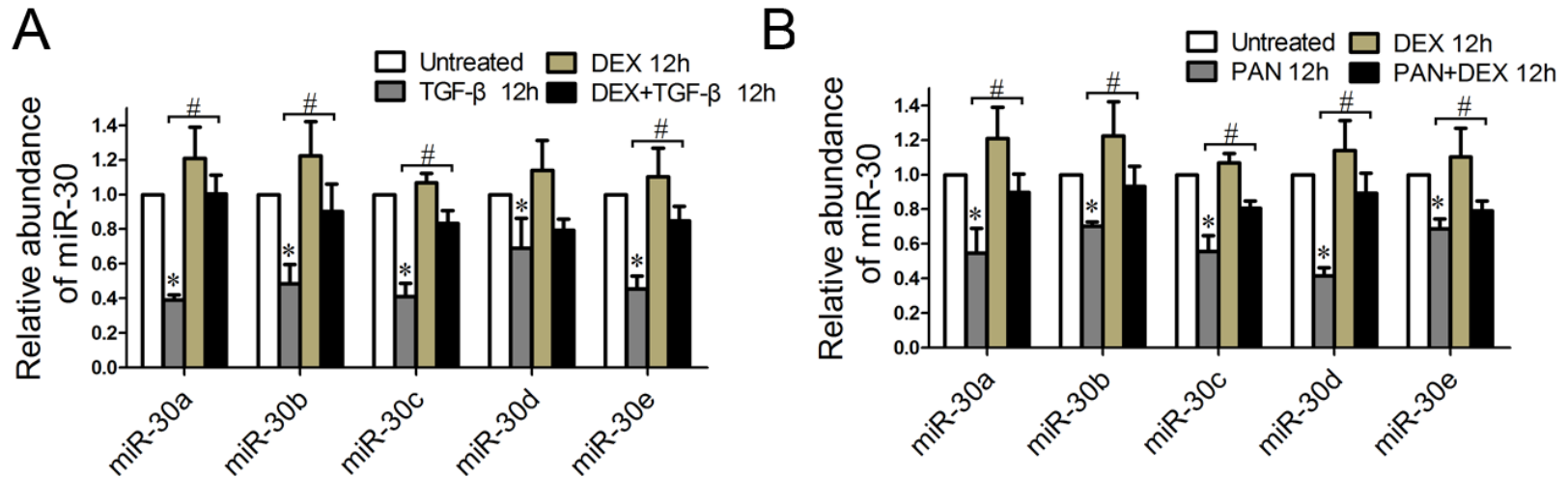


Western blot: 足细胞NICD检测 (图B)

足细胞转染miR-30a质粒后可显著拮抗TGF-β 所致的 p53通路激活



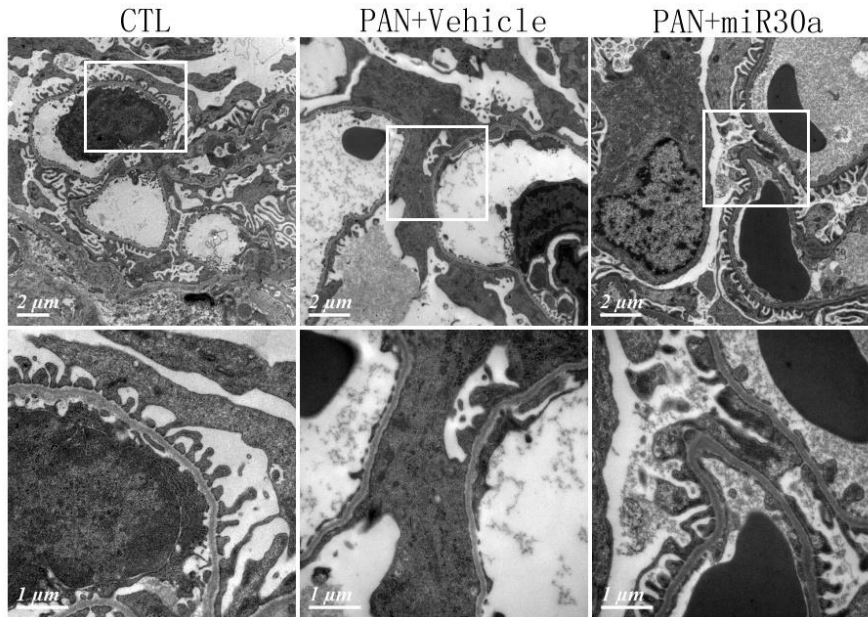
糖皮质激素（DEX）可以逆转TGF- β /PAN诱导的足细胞miR-30s表达下降



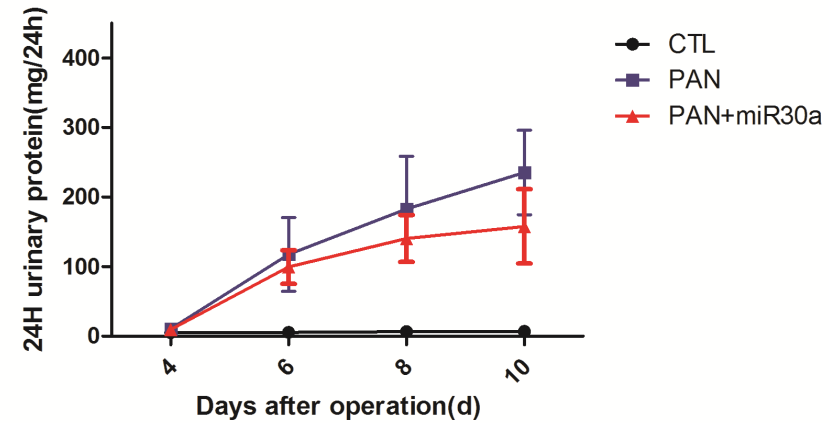
qPCR: 足细胞miR-30s表达 (图A、B)

miR-30a治疗可逆转PAN大鼠肾小球损伤

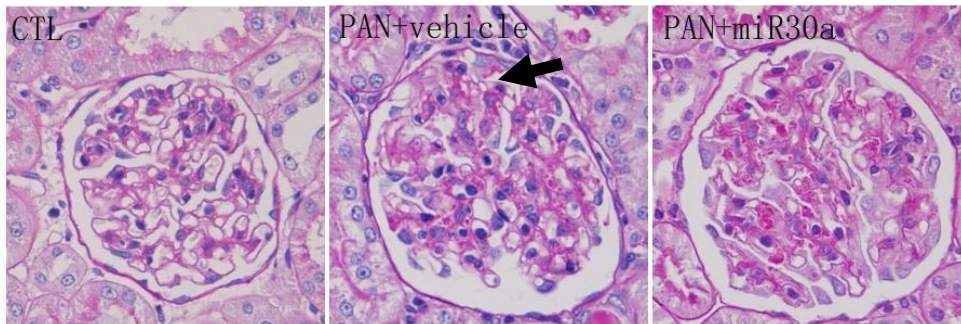
A



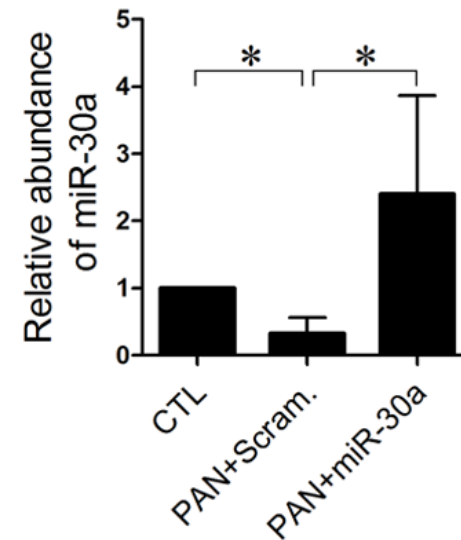
B



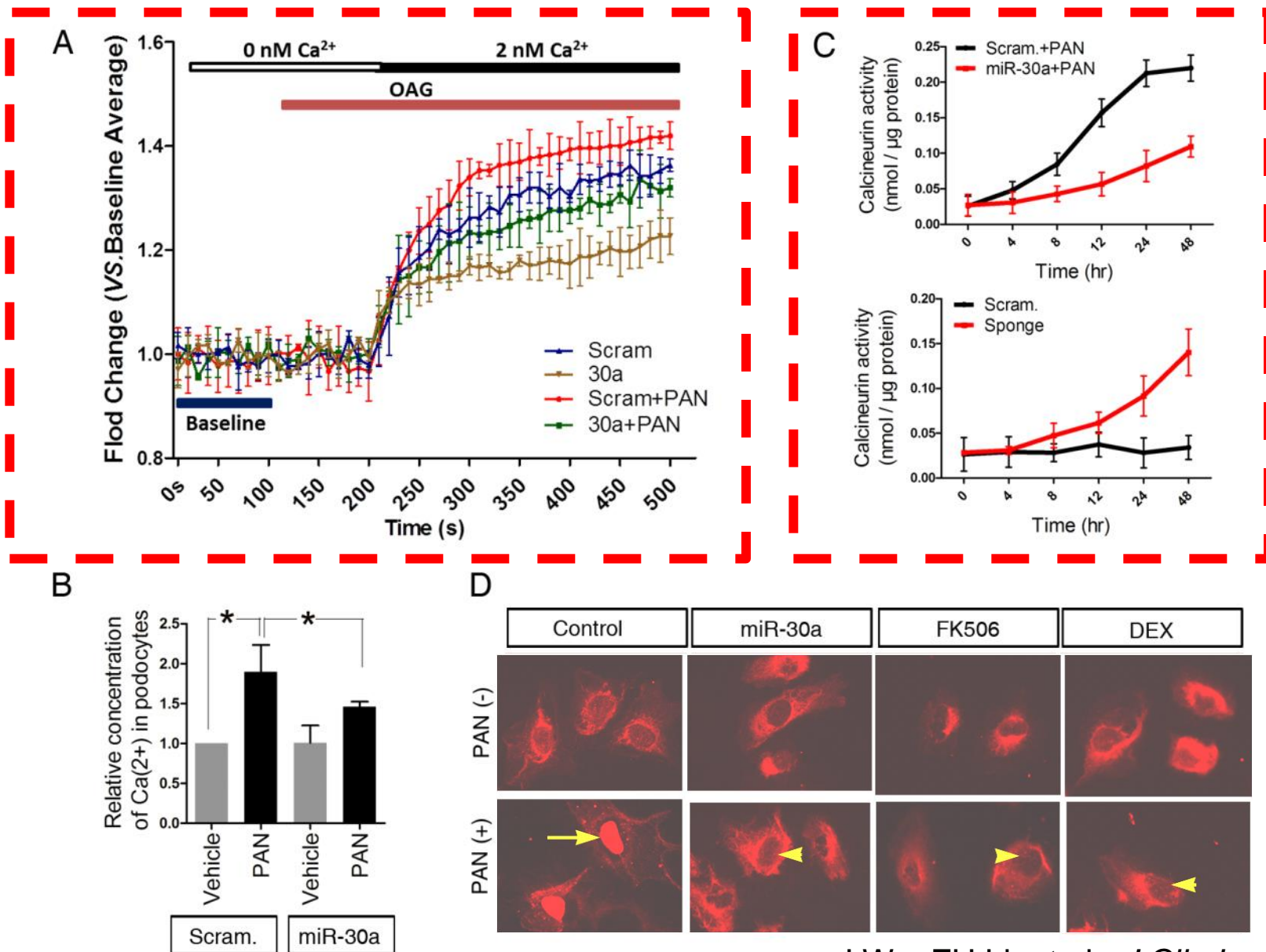
C



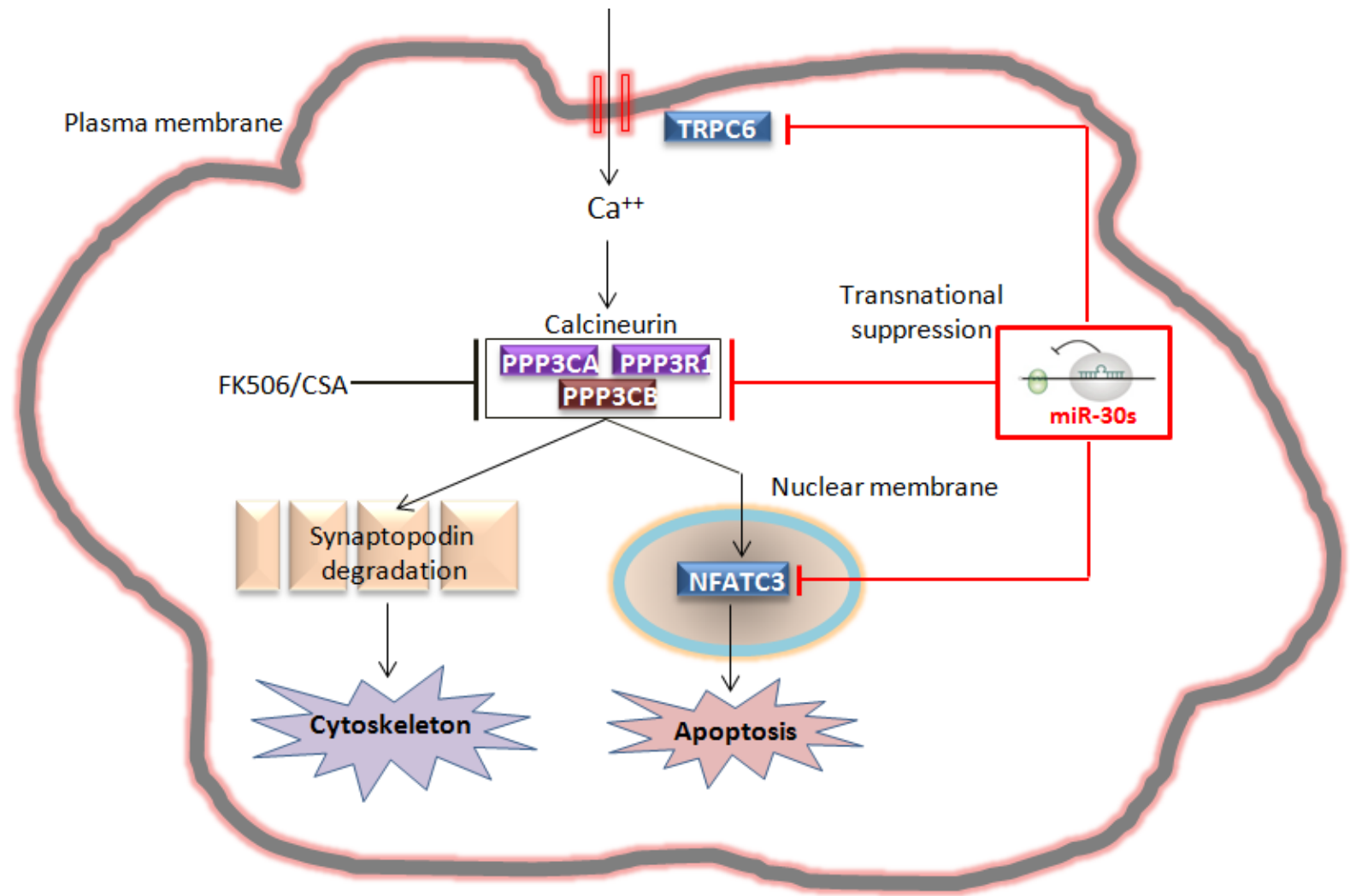
D



miR-30s调控足细胞钙内流及calcineurin活性



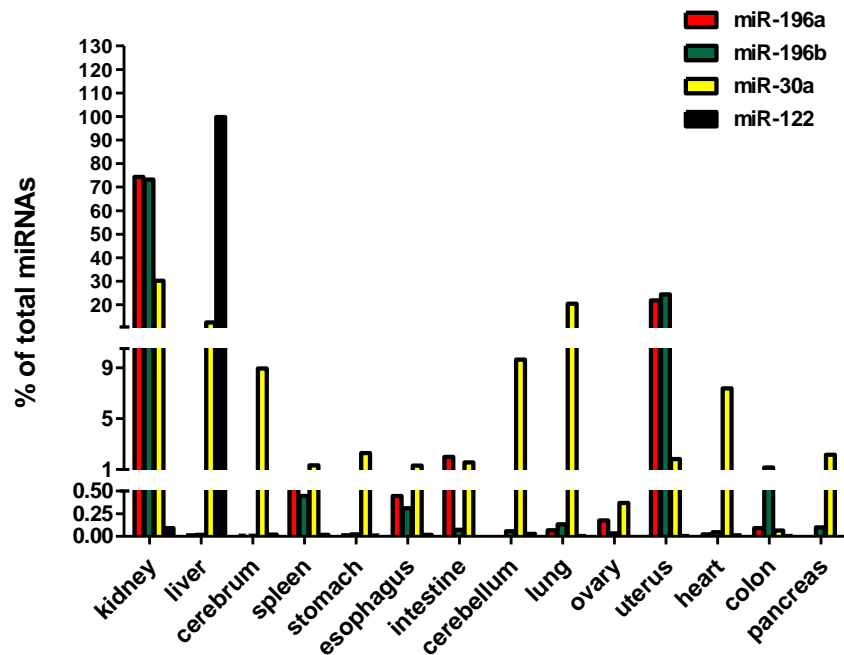
miR-30s调控Ca²⁺-Calcineurin信号通路



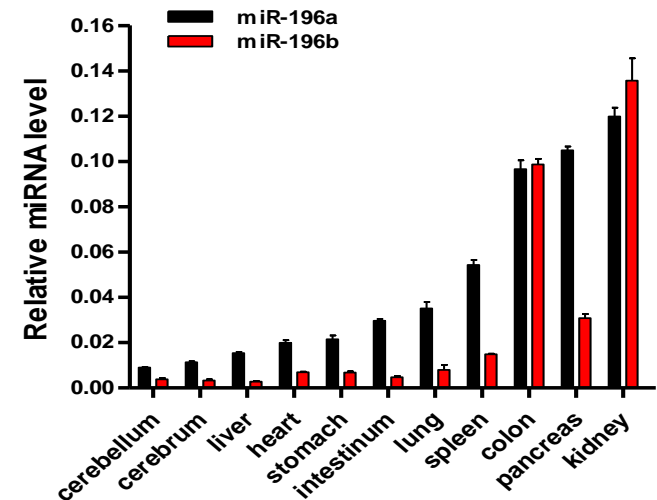
FSGS患者尿液miR-196变化的意义

miR-196a和miR-196b在小鼠肾脏中高表达

A



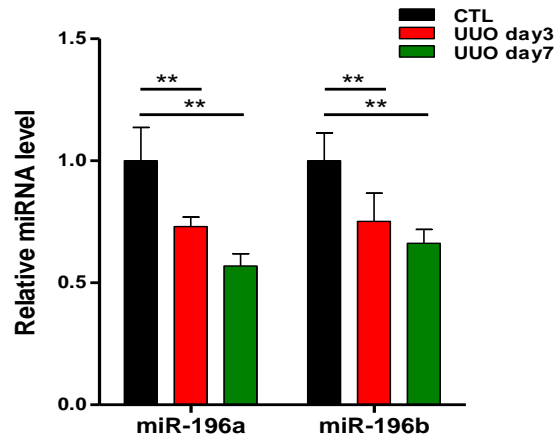
B



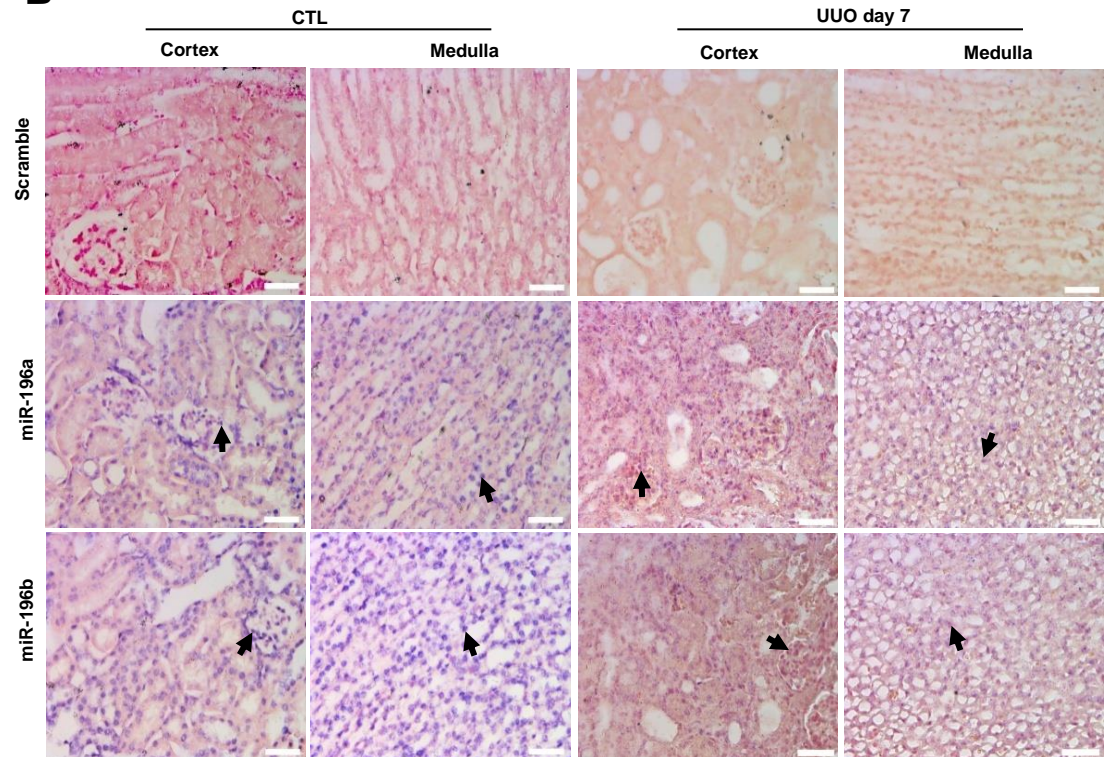
Using Solexa deep sequencing and identified that miR-196a is highly expressed in kidney

UUO模型小鼠肾脏miR-196a和miR-196b表达下调

A

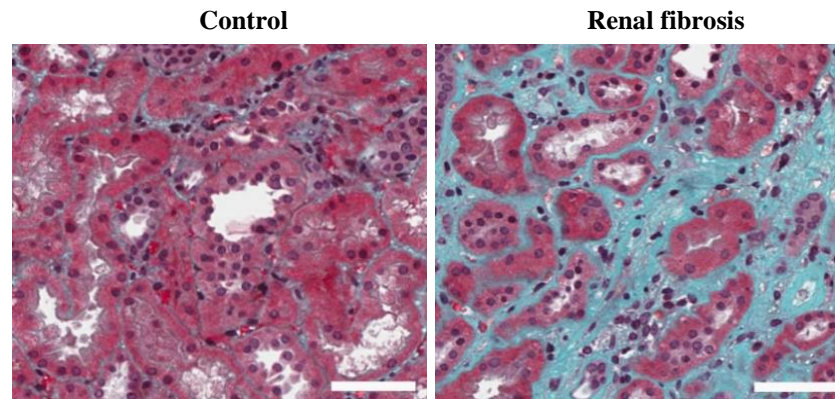


B

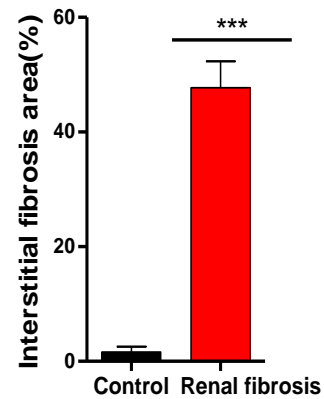


肾脏纤维化患者肾组织中miR-196a和miR-196b表达下调

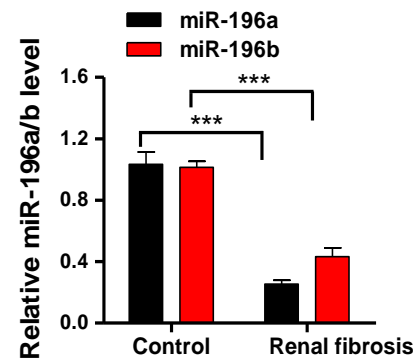
A



B



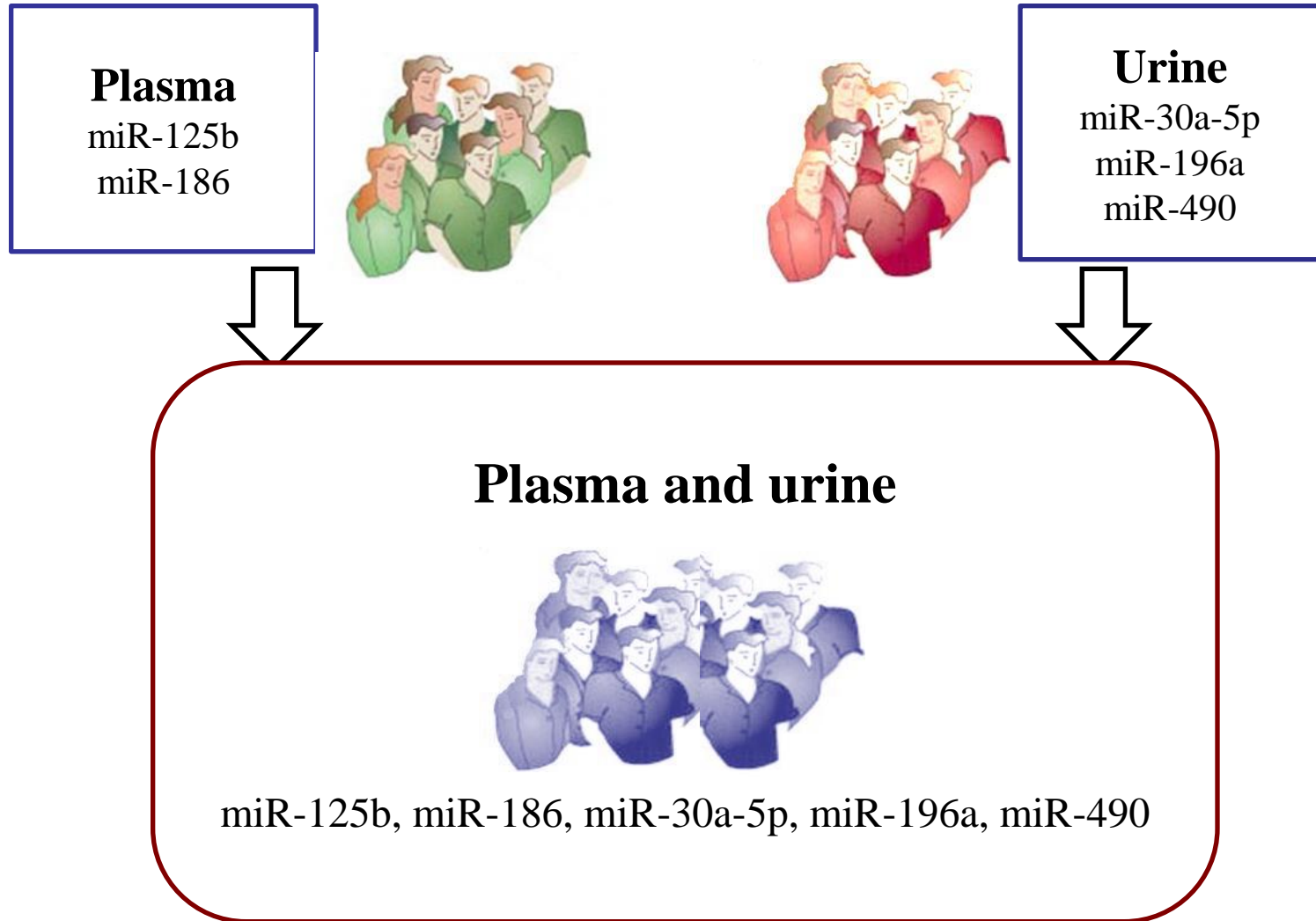
C



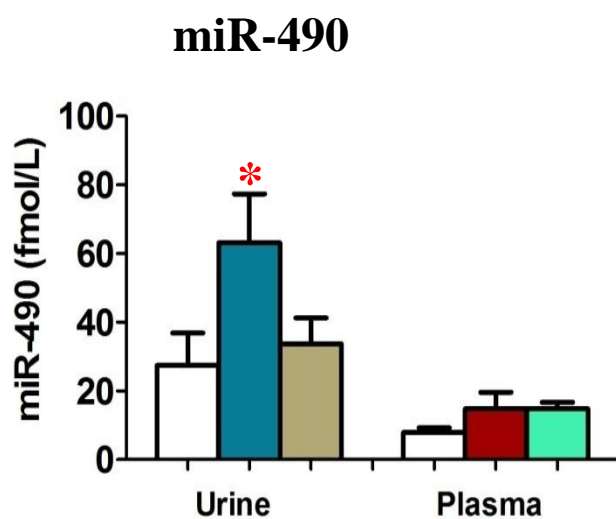
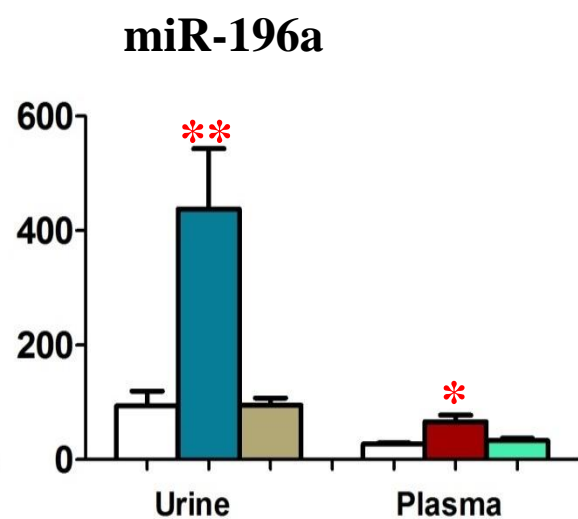
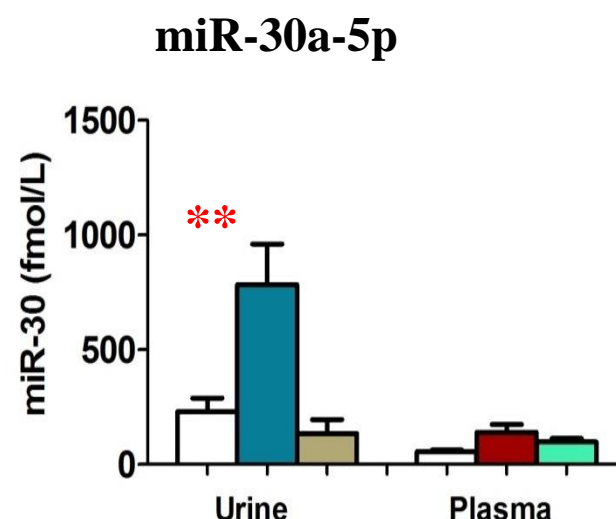
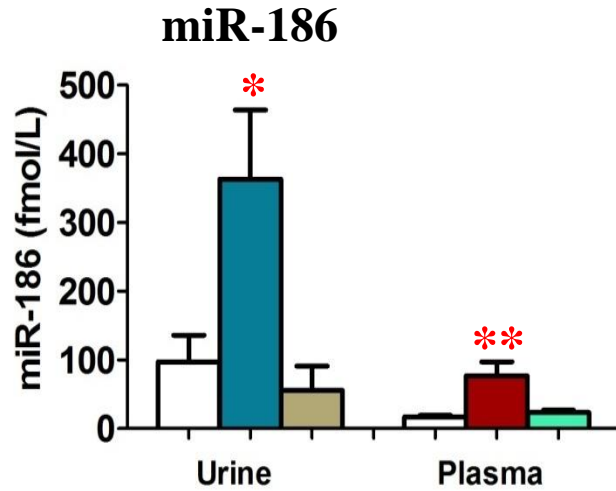
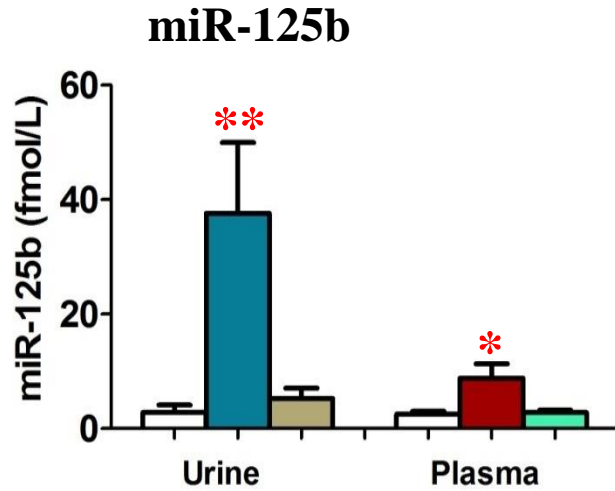
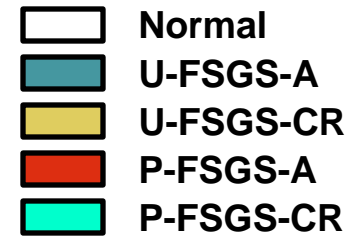
体内外研究证实miR-196a/b通过TGF β -Smad信号通路抑制肾脏纤维化。

肾脏富含miR-196，FSGS患者尿液中高表达的miR-196可能源于损伤肾脏的被动泄漏或主动分泌。

血浆和尿液miRNAs作为FSGS生物标志物的联合检测



Testing (20 cases for each group)



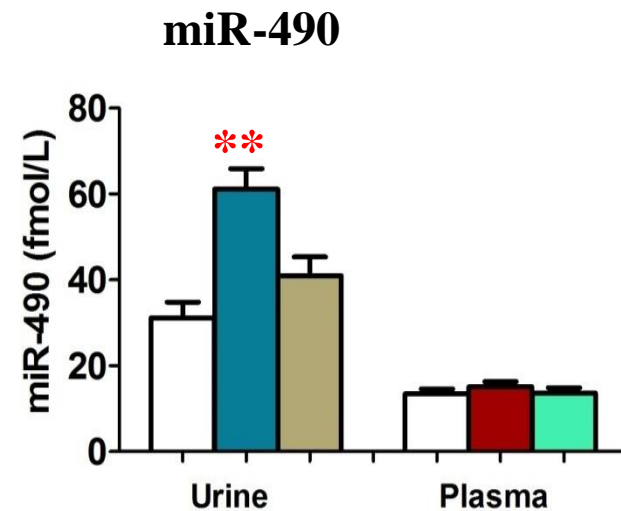
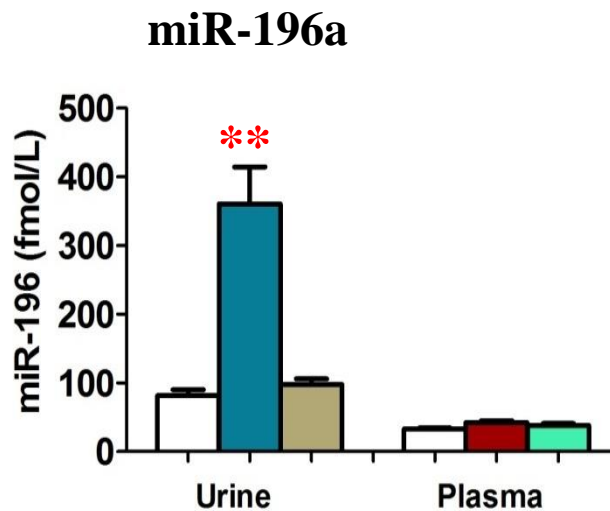
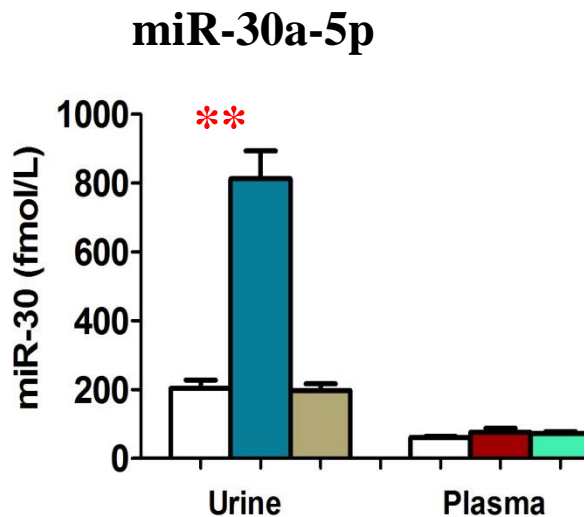
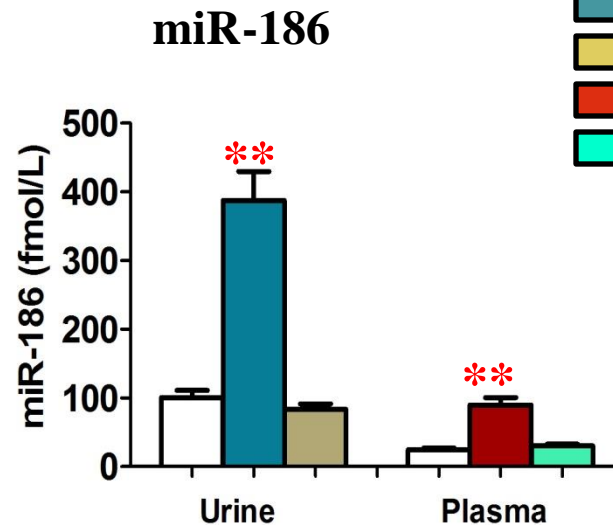
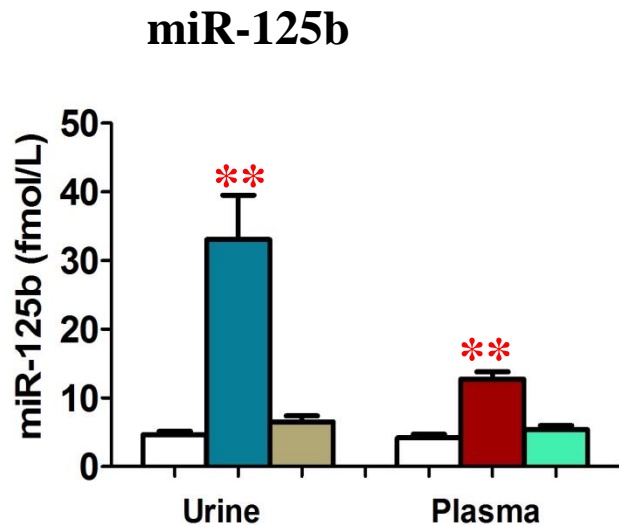
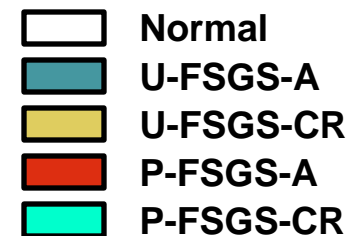
➤ Plasma: miR-125b, miR-186, miR-196a

➤ Urine: miR-30a-5p, miR-196a, miR-490, miR-125b, miR-186

*p < 0.05 vs Normal, FSGS-CR

**p < 0.01 vs Normal, FSGS-CR

Confirmation (60 cases for each group)



➤ Plasma: miR-125b, miR-186

➤ Urine: miR-30a-5p, miR-196a, miR-490, miR-125b, miR-186

**p < 0.01 vs Normal, FSGS-CR

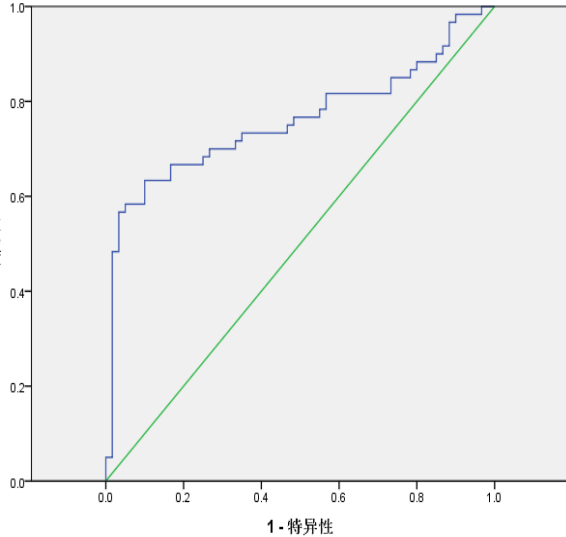
血浆和尿液miR-125b区分活动性FSGS和完全缓解FSGS

Plasma miR-125b

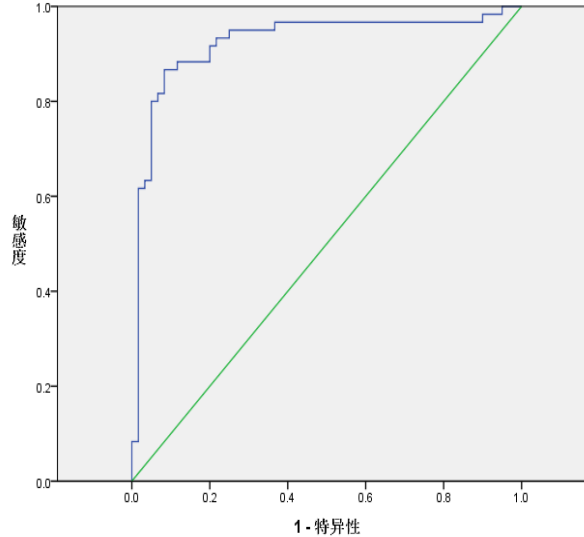
Urinary miR-125b

2 panel

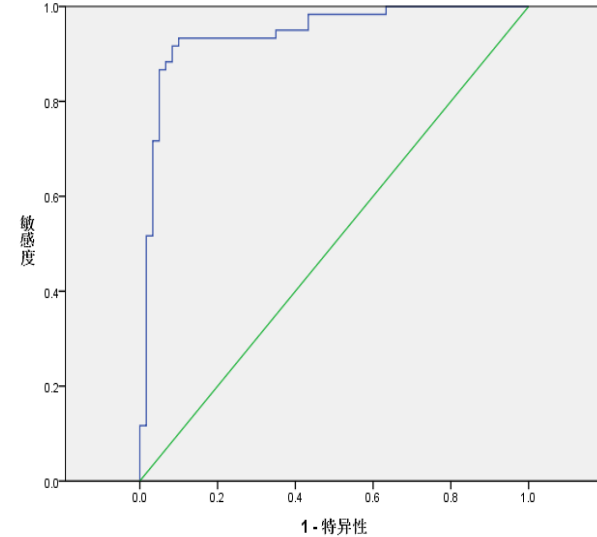
ROC 曲线



ROC 曲线



ROC 曲线



AUC

95% CI

Specificity

Sensitivity

P value

P-miR-125b

0.760

0.669-0.851

0.967

0.567

8.98×10^{-7}

U-miR-125b

0.924

0.869-0.978

0.917

0.867

1.20×10^{-15}

2 panel

0.943

0.899-0.987

0.917

0.917

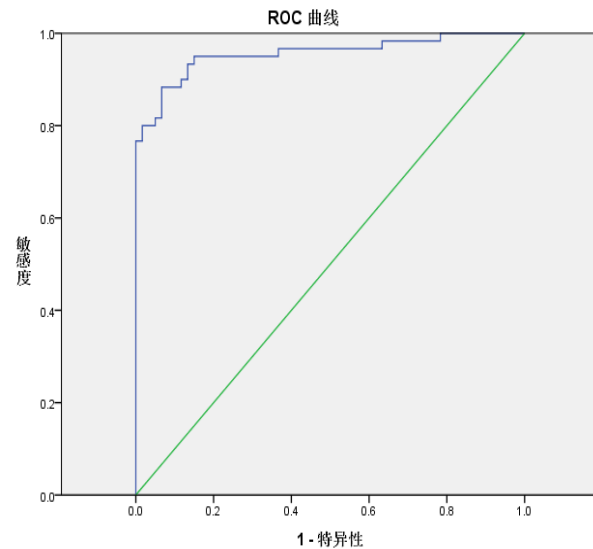
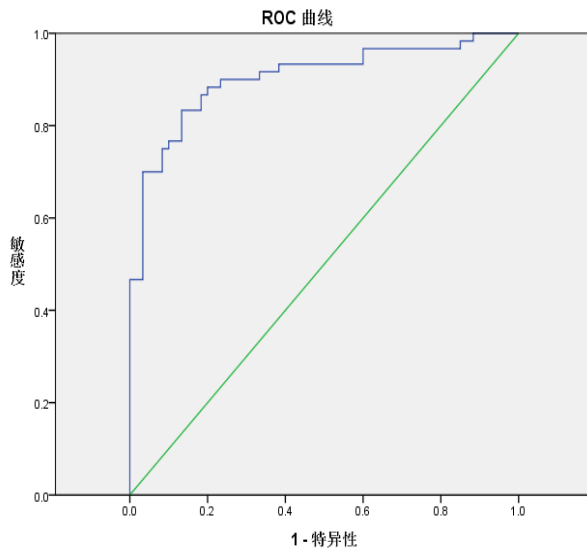
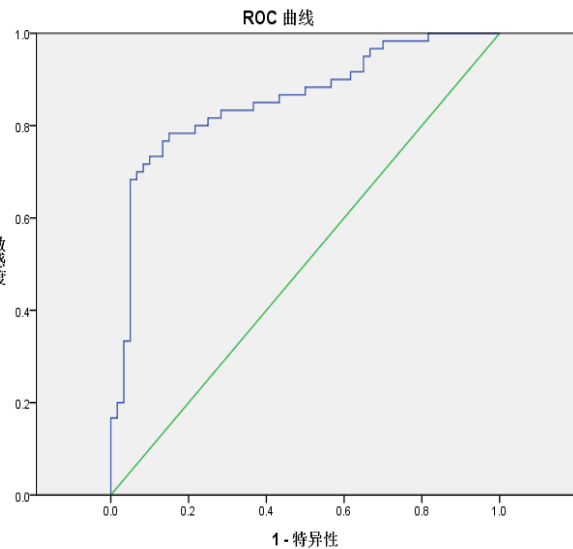
5.94×10^{-17}

血浆和尿液miR-186区分活动性FSGS和完全缓解FSGS

Plasma miR-186

Urinary miR-186

2 panel



	AUC	95% CI	Specificity	Sensitivity	P value
P-miR-186	0.854	0.785-0.924	0.867	0.767	2.12×10^{-11}
U-miR-186	0.903	0.847-0.960	0.867	0.833	2.52×10^{-14}
2 panel	0.956	0.918-0.993	0.933	0.883	7.45×10^{-18}

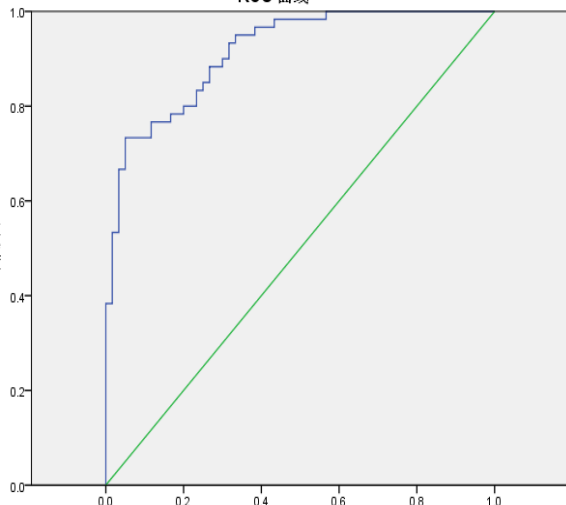
尿液miR-30a, miR-196a 和miR-490区分活动性FSGS和完全缓解FSGS

Urinary miR-30a

Urinary miR-196a

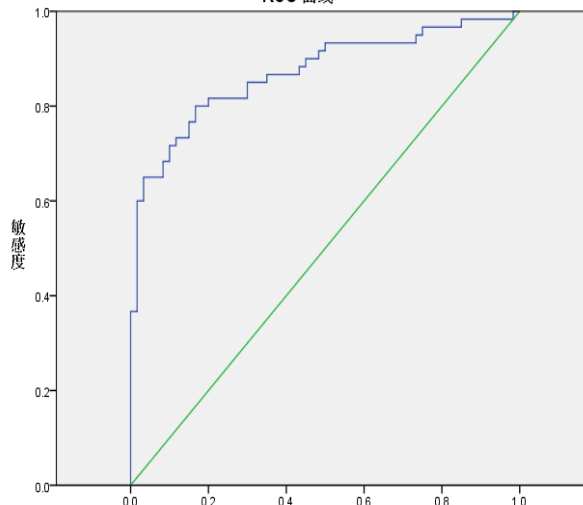
Urinary miR-490

ROC 曲线



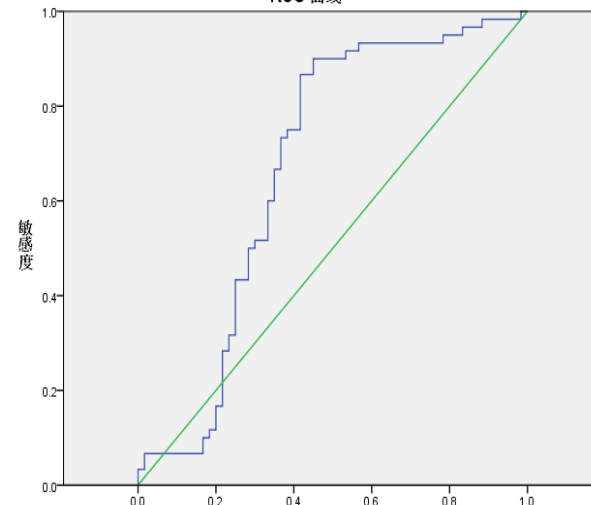
1-特异性

ROC 曲线



1-特异性

ROC 曲线



1-特异性

AUC

95% CI

Specificity

Sensitivity

P value

U-miR-30a

0.915

0.867-0.962

0.950

0.733

4.64×10^{-15}

U-miR-196a

0.870

0.805-0.936

0.833

0.800

2.63×10^{-12}

U-miR-490

0.673

0.571-0.775

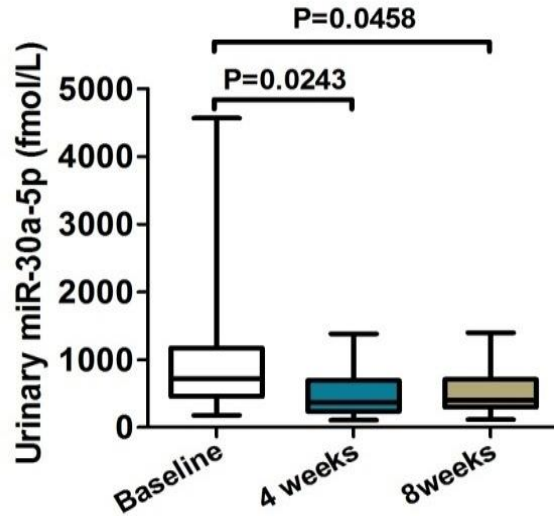
0.550

0.900

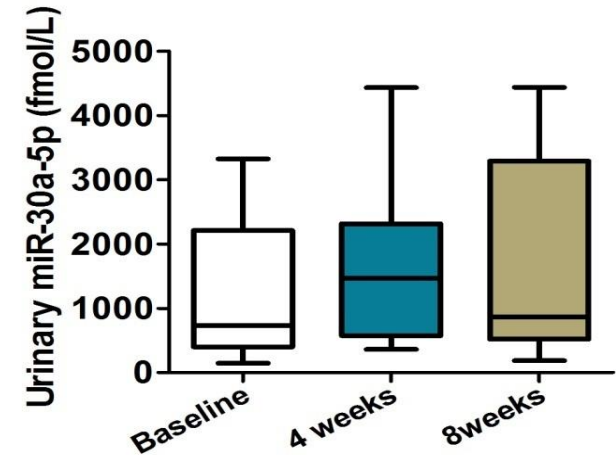
0.001

激素治疗前后尿液miR-30a-5p水平

Steroid-sensitive patients



Steroid-resistant patients



激素敏感患者，治疗4周后尿液miR-30a-5p水平较治疗前已有明显下降，早于尿蛋白的变化，激素抵抗患者在治疗前后则无明显差异。

miR-125b

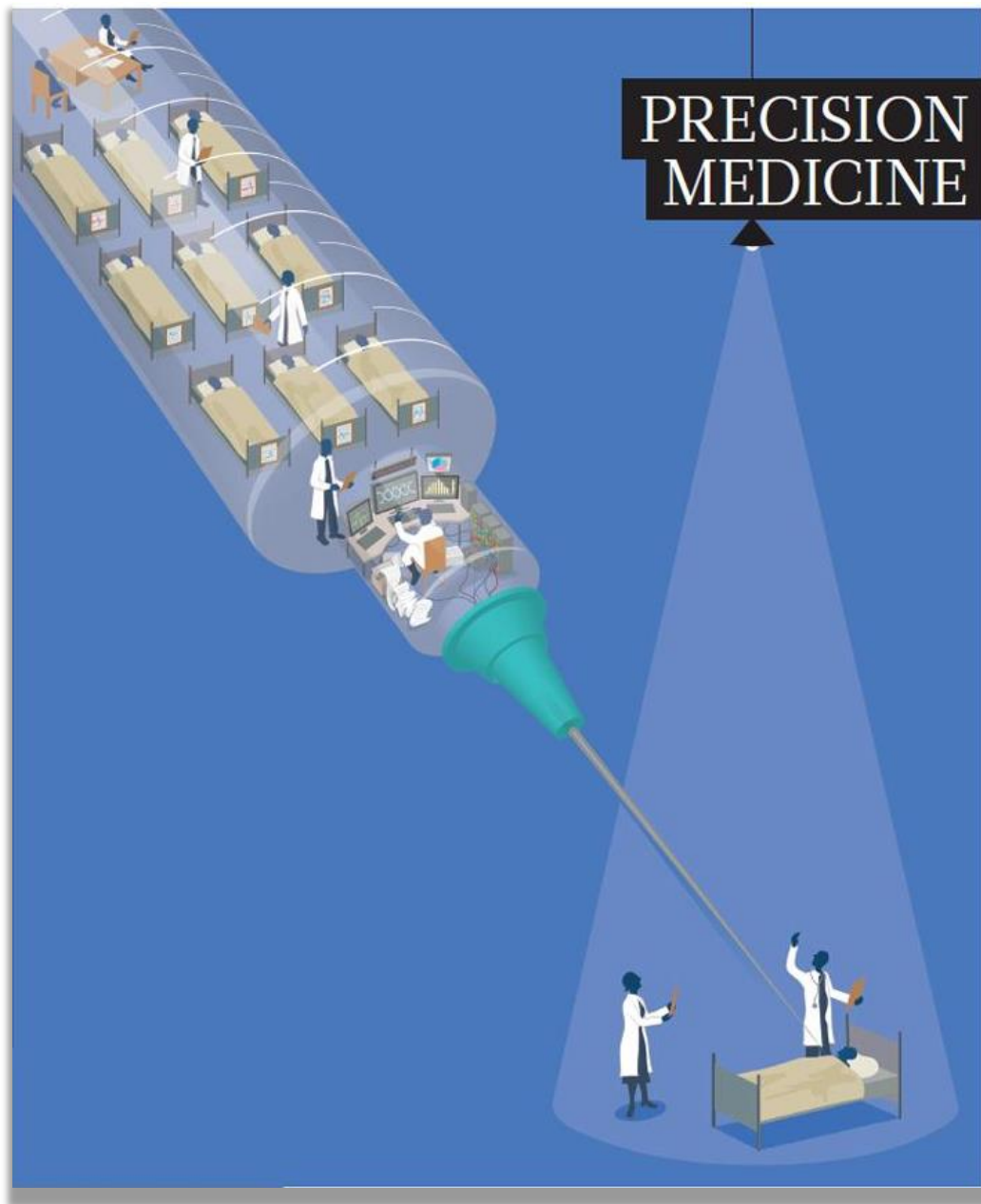
血浆和尿液miR-125b均与FSGS疾病活动相关，尿液miR-125b呈现更好的预测效力，尿液miR-125b可以作为FSGS诊断和监测的生物标志物。

miR-186

血浆和尿液miR-186均与FSGS疾病活动相关，尿液miR-186呈现更好的预测效力，尿液miR-186可以作为FSGS诊断和监测的生物标志物。

miR-30a

监测尿液miR-30a-5p的变化可用于预测FSGS患者对激素治疗的反应。



PRECISION MEDICINE

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为病人提供更好地服务

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前瞻性研究队列
生物样本库
基于组学的系统研究
大数据的生物信息学分析

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Pharmaceutical Biotechnology
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Ke Zeng, PhD

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