

# 开花调控基因CONSTANS的生物信息学分析

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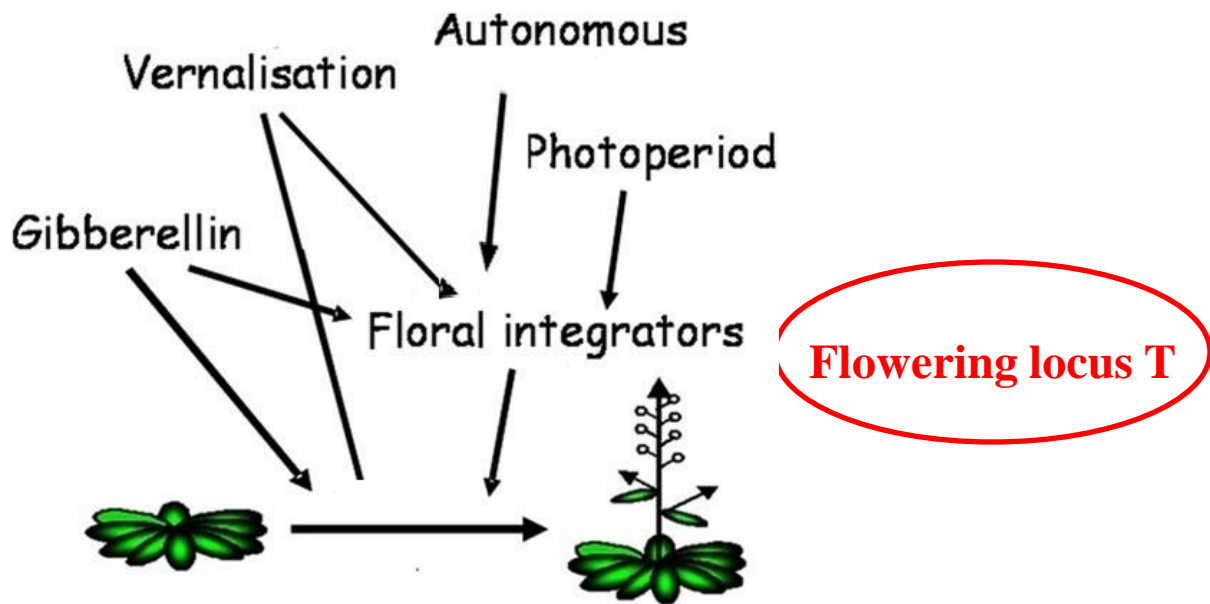
**背景介绍**

**CO序列及结构分析**

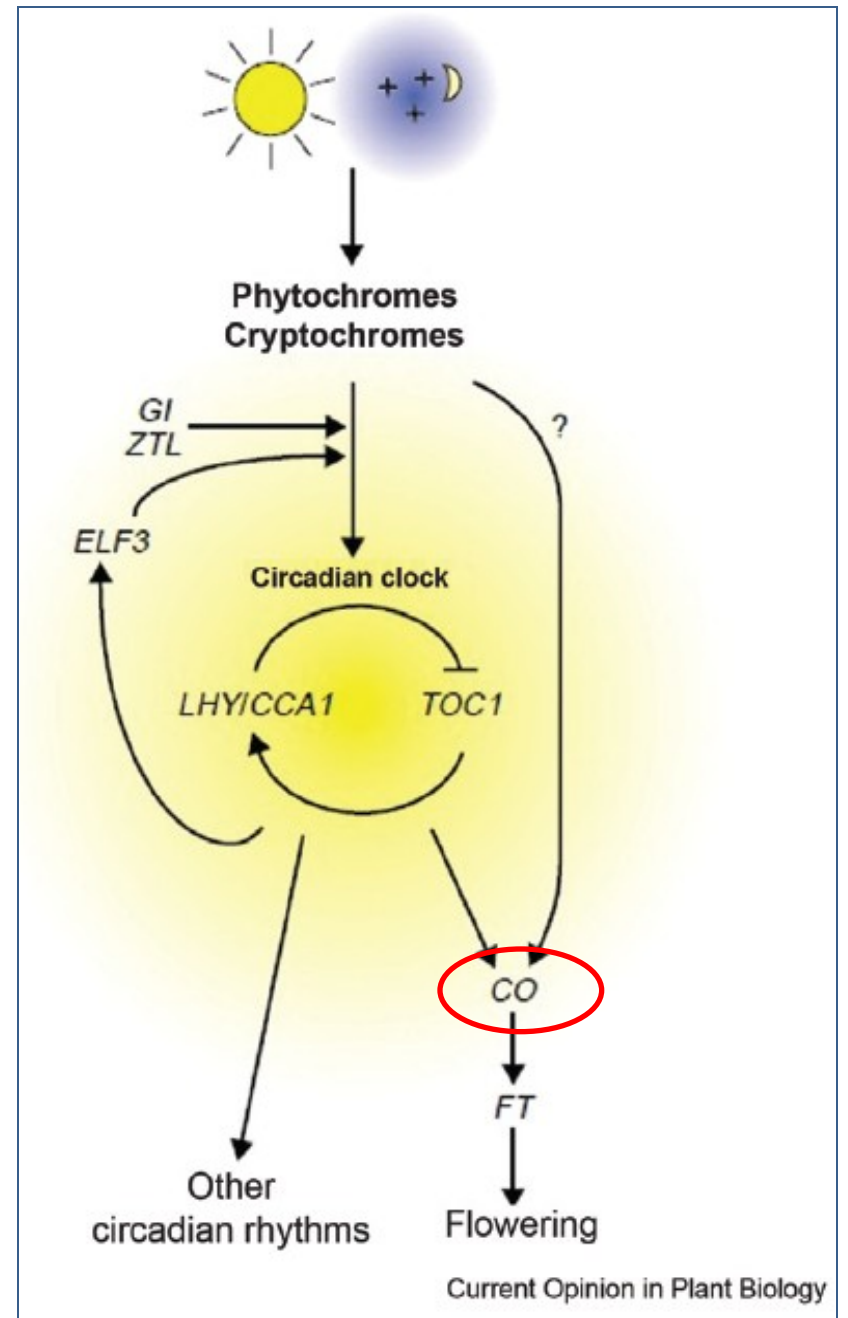
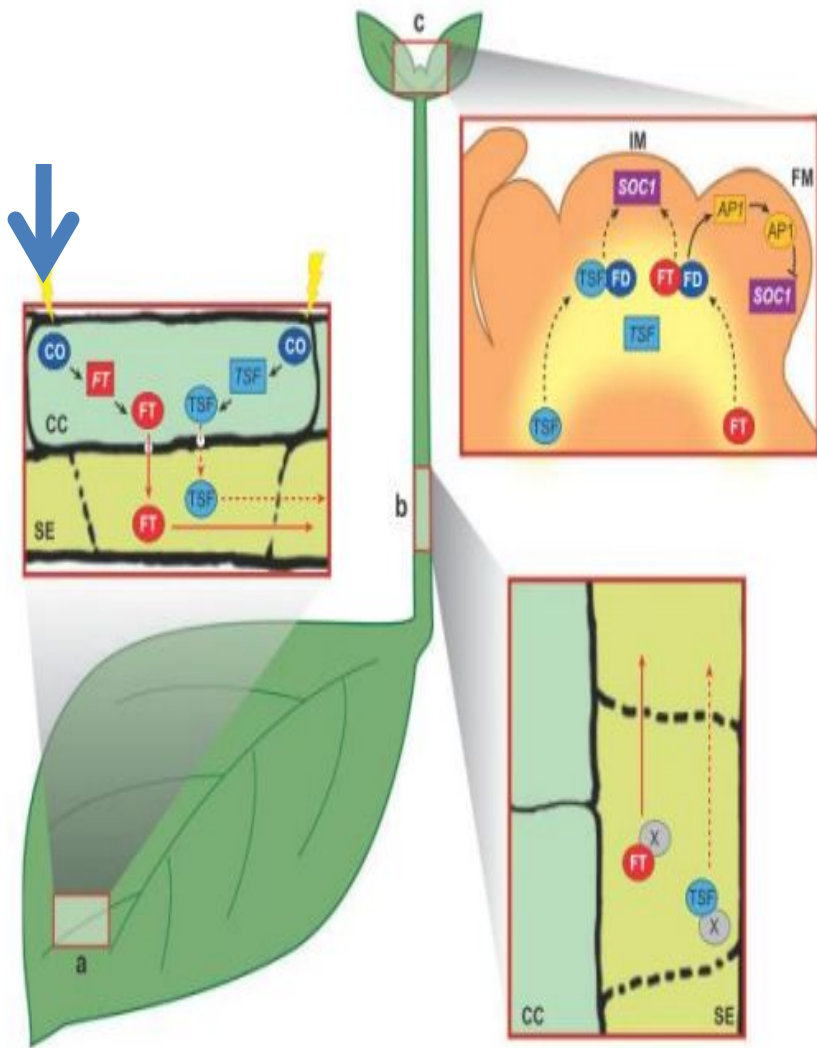
**蛋白质结构预测**

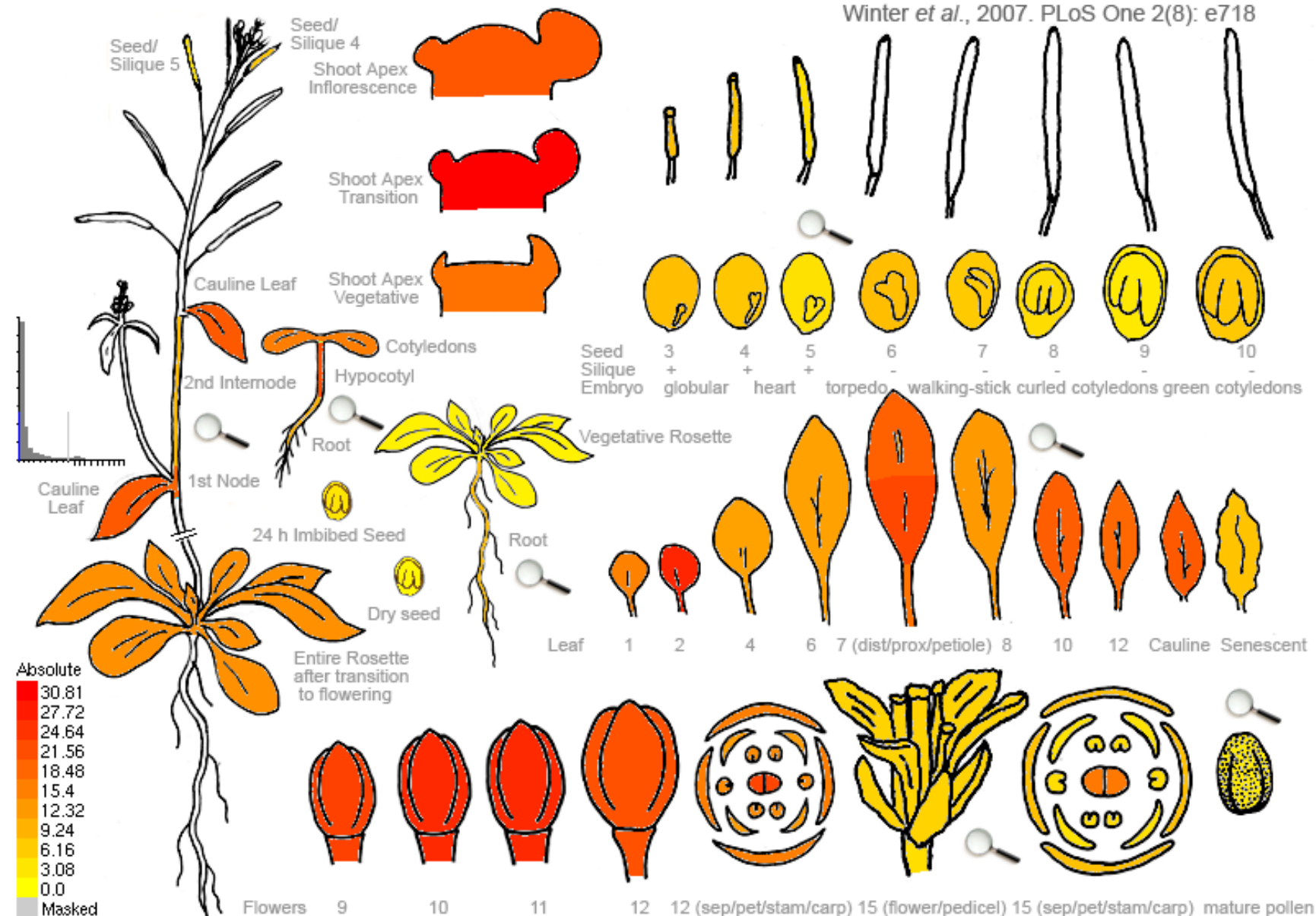
# 背景介绍

CONSTANS: 1995年从拟南芥晚花突变体中发现。  
可以整合光信号和生物钟信号，节律性地激活  
Flowering locus T 的表达，从而诱导开花。

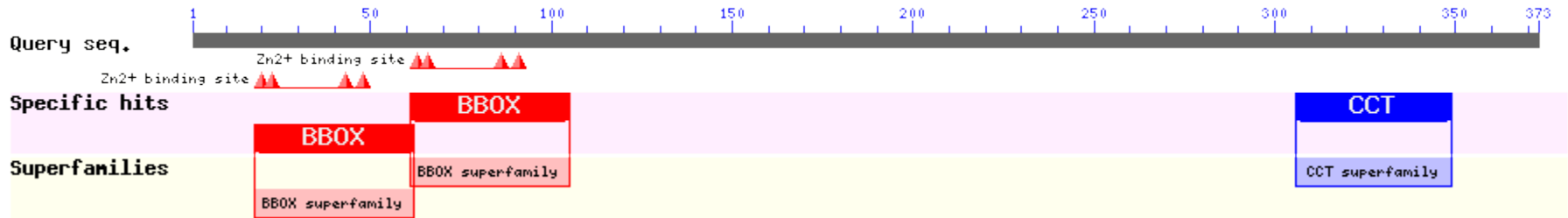


调控开花的四种途径：





# CDD

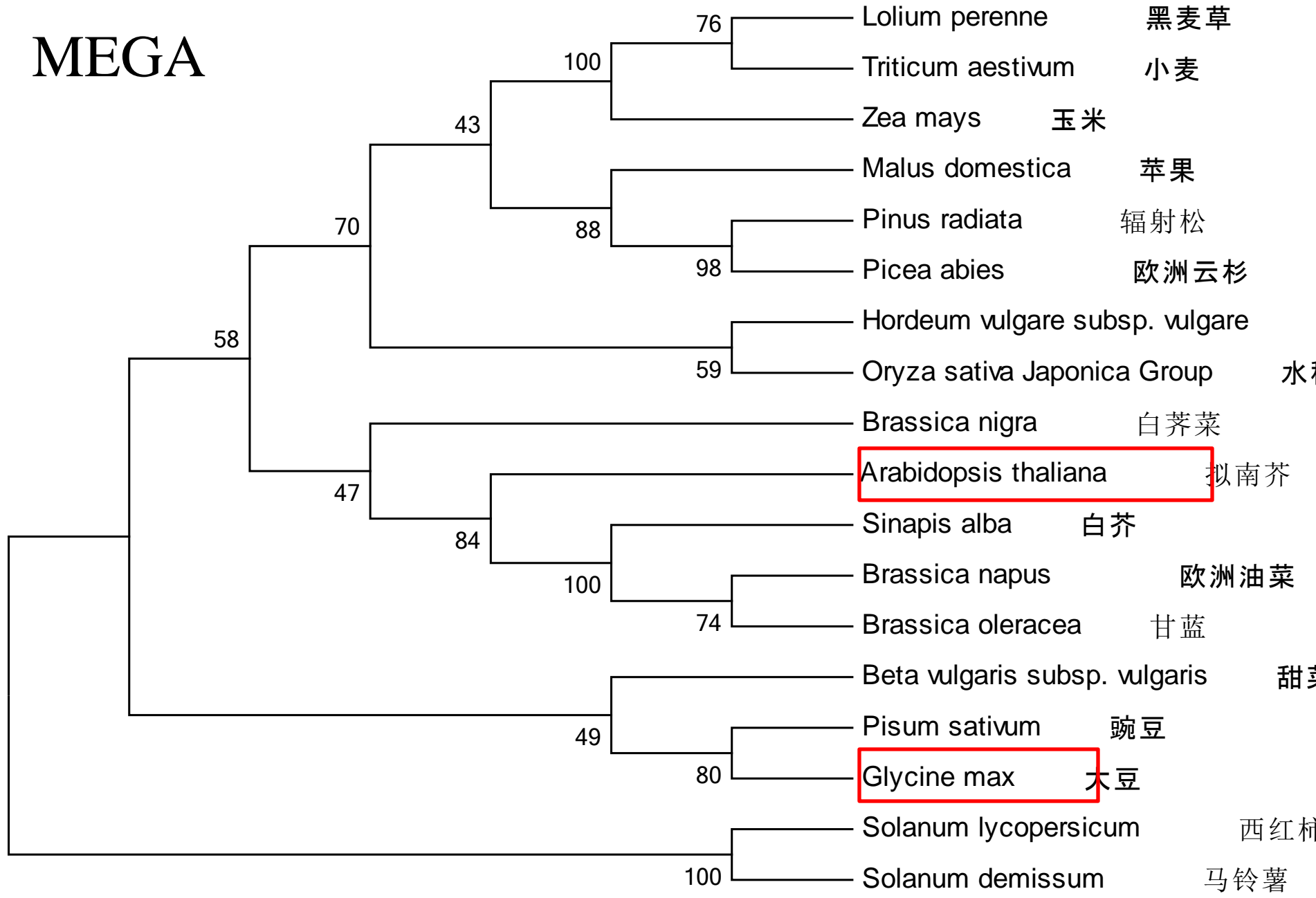


B-BOX： 锌指蛋白。

结合锌离子后能够稳定  $\alpha$ -螺旋结构，使  $\alpha$ -螺旋镶嵌于DNA的大沟中，因此含锌指结构的蛋白质都能与DNA或RNA结合。

CCT： 具有核定位信号

# MEGA



# 拟南芥CO家族成员序列比对

## N 端

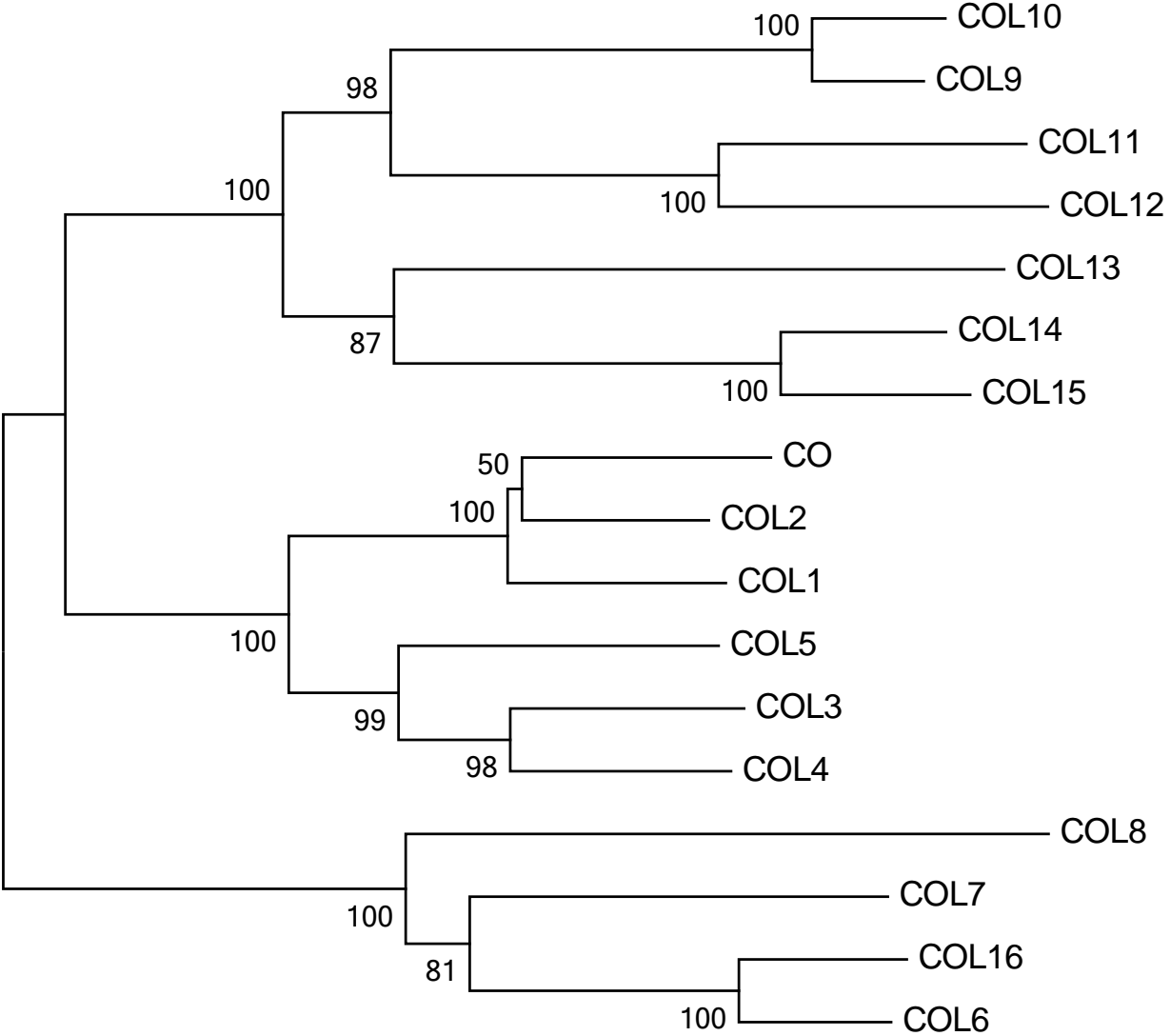
MLKQESNDIGSGEN	NRARPCDTCRSN	ACTVYCHADSAYLCMSQDAQVHSANRVA SRHKRVRCESCEERAPAAFLCEADDA	LCTACDSEVHSAN	PLARRHQVRPIL	106	Q39057	CONS_ARATH			
MEPEKCDHCATS	QAL IYCKSDLAKLCLMCDVHVHSANPLSHRH IRLSLICEKCFSPQAA I RCLDERVS	YCQGCHWHE SMC	SELGHRVQSLN	89	Q9LJ44	COL12_ARATH				
ML	XVESNWAQA	CDTCRSA	ACTVYCRADSAYLCSSCDAQVHAANRLA SRHERVRCQSCERAPAAFFCKADAAS	LCTTCDSEIHSAN	PLARRHQVRPIL	98	050055	COL1_ARATH		
MLKEESNESG	TWARA	CDTCRSA	ACTVYCEAD SAYLCTTCDARVHAANRVA SRHERVRCQSCESAPAAFLCKADAAS	LCTACDAEIHSAN	PLARRHQVRPIL	102	Q96502	COL2_ARATH		
MGYM	CDFCGEQ	RSVVYCRSDAACLCLCDNRVHSANALSKRH SRTLVCERCNAQPA SVRCSDERSV	LCQNCDSWGHGKNS	TTTSHKRRQTI N	93	Q9LUA9	COL10_ARATH			
MEAEEGHQDRDL	CDYCDSS	VALVYCKADSAKLCLACDKQVHVANQLPAKHFRSLLCDSCNE SPSSLFCETERSV	LCQNCDSWGHHTAS	SSLHSRRPFE	97	082256	COL13_ARATH			
MGTSTTESVVA	CEFCGER	TAVLFCRADTAKLCLPCDQVHSANLLSRKH VRSQICDNC SKEPV SVRCFTDNLV	LCQECDWDVHG SC	SSSATHERSAVE	98	022800	COL14_ARATH			
MASSSRL	CDSCKST	AATLFCRADAAPFCGDCDGKIHTANKLA SRHERVWLCVEVCEQAPAHVTCADAAA	LCVTCDRDIHSAN	PLSRRHERVPIT	94	Q9SK53	COL3_ARATH			
MGFGLSEIKSISGGWGAAARS	CDACKSV	TAAVFCRVD SAFLCIACDTRIHSE	TRHERVWVCEVCEQAPAAVTCADAAA	LCVSCDADIHSAN	PLASRHERVPVE	104	Q9FHH8	COL5_ARATH		
MFCAEIMI SKYQEDVQAPRA	CELCLNK	HAVVYCA SDDAFLCHVCDSEVHSANHVATKHERVCLRTNEI S	NDVR	GGTTLT SVWH	SGFRKART	92	Q9M9B3	COL8_ARATH		
MEAR	CDFCGTE	KAL IYCKSDSAKLCLMCDVNVHSANPLSQRHTRSLLECKCSLQPTAVHGMNENVS	LCQGCQWTA SMC	TGLGHLQSLN	89	023379	COL11_ARATH			
MSSSERVP	CDFCGER	TAVLFCRADTAKLCLPCDQVHTANLLSRKH VRSQICDNC NEPV SVRCFTDNLV	LCQECDWDVHG SC	SVSDAHVRSAVE	95	Q9C7E8	COL15_ARATH			
MM	KSLANAVG	AKTARA	CDSCVKR	RARVYCAADDAFLCQSCDSL VHSANPLARRHERVRLKTA SPA	VVKHSMHSSA SPPHEVATWH	HGFRKART	94	Q8RWD0	COL16_ARATH	
MA SKL	CDSCKSA	TAALYCRPDA AFLCLCD SKVHAANKLA SRHARVWVCEVCEQAPAHVTCADAAA	LCVTCDRDIHSAN	PLARRHERVPVT	92	Q940T9	COL4_ARATH			
MM	KSLA SAVG	AKTARA	CDSCVKR	RARVYCAADDAFLCHACDC SVHSANPLARRHERVRLK SASAG	KYRHA	S PPHQATWH	QCFRKART	88	Q8LG76	COL6_ARATH
MVVDVESRTA SVTGEKMAARG	CDACMKRSRA	SWYCPADDAFLCQSCDA SIHSANHLAKRHERVRLQSSSPT	ETA	DKTTSVWY	EGFRKART	91	Q9C9A9	COL7_ARATH		
MGYM	CDFCGEQ	RSVVYCRSDAACLCLCDNRVHSANALSKRH SRTLVCERCNAQPA TVRCVEERVS	LCQNCDSWGHNSNNNNSSSSTSPQQHKKRQTI S		100	Q9SSE5	COL9_ARATH			

VEQQP	DPASQ	MITVTIQLSPND	REARVRLYREKRETRREFEET IRYASREKAYAE I RPRVNGRFAKREIEA	FEQGF	NTMLNWTG	YGVPSF	373	Q39057	CONS_ARATH			
MEEA	P	WEIN	FEVSC	PQARWFAKLRYEPEKELERS	FGDQ	IRYASREKARADITRERVEGRFVVEAGISYDVPSSPT	---	TNN	364	Q9LJ44	COL12_ARAI	
TIQPP	YPPAQ	---	LSPRD	REARVRLYREKRETRREFEET IRYASREKAYAEERPRI	EGRFAKRE	EDVDEFAWQAF	---	STMIIFDTG	YGVPSF	355	050055	COL1_ARATH
IDQLS	GPPTQV	---	VQQLTP	REARVRLYREKRETRREFDET IRYASREKAYAE I RPRI	EGRFAKRE	IETFAAEFIIF	---	STLSLSEITG	YGVPSF	347	Q96502	COL2_ARATH
LSREPQ	WCHPTAQDI	IASSHATT	WNAVNRVYREKREKAREFDERVRYVS	REERADVRRRVEGRFVVEAG	GEAYDYDPE	SPSPT	---	RSY	373	Q9LUA9	COL10_ARAI	
TVNVP	VITSTR	WHEIN	LSLE	WNSALSRYEPEKESRRYERH IRYESREVRAPESRIRI	EGRFAKRE	ADP	---	---	332	082256	COL13_ARAI	
IAITSWRATRLVAVINADLE	QWAGN	WNAVNRVYREKREKTRRYDET IRYETRKAETRL	RVEGRFVVEAG	TDF	---	---	402	022800	COL14_ARAI			
---	---	---	---	---	---	---	294	Q9SE53	COL5_ARATH			
T	---	---	---	---	---	---	355	Q9FHH8	COL5_ARATH			
PG	---	---	---	---	---	---	319	Q9M9B3	COL8_ARATH			
IGIS	P	WESN	---	VEVSNPEL	RDEAKERYEQEESERNFGEQ	IRYASREKARADITRERVEGRFVVEAG	GETIFEYDPSL	---	330	023379	COL11_ARAI	
IAGTSCETITRLVA	TKADLERLAGN	PCDA	WNAVNRVYREKREKTRRYDET IRYESREKARADITRL	RVEGRFVVEAG	SEAPYP	---	433	Q9C7E8	COL15_ARAI			
MGGESIHQEQYVGGCLPSSGFGDGG	REARVRS	RYREKRETRREFEET IRYASREKAYAEERPRI	EGRFAKRE	ASLAAAAS	PLGVWY	---	417	Q8RWD0	COL16_ARAI			
SGADP	G	---	---	---	---	---	362	Q940T9	COL4_ARATH			
SGAAHHHHHFRGLGHLGDAGDGG	REARVRS	RYREKRETRREFEET IRYASREKAYAEERPRI	EGRFAKRE	SSIGVAH	---	---	406	Q8LG76	COL6_ARATH			
YEKLSSSDGSVTRQGRDGGGSDGE	REARVRLYREKRETRREFEET IRYASREKAYAEERPRI	EGRFAKRE	ISLIT	---	---	---	392	Q9C9A9	COL7_ARATH			
LSGEP	P	WYPPTLQINWACSB	WNAVNRVYREKREKAREFDERVRYAS	REKARADVRRRVEGRFVVEAG	GEAYDYDPLIPT	---	---	RSY	372	Q9SSE5	COL9_ARATH	

## C 端



# 拟南芥CO家族成员进化树分析

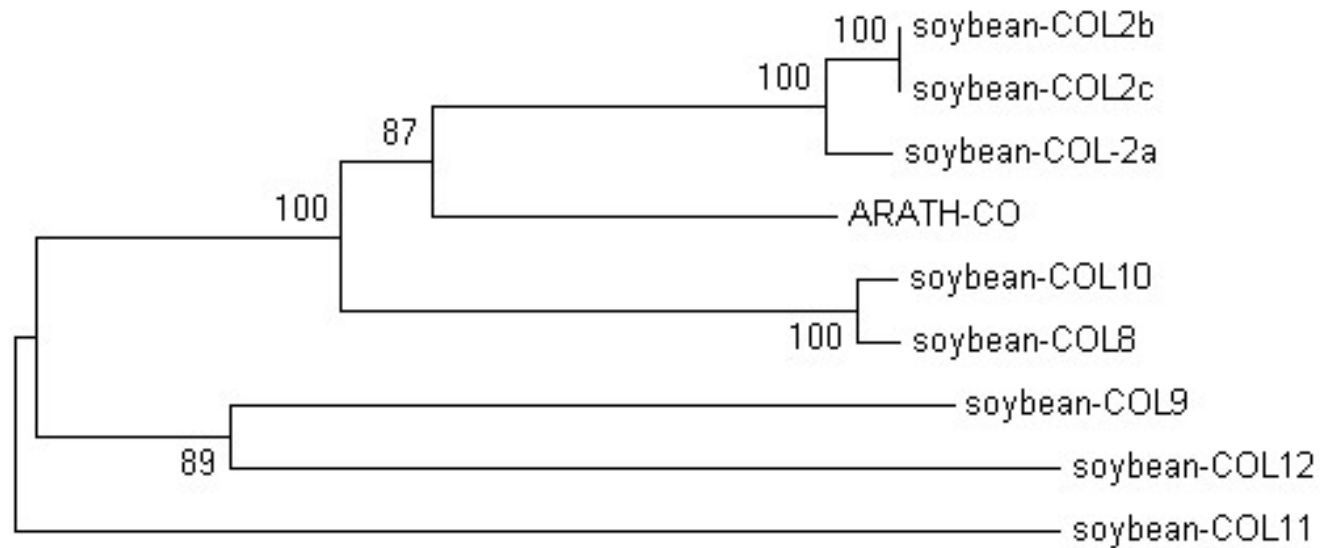


0.1

- 大豆是典型的短日作物，不同品种早晚有较大的差异。
- 影响大豆开花候选基因的克隆，表达分析，转基因研究，鉴定其在成花转变中的作用，对探究基因与开花整合因子FT的相互关系以及调节大豆开花的分子网络的描绘有重要意义。

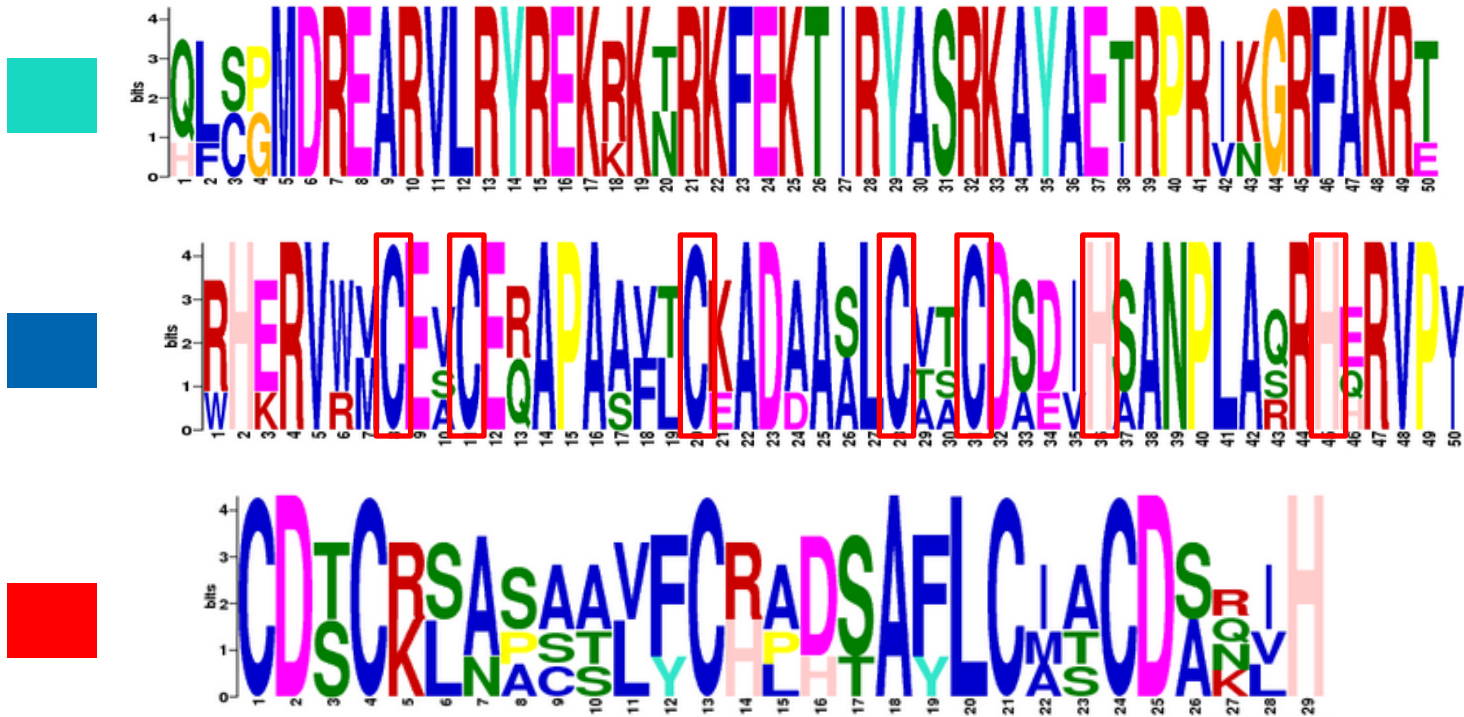
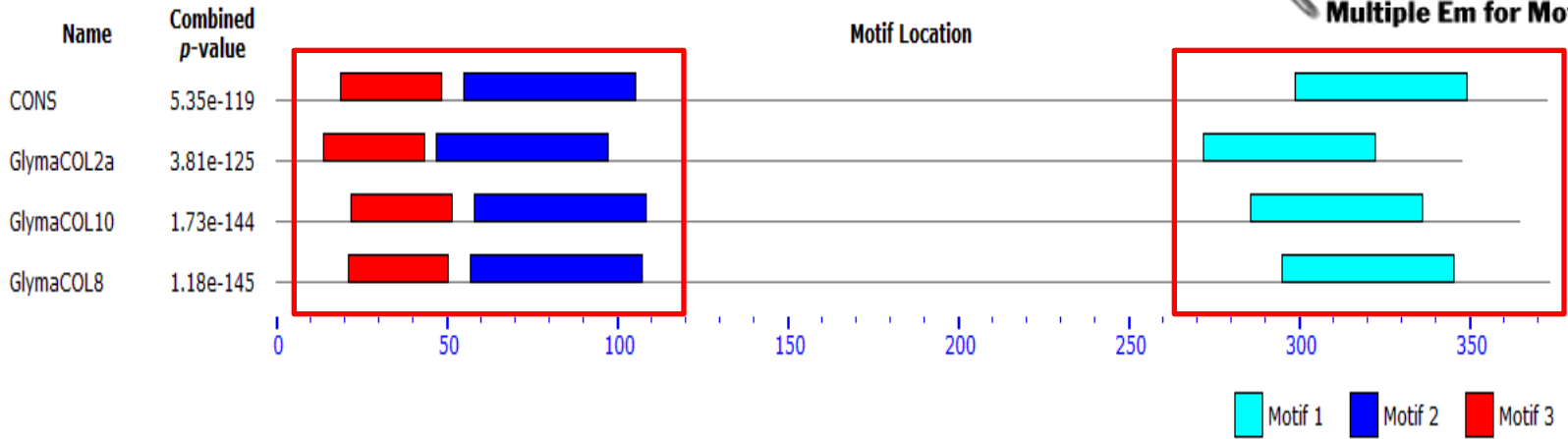
# 大豆CO-Like基因分析

Entry	Entry name	Status	Protein names	Gene names	Organism	Length
<input checked="" type="checkbox"/> B9MSS5	B9MSS5_SOYBN	★	CONSTANS-like 2a	COL2a	Glycine max (Soybean) (Glycine hispida)	348
<input checked="" type="checkbox"/> D0EP05	D0EP05_SOYBN	★	CONSTANS-like zinc finger protein	COL10	Glycine max (Soybean) (Glycine hispida)	365
<input checked="" type="checkbox"/> D0EP04	D0EP04_SOYBN	★	CONSTANS-like zinc finger protein	COL9	Glycine max (Soybean) (Glycine hispida)	351
<input checked="" type="checkbox"/> D0EP03	D0EP03_SOYBN	★	CONSTANS-like zinc finger protein	COL8	Glycine max (Soybean) (Glycine hispida)	374
<input checked="" type="checkbox"/> B9MSS6	B9MSS6_SOYBN	★	CONSTANS-like 2b	COL2b	Glycine max (Soybean) (Glycine hispida)	328
<input checked="" type="checkbox"/> D0EP07	D0EP07_SOYBN	★	CONSTANS-like zinc finger protein	COL12	Glycine max (Soybean) (Glycine hispida)	418
<input checked="" type="checkbox"/> B9MSS7	B9MSS7_SOYBN	★	Zinc finger protein CONSTANS-LIKE 2	COL2c	Glycine max (Soybean) (Glycine hispida)	352
<input checked="" type="checkbox"/> D0EP06	D0EP06_SOYBN	★	CONSTANS-like zinc finger protein	COL11	Glycine max (Soybean) (Glycine hispida)	366



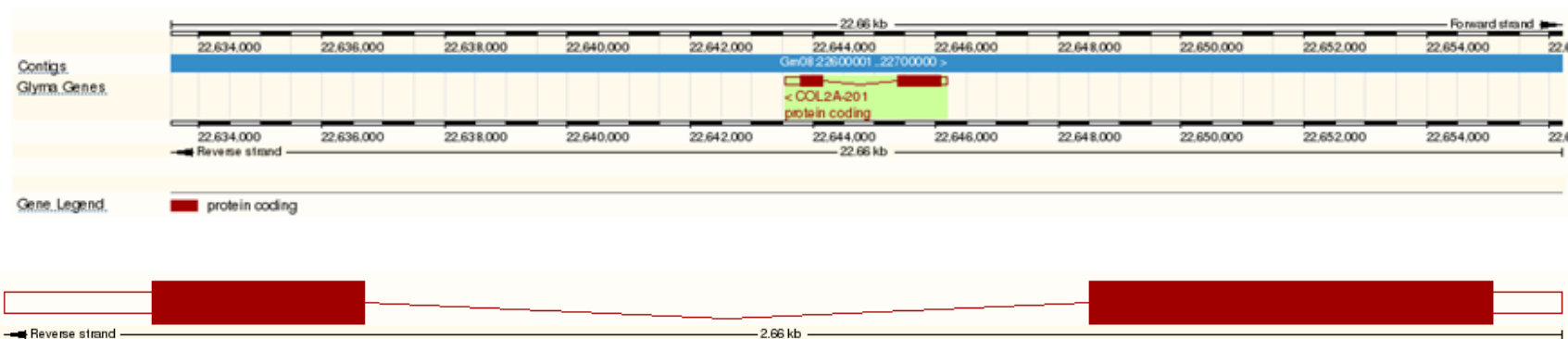
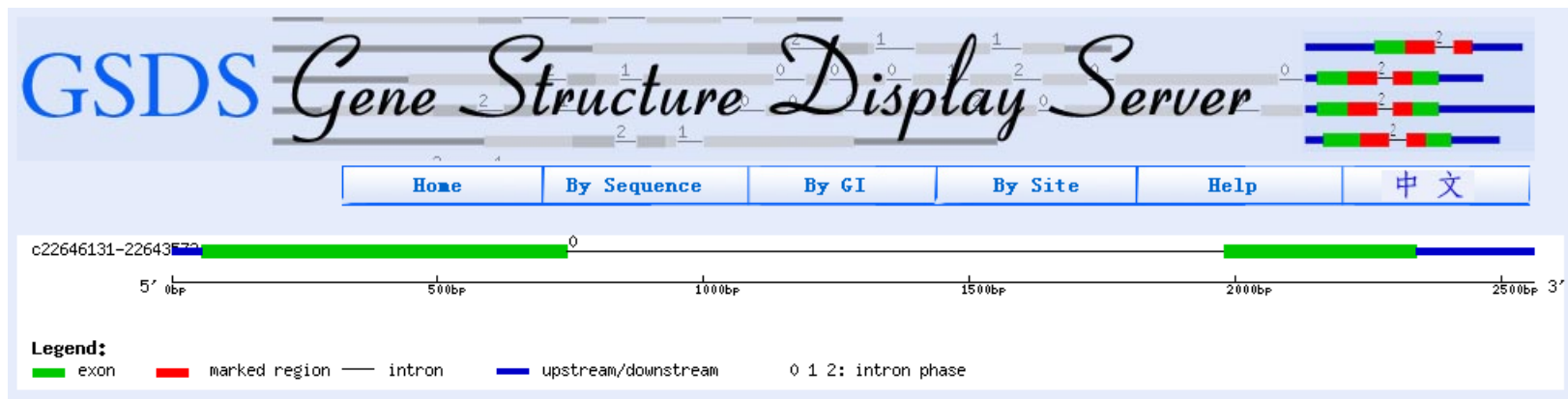
0.1





**B-box :  $CX_2CX_8CX_7CX_2CX_4HX_8H$**

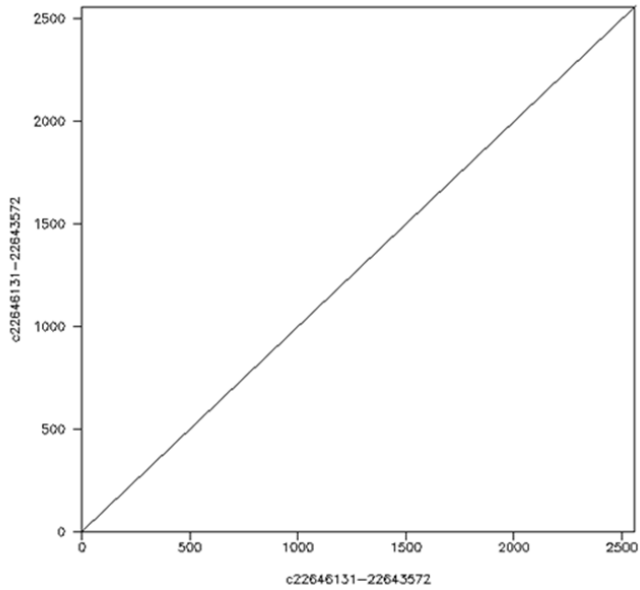
# GmCOL2a的基因结构



# GmCOL2a的重复序列分析

Dotpath: fasta::6039:c22646131-22643572 vs fasta::6040:c

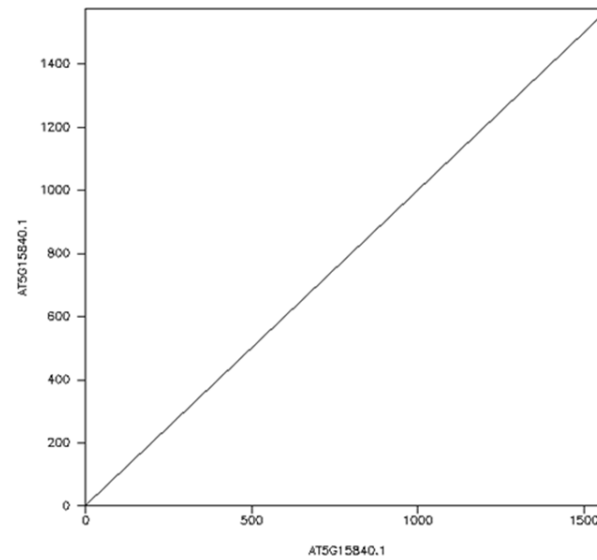
Sat 30 Nov 2013 13:51:05



大豆

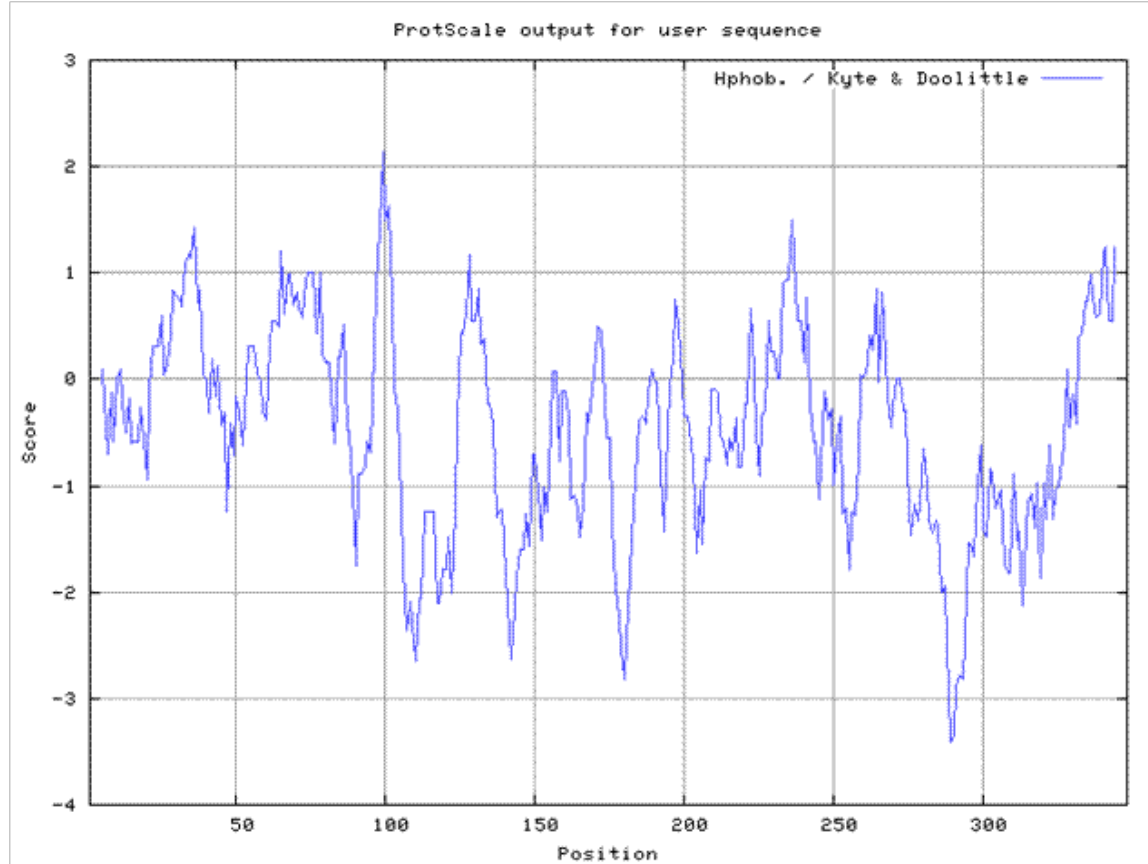
Dotpath: fasta::6019:AT5G15840.1 vs fasta::6020:AT5G1584.

Sat 30 Nov 2013 12:47:13

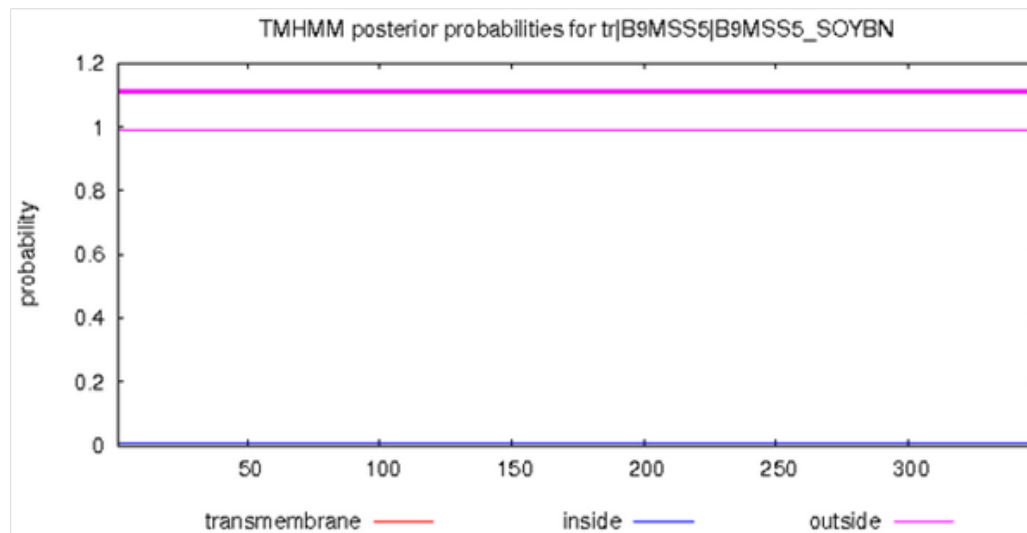


拟南芥

# ProScale



# TMHMM





# GmCOL2a的亚细胞定位预测



## TargetP 1.1 Server - prediction results

Technical University of Denmark

```
### targetp v1.1 prediction results #####  
Number of query sequences: 1  
Cleavage site predictions not included.  
Using PLANT networks.
```

Name	Len	cTP	mTP	SP	other	Loc	RC
Soybean_COL_2a	348	0.144	0.262	0.018	0.680	_	3
cutoff		0.000	0.000	0.000	0.000		

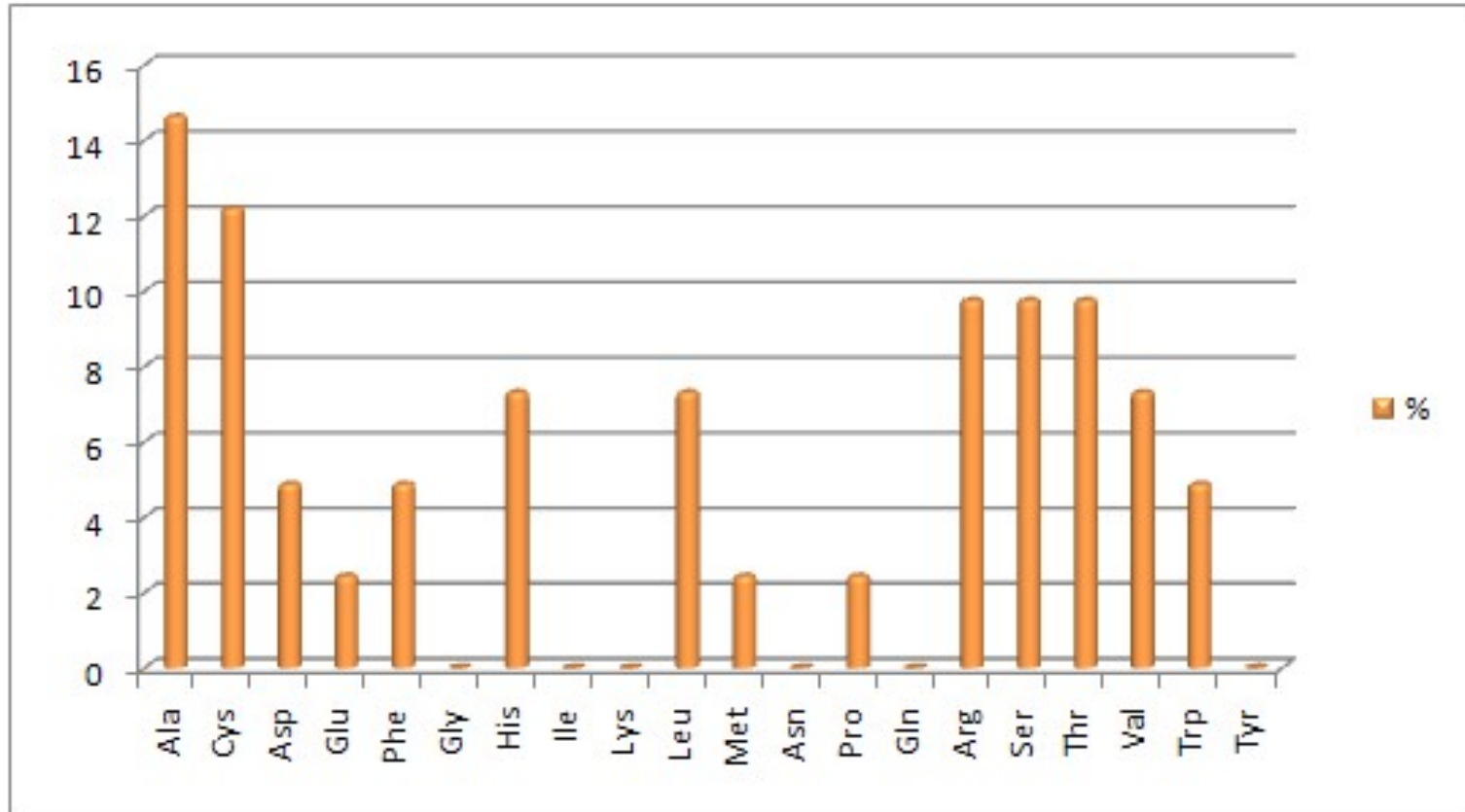
# GmCOL2a的氨基酸组分分析

Pepstats

AA	SwissProt*	Glymax-COL2a	
	%	number	%
Ala	8.25	38	10.92
Cys	1.37	11	3.161
Asp	5.45	29	8.333
Met	2.42	7	2.011
Phe	3.86	15	4.31
Gln	3.93	8	2.299
His	2.27	18	5.172
Tyr	2.92	9	2.586
Ile	5.95	11	3.161
Lys	5.84	13	3.736
Gly	7.07	15	4.31
Asn	4.06	17	4.885
Pro	4.7	18	5.172
Glu	6.75	16	4.598
Arg	5.53	21	6.034
Ser	6.57	31	8.908
Thr	5.34	21	6.034
Val	6.87	27	7.759
Trp	1.08	4	1.149
Leu	9.66	19	5.46

# Pepstats

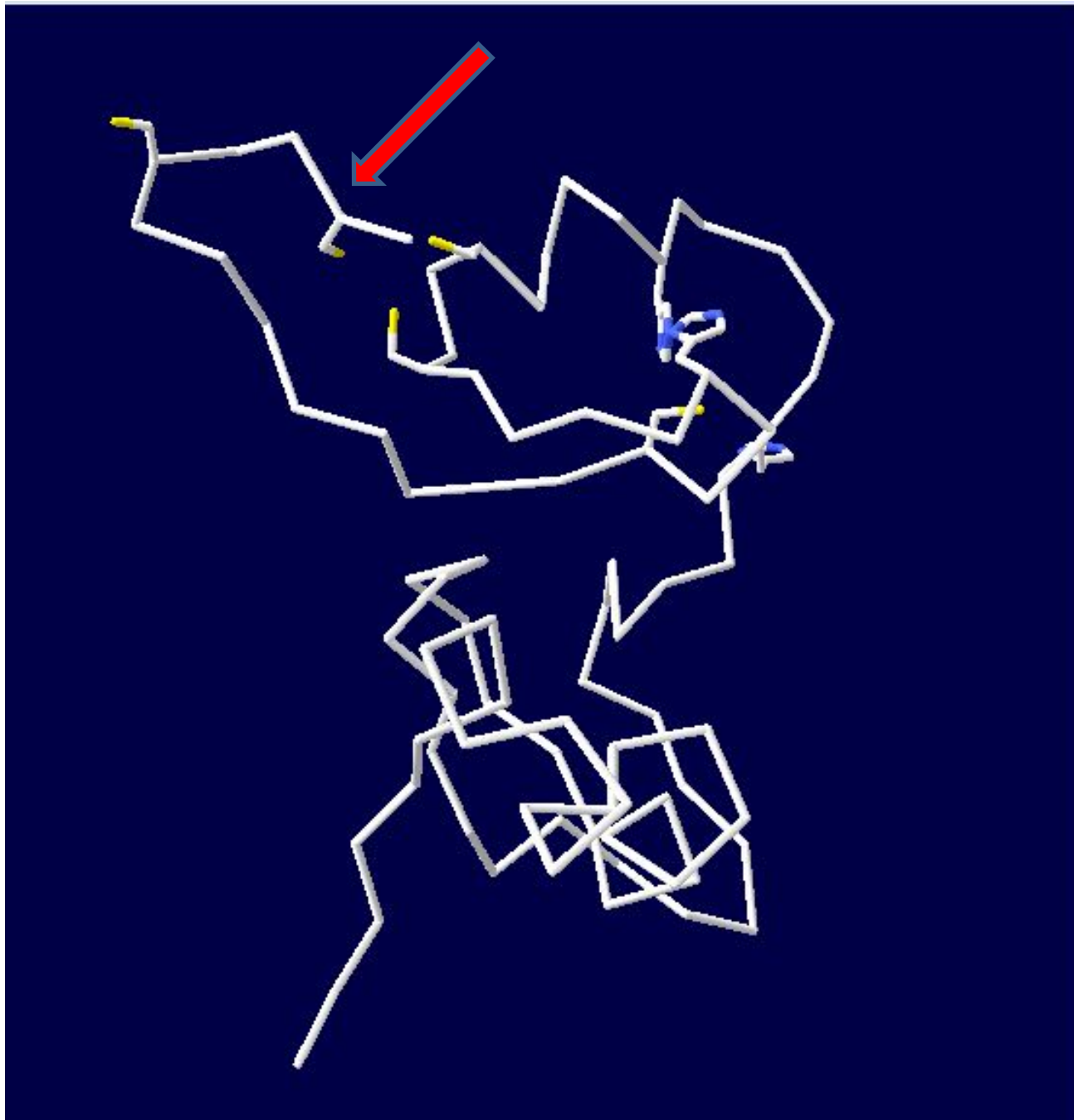
## GmCOL2a B-box区氨基酸组分分析







$CX_2CX_8CX_7CX_2CX_4HX_8H$

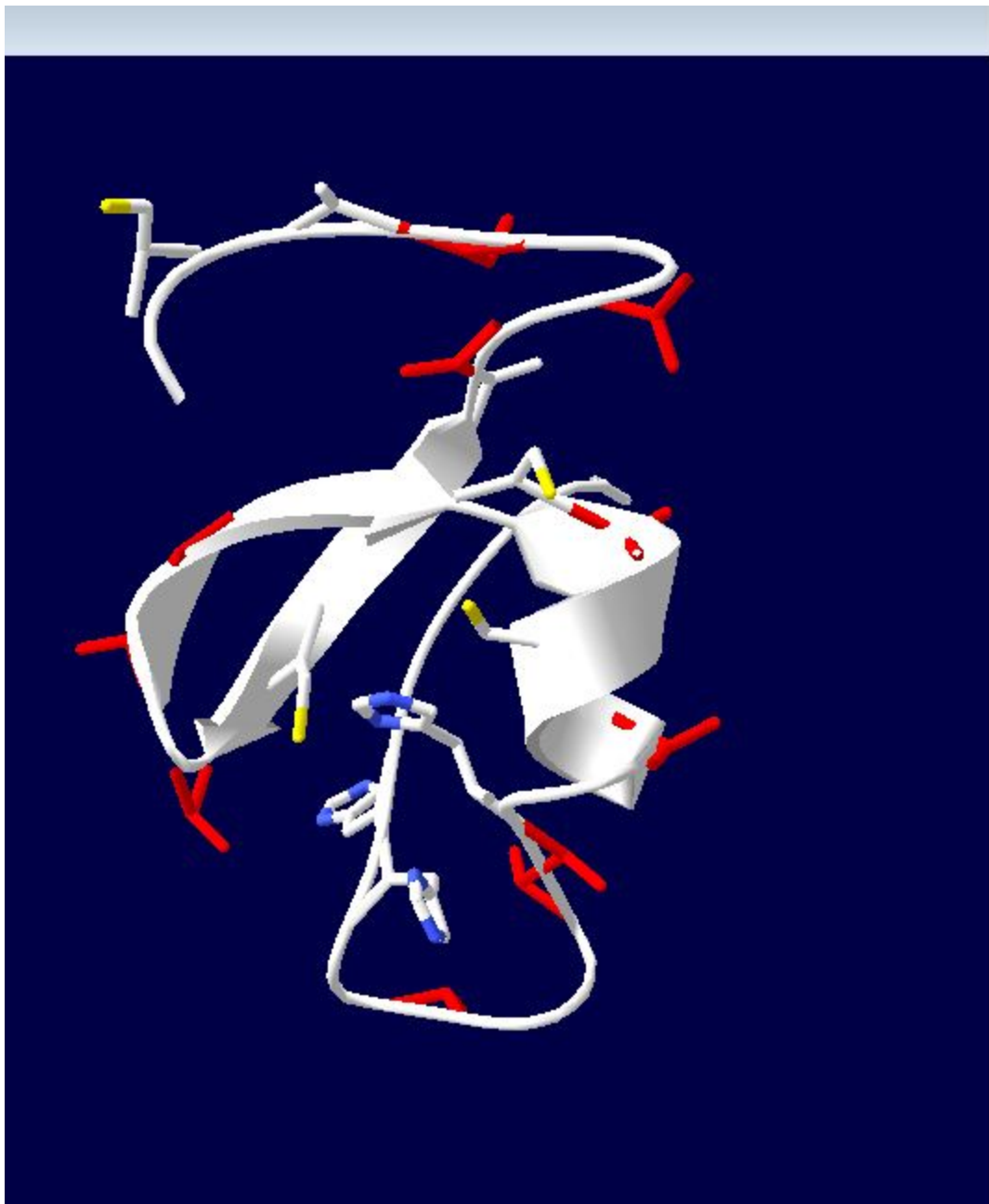


ool2		visible	?	can move	
group	show	side	labl	ribn	colgs
MET14	v				<input type="checkbox"/>
CYS15	v	v			<input type="checkbox"/>
ASP16	v				<input type="checkbox"/>
THR17	v				<input type="checkbox"/>
CYS18	v	v			<input type="checkbox"/>
ARG19	v				<input type="checkbox"/>
SER20	v				<input type="checkbox"/>
ALA21	v				<input type="checkbox"/>
PRO22	v				<input type="checkbox"/>
SER23	v				<input type="checkbox"/>
SER24	v				<input type="checkbox"/>
s VAL25	v				<input type="checkbox"/>
s PHE26	v				<input type="checkbox"/>
s CYS27	v	v			<input type="checkbox"/>
s ARG28	v				<input type="checkbox"/>
ALA29	v				<input type="checkbox"/>
HIS30	v	v			<input type="checkbox"/>
s THR31	v				<input type="checkbox"/>
s ALA32	v				<input type="checkbox"/>
s PHE33	v				<input type="checkbox"/>
s LEU34	v				<input type="checkbox"/>
CYS35	v	v			<input type="checkbox"/>
h ALA36	v				<input type="checkbox"/>
h THR37	v				<input type="checkbox"/>
h CYS38	v	v			<input type="checkbox"/>
h ASP39	v				<input type="checkbox"/>
h ALA40	v				<input type="checkbox"/>
h ARG41	v				<input type="checkbox"/>
h LEU42	v				<input type="checkbox"/>
HIS43	v	v			<input type="checkbox"/>
ALA44	v				<input type="checkbox"/>
SER45	v				<input type="checkbox"/>
LEU46	v				<input type="checkbox"/>
THR47	v				<input type="checkbox"/>
TRP48	v				<input type="checkbox"/>
HIS49	v	v			<input type="checkbox"/>
GLU50	v				<input type="checkbox"/>
ARG51	v				<input type="checkbox"/>

**CX<sub>2</sub>CX<sub>8</sub>CX<sub>7</sub>CX<sub>2</sub>CX<sub>4</sub>HX<sub>8</sub>H**



group	+	+	+	+	+
	show	side	labl	ry	ribn
					oolg
VAL52	v				<input type="checkbox"/>
TRP53	v				<input type="checkbox"/>
VAL54	v				<input type="checkbox"/>
CYS55	v	v			<input type="checkbox"/>
GLU56	v				<input type="checkbox"/>
ALA57	v				<input type="checkbox"/>
CYS58	v	v			<input type="checkbox"/>
GLU59	v				<input type="checkbox"/>
ARG60	v				<input type="checkbox"/>
ALA61	v				<input type="checkbox"/>
PRO62	v				<input type="checkbox"/>
ALA63	v				<input type="checkbox"/>
ALA64	v				<input type="checkbox"/>
s PHE65	v				<input type="checkbox"/>
s LEU66	v				<input type="checkbox"/>
s CYS67	v	v			<input type="checkbox"/>
s LYS68	v				<input type="checkbox"/>
ALA69	v				<input type="checkbox"/>
ASP70	v				<input type="checkbox"/>
s ALA71	v				<input type="checkbox"/>
s ALA72	v				<input type="checkbox"/>
s SER73	v				<input type="checkbox"/>
s LEU74	v				<input type="checkbox"/>
CYS75	v	v			<input type="checkbox"/>
h ALA76	v				<input type="checkbox"/>
h SER77	v				<input type="checkbox"/>
h CYS78	v	v			<input type="checkbox"/>
h ASP79	v				<input type="checkbox"/>
h ALA80	v				<input type="checkbox"/>
ASP81	v				<input type="checkbox"/>
ILE82	v				<input type="checkbox"/>
HIS83	v	v			<input type="checkbox"/>
ALA84	v				<input type="checkbox"/>
ALA85	v				<input type="checkbox"/>
ASN86	v				<input type="checkbox"/>
PRO87	v				<input type="checkbox"/>
LEU88	v				<input type="checkbox"/>
ALA89	v				<input type="checkbox"/>
SER90	v				<input type="checkbox"/>
ARG91	v				<input type="checkbox"/>
HIS92	v	v			<input type="checkbox"/>
HIS93	v	v			<input type="checkbox"/>
ARG94	v				<input type="checkbox"/>



visible	?		can move	
group	+	+	+	+
	show	side	labl	ribn
			oolg	
CYS55	v	v		v
GLU56				v
ALA57	v	v		v
CYS58	v	v		v
GLU59				v
ARG60				v
ALA61	v	v		v
PRO62				v
ALA63	v	v		v
ALA64	v	v		v
s PHE65				v
s LEU66				v
s CYS67	v	v		v
s LYS68				v
ALA69	v	v		v
ASP70				v
s ALA71	v	v		v
s ALA72	v	v		v
s SER73				v
s LEU74				v
CYS75	v	v		v
h ALA76	v	v		v
h SER77				v
h CYS78	v	v		v
h ASP79				v
h ALA80	v	v		v
ASP81				v
ILE82				v
HIS83	v	v		v
ALA84	v	v		v
ALA85	v	v		v
ASN86				v
PRO87				v
LEU88				v
ALA89	v	v		v
SER90				v
ARG91				v
HIS92	v	v		v
HIS93	v	v		v
ARG94				v
VAL95				v
PRO96				v
ILE97				v



**Thank you !**