

进化和结构分析预测拟南芥**PRK6**的共受体

G01成员： **G01A 宁妮娜**

G01B 路菡

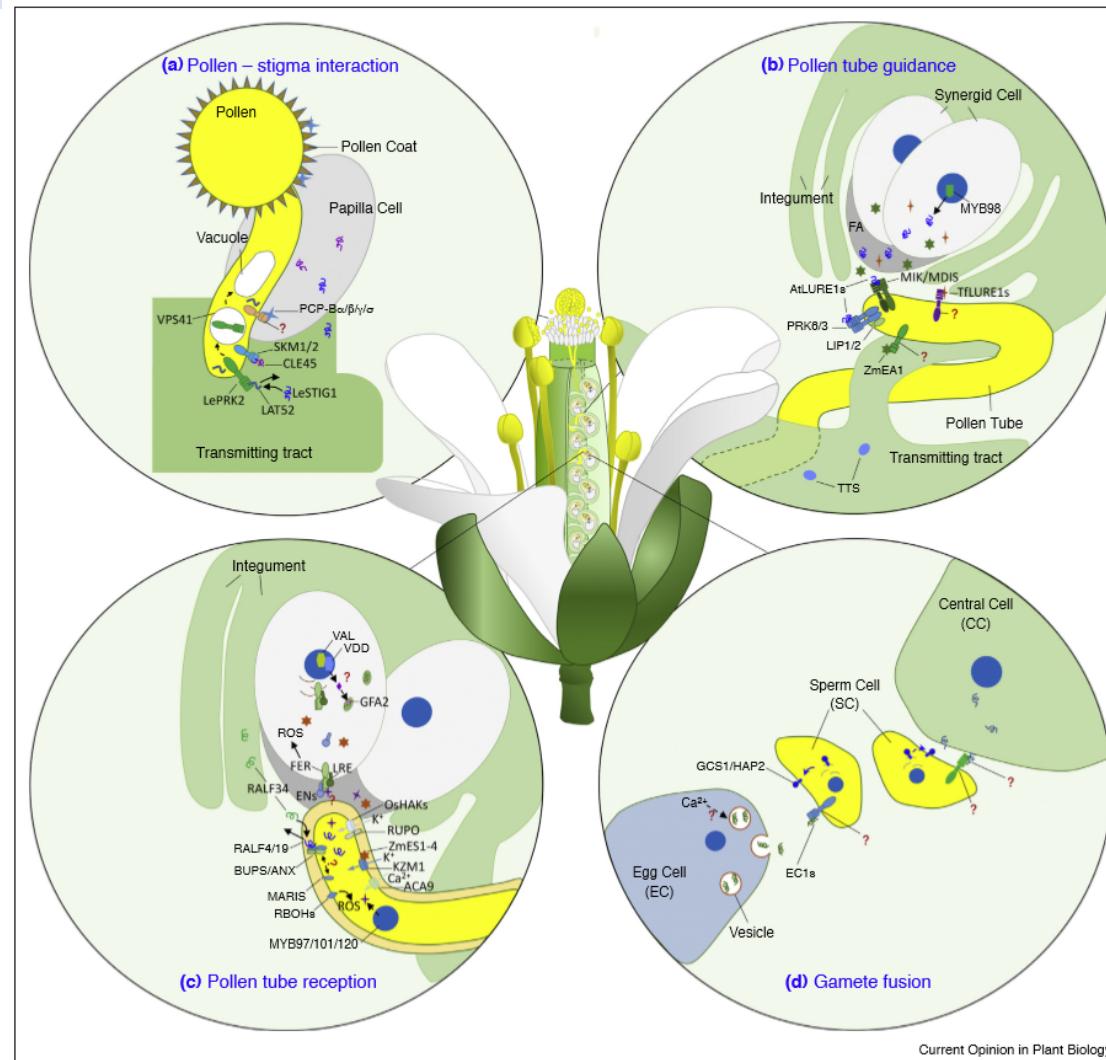
G01C 王鼎岳

G01D 李其昀

汇报人： **G01B 路菡**

BACKGROUND

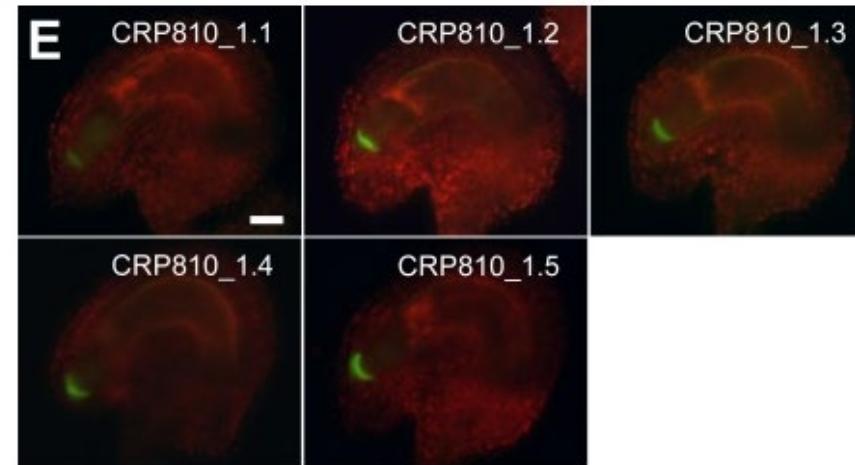
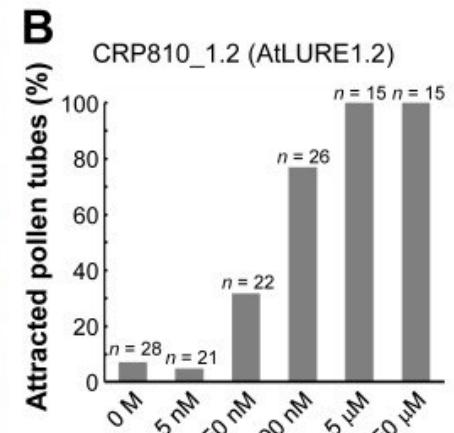
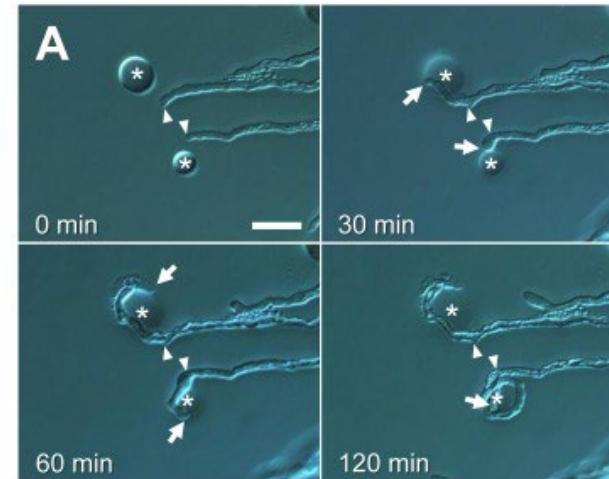
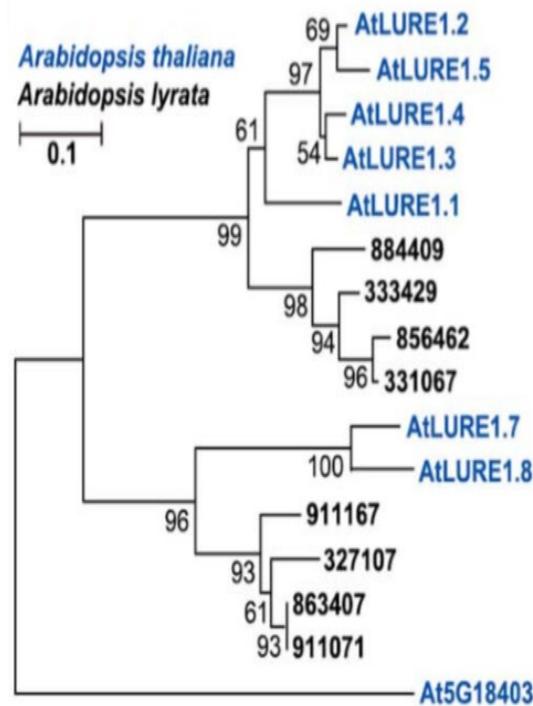
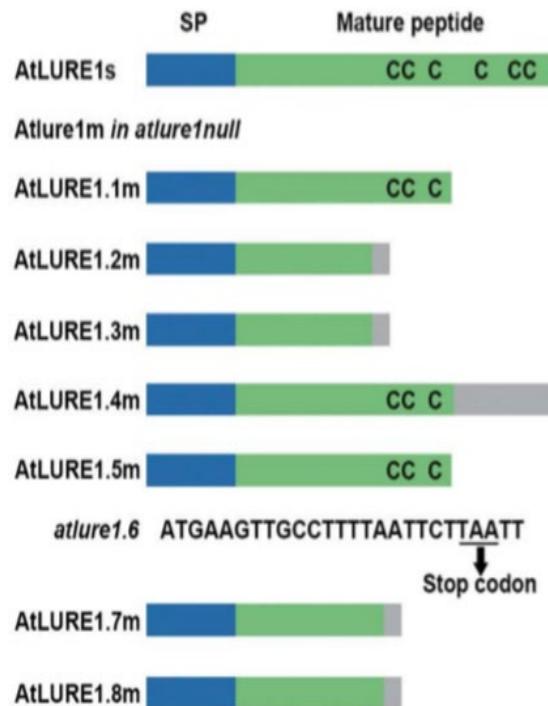
BACKGROUND



- 花粉粘附, 水合, 萌发
- 花粉管生长, 导向, 接受
- 配子激活, 融合

ATTRCTANT

AtLURE1s

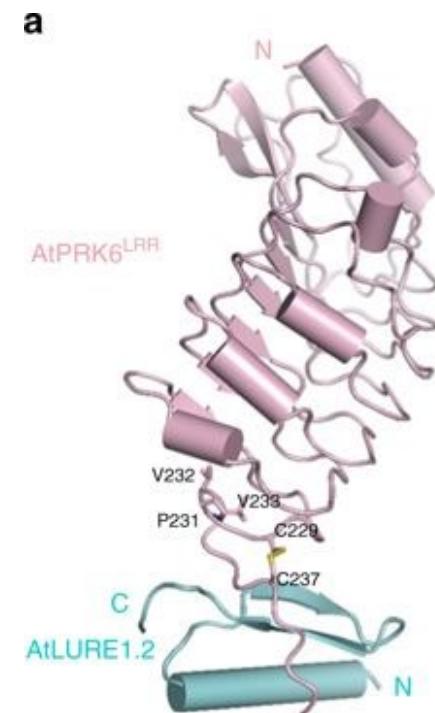
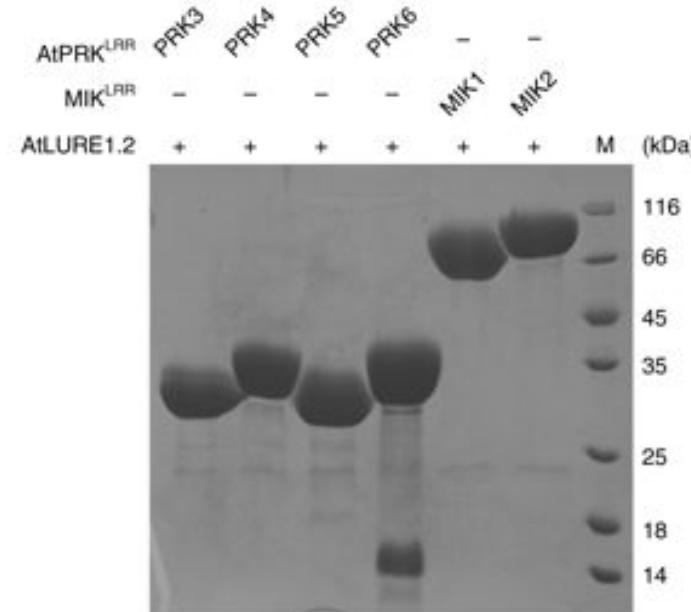
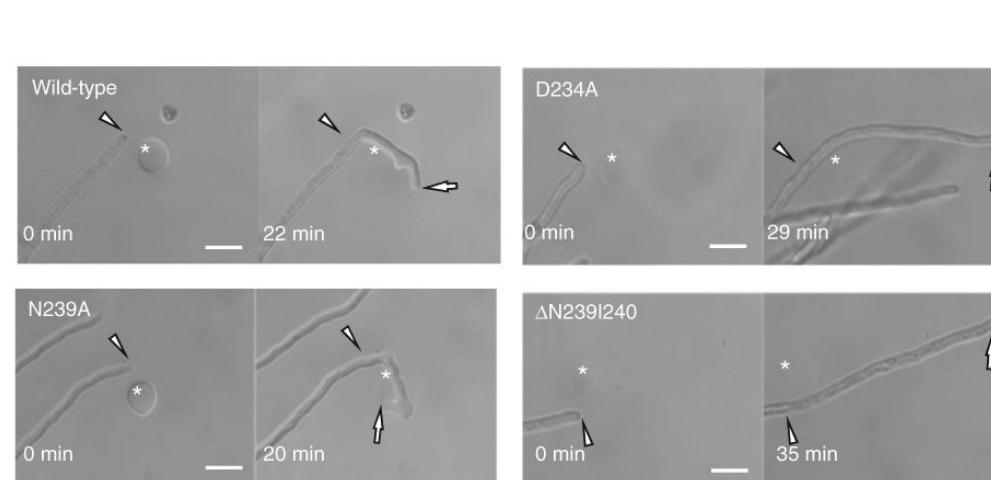


Zhong et al. *Science* 2019

Takeuchi et al. *PloS Biol* 2012

ATTRACTANT RESPONSE-PRK6 (pollen-specific receptor kinase 6)

AtLURE1s receptor PRK6 (pollen-specific receptor kinase 6)



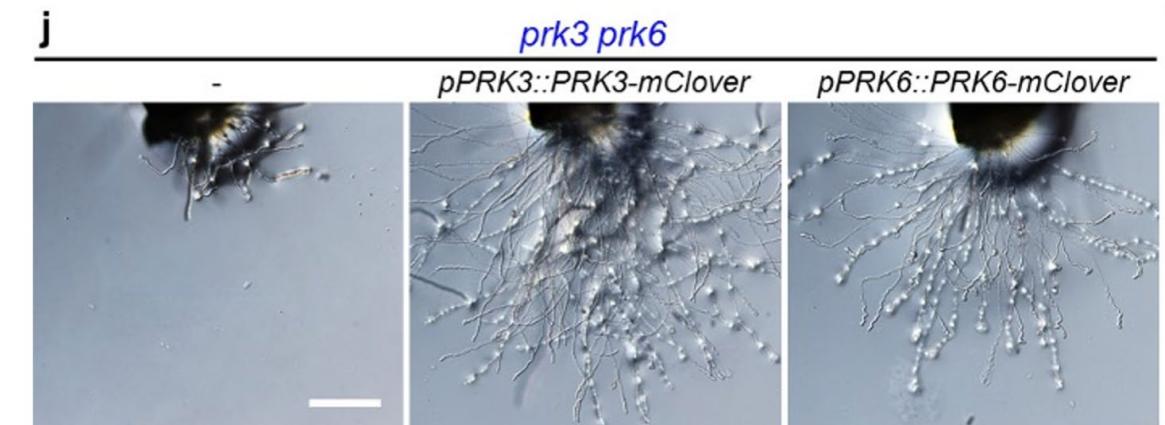
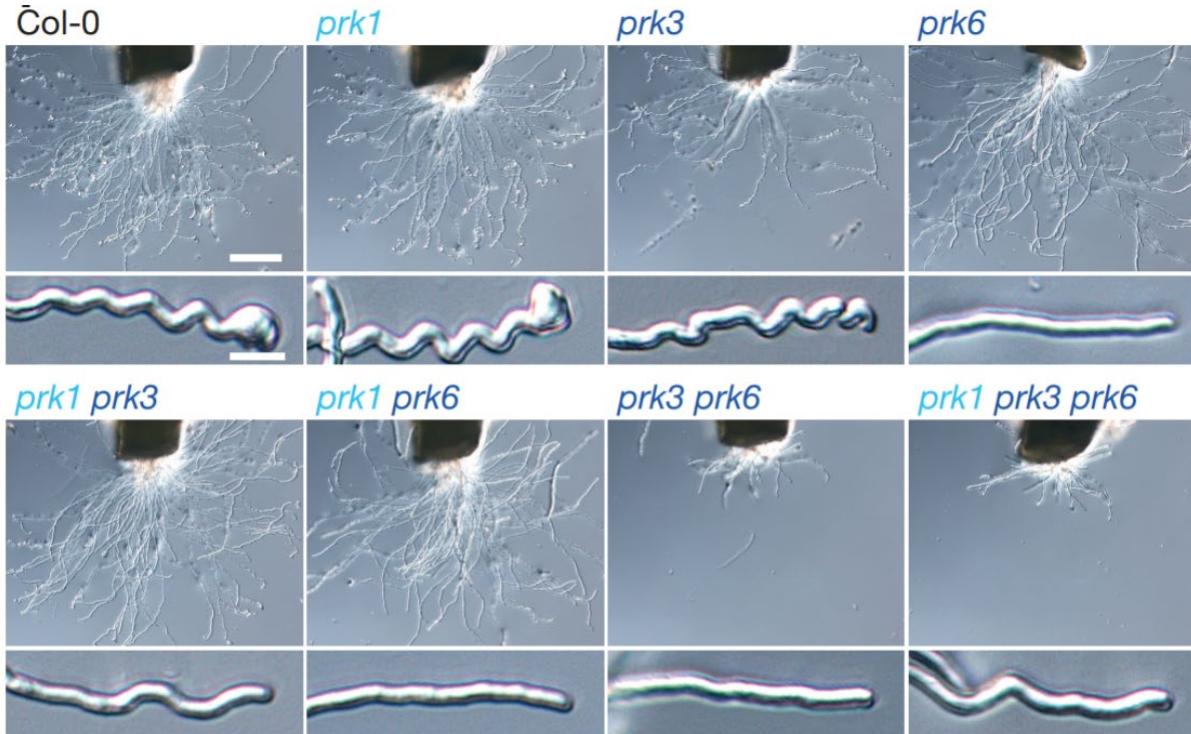
- AtLURE1.2 specifically interacts with AtPRK6^{LRR}
- Interaction between AtLURE1.2 and AtPRK6^{LRR} is mediated by a combination of polar and hydrophobic contacts

Takeuchi et al. *Nature* 2016

Zhang et al. *Nat Comm* 2017

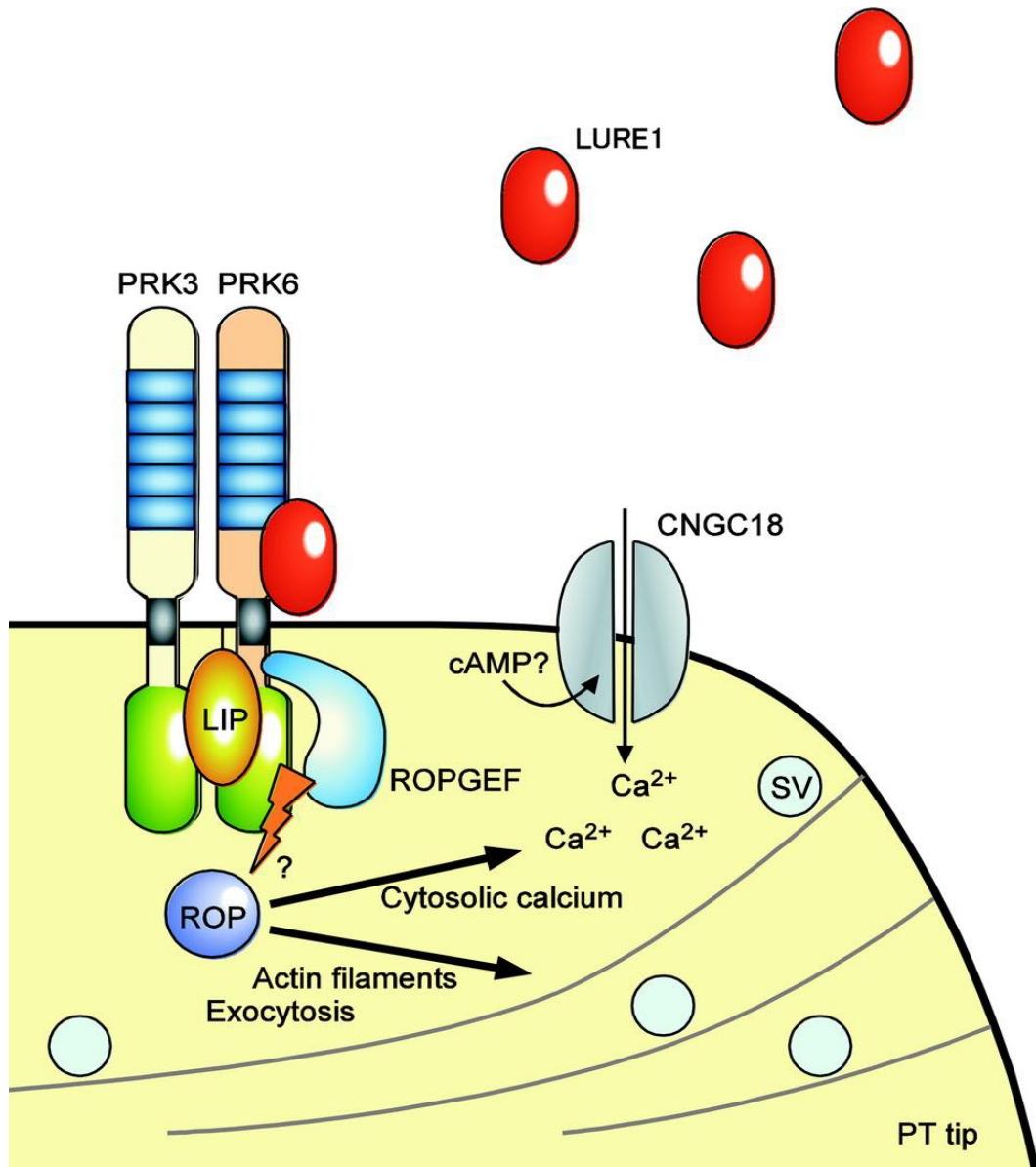
PRK FAMILY

PRKs in ovular guidance——PRK1/3/6



- *prk1 prk3* double mutations as well as a *prk6* single mutation **impaired the response to AtLURE1.2**
- **Expression of PRK3 in *prk3 prk6***, which showed similar tip localization to that of PRK6, restored the growth defect but **not the wavy response to AtLURE1**

Takeuchi et al. *Nature* 2016

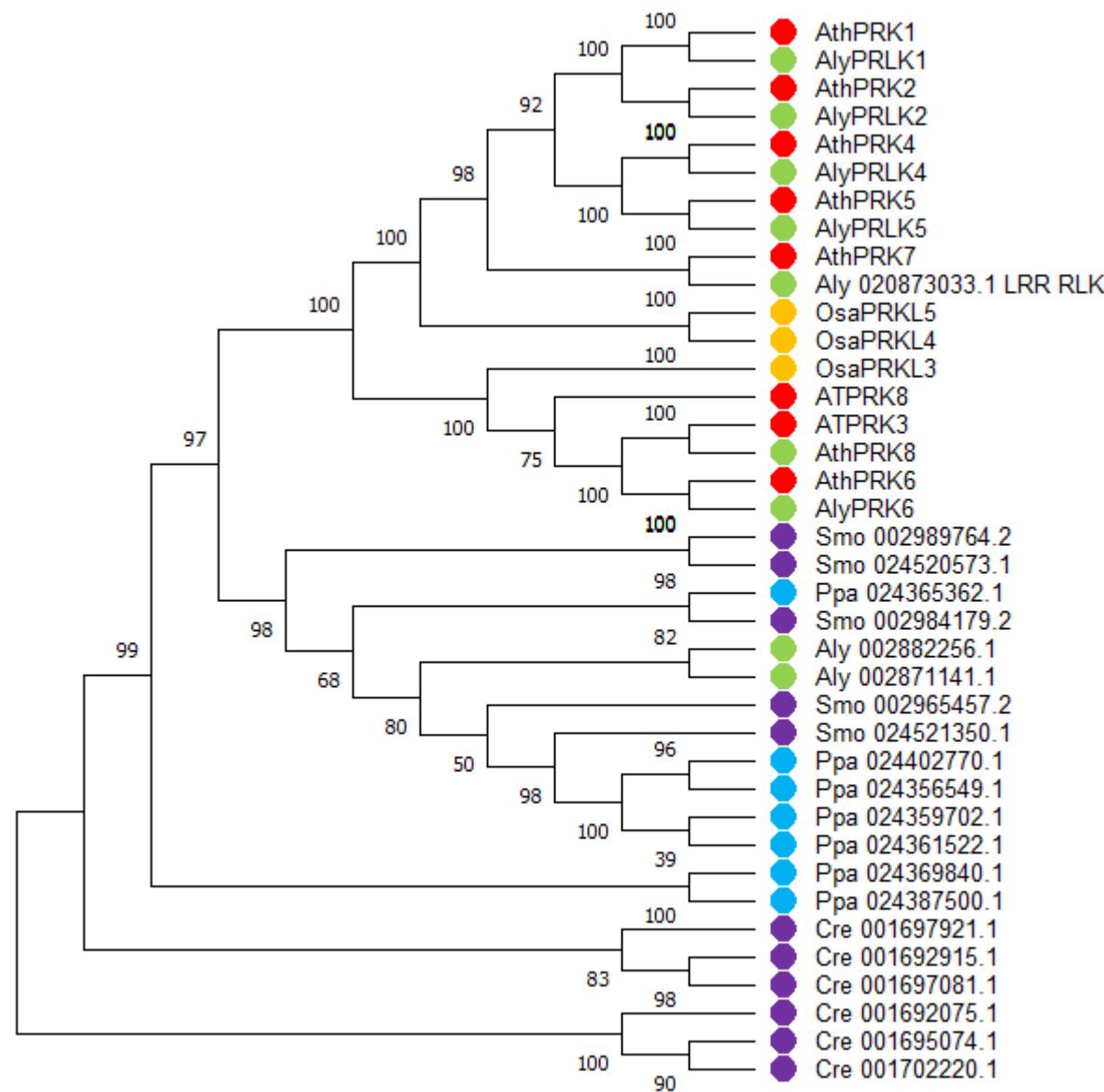


- 1. PRK6二聚化起作用
- 2. PRK3似乎可增强PRK6的作用

- PRK6结合LURE是否有共受体?
- 进化分析+结构预测

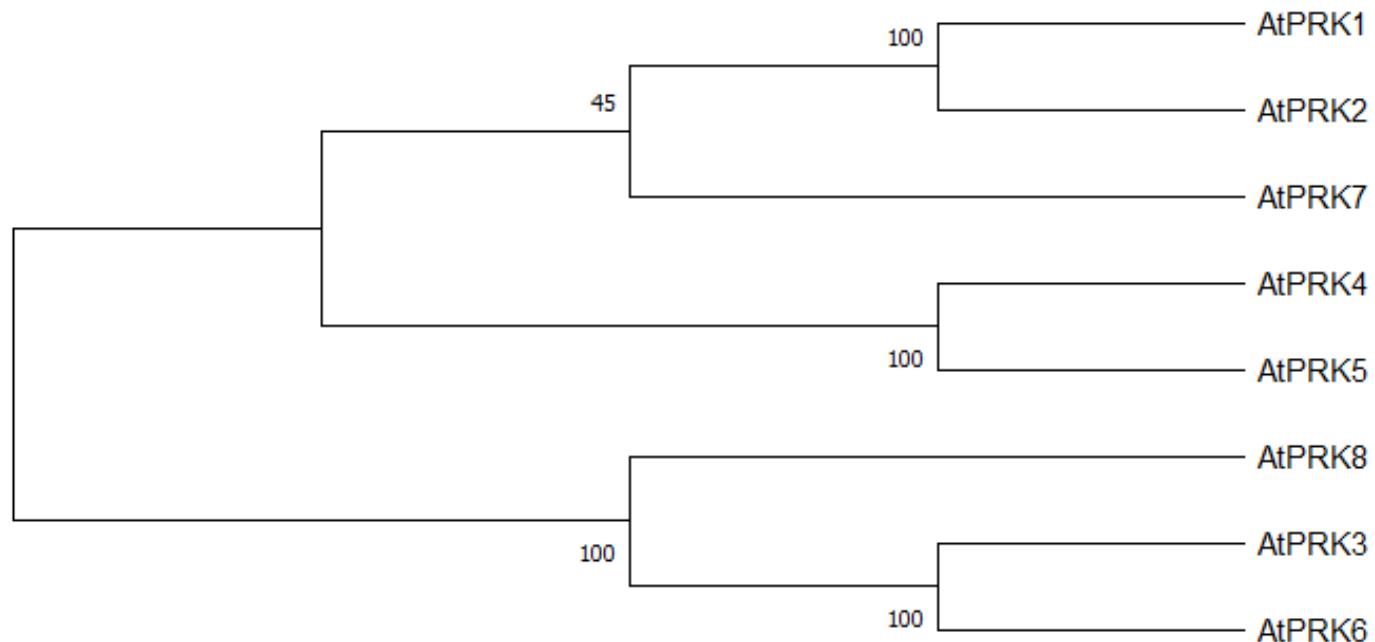
Phylogenetic analysis

Phylogenetic tree of plant PRKs



Ath - *Arabidopsis thaliana*, 拟南芥
Aly – *Arabidopsis, lyrata*, 琴叶拟南芥
Osa - *Oryza sativa*, 水稻
Smo - *Selaginella moellendorffii*, 江南卷柏
Ppa – *Physcomitrella patens*, 小立碗藓
Cre-*Chlamydomonas reinhardtii* , 莱茵绿藻

Phylogenetic tree of AtPRKs



Multi sequence alignment of AtPRKs

LRR domain

PRK6-AtLURE1 binding site

kinase domain

AtPRK7/1-676	313 RKLNEATVQRDSTATSGAISVGGLS-PDEDKRGDQRKLHFRVNDQERFTLQDMLRASAEVLGS-----GGFGSSYKAALSSGRAVVVKRFRFSNIGREEFYDHMKKIGRLSHPNLLPLIAFYRKEEKLLVTNYISNGSLANLLHANRTPGQVVLDPWIRLKIVRGVTRGLAYLYRVFPDLN489
AtPRK4/1-679	334 KAA-----DSVTSYTS--R-RGA-VPD-QNKLLFLQDDIQRFDLQDLLRASAEVLGS-----GSFGSSYKTGINSQMLVVVKRYKHMNNSVGRDEFHEHMRLRLKHPNLLPIVAYYYYREEKLLIAEFMPNRSLASHLHANHSVDQPGLDPWTRLKIQGVAKGLGYLFNELTTLT495
AtPRK5/1-686	337 NKP-----AESVNHT--R-RGS-MDPGGRLLFVRDDIQRFDLQDLLRASAEVLGS-----GTFGASYYKAIASSGQTLVVVKRYKHMNNSVGRDEFHEHMRLRLKHPNLLPILPLVAYYYYREEKLLVTEFMPNSSLASHLHANN--SAGLDWITRLKIKGVAKGLSYLFDELPTLT495
AtPRK1/1-662	315 KKAD-----HRKGSGTTK--R-MGAAAGVENTKLSFLREDREKFQDQLLKASAEILGS-----GCFGASYYKAVLSSGQMVMVVKRFQKOMNNAGRDEFQEHMKRLRLMHNNLLSVAYYYYREEKLLVCDFAERGSLAINLHSNQS LGKPSLDWPTRLKIVKGVAKGFLYHQLDPLSLM480
AtPRK2/1-647	306-----HSQ-----N-RAAKMIHTTKLFLRDDKGKFELQDLLKASAEILGS-----GCFGASYYKTLLSNGSVMVVKRFKHMNSAGIDFQEHMKRLRLHNHENLLPVAYYYYKEEKLFVSDFVANGSLA AHLHGHSLGQPSLDWPTRFLNIVKGVGRGLLYLHKNLPSLM461
AtPRK8/1-644	310 QS-----RSSRSG-----ELNKVGAVGTSQDLMVNKEKGVFRLSDLMKAIAAHVLGNPGGGSNRPRSSGGVGSAYKAVLSNGVTVVVKRVTVMNOVSVDFTKEIRKLGSLQHKNVLTPLAYHFRODEKLLVFEFPVNLLNHLRHDGEFF--QLDWPSSRLKIQGIARGMVYLHRELGFN478
AtPRK6/1-659	330 RKRGHSDDGSTKKGSNSIKGGNGGGGGALGGGGMDIIMVNTDKGSFLPDLMAAAEVLGN-----GSLGSAYKAVMTTGLSVVVKRIRDMNQLAREPFMRRFGKLHPRNIPITPLAYHYRREEKLVSEYMPKSSLLYVLHGDRGIVYHSELTWATRKLQI QGVAHGMLKFHEE FASYD507
AtPRK3/1-633	310 KKRS---NAEGSSKKGSNSHNGKG---AGGGPSSGGMGDJIMVNSEKGSFGFLPDLMKAAAEVLGN-----GSLGSAYKAVMANGLISVVKRIRDMNKLAREAFDTMREGFKLHPRNIPITPLAYHYRREEKLVSEYMPKSSLLYVLHGDRGIVYHSELTWATRKLQI QGVAHGMLKFHEE FASYD481

AtPRK7/1-676	490 LPHGHLKSSNVLLDPNFEPPLTDYALPVVNQRDSQQFMVAYKAPEFTQQ--DRTSRSDWSLGLIILEI LTGKFPA NYLRQKGADDELA AAWESVARTEWTADVFDEKMAGKEH---EAQMLKLK IGLRCDDW DIEKRIELHEAVDRIEEVDRDAGGGQESVRSSYVTASDGDRSSRAMTEE 672
AtPRK4/1-679	496 IPHGHLKSSNVLLDESFEPLLTDYALRPVMNSEQSHNLMSYKSPEYSLK--GHLT KKT DWCLGV LLELTGRFPE NYLSQGYDANMSLTVWSNMVK EKKTGDVF DKE MTGKNC---KAEMLNLLK IGLSCCEEDEERRM ERM RDAVEKIERLK-EGEF-DNDFAST----THNVFASRLIDDD 671
AtPRK5/1-686	496 IPHGHMKSSNIVLDDSFEPPLTDYALRP MMSSEAHNFMTAYKSPEYRPSKGQ I T KKT DWCLGV LLELTGRFPE NYLSQGYDANMSLTVWSNMVK EKKTGDVF DKE MTGKNC---KAEMLNLLK IGLSCCEEDEERRM ERM RDAVEKIERLK-EGEF-DNDFAST----THNVFASRLIDDD 678
AtPRK1/1-662	481 APHGHKLSSNVLLTKTFEPLTDYGLIPLI NQEKAQM HMAA YRSPEYLQH--RRITKKT DWCLGV LLELTGKFPA NFQ S---SSEEDLASWNSGFHGWAPS LFDKGGMGKTSHC---EGQILKLLTIGLNCCEP DV EKRLD I GQAVEKIEEL-KEREGDDDDFYSTYSETDG--RSSKGESCE 657
AtPRK2/1-647	462 APHGHKLSSNVLLSEKFEPPLMDYGLIPMI NEESAQELMVAYKSPEYVKQ--SRVT KKT DWCLGV LLELTGKLLESFSQV DK ESEEDLASW RSSFKGEWT QEL F DQE MGKTSNC---EAHI LNLMR I GLSCCEV D V EKRLD I REAVEK MEDLMKEREQGDDDFYSTYASEADG--RSSRGLSSE 642
AtPRK8/1-644	479 LP HGNLKSSNIFLAEDGEPLI SEFGLQKL INPDQASQSLVAFKSP EADRD--GTVSAKSDVFSFGVV LLELTGKFPSOYAGLNRAGGANL VEWLGS AL E QGGWM DLLH PMPV TAAEDKIMEEEIENVL RIGVRCTREDP DORPNMTEV D E LTI EDSN----DDF ITI ET----- 644
AtPRK6/1-659	508 LP HGNLKSSNVL SETYEPLISDYAFLPLQPSNASQALFAKTFPEFAQT--QVQVHSKSDVYCLG I LLELTGKFPSOYLN N-GKGGD I VQWQSSVAE QKEEL IDPEI VNNTESM--RQM VELL RIGVACI ASNP DERLDLMR EAVR RLIEQVKT----- 659
AtPRK3/1-632	482 LP HGNLKSSNVLL SETYEPLISDYAFLPLIOPNNAQSALFAKTFPEFAQVON--D0VSPKSDVYCLG I LLELTGKFPSOYLN N-GKGGD I VQWQSSVAE QKEEL IDPEI VNNTESM--RQM VELL RIGVACI ASNP DERLDLMR EAVR RLIEQVKT----- 623

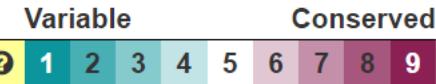
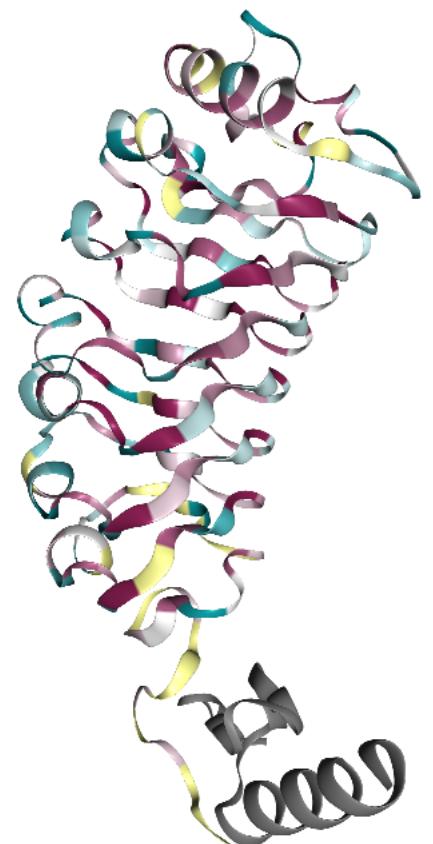


北京大學
PEKING UNIVERSITY

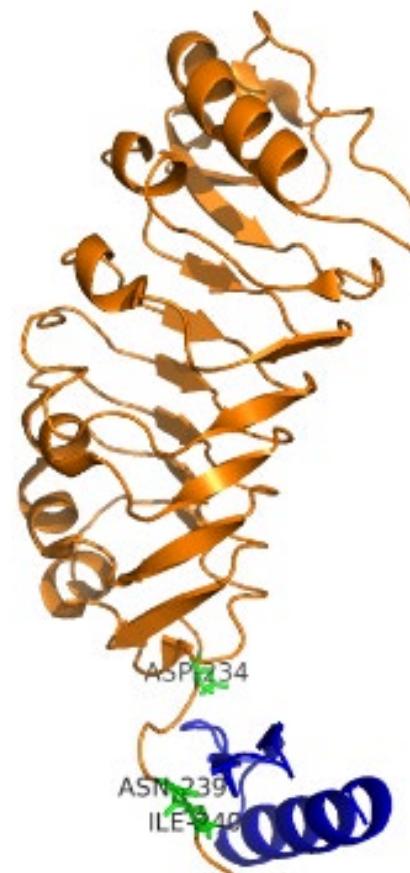
Structure analysis

Mutant structure prediction of PRK6

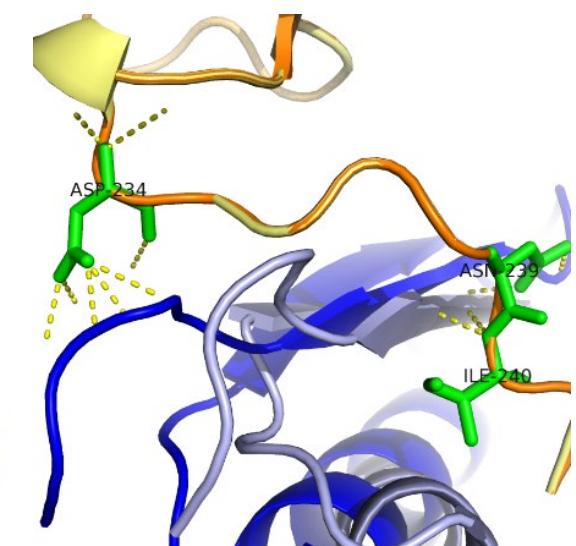
WT PRK6 保守度分析



WT PRK6-AtLURE1.2



WT PRK6-ZDOCK PRK6 结构对比



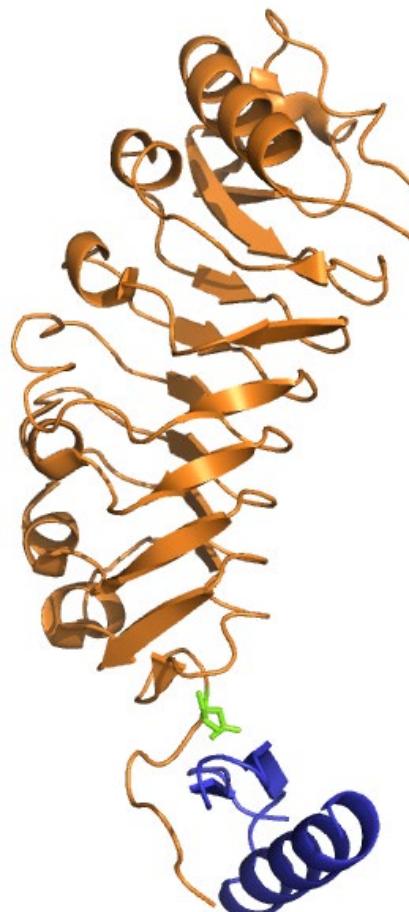
北京大学
PEKING UNIVERSITY

Mutant structure prediction of PRK6

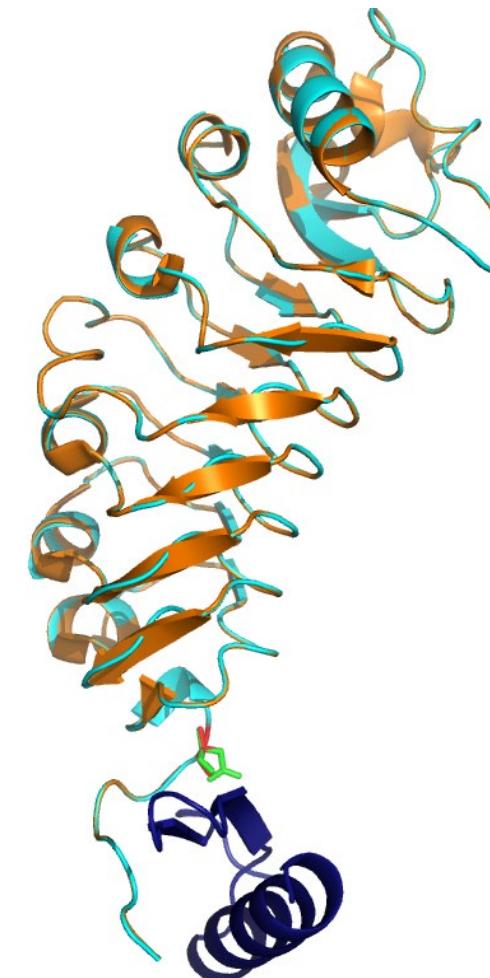
WT PRK6 保守度分析



WT PRK6-AtLURE1.2

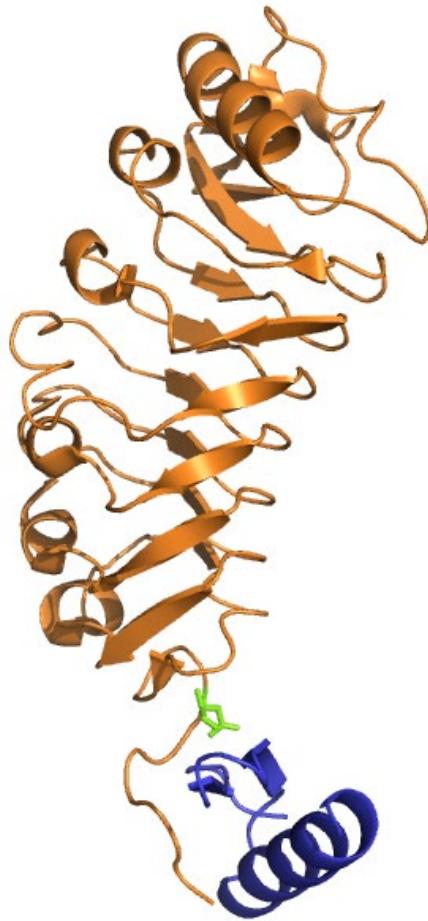


D234A PRK6-WT PRK6 结构对比

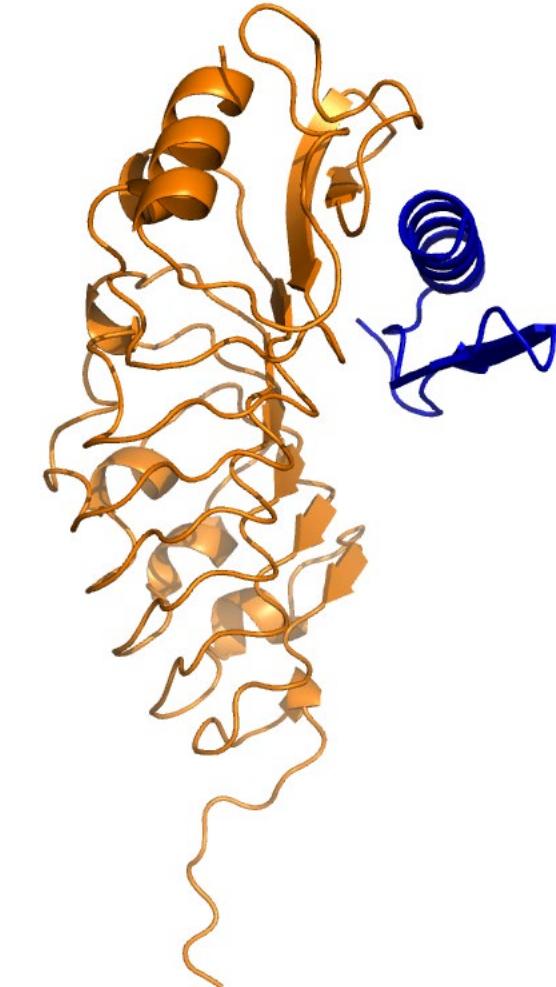
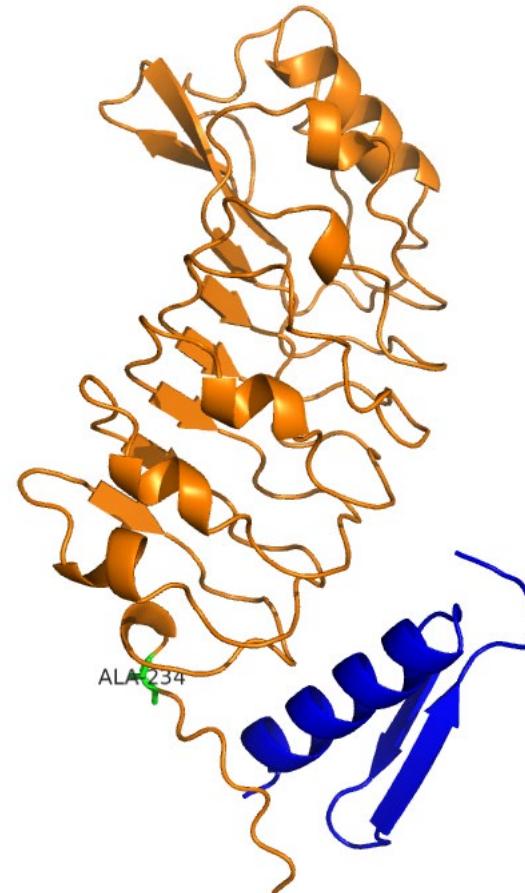


Molecular docking prediction of mutant PRK6 and AtLURE1.2

WT PRK6-AtLURE1.2



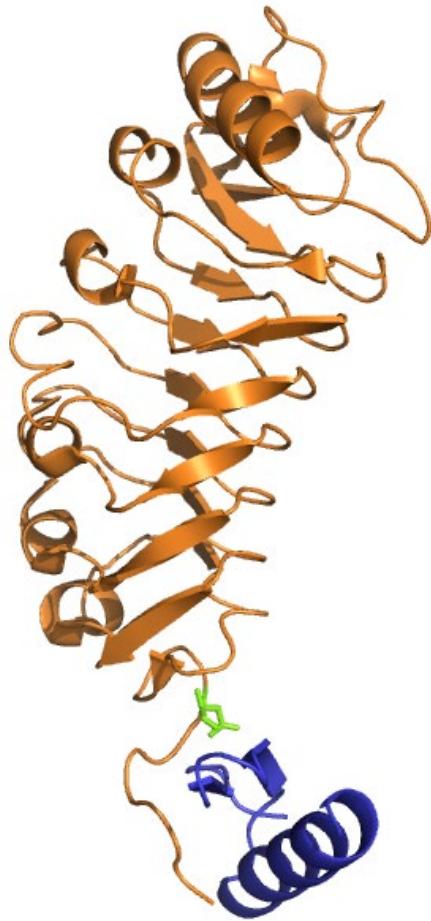
D234A PRK6-AtLURE1.2 ZDOCK对接预测



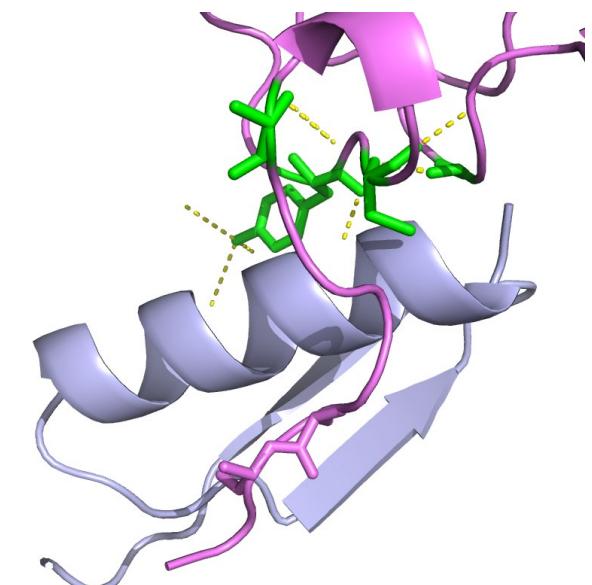
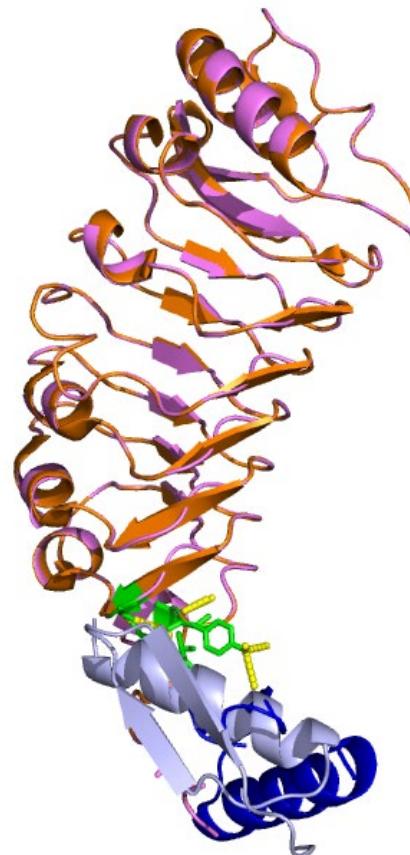
➤ 初步认为突变结合位点后的PRK6丧失与LURE的结合能力

Molecular docking prediction of mutant PRK6 and AtLURE1.2

WT PRK6-AtLURE1.2



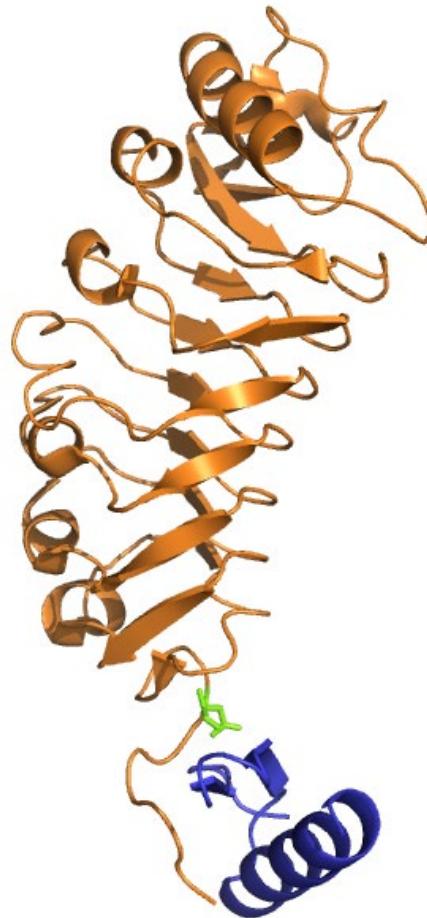
D234A N239A I240A PRK6-AtLURE1.2 ZDOCK对接预测



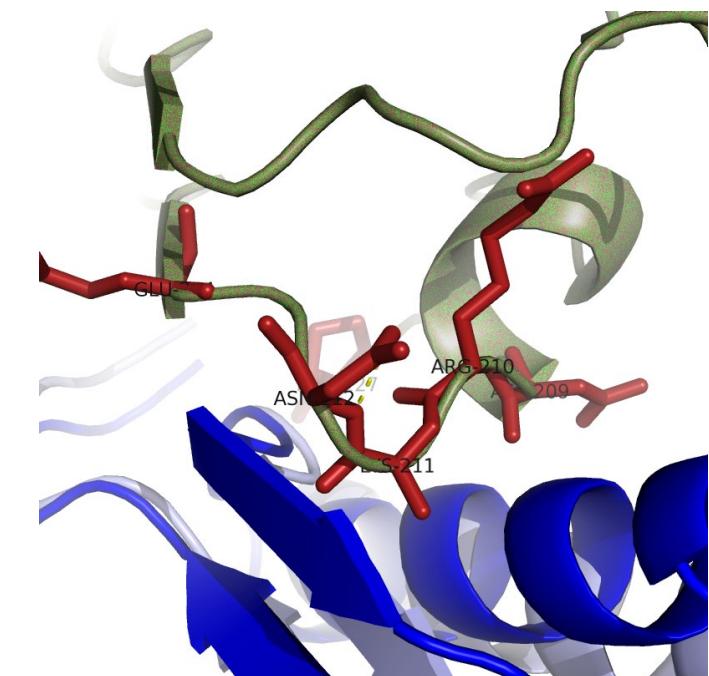
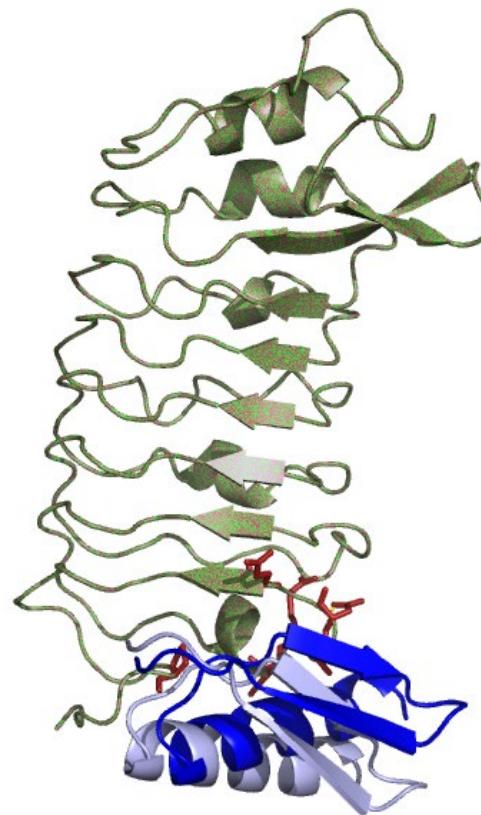
- 初步认为突变结合位点后的PRK6丧失与LURE的结合能力，无有效氢键

Molecular docking prediction of PRK3 and AtLURE1.2

WT PRK6-AtLURE1.2



WT PRK3-AtLURE1.2 ZDOCK对接预测



PRK3-AtLURE1.2实际上无有效氢键联系

- 初步认为PRK3不具备与LURE的结合能力

总结

- 1.与PRK6亲缘关系最近的 PRK3或许不是PRK6结合AtLURE1的共受体，或者是共受体但不负责结合AtLURE1。
- 2.应通过生化、遗传实验进一步证实
- 3.应考虑其他家族受体作为PRK6共受体的可能性。

Thanks!