



# 自闭症风险基因 *FMR1* (*Fmr1*) 的生物信息学分析

—— | G4小组成员：毕金龙、李戈、程佳琪 | ——



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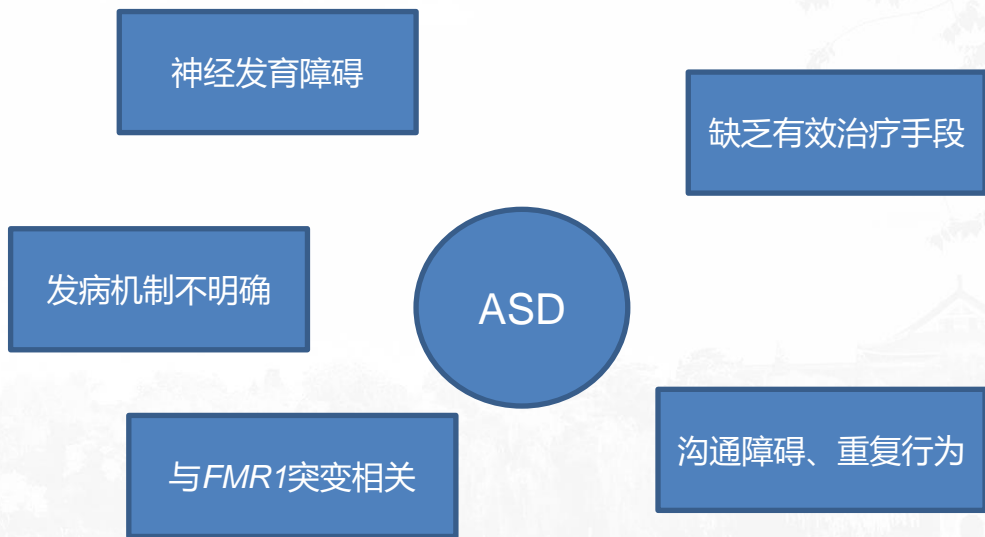
探究问题及结论

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小组成员总结

# 自闭症简介



- 自闭症 (Autism) , 又称孤独症, 是一种广泛的神经起源的发育障碍。它的谱系很广, 统称为自闭症谱系障碍 (Autism Spectrum Disorder, ASD) 。
- 自闭症的特点: ①在多种背景下的社交沟通和社交互动持续存在缺陷; ②有限的、重复的行为、兴趣或活动模式
- 自闭症障碍全世界人口患病率约为1~2%, 大约每100个儿童中就有一人患自闭症

# 自闭症动物模型

小鼠模型

单基因模型

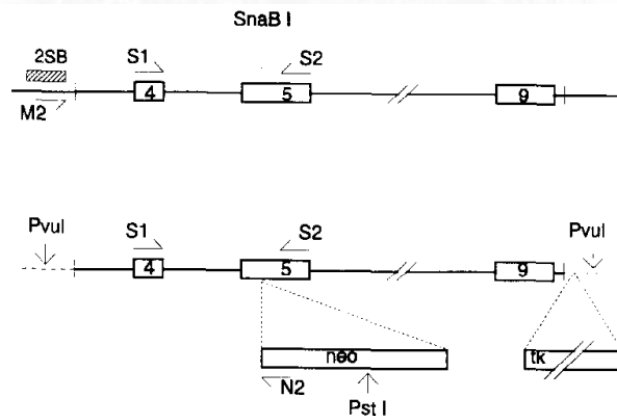
*Fmr1* 敲除模型

多基因模型

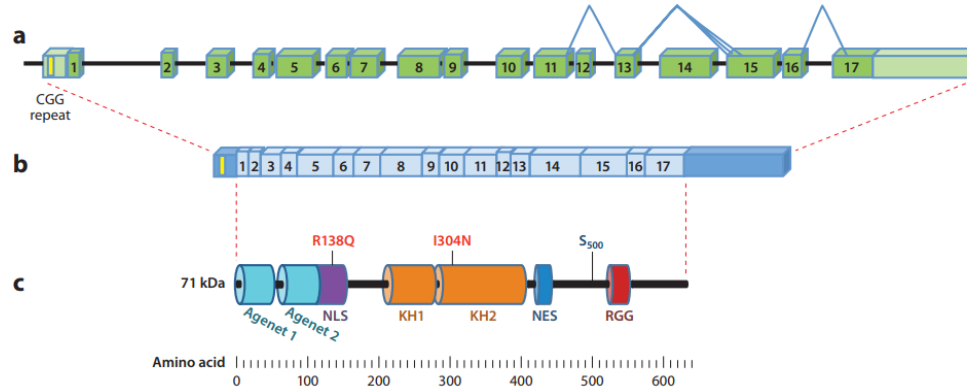
动物模型

果蝇模型

犬等其他模型



# Fmr1及FMRP结构

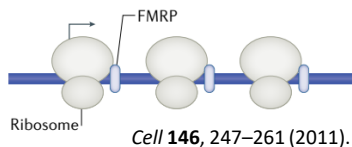


*Annu Rev Pathol.* 2012;7:219-45.

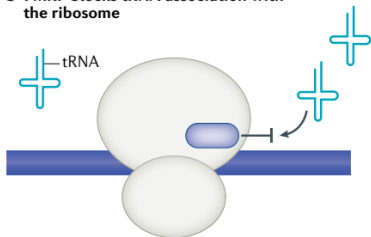
- Agenet 1 and Agenet 2, tandem Agenet domains, **chromatin binding**;
- NLS, nuclear localization signal; NES, nuclear export **signal**;
- RGG, arginine-glycine-glycine box, **RNA-binding**;
- KH1 and KH2, K homology domains 1 and 2, **RNA-binding**;

# FMRP具有和mRNA结合

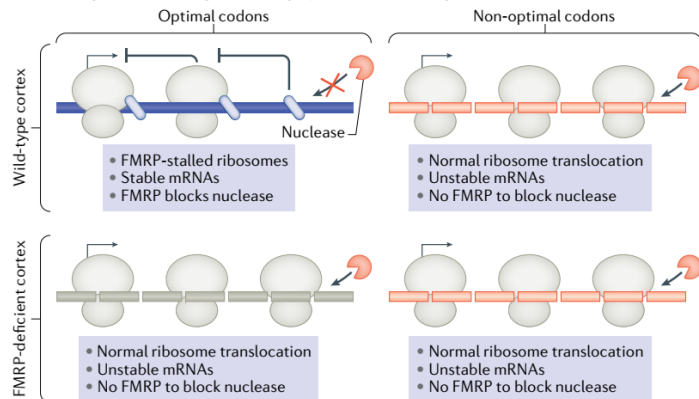
**a FMRP blocks ribosome translocation**



**b FMRP blocks tRNA association with the ribosome**



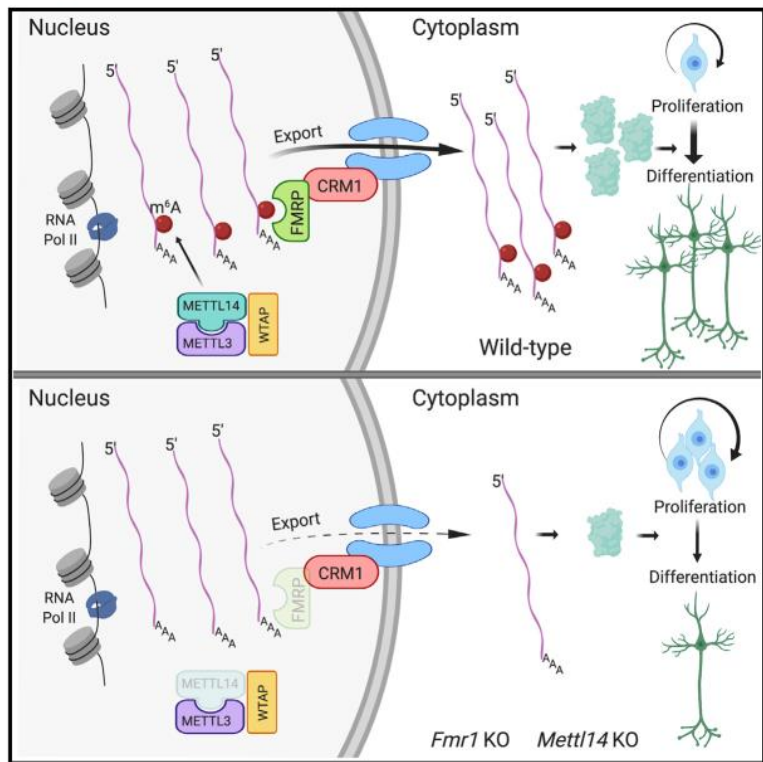
**c FMRP regulates RNA degradation by optimal codon recognition**



*Nature Reviews Neuroscience* 22, pages209-222 (2021)

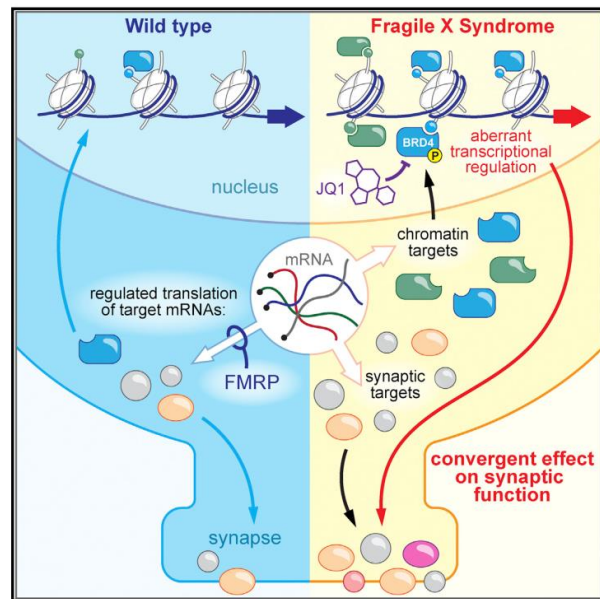
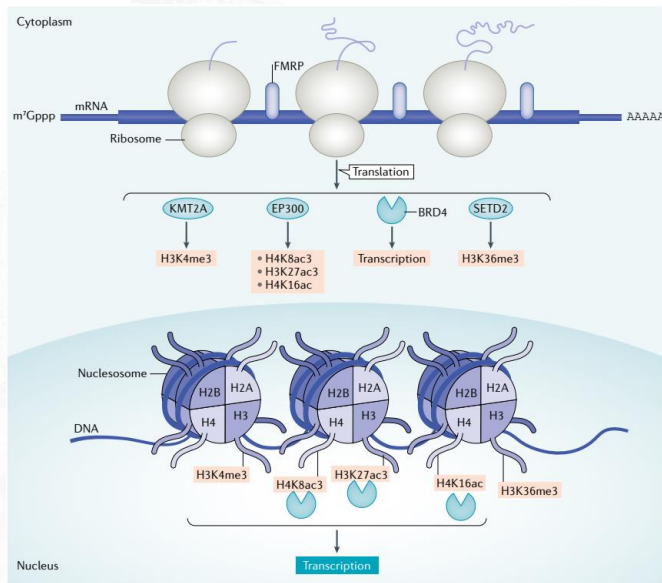
- FMRP与mRNA结合可能会阻碍核糖体易位，从而导致多肽延伸率降低。
- FMRP是一种抑制tRNA与核糖体结合的核糖体蛋白。
- FMRP直接或间接地与野生型小鼠皮质mRNA的翻译机制相关，这些mRNA含有最佳密码子并阻止这些RNA上的核糖体。

# FMRP与核质转运



- FMRP优先结合m6A甲基化mRNA并促进其核输出，这对于神经元分化至关重要。

# FMRP与染色质调控



Cell 170, 1209–1223 (2017)

- FMRP控制表观遗传修饰酶的合成，进而改变染色质景观，导致转录激活
- FMRP调控转录因子的翻译，进而调控基因表达

## 探究目的

- 是否鼠源*Fmr1* 转录的蛋白质功能研究能够对人源FMRP功能有参考意义？
- FMRP和哪些蛋白质可以发生相互作用？

# FMR1基因序列的保守性

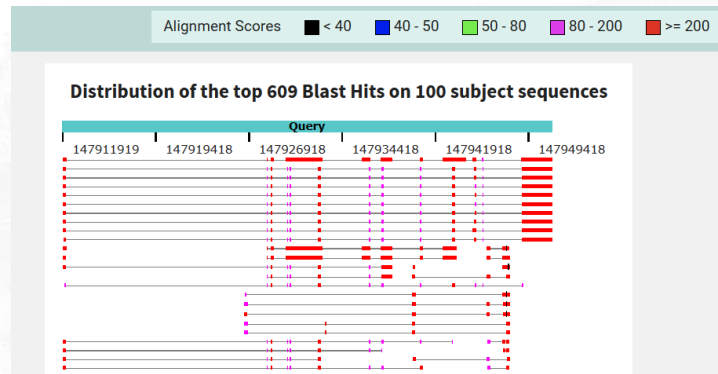
## 基因序列

Descriptions Graphic Summary Alignments Taxonomy

Sequences producing significant alignments Download Select columns Show 100

select all 100 sequences selected

Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
<input checked="" type="checkbox"/> Mus musculus X BAC RP23-149C7 (Roswell Park Cancer Institute Mouse BAC Library) complete seque...	Mus musculus	3048	9231	26%	0.0	88.46%	222658	AC055766.31
<input checked="" type="checkbox"/> Mus musculus fragile X mental retardation syndrome 1 homolog_mRNA (cDNA clone MGC 90722 IMAG...	Mus musculus	2946	5555	10%	0.0	89.03%	4412	BC079671.1
<input checked="" type="checkbox"/> Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant 3_mRNA	Mus musculus	2942	5546	10%	0.0	89.02%	4305	NM_001374719.1
<input checked="" type="checkbox"/> Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant 6_mRNA	Mus musculus	2942	5347	10%	0.0	89.02%	4230	NM_001412091.1
<input checked="" type="checkbox"/> Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant 4_mRNA	Mus musculus	2942	5485	10%	0.0	89.02%	4332	NM_001412089.1
<input checked="" type="checkbox"/> Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant 2_mRNA	Mus musculus	2942	5353	10%	0.0	89.02%	4293	NM_001290424.2
<input checked="" type="checkbox"/> Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant 5_mRNA	Mus musculus	2942	5479	10%	0.0	89.02%	4269	NM_001412090.1
<input checked="" type="checkbox"/> Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant 1_mRNA	Mus musculus	2942	5551	10%	0.0	89.02%	4368	NM_008031.4
<input checked="" type="checkbox"/> PREDICTED: Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1) transcript variant X4_misc...	Mus musculus	2942	5666	10%	0.0	89.02%	5147	XR_001782738.3
<input checked="" type="checkbox"/> Mus musculus fragile X mental retardation syndrome protein (Fmr1) (homologue) mRNA, complete cds	Mus musculus...	2726	5319	10%	0.0	87.57%	4240	L23971.1



# FMR1转录本和蛋白序列的保守性

## 转录本序列

Descriptions | Graphic Summary | Alignments | Taxonomy

Sequences producing significant alignments

Download Select columns Show 100

select all 7 sequences selected

GenBank Graphics Distance tree of results MSA Viewer

Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant 3, mRNA	Mus musculus	2942	5252	94%	0.0	89.02%	4305	NM_001374719.1
Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant 6, mRNA	Mus musculus	2942	5252	94%	0.0	89.02%	4230	NM_001412091.1
Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant 4, mRNA	Mus musculus	2942	5336	96%	0.0	89.02%	4332	NM_001412089.1
Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant 2, mRNA	Mus musculus	2942	5336	96%	0.0	89.02%	4293	NM_001290424.2
Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant 5, mRNA	Mus musculus	2942	5252	94%	0.0	89.02%	4269	NM_001412090.1
Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant 1, mRNA	Mus musculus	2942	5336	96%	0.0	89.02%	4368	NM_008031.4
PREDICTED: Mus musculus fragile X messenger ribonucleoprotein 1 (Fmr1), transcript variant X4, misc RNA	Mus musculus	2942	5336	96%	0.0	89.02%	5147	XR_001782738.3

## 蛋白质序列

Descriptions | Graphic Summary | Alignments | Taxonomy

Sequences producing significant alignments

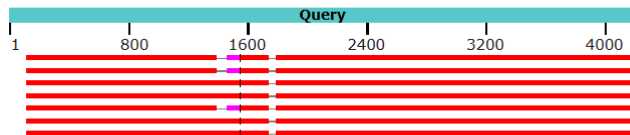
Download Select columns Show 100

select all 49 sequences selected

GenPept Graphics Distance tree of results Multiple alignment MSA Viewer

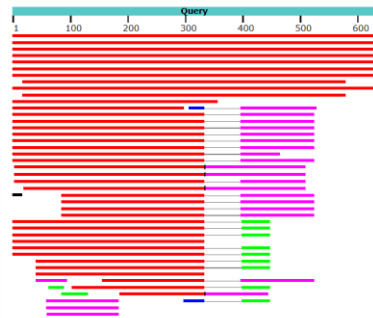
Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
fragile X messenger ribonucleoprotein 1 isoform 1 [Mus musculus]	Mus musculus	1221	1221	100%	0.0	94.15%	614	NP_032057.2
RecName: Full=Fragile X messenger ribonucleoprotein 1; AltName: Full=Fragile X messenger ribonucleoprotein 1 [Mus musculus]	Mus musculus	1218	1218	100%	0.0	93.99%	614	P35922.1
fragile X messenger ribonucleoprotein 1 isoform 2 [Mus musculus]	Mus musculus	1190	1190	100%	0.0	92.25%	602	NP_001399018.1
fragile X messenger ribonucleoprotein 1 isoform 3 [Mus musculus]	Mus musculus	1179	1179	100%	0.0	91.61%	593	NP_001361648.1
fragile X messenger ribonucleoprotein 1 isoform 2 [Mus musculus]	Mus musculus	1157	1157	100%	0.0	90.19%	589	NP_001277353.1
fragile X mental retardation protein FMRP [Mus musculus]	Mus musculus	1153	1153	100%	0.0	90.03%	589	AAL66364.1
fragile X messenger ribonucleoprotein 1 isoform 5 [Mus musculus]	Mus musculus	1147	1147	100%	0.0	89.72%	561	NP_001399019.1
fragile X mental retardation syndrome 1 homolog, isoform CRA_a [Mus musculus]	Mus musculus	1132	1132	88%	0.0	96.98%	561	EDL01213.1
fragile X messenger ribonucleoprotein 1 isoform 6 [Mus musculus]	Mus musculus	1114	1114	100%	0.0	87.66%	568	NP_001399020.1

Distribution of the top 24 Blast Hits on 7 subject sequences

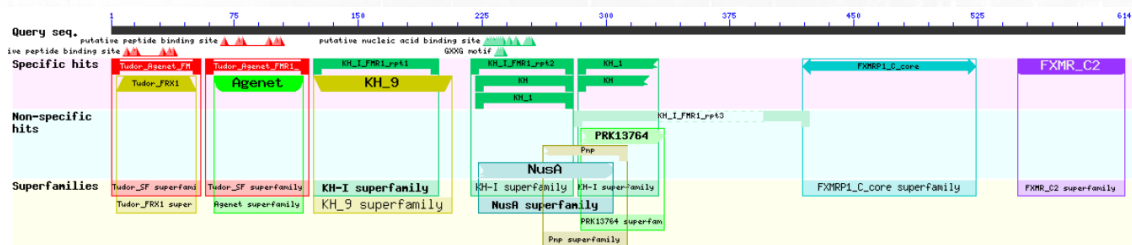
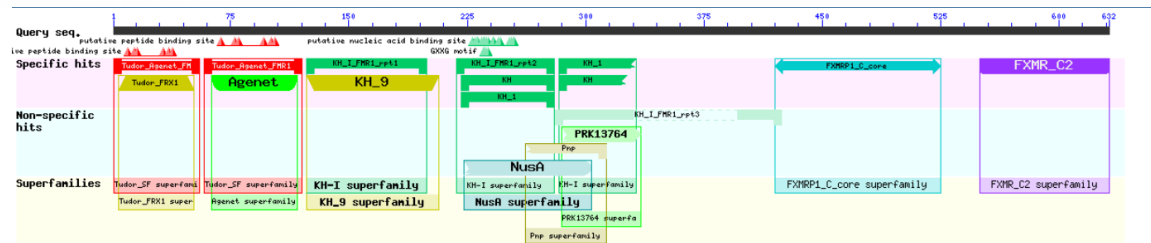
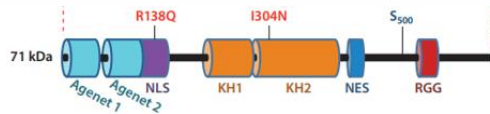
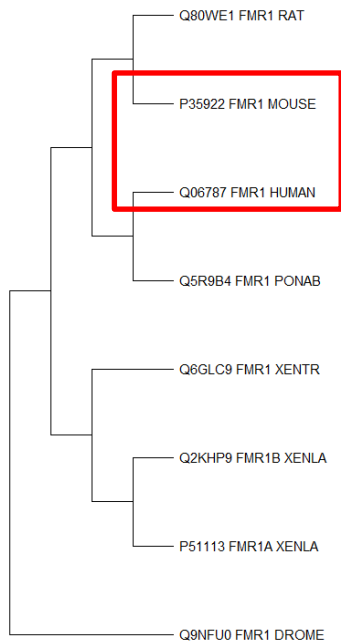


Alignment Scores ■ < 40 ■ 40 - 50 ■ 50 - 80 ■ 80 - 200 ■ >= 200

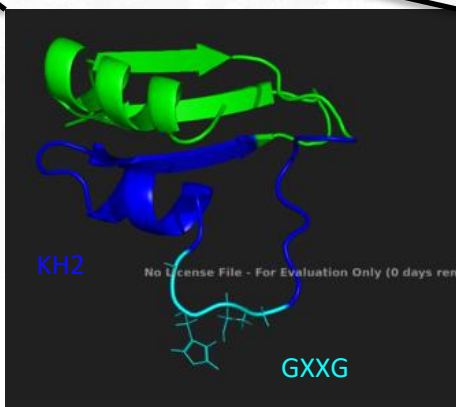
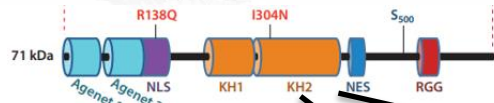
Distribution of the top 77 Blast Hits on 49 subject sequences



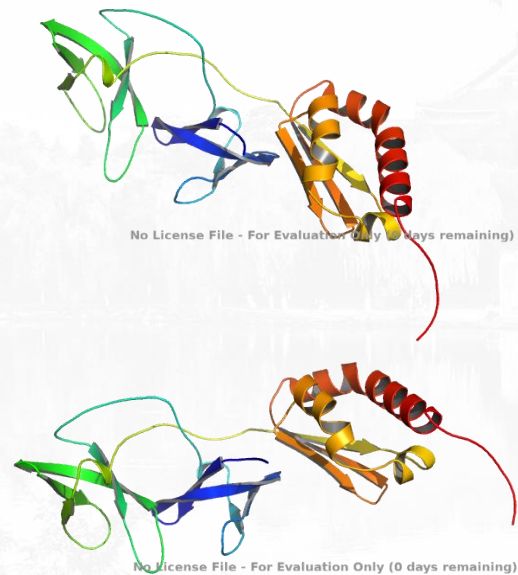
# FMRP发育树和保守结构域分析



# FMRP已表征结构域及全长结构预测



Mouse FMRP prediction



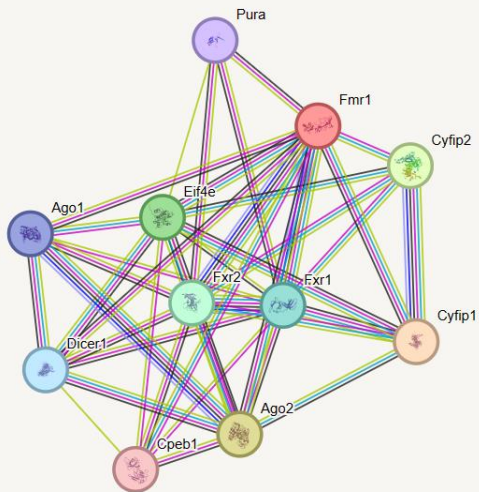
Human FMRP prediction

## 探究目的

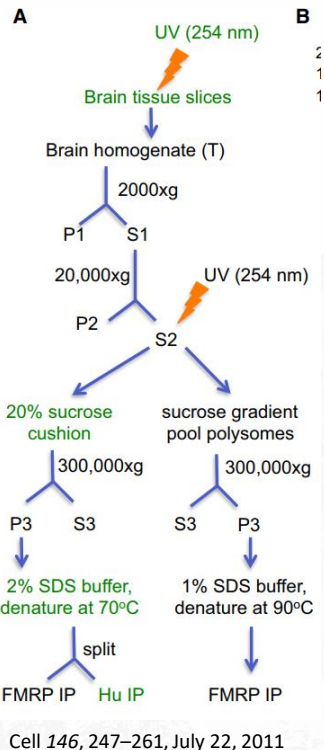
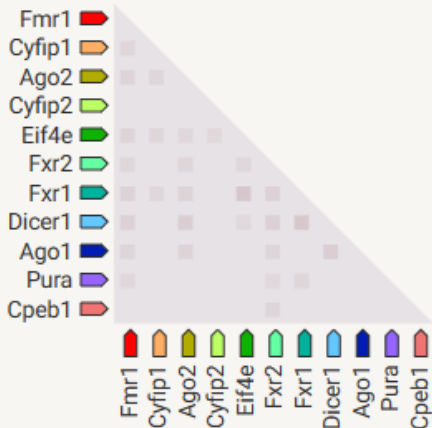
- 是否鼠源*Fmr1* 转录的蛋白质功能研究能够对人源FMRP功能有参考意义？
- FMRP和哪些蛋白质可以发生相互作用？

# 鼠源FMRP蛋白相互作用分析及共表达情况

STRING数据库:



observed Coexpression in  
Mus musculus:



- 利用交联免疫沉淀分离RNA，之后进行高通量测序，
- 鉴定出842个FMRP靶标mRNA

- FMRP可以和多种基因、蛋白质相互作用，主要包括但不局限于蛋白稳态、信号转导、细胞骨架、囊泡运输、细胞周期、染色质调控等。



## 课程感悟

G4A毕金龙:

G4B李戈:

G4C程佳琪:



**THANKS**