у									
y Mtype	PRT						Species	Rattus norvegicus (Pat)
		IDAD	D						rdi)
Class	STAN						Org	Entry	
Id	100K	_RAI						Eukaryota	
Seglen	889							Metazoa	
Ac	Entry	_						Chordata	
					•			Craniata	
Mod	Date				el Type			Vertebrata	
	01-N	OV-1	997	3	5 Created			Euteleostomi	
	01-N			3!			1	Mammalia	
	15-Jl			3			1	Eutheria	
Descr				(EC 6.3.2.				Rodentia	
Desci	100 1	(D) (I	ROTEIN	(LO 0.5.2.	,			Sciurognathi	
								Muridae	
								Murinae	
								Rattus	
Ref	Num	Pos			Comment	Db	Medlineid	Author	Cite
					Entry			Entry	
	1	SFO	UFNCF F	ROM N.A	STRAIN=WISTAR	MEDLINE	92253337	Mueller D	Nucleic Acids Res. 20:1471-1475(1992
					TISSUE=TESTIS			Rehbein M	
					11330E=1E3113			Baumeister H	
	_							Richter D	
	2	ERR	MUTA					Mueller D	Nucleic Acids Res. 20:2624-2624(1992
								Rehbein M	
								Baumeister H	
								Richter D	
Embl	Sec I	d	Prim Id	Status	-		Pir	Sec Id Prim Id	
			X64411						
Interpro	Sec						Mendel		
interpro	Id	-11111	iu				Hssp		
		IDD04	205/0					0	I-
			00569				Prosite	Status Sec Id	F
			02004						
Pfam	Statu					Prim Id	Gene		
	1	HE	CT			PF00632	Prints		
	1	PA	BP			PF00658	Flybase		
Keyword	Entry	,					Maizedb		
.,			conjugati	on			Wormpep		
	Ligas		oonjugan	011					
Features	Ligas	C							
	Ir		T- D						
Doma	in F	rom							
	_			ntry					
	7	7	88						
	1	27	150						
	4	20	439						
		48	457						
		85	514						
		79	590						
		86	889						
		27	847						
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F	rom 8	358					-		
Т		358							
			IIITIN (R	Y SIMILAR	ITY)		•		
Signa		שוע	51111V (D	. JIMILAN	,				
			T- ID						
Chain	L L	rom	To Des	CI					
Disulf	id F	rom	To Des	cr					
Non 7	Ter								
Mod F									
Confli		rom	To Des	cr					
301111		. 5111	. 0 003						
Carbo	hyd F	rom	To De	ecr					
Carbo	niyu F	TOTT	TO DE	361					

Np Bind From To Descr

Similar

Prim Id

Mtype	PRT		Species	Theileria parva	
Class	STANDARD			Entry	
Id	104K THEPA			Eukaryota	
Seglen	924			Alveolata	
Ac	Entry			Apicomplexa	
				Piroplasmida	
Mod	Date Rel Type			Theileriidae	
	01-APR-1990 14 Created			Theileria	
	01-APR-1990 14 Last sequence update				
	01-AUG-1992 23 Last annotation update				
Descr	104 KDA MICRONEME-RHOPTRY ANTIGEN				
Ref	Num Pos Comment	Db	Medlineid	Author	Cite
	Entry			Entry	
	1 SEQUENCE FROM N.A	MEDLINE	90158697		Mol. Biochem. Parasitol. 39:47-60(1990)
				Young J.R	
				Nene V	
				Desai J	
				Webster P	
				Ole-Moiyoi	
				O.K	
				Musoke A.J	
Embl	Sec Id Prim Id Status	_	Pir	Sec Id Prim Id	•
	AAA18217 M29954			A44945 A44945	
Interpro	Sec Prim Id		Mendel		
	Id		Hssp		
				Status Sec Id	Prim Id
Pfam	Status Sec Id	Prim Id			
			Gene		
Keyword	Entry		Prints		
	Antigen		Flybase		
	Sporozoite		Maizedb		
	Repeat		Wormpep		
Features					
Doma	in From To Descr				
	Entry				
	1 19				
	905 924				
Bindir	ng				
Signa					
Chain					
Disulf	id From To Descr				

Np Bind

Similar

Carbohyd From To Descr

From To Descr

Mtype	PRT	DAR		_						Lycopersicon escule	intum (Tomato)
Class	STANI								Org	Entry	
d	108_L	YCE:	Ś							Eukaryota	
Seglen	102									Viridiplantae	
Ac .	Entry									Embryophyta	
										Tracheophyta	
Mod	Date						Туре			Spermatophyta	
		15-JUL-1999 38 Created 15-JUL-1999 38 Last sequence update								Magnoliophyta	
										eudicotyledons	
		15-JUL-1999 38 Last annotation update								Asteridae	
Descr	PROT	PROTEIN 108 PRECURSOR								euasterids I	
										Solanales	
										Solanaceae	
										Solanum	
Ref	Num	Pos				С	Comment	Db	Medlineid	Author	Cite
							Entry			Entry	
	1	SEQI	JENO	E FRO	M N.A		STRAIN=CV. VF36	MEDLINE	94143497		Plant Physiol. 101:1413-1413(1993)
							TISSUE=ANTHER			Smith A.G	
Embl	Sec Io	i	Prin	n Id	Status				Pir	Sec Id Prim Id	
	CAA78			088							
nterpro	Sec P	rim I	d						Mendel		
	Id								Status	1133	
									Sec Id	LYCes	
Pfam	Status	Sec	: Id					Prim Id	Prim I	d 8853	
									Hssp		
Keyword	Entry								Prosite	Status Sec Id	Prim Io
eatures									Gene		
Doma	ain Fr	om	To	Descr					Prints		
				Ent	ry				Flybase		
									Maizedb		
Bindir									Wormpep		
Signa	al										
F	rom 1										
T	To 3	0									
D	Descr P	OTE	AITI	L							
Chain	n Fr	om	To	Descr							
	31	1	102	PROTE	EIN 108						
Disulf	fid Fr	om		Descr							
	4				ЛІLARIT	Υ					
	51	1			ЛІLARIT				•		
	_				ИLARIT						
	67	/									

Np Bind Similar

 Conflict
 From To Descr

 Carbohyd
 From To Descr

 Np Bind
 From To Descr

Mtype	PRT	_	_					Vigna unguiculata (0	Cowpea)	
Class	STANDAR							Entry		
ld	10KD_VIC	SUN						Eukaryota		
Seqlen	75							Viridiplantae		
Ac	Entry							Embryophyta		
								Tracheophyta		
Mod	Date				Туре			Spermatophyta		
	01-NOV-1				Created			Magnoliophyta		
	01-NOV-1				Last sequence update			eudicotyledons		
	01-FEB-19				Last annotation update			Rosidae		
Descr	10 KDA P	ROTE	IN PRECU	URSOR	(CLONE PSAS10)			eurosids I		
								Fabales		
								Fabaceae		
								Papilionoideae		
								Vigna		
Ref	Num Pos			(Comment	Db	Medlineid		Cite	
					Entry			Entry		
	1 SEC	UENC	E FROM	N.A		MEDLINE	91355865	Ishibashi N	Plant Mol. Biol. 15:59-64(1990)	
								Yamauchi D		
								Miniamikawa		
								T		
Embl	Sec Id	Prim		tatus				Sec Id Prim Id		
	CAA34760		877					S11156 S11156		
Interpro	Sec Prim	ld					Mendel			
	Id						Hssp			
		02118						1CHL		
Pfam	Status Se					Prim Id		d P45639		
		ımma-	-thionin			PF00304	Prosite	Status Sec Id		Prim Id
Keyword								1 GAMMA_THI	IONIN	PS00940
	Germinati	on					Gene			
	Signal						Prints			
							Flybase			
Features			Descr				Maizedb			
Features Doma	ain From	10	Desci							
	ain From	10	Entry				Wormpep			
Doma		10								
Doma	ng	10								
Doma Bindir Signa	ng al	10								
Doma Bindir Signa F	ng al	10								
Bindir Signa F	ng al From 1 Fo ?		Entry							
Bindir Signa F T	ng al From 1 Fo ? Descr POTE	NTIAL	Entry							
Bindir Signa F	ng al From 1 Fo ? Descr POTE	NTIAL To	Entry							
Bindir Signa F T Chain	ng al From 1 Fo ? Descr POTE From ?	NTIAL To 1	Entry Descr	PROTEI	N					
Bindir Signa F T	ng al From 1 Fo ? Descr POTE n From ?	NTIAL To 75 To	Entry Descr 10 kDa P Descr		V					
Bindir Signa F T Chain	ng al From 1 Fo ? Descr POTE From ?	NTIAL To 75 To 75	Descr 10 kDa P Descr BY SIMIL	LARITY	N					
Bindir Signa F T Chain	ng From 1 Fo ? Descr POTE From ? fid From	TO 175 163 163	Descr 10 kDa P Descr 8Y SIMIL BY SIMIL	LARITY LARITY	N					
Bindir Signa F T Chain	ng al From 1 Fo ? Descr POTE From ? fid From 31	TO 175 163 163	Descr 10 kDa P Descr BY SIMIL	LARITY LARITY	N					

Similar

 Conflict
 From To Descr

 Carbohyd
 From To Descr

 Np Bind
 From To Descr

Mtype Class Id Seglen	PRT						Species	Plasmodium	knowle	si
Id Seqlen	STAND	ARD					Org	Entry		
	110K_F						. 3	Eukaryota		
	296							Alveolata		
Ac	Entry	Ī						Apicomplexa	a	
								Haemospori		
Mod	Date			Rel	Туре			Plasmodium		
	01-JAN	-1990		13				. idomodiam		
	01-JAN			13						
	01-FEB				Last annotation update					
Descr			GEN (PK110)							
Ref	Num P		OLIV (FRTTO)		Comment	Db	Medlineid	Author		Cite
Kei	INUITIF	US		Ľ	Entry	100	Medilileid	Entry		Cite
	1 5	FOLIENI	CE FROM N.A	-	Entry	MEDITAL	88039002		<u> </u>	Mol. Biochem. Parasitol. 25:185-193(1987)
	1 2	EQUEN	JE FROM N.A			MEDLINE	88039002			Moi. Biochem. Parasitoi. 25:185-193(1987)
								Moon A.		
								Qiang B		
								Meda M		
								Dalton N	Λ	
								Card C		
								Schmidt		
								Ullrich R		
								Wallach		
								Lynch J		
								Donelso		
Embl	Sec Id		n Id Statu	s					im Id	
	AAA294		9152					A54527 A5	54527	
Interpro	Sec Pri	m Id					Mendel			
	Id						Hssp			
							Prosite	Status Sec	Id	Prim Id
Pfam	Status	Sec Id				Prim Id				
							Gene			
Keyword							Prints			
	Malaria						Flybase			
	Antiger	1					Maizedb			
	Repeat						Wormpep			
Features										
Doma	ain Fro	m To	Descr							
			Entry							
	131	296	13.5 X 12	AA	TANDEM REPEATS OF E-E	-T-Q-				
			K-							
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Mod Res Conflict

Np Bind

Similar

From To Descr Carbohyd From To Descr

From To Descr

Mtype	PRT							Species	Helianthus annuus (Common sunflower)	
Class	STAN	IDARD						Org	Entry	·	
ld	11S3	HELAN						3	Eukaryota		
Seglen	493								Viridiplantae		
Ac	Entry	,							Embryophyta		
									Tracheophyta		
Mod	Date			Re	I Type				Spermatophyta		
		OV-1990		16	Created				Magnoliophyta		
		OV-1990		16		ate			eudicotyledons		
		AY-2000			Last annotation up				Asteridae		
Descr			I SEED ST		E PROTEIN G3 PREC				euasterids II		
		IANTHIN							Asterales		
	`		,						Asteraceae		
									Asteroideae		
									Heliantheae		
									Helianthus		
Ref	Num	Doc			Comment		Db	Medlineid		Cite	
101	Nulli	1 03			Entry		00	weumelu	Entry	- Citic	
	1	SECHEN	CE FROM	N A	Liiu y		MEDLINE	89232734		Gene 74:433-443(1988)	
	1	SEQUEN	CE FROIVI	IV.A			IVIEDLINE	09232734	R.A	Gene 74.455-445(1966)	
									Allen R.D		
									Cohen E.A		
									Nessler C.L		
									Thomas T.L		
Embl	Sec I	al Dei	m Id St	tatus				Pir	Sec Id Prim Id		
LIIIDI		3374 M2		latus				rii	JA0089 JA0089		
Interpro		Prim Id	.0032					Mendel	JAUU09 JAUU09		
interpro	Id	riiii iu									
		IPR00045	0					Hssp Prosite	Status Sec Id		Prim I
Pfam		s Sec Id	7				Prim Id	riusite	1 11S_SEED_S	TORACE	PS003
laili	1		ore_11s					Gene	HAG3	TORAGE	P3003
Keyword	Entry		010_113				1100170	Prints	TIAGS		
toy word		storage	orotein						11SGLOBULIN		
		gene fam							d PR00439		
	Signa		y					Flybase	u F K00437		
eatures	Jigino							Maizedb			
Doma	in F	rom To	Descr					Wormpep			
Doma	I.	10111 10	Entry					wormpep			
	2	3 35	Linay								
		11 127									
		91 297									
Bindir		71 277									
Signa											
	rom										
T		20									
	escr	20									
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Chain	_	rom To	Descr								
		1 305									
- ·		06 493		HAIN							
Disulf	_	rom To	Descr								
		03 312	INTERCH	AIN (A	ACIDIC-BASIC) (POT	ENTIAL)				
Non T											
Mod F			-								
Confli	ct F	rom To	Descr								
Carbo	hyd F	rom To	Descr								
NI- DI-	nd F	rom To	Descr								
Np Bii											

	PRT						Cucurbita maxima (F	rumpkin) (winter squasn)	
Class		NDARD				Org	Entry		
ld		B_CUCMA					Eukaryota		
Seqlen	480						Viridiplantae		
Ac	Entry	У					Embryophyta		
							Tracheophyta		
Mod	Date			el Type			Spermatophyta		
	01-J	AN-1990	13	3 Created			Magnoliophyta		
	01-J	AN-1990	13	Last sequence upda	ate		eudicotyledons		
	01-N	NOV-1990	16	Last annotation up	date		Rosidae		
Descr			I BETA SUBUNI				eurosids I		
							Cucurbitales		
							Cucurbitaceae		
							Cucurbita		
Ref	Num	Pos		Comment	Db	Medlineid		Cite	
Kei	Nulli	11105		Entry		weumeiu	Entry	cite	
	1	OFOLIENI	OF FROM N. A	Entry	MEDIANE	004//744		E	
	1	SEQUEN	CE FROM N.A		MEDLINE	88166744		Eur. J. Biochem. 172:627-632(1988)	
							Mori H		
							Nishimura M		
							Akazawa T		
							Hara-		
							Nishimura I		
	2	SEOUEN	CE OF 22-30				Ohmiya M	Plant Cell Physiol. 21:157-167(1980))
	_	AND 297					Hara I	Tank ocii 111y3ioi. 21:107 107(1700)	
		AND 277	-302						
							Mastubara H		
Embl	Sec		n Id Status			Pir	Sec Id Prim Id		
		33110 M3	6407				FWPU1B S00366		
Interpro		Prim Id				Mendel			
	Id					Hssp			
	-	IPR00045	9			Prosite	Status Sec Id		Prim Id
Pfam	Statu	us Sec Id			Prim Id		1 11S_SEED_S	TORAGE	PS0030
	1	Seedsto	ore_11s		PF00190	Gene	1		1
Keyword	Entry				1	Prints			
icey word		d storage p	rotoin				11SGLOBULIN		
			n otcii i						
	Sign						d PR00439		
Features		al	D			Flybase	d PR00439		
Features Doma		al	Descr			Flybase Maizedb	a PR00439		
		al	Descr Entry			Flybase	a <u> PR00439</u>		
Doma	ain [al				Flybase Maizedb	a PR00439		
Doma	ain I	al				Flybase Maizedb	a <u>PR00439</u>		
Doma	ain I	al				Flybase Maizedb	a PR00439		
Doma Bindir Signa	ain I	From To				Flybase Maizedb	a PR00439		
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Bindir Signa F	ain Fing From Fo	From To				Flybase Maizedb	a PR00439		
Bindir Signa F T	ain F ng al From F Oescr	From To 1 21	Entry			Flybase Maizedb	a PR00439		
Bindir Signa F	ng all From Fo Descr	From To 1 21 From To	Entry	I DETA CUDUNIT		Flybase Maizedb	a PR00439		
Bindir Signa F T	ng all From Fo Descr	From To 1 21 From To 22 480	Entry Descr 11S GLOBULIN	J BETA SUBUNIT		Flybase Maizedb	a PR00439		
Bindir Signa F T	ng all From Fo Descr	From To 1 21 From To 22 480 22 296	Entry Descr 11S GLOBULIN GAMMA CHAIN	I (ACIDIC)		Flybase Maizedb	a PR00439		
Bindir Signa F T Chain	ng all From Fo Descr	From To 1 21 From To 22 480 22 296 297 480	Descr 11S GLOBULIN GAMMA CHAIN DELTA CHAIN	I (ACIDIC)		Flybase Maizedb	a PR00439		
Bindir Signa F T	ng all From Fo Descr	From To 122 480 22 296 297 480 From To To 150 150 150 150 150 150 150 150 150 150	Descr 11S GLOBULIN DELTA CHAIN DESCR	(ACIDIC) (BASIC)		Flybase Maizedb	d PR00439		
Bindir Signa F T Chain	ain fid fid fid	From To 122 480 22 296 297 480 From To To 150 150 150 150 150 150 150 150 150 150	Descr 11S GLOBULIN DELTA CHAIN DESCR	I (ACIDIC)	TENTIAL)	Flybase Maizedb	a PR00439		
Bindir Signa F T Chain	ain F ng al From F O Descr	From To 122 480 22 296 297 480 From To To 150 150 150 150 150 150 150 150 150 150	Descr 11S GLOBULIN DELTA CHAIN DESCR	(ACIDIC) (BASIC)	TENTIAL)	Flybase Maizedb	a PR00439		
Bindin Signa F T C Chain	ain Find Find Find Find Find Find Find Fi	From To 122 480 22 296 297 480 From To To 150 150 150 150 150 150 150 150 150 150	Descr 11S GLOBULIN DELTA CHAIN DESCR	(ACIDIC) (BASIC)	TENTIAL)	Flybase Maizedb	a PR00439		
Bindin Signa F T C Chain Disulf	ng all From Fo Descr n fid Frer Res	From To 1 21 From To 22 480 22 296 297 480 From To 124 303	Descr 11S GLOBULIN DELTA CHAIN DESCR	(ACIDIC) (BASIC)	TENTIAL)	Flybase Maizedb	a PR00439		
Bindin Signa F T C Chain Disulf	ng all From Fo Descr fid Frer Res From	From To 1 21 From To 22 480 297 480 From To 124 303	Descr 11S GLOBULIN DELTA CHAIN DESCR	(ACIDIC) (BASIC)	TENTIAL)	Flybase Maizedb	a PR00439		
Bindin Signa F T C Chain Disulf	ng all From Fo Descr Ifid Frem Res From Fo	From To 1 21 From To 22 480 22 296 297 480 From To 124 303	Descr 11S GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PR00439		
Bindir Signa F T Chain Disulf Non 1 Mod I F T T T	ng ala ala ala ala ala ala ala ala ala al	1 21 From To 22 480 22 296 297 480 2124 303 22 22 PYRROLID	Descr 115 GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PR00439		
Bindin Signa F T C Chain Disulf	ng ng lal r-rom ro Obescr lal reservation lal	From To 1 21 From To 22 480 22 296 297 480 From To 124 303 22 PYRROLID From To	Descr 115 GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PR00439		
Bindir Signa F T Chain Disulf Non 1 Mod I F T T T	ng Ing Ing Ing Ing Ing Ing Ing Ing Ing I	From To 1	Descr 11S GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PROU439		
Bindii Signa F T C Chain Disulf Non 7 Mod I F T C Confil	ng lal lal lar lar lar lar lar lar lar lar	From To 1 21 From To 22 480 22 296 297 480 To 124 303 22 22 PYRROLID From To 22 27 27 30 30	Descr 115 GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PR00439		
Bindir Signa F T C Chain Disulf Non 1 Mod I F T C Confil	ng lal lal lar lar lar lar lar lar lar lar	From To 1 21 From To 22 480 22 296 297 480 To 124 303 22 22 PYRROLID From To 22 27 27 30 30	Descr 11S GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PR00439		
Bindir Signa F T C Chain Disulf Non 1 Mod I F T C Confil	ng lal lal lar lar lar lar lar lar lar lar	From To 1	Descr 11S GLOBULIN GAMMA CHAIN DELTA CHAIN DESCR INTERCHAIN (ONE CARBOXY Descr S -> E (IN REF E -> S (IN REF	I (ACIDIC) (BASIC) GAMMA-DELTA) (POT	TENTIAL)	Flybase Maizedb	a PR00439		

Similar

• Mtype PRT Species Rickettsia rickettsii

Class	STAN	IDARD				Org	Entry	
Id	120K	_RICRI				-	Bacteria	
Seglen	1300						Proteobacteria	
Ac	Entry						alpha subdivision	
							Rickettsiales	
Mod	Date		Rel	Туре			Rickettsiaceae	
	01-AF	PR-1990	14	Created			Rickettsieae	
	01-FE	B-1996	33	Last sequence update			Rickettsia	
	01-FE	B-1996	33	Last annotation update				
Descr	120 k	CDA SURFACE-EXPOSE	DΡ	ROTEIN				
Ref	Num	Pos	(Comment	Db	Medlineid	Author	Cite
				Entry			Entry	
	1	SEQUENCE FROM N.A	١		MEDLINE	90136087	Gilmore R.D.	Mol. Microbiol. 3:1579-1586(1989)
							Jr	
							Joste N	
							McDonald G.	
							A	

Embl	Sec		Prim		Status	S					Pir
		134402		353							
Interpro		Prim	ld								Mendel
	Id										Hssp
									1		Prosite
Pfam	Sta	tus Se	c Id						Prim	ld	
	_										Gene
Keyword	Ent	_									Prints
		igen									Flybase
		coprot	ein								Maizedb
	S-la	iyer									Wormpep
Features		_	т.								
Doma	iin	From	10	Descr							
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Chain		From	lo l	Descr							
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Disulf	ıa	From	To I	Jescr							
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Carbo	hyd	From		Descr							
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		66	66		NTIAL						
		86	86		NTIAL						
		103	103		NTIAL						
		147	147		NTIAL						
		268	268		NTIAL						
		330	330		NTIAL						
		375	375		NTIAL						
		415	415		NTIAL						
		424	424		NTIAL						
		430	430		NTIAL						
		436	436		NTIAL						
		444	444		NTIAL						
		515	515		NTIAL						
		547	547		NTIAL						
		593	593		NTIAL						
		655	655		NTIAL						
		698	698		NTIAL						
		710	710		NTIAL						
		799	799		NTIAL						
		800	800		NTIAL						
		826	826		NTIAL						
		844	844		NTIAL						
		861	861	POTE	NTIAL	_					
		879	879		NTIAL						
		920	920	POTE	NTIAL	_					
		926	926		NTIAL						
		1116	1116	POTE	NTIAL	_					
		1128	1128	POTE	NTIAL						
				POTE	NTIAL						
		1146	1146	POTE	NTIAL						
		1211		POTE	NTIAL						
Np Bir	nd	From	To I	Descr							
Simila	ır										

 Sec Id
 Prim Id

 S07575
 S07575

Status Sec Id

P120

Prim Id

Mtype	PR	г							Species	Drosonh	nila malanon	aster (Fruit fly)	
Class		ANDAR	n						Org	Entry	ilia melanog	dater (Fruit hy)	
Id		BU DR							org	Eukaryo	ıta		
Seglen	368	_	OIVIL							Metazoa			
Ac	Ent									Arthropo			
AC	LIII	ı y								Trachea			
Mod	Dat	.0			Do	I Type				Нехаро			
IVIOU		OCT-1	വാ			Created				Insecta	ua		
		OCT-1					uence update			Pterygo	to		
		FEB-19					otation update			Neopter			
Descr				PPOTE	IN 128U		otation apaate			Endopte			
DC3CI	1011	-DIIVL	/IIVO I	IKOIL	114 1200					Diptera	rygota		
										Brachyc	ora		
										Muscom			
										Ephydro			
										Drosoph			
										Drosoph			
Ref	Niur	n Pos				Comment		Db	Medlineid		ilia	Cite	
Kei	ivui	11 1 03				Entry		- 00	Medilileid	Entr	V	Cite	
	1	SEO	HENC	CE FRO	Λ Ν Μ	Liitiy		MEDI INE	94166747		mer K.A	Mol. Gen. Genet. 242:391-398(1	004)
	Ι'	JLU	OLIVO	LIKO	IVI IV.A			IVILDLINL	74100747		ersen G	Wol. Gen. Genet. 242.371-370(1	774)
											tz E.K.F		
Embl	Sec	14	Prin	o Id	Status				Pir	Sec Id			
LIIIDI		A50701			Status				r II	S33467			
Interpro		Prim		000	ļ					S42582			
interpro	Id	1 11111	Iu						Mendel	342302	342302		
	-	IPR0	00765	5					Hssp				
Pfam	Sta	tus Se						Prim Id	Prosite	Status S	Sec Id		Prim Id
	1		P1_C	DBG				PF01018	1100110		GTP1_OBG		PS00905
Keyword	Ent								Gene		OR GTP-BP		1.000700
.,									Prints	12001	J. C. I. D.		
Features										GTP10	OBG		
Doma	ain	From	To	Descr						d PR003			
				Entr	rv				Flybase				
										128up)		
Bindi	ng									d FBqn0			
Signa	al								Maizedb				
Chair	1	From	To	Descr					Wormpep				
Disult	fid	From	To	Descr									
Non ⁻													
Mod I													
Confl	ict												
Carbo	ohyd	From	To	Descr	r								
Np Bi	ind	From		Descr									
		71			BY SIMIL								
		117			BY SIMIL								
		248	251	GTP (E	BY SIMIL	ARITY)							
Simila	~ "												

Mtype	PRT				Species	Clostridium sp. (stra	nin C 48-50)
Class	STANDARD					Entry	·
Id	12AH_CLOS4				Ĭ	Bacteria	
Seglen	29					Firmicutes	
Ac	Entry					Bacillus/Clostridium	group
						Clostridiaceae	
Mod	Date	Rel Type				Clostridium	
	01-AUG-1991	19 Create					
	01-AUG-1991		equence update				
	01-AUG-1991		nnotation update				
Descr	12-ALPHA-HYDROXYSTERC (FRAGMENT)	ID DEHYDF	ROGENASE (EC 1.1.				
Ref	Num Pos	Comme		Db	Medlineid		Cite
		Entry				Entry	
	1 SEQUENCE			MEDLINE	91177018		Eur. J. Biochem. 196:439-450(1991)
						Luensdorf H	
						Bueckmann	
					L .	A.F	
Embl	Sec Id Prim Id Statu	IS				Sec Id Prim Id \$14099 \$14099	
Interpro	Sec Prim Id				Mendel	314099 314099	
merpro	Id Id				Hssp		
	14					Status Sec Id	Prim Id
Pfam	Status Sec Id			Prim Id	FIUSILE	Status Sec Tu	FIIII IQ
					Gene		
Keyword	Entry				Prints		
	Oxidoreductase				Flybase		
	NADP				Maizedb		
Features					Wormpep		
Doma							
	Entry						
Bindin							
Signal							
Chain	From To Descr						
Disulf	id From To Descr						
Non T							
	rom 29						
T	o <mark>29</mark>						

Mod Res Conflict

Np Bind Similar

From To Descr Carbohyd From To Descr From To Descr

Mtype	PRT								Species	Fragaria ananassa (S	Strawberry)	
Class	STA	NDAF	RD						Org	Entry		
Id	12K	D_FR	AAN						Ü	Eukaryota		
Seglen	111									Viridiplantae		
Ac	Entr	y								Embryophyta		
										Tracheophyta		
Mod	Date	е			Re	el Type				Spermatophyta		
	01-0	OCT-1	996		34	1 Create	d			Magnoliophyta		
	01-0	OCT-1	996		34	1 Last se	equence update			eudicotyledons		
		NOV-1					nnotation update			Rosidae		
Descr	AUX	(IN-RI	EPRE:	SSED 12	2.5 KDA	PROTEIN	l			eurosids I		
										Rosales		
										Rosaceae		
										Fragaria		
Ref	Nun	n Pos				Comme	nt	Db	Medlineid	Author	Cite	
						Entry				Entry		
	1	SEC	QUEN	CE FRO	M N.A	STRA	IN=CV. OZARK	MEDLINE	91329668	Reddy A.S.N	Plant Mol. Biol. 14:127-136(1990)	
						BEAU				Poovaiah B.		
						TISSI	JE=FLOWER			W		
Embl	Sec			m Id	Status				Pir	Sec Id Prim Id		
		3667										
		7387		<u>41</u> 42					Mendel			
Interpro		Prim	ld						Hssp			
	Id								Prosite	Status Sec Id		Prim Id
									_			
Pfam	Stat	us Se	ec Id					Prim Id	Gene			
									Prints			
Keyword	Entr	У							Flybase			
Features									Maizedb			
Doma	nim.	From	То	Descr					Wormpep			
Doma	4111	FIOIII	10	Enti	r.							
		43	57	EIIU	ıy							
Bindir		43	37									
Signa												
Chain		From	То	Descr								
Cridiii	'	TOIT	10	Desci								
Disulf	fid	From	To	Descr								
Distail		110111	10	Desci								
Non 1	Ter											
Mod I												
Confli												
Carbo	ohyd	From	То	Descr	r							
Np Bi	ind	From	То	Descr								

Similar

Mtype	PRT				Species	Mycobacterium sme	nmatis
Class	STANDARD					Entry	gridus
Id	12KD_MYCSM					Bacteria	
Seglen	24					Firmicutes	
Ac	Entry					Actinobacteria	
AC	Lift y					Actinobacteridae	
Mod	Date	Rel	Туре			Actinomycetales	
WIOG	01-NOV-1995	32	Created			Corynebacterineae	
	01-NOV-1995		Last sequence update			Mycobacteriaceae	
	01-NOV-1995		Last annotation update			Mycobacterium	
Descr	12 KDA PROTEIN (FRAGME		Last armotation apacto			my oob a or or rain	
Ref	Num Pos		Comment	Db	Medlineid	Author	Cite
			Entry			Entry	
	1 SEQUENCE					Pahl A	Submitted (MAR-1995) to the SWISS-PROT
						Keller U	data bank
Embl	Sec Id Prim Id Statu	IS			Pir	Sec Id Prim Id	
						•	
Interpro	Sec Prim Id				Mendel		
	Id				Hssp		
					Prosite	Status Sec Id	Prim Id
Pfam	Status Sec Id			Prim Id			·
					Gene		
Keyword	Entry				Prints		
					Flybase		
Features					Maizedb		
Doma					Wormpep		
	Entry						
Dia dia							
Bindin							
Signal Chain							
Criairi	FIOIII TO Desci						
Disulfi	id From To Descr						
Distill	id Trom To Descr						
Non T	er						
	rom 24						
To							
Mod R							
Confli							
Carbo	hyd From To Descr						
Np Bir	nd From To Descr						
Simila	ir						

Mtype	PRT							Species	Arabidopsis thaliana	(Mouse-par cross)	
Class		NDARD	١					Org	Entry	(Mouse-ear cress)	
Id		_ARA						org	Eukaryota		
Seglen	472	_ANA							Viridiplantae		
Ac	Entry	,							Embryophyta		
AC	LIII								Tracheophyta		
Mod	Date			D	ام	Туре			Spermatophyta		
IVIOU		PR-19	90			Created			Magnoliophyta		
		PR-19				Last sequence update			eudicotyledons		
		EC-19				Last annotation update			Rosidae		
Descr				AGE PROTEII					eurosids II		
<i>B</i> 050.	120	JEED (0.0.0		•	TL CONCORT			Brassicales		
									Brassicaceae		
									Arabidopsis		
Ref	Num	Pos			(Comment	Db	Medlineid		Cite	
Itoi	Ivaiii	1 03			F	Entry		Wicamicia	Entry	- Once	
	1	SEOL	IENCE	FROM N.A	Ť	z.m.j			Pang P.P	Plant Mol. Biol. 11:805-820(1988)	
	•	JEQU	LIVOL	TROWN N.A					Pruitt R.E	Tidit Moi. Biol. 11.000 020(1700)	
									Meyerowitz		
									E.M		
	2	SEOL	IENCE	OF 420-472	,				Raynal M	Submitted (OCT-1992) to the	
	_		ЛN.A	0. 120 172					Grellet F	EMBL/GenBank/DDBJ databases	
									Laudie M		
									Meyer Y		
									Cooke R		
									Delseny M		
Embl	Sec	d	Prim	Id Status				Pir	Sec Id Prim Id		
211101		32777							S08509 S08509		
		32493						Mendel	000007		
		79005						Hssp			
Interpro		Prim I						Prosite	Status Sec Id		Prim Id
	Id								1 11S_SEED_S	TORAGE	PS00305
	-	IPR00	0459					Gene	CRA1		
Pfam	Statu	ıs Sec	Id				Prim Id	Prints			
	1		dstore	e_11s			PF00190		11SGLOBULIN		
Keyword	Entry	,						Prim I	d PR00439		
-	Seed	stora	ge pro	tein				Flybase			
	Multi	gene f	family					Maizedb			
	Signa	al						Wormpep			
Features											
Doma	ain F	rom	Γo D	escr							
				Entry							
Bindir											
Signa											
	rom										
		24									
		POTEN									
Chain		rom									
						BY SIMILARITY)					
					(B	Y SIMILARITY)					
Disulf		rom		escr							
		112 2	11 985	NTERCHAIN	(A	LPHA-BETA) (POTENTIAL)					
Non 7	Tor										

Np Bind Similar

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 Carbohyd
 From To
 Descr

 Np Bind
 From To
 Descr

Mtype	PRT						Species	Arabidopsis thaliana	(Mouse-ear cress)	
Class	STAN	IDARD					Org	Entry	,	
Id	12S2	ARATH						Eukaryota		
Seglen	455							Viridiplantae		
Ac	Entry	1						Embryophyta		
								Tracheophyta		
Mod	Date			Rel	Туре			Spermatophyta		
	01-A	PR-1990		14	Created			Magnoliophyta		
		PR-1990			Last sequence update			eudicotyledons		
		EC-1998			Last annotation update			Rosidae		
Descr	12S S	SEED STO	RAGE PROTI	EIN	PRECURSOR			eurosids II		
								Brassicales		
								Brassicaceae		
								Arabidopsis		
Ref	Num	Pos		L	Comment	Db	Medlineid		Cite	
					Entry			Entry		
	1	SEQUEN	CE FROM N.A	A				Pang P.P	Plant Mol. Biol. 11:805-820(1988)	
								Pruitt R.E		
								Meyerowitz		
								E.M		
	2		CE OF 240-3	60				Raynal M	Submitted (OCT-1992) to the	
		FROM N.	A					Grellet F	EMBL/GenBank/DDBJ databases	
								Laudie M		
								Meyer Y		
								Cooke R Delseny M		
Embl	Sec I	d Drir	n Id Stati	ıc		1	Pir	Sec Id Prim Id		
EIIIDI		32778 M3		12			FII	S08510 S08510		
		32494 X14					Mendel	300310 300310		
		9024 Z17					Hssp			
Interpro		Prim Id					Prosite	Status Sec Id		Prim Id
orpro	Id							1 11S_SEED_S	STORAGE	PS00305
	-	IPR00045	9				Gene	CRB		1
Pfam	Statu	s Sec Id				Prim Id	Prints	1 -		
	1	Seedsto	ore_11s			PF00190	Sec Id	11SGLOBULIN		
Keyword	Entry	1					Prim I	d PR00439		
	Seed	storage p	rotein				Flybase			
	Multi	gene fami	ily				Maizedb			
	Signa	al					Wormpep			
Features										
Doma	ain F	rom To	Descr							
			Entry							
Bindir										
Signa										
	rom									
		24								
		POTENTIA								
Chain	_	rom To		INI	(BY SIMILARITY)					
					(BY SIMILARITY) BY SIMILARITY)					
Disulf			Descr	14 (E	DI SIIVILAKIIY)					
Disuil				N (A	LPHA-BETA) (POTENTIAL)					
		270	LINOI IAII	. (/	L DE IM (I O I ENTIAL)					

Np Bind Similar

 Conflict
 From To
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 Carbohyd
 From To
 Descr

 Np Bind
 From To
 Descr

	PRT					Species	Entam	oeba histolyti	ca	
		DARD				Org	Entry	and the state of the		
Id		_ENTHI					Eukary	ota		
Seglen	239							oebidae		
Ac	Entry						Entam	oeba		
				.1_						
	Date			Туре						
		OV-1995	32							
		OV-1995	32							
		OV-1997		Last annotation update						
Descr		3 PROTEIN 1 (14-3-3-	-1)		1					
Ref	Num	Pos		Comment	Db	Medlineid			Cite	
	_	OF OUT NOT FROM N		Entry			En		0.1. 11. 1 (4110.400.4) 1. 11	
	1	SEQUENCE FROM N.A	4					muelson J	Submitted (AUG-1994) to the EMBL/GenBank/DDBJ databases	
								en P eckler G	EWBL/Geribarik/DDBJ databases	
								scoteaux		
							S	Scoleaux		
							Fu	н		
								hia A		
Embl	Sec Io	d Prim Id Statu	ıs			Pir		Prim Id		
Lilloi		0185 U13418	45				occ iu	i i i i i i		
Interpro		Prim Id				Mendel				
	Id					Hssp				
		PR000308					Status	Sec Id		Prim Id
Pfam	Statu	s Sec Id			Prim Id		1	1433_1		PS0079
	1	14-3-3			PF00244		1	1433_2		PS0079
Keyword	Entry					Gene				
_						Prints				
Features						Sec Id	1433	BZETA		
						Prim I	d PRO	0305		
						Flybase				
						Maizedb				
						Wormpep				
	DDT					C!				
	PRT	DARD				Species Org		ersicon escule	ntum (Tomato)	
		LYCES				org	Entry	oto		
	249	LICES					Viridip			
_	Entry	_						ophyta		
AC	Liitiy							eophyta		
Mod	Date		Re	Type				atophyta		
		L-1998	36					liophyta		
		L-1998		Last sequence update				tyledons		
		L-1998		Last annotation update			Asterio			
Descr		3-LIKE PROTEIN 1	-	Last annotation apacto			euaste			
							Solana			
							Solana			
							Solanu			
Ref	Num	Pos		Comment	Db	Medlineid			Cite	
	IVUIII			Entry			En			
	IVaiii								Submitted (JAN-1997) to the	
	1	SEQUENCE FROM N.A	4	STRAIN=CV.			Ro	berts M.R		
			4	MONEYMAKER				wles D.J	EMBL/GenBank/DDBJ databases	
	1	SEQUENCE FROM N.A					Во	wles D.J	EMBL/GenBank/DDBJ databases	
	1 Sec Io	SEQUENCE FROM N. /		MONEYMAKER		Pir		wles D.J	EMBL/GenBank/DDBJ databases	
Embl	1 Sec Io	SEQUENCE FROM N.A		MONEYMAKER			Во	wles D.J	EMBL/GenBank/DDBJ databases	
Embl Interpro	1 Sec Id CAA6 Sec F	SEQUENCE FROM N. /		MONEYMAKER		Mendel	Во	wles D.J	EMBL/GenBank/DDBJ databases	
Embl Interpro	Sec Io CAA6 Sec Id	SEQUENCE FROM N./ d Prim Id Statu 5145 X95900		MONEYMAKER		Mendel Hssp	Sec Id	Prim Id	EMBL/GenBank/DDBJ databases	
Embl Interpro	Sec Id CAA6 Sec Id	d Prim Id Statu 5145 X95900 Prim Id		MONEYMAKER		Mendel Hssp Prosite	Sec Id	Prim Id Sec Id	EMBL/GenBank/DDBJ databases	Prim Id
Embl Interpro	Sec Id CAA6 Sec Id	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER	Prim Id	Mendel Hssp Prosite	Sec Id Status	Prim Id Sec Id 1433_1	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam	Sec Id CAA6 Sec Id Id Statu:	d Prim Id Statu 5145 X95900 Prim Id		MONEYMAKER	Prim Id PF00244	Mendel Hssp Prosite	Sec Id Status 1	Prim Id Sec Id	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam	Sec Id CAA6 Sec Id	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene	Sec Id Status	Prim Id Sec Id 1433_1	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints	Sec Id Status 1 TFT1	Prim Id Sec Id 1433_1 1433_2	EMBL/GenBank/DDBJ databases	
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I Flybase	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I Flybase Maizedb	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I Flybase	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I Flybase Maizedb	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I Flybase Maizedb	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079
Embl Interpro Pfam Keyword	Sec Id CAA6 Sec Id Id Statu:	d Prim Id State 5145 X95900 Prim Id PR000308		MONEYMAKER		Mendel Hssp Prosite Gene Prints Sec Id Prim I Flybase Maizedb	Status 1 TFT1 1433	Prim Id Sec Id 1433_1 1433_2 SZETA	EMBL/GenBank/DDBJ databases	PS0079

Mtype	PRT				Species	Zea mays (Maize)		
Class	STANDARD				Org	Entry		
ld	1431_MAIZE				ŭ	Eukaryota		
Seglen	261					Viridiplantae		
Ac	Entry					Embryophyta		
						Tracheophyta		
Mod	Date	Rel	Туре			Spermatophyta		
	01-FEB-1996	33	Created			Magnoliophyta		
	01-FEB-1996	33	Last sequence update			Liliopsida		
	01-OCT-1996	34	Last annotation update			Poales		
Descr	14-3-3-LIKE PROTEIN GF14	-6				Poaceae		
						Zea		
Ref	Num Pos	(Comment	Db	Medlineid	Author	Cite	
			Entry			Entry		
	1 SEQUENCE FROM N.A	١		MEDLINE	95148741		Plant Physiol. 106:1593-1604(1994)	
						<u>C</u>		
						Ferl R.J		
Embl	Sec Id Prim Id Statu	IS			Pir	Sec Id Prim Id		
	AAB33304 S77133							
Interpro	Sec Prim Id				Mendel			
	Id				Hssp	a la		
	- IPR000308				Prosite	Status Sec Id		Prim Id
Pfam	Status Sec Id			Prim Id		1 1433_1		PS00796
14	1 14-3-3			PF00244	•	1 1433_2		PS00797
Keyword	Entry					GRF1		
F. strees					Prints Sec Id	1433ZETA		
Features						1433ZETA d PR00305		
					Flybase	1 PR00305		
					Maizedb			
					Sec Id	d 65832		
					Wormpep	u 03032		
				l	wompep			

	DD=									0 1 1 1		. (0) (1.)
Mtype	PRT	10.45									osoma mansoi	ni (Blood fluke)
Class		NDAR							Org	Entry		
Id		_SCF	IMA							Eukary		
Seglen	252									Metaz		
Ac	Entr	У									elminthes	
											ermata	
Mod	Date					I Type				Trema		
		UL-19				Created				Digene		
		UL-19				Last sequence				Strigei		
		UL-19				Last annotati	ion update				osomatoidea	
Descr	14-3	-3 PR	OTE	IN HOM	10L0G 1						osomatidae	
											osoma	
Ref	Num	Pos				Comment		Db	Medlineid	Autho	or	Cite
						Entry				En	itry	
	1	SEQ	UEN	CE FRO	M N.A			MEDLINE	96123403	Sc	hechtman	Mol. Biochem. Parasitol. 73:275-278(1995)
										D		
										Ra	ım D	
										Ta	rrab-	
										Ha	izdai R	
										Arı	non R	
										Sc	hechter I	
Embl	Sec	ld	Prir	n Id	Status				Pir	Sec Id	l Prim Id	
	AAC	46983	U24	4281								
Interpro	Sec	Prim	ld						Mendel			
·	Id								Hssp			
	-	IPR00	0030	8					Prosite	Status	Sec Id	Prim Id
Pfam	Statu	ıs Se	c Id					Prim Id		1	1433_1	PS00796
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Keyword	Entr	У							Gene			<u> </u>
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Features									Sec Id	1433	3ZETA	
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From To Descr

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To 149
Descr TO THE C-TERMINUS OF ANNEXINS

Np Bind Similar

Mtype	PRT							oeba histolyt	ica	
Class	STAN	DARD					Entry			
Id		ENTHI					Eukary	nta		
Seglen	238	_CIVIIII						oebidae		
Ac							Entam			
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Mtype										
	PRT						Lycope	ersicon escule	entum (Tomato)	
Class		DARD				Species	Lycope Entry	ersicon escule	entum (Tomato)	
Class	STAN					Species Org	Entry		entum (Tomato)	
Class Id	STAN 1432	DARD _LYCES				Species Org	Entry Eukary	ota	entum (Tomato)	
Class Id Seqlen	STAN 1432 254	LYCES				Species Org	Entry Eukary Viridip	ota antae	entum (Tomato)	
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- 71	PRT							Zea mays (Maize)	
		IDARD					5	Entry	
Id	1432	<u>_M</u> A12	ZE					Eukaryota	
	261							Viridiplantae	
Ac	Entry	1						Embryophyta	
								Tracheophyta	
	Date				Type			Spermatophyta	
	01-Jl	JL-199	93	26				Magnoliophyta	
		EB-199		33				Liliopsida	
		EB-199			Last annotation update			Poales	
Descr	14-3-	-3-LIK	E PROTEIN	GF14-12	2			Poaceae	
								Zea	
Ref	Num	Pos			Comment	Db	Medlineid	Author	Cite
		SEQUENCE FROM N.A.			Entry			Entry	
	1	1 SEQUENCE FROM N.		M N.A		MEDLINE	95148741	de Vetten N.	Plant Physiol. 106:1593-1604(1994)
								С	
								Ferl R.J	
	2	SEQL	JENCE OF 1	14-261		MEDLINE	93076113	de Vetten N.	Plant Cell 4:1295-1307(1992)
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	-		S77135	NOT_AN	INOTATED_CDS			JQ1680 JQ1680	
	AAA3	3505	M96856				Mendel		
Interpro	Sec I	Prim I	d				Hssp		
	Id						Prosite	Status Sec Id	Prim Id
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Pfam	Statu	is Sec	ld			Prim Id		1 1433_2	PS00797
	1	14-	3-3			PF00244	Gene	GRF2	
Keyword	Entry	1					Prints		
							Sec Id	1433ZETA	
Features							Prim I	PR00305	
							Flybase		
							Maizedb		
							Sec Id	-	
							Prim I	d 65832	

Mtype	PRT		_					Species	Schisto	soma mansoi	ni (Blood fluke)	
Class		DARD						Org	Entry			
Id	1432	SCHMA							Eukary	ota		
Seglen	214								Metazo	oa		
Ac	Entry								Platyhe	elminthes		
									Neode	rmata		
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	15-JL	L-1998		3	6 La	ast sequence update			Strigeio	dida		
	15-JL	L-1998		3	6 La	ast annotation update			Schisto	somatoidea		
Descr	14-3-	3 PROTE	IN HON	10LOG 2	(14	-3-3-2) (FRAGMENT)			Schisto	somatidae		
									Schisto	soma		
Ref	Num	Pos			Cor	mment	Db	Medlineid	Autho	r	Cite	
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	1	SEQUEN	CE FRO	M N.A					Lee	e K.W	Submitted (JUN-1996) to the	
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									Lov	verde P.T		
Embl	Sec I	d Pri	m Id	Status				Pir	Sec Id	Prim Id		
	AAB0	2100 L78	3441									
Interpro	Sec F	Prim Id		•				Mendel				
	Id							Hssp				
	- I	PR00030	8					Prosite	Status	Sec Id		Prim Id
Pfam	Statu	s Sec Id					Prim Id		1	1433_1		PS00796
	1	14-3-3					PF00244		1	1433_2		PS00797
Keyword	Entry							Gene				
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A	Mtype	PRT							Species	Caoro	rhabditis ele	nane	
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_	d		CAE						Org	Eukary	nta		
	Seglen	248	_CAE	EL						Metazo			
	Ac	Entr	.,							Nemat			
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		Q21								Rhabd			
	Mod	Date				Dol	Туре				itoidea		
	viou		IOV-19	005			Created			Rhabd			
			IOV-19				Last sequence update			Pelode			
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Г	Descr			E PROTEI		5,	Last annotation apacte			oderio	mabantis		
	Ref		Pos	L I NOTE			Comment	Db	Medlineid	Autho	r	Cite	
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		1	SEO	JENCE FR	OM N A		z.m.y	MEDI INF	95011616		ing W	Gene 147:215-218(1994)	
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		AAA	61872	U05038									
		CAA	98138	Z73910					Mendel				
Ī	nterpro	Sec	Prim I	d					Hssp				
		Id							Prosite	Status	Sec Id		Prim Id
		-	IPR00	0308						1	1433_1		PS00796
F	Pfam	Stat	us Sec					Prim Id		1	1433_2		PS00797
		1		3-3				PF00244		FTT-1	OR M117.2		
k	Keyword	Entr	у						Prints				
										1433			
F	eatures									d PRO	305		
	Doma	in	From						Flybase				
		L		En	itry				Maizedb				
									Wormpep				
	Bindir								Prim I	d M11	7		
	Signa												
	Chain	' <u> </u>	From	To Desci	ſ								
	D: 16			T D									
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	COIIII	-		118 A ->		FF	2)						
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	oui bo	,		.0 DC3	·.	•							
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	Simila	ar											

Mtype	PRT				Species	Candida albicans (Y	'east)	
Class	STANDARD				Org	Entry	•	
Id	1433_CANAL				Ĭ	Eukaryota		
Seglen	264					Fungi		
Ac	Entry					Ascomycota		
						Saccharomycetes		
Mod	Date	Re	el Type			Saccharomycetales		
	15-JUL-1998	36			Ī	anamorphic Saccha	romycetales	
	15-JUL-1998		Last sequence update		-	Candida	,	
	15-JUL-1999		Last annotation update					
Descr	14-3-3 PROTEIN HOMOL		•					
Ref	Num Pos		Comment	Db	Medlineid	Author	Cite	
			Entry			Entry	1	
	1 SEQUENCE FROM	V.A				Cognetti D	Submitted (DEC-1997) to the	
						Sturtevant J	EMBL/GenBank/DDBJ databases	
Fmbl	Sec Id Prim Id St	atus			Pir	Sec Id Prim Id		
	AAB96910 AF038154	atus			1	000 14 111111 14		
Interpro	Sec Prim Id				Mendel			
p. 5	Id				Hssp			
	- IPR000308					Status Sec Id		Prim Id
Pfam	Status Sec Id			Prim Id		1 1433_1		PS007
	1 14-3-3			PF00244		1 1433_2		PS007
Keyword	Entry				Gene	BMH1 OR BMH		
.,					Prints			
Features	•				Sec Io	1433ZETA		
						d PR00305		
					Flybase			
					Maizedb			
					Wormpep			
Mtype	PRT					Chlamydomonas re	inhardtii	
Class	STANDARD				Org	Entry		
ld	1433_CHLRE							
Seglen	259							
Ac	Entry							
			_		_			
Mod	Date		l Type					
	01-OCT-1996	34						
	01-OCT-1996	34						
	01-OCT-1996	34	Last annotation update					
Descr	14-3-3-LIKE PROTEIN				Ļ			
Ref	Num Pos		Comment	Db	Medlineid		Cite	
			Entry			Entry		
Embl	Sec Id Prim Id St	atus			Pir	Sec Id Prim Id		
Interpro	Sec Prim Id				Mendel			
	Id				Hssp			
D.C.	0			D: 1:	Prosite	Status Sec Id		Prim Id
Pfam	Status Sec Id			Prim Id				
Vararia	Futu.				Gene			
Keyword	Entry				Prints			
F. et					Flybase			
Features					Maizedb			
					Wormpep			
					rvvormpep			