Entry

- Mtype: PRT
  Class: STANDARD
  Id: 100K_RAT
  Seqlen: 889
  Ac
  - Entry:
  Mod
  - Date: 01-NOV-1997
    Rel: 35
    Type: Created
  - Date: 01-NOV-1997
    Rel: 35
    Type: Last sequence update
  - Date: 15-JUL-1999
    Rel: 38
    Type: Last annotation update

  Descr: 100 KDA PROTEIN (EC 6.3.2.-)
  Species: Rattus norvegicus (Rat)
  Org
  - Entry: Eukaryota
  - Entry: Metazoa
  - Entry: Chordata
  - Entry: Craniata
  - Entry: Vertebrata
  - Entry: Euteleostomi
  - Entry: Mammalia
  - Entry: Eutheria
  - Entry: Rodentia
  - Entry: Sciurognathi
  - Entry: Muridae
  - Entry: Murinae
  - Entry: Rattus

  Ref
  - Num: 1
    Pos: SEQUENCE FROM N.A
    Comment
    - Entry: STRAIN=WISTAR
    - Entry: TISSUE=TESTIS

    Db: MEDLINE
    Medlineid: 92253337
    Author
    - Entry: Mueller D
    - Entry: Rehbein M
    - Entry: Baumeister H
    - Entry: Richter D

    Cite: Nucleic Acids Res. 20:1471-1475(1992)
  - Num: 2
    Pos: ERRATUM
    Comment
    Db:
    Medlineid:
    Author
    - Entry: Mueller D
    - Entry: Rehbein M
    - Entry: Baumeister H
    - Entry: Richter D

    Cite: Nucleic Acids Res. 20:2624-2624(1992)

Embl

- Sec Id: CAA45756
  Prim Id: X64411
  Status:

Interpro

- Sec Id: -
  Prim Id: IPR000569

- Sec Id: -
  Prim Id: IPR002004

Pfam

- Status: 1
  Sec Id: HECT
  Prim Id: PF00632

- Status: 1
  Sec Id: PABP
  Prim Id: PF00658

Keyword

- Entry: Ubiquitin conjugation

- Entry: Ligase

Features

Domain

- From: 77
  To: 88
  Descr
  - Entry:

- From: 127
  To: 150
  Descr
  - Entry:

- From: 420
  To: 439
  Descr
  - Entry:

- From: 448
  To: 457
  Descr
  - Entry:

- From: 485
  To: 514
  Descr
  - Entry:

- From: 579
  To: 590
  Descr
  - Entry:

- From: 786
  To: 889
  Descr
  - Entry:

- From: 827
  To: 847
  Descr
  - Entry:

Binding

From: 858
To: 858
Descr: UBIQUITIN (BY SIMILARITY)

Signal

Chain

Disulfid

Non Ter

Mod Res

Conflict

Carbohyd

Np Bind

Similar

- Pir
- Mendel
- Hssp
- Prosite
- Gene
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 104K_THEPA
- Seqlen: 924
- Ac
  - Entry
- Mod
  - Date: 01-APR-1990
    Rel: 14
    Type: Created
  - Date: 01-APR-1990
    Rel: 14
    Type: Last sequence update
  - Date: 01-AUG-1992
    Rel: 23
    Type: Last annotation update
- Descr: 104 KDA MICRONEME-RHOPTRY ANTIGEN
- Species: Theileria parva
- Org
  - Entry: Eukaryota
  - Entry: Alveolata
  - Entry: Apicomplexa
  - Entry: Piroplasmida
  - Entry: Theileriidae
  - Entry: Theileria
- Ref
  - Num: 1
    Pos: SEQUENCE FROM N.A
    Comment
    - Entry
    Db: MEDLINE
    Medlineid: 90158697
    Author
    - Entry: Iams K.P
    - Entry: Young J.R
    - Entry: Nene V
    - Entry: Desai J
    - Entry: Webster P
    - Entry: Ole-Moiyoi O.K
    - Entry: Musoke A.J

    Cite: Mol. Biochem. Parasitol. 39:47-60(1990)
- Embl
  - Sec Id: AAA18217
    Prim Id: M29954
    Status
- Interpro
- Pfam
- Keyword
  - Entry: Antigen
  - Entry: Sporozoite
  - Entry: Repeat

- Features
  - Domain
    - From 1
      To 19
      Descr
      - Entry
    - From 905
      To 924
      Descr
      - Entry
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
  - Sec Id A44945
    Prim Id A44945
- Mendel
- Hssp
- Prosite
- Gene
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype PRT
- Class STANDARD
- Id 108_LYCES
- Seqlen 102
- Ac
  - Entry
- Mod
  - Date 15-JUL-1999
    Rel 38
    Type Created
  - Date 15-JUL-1999
    Rel 38
    Type Last sequence update
  - Date 15-JUL-1999
    Rel 38
    Type Last annotation update
- Descr PROTEIN 108 PRECURSOR
- Species Lycopersicon esculentum (Tomato)
- Org
  - Entry Eukaryota
  - Entry Viridiplantae
  - Entry Embryophyta
  - Entry Tracheophyta
  - Entry Spermatophyta
  - Entry Magnoliophyta
  - Entry eudicotyledons
  - Entry Asteridae
  - Entry euasterids I
  - Entry Solanales
  - Entry Solanaceae
  - Entry Solanum

- Ref
  - Num: 1
  - Pos: SEQUENCE FROM N.A
  - Comment
    - Entry: STRAIN=CV. VF36
    - Entry: TISSUE=ANTHER
  - Db: MEDLINE
  - Medlineid: 94143497
  - Author
    - Entry: Chen R
    - Entry: Smith A.G
  - Cite: Plant Physiol. 101:1413-1413(1993)
- Embl
  - Sec Id: CAA78466
  - Prim Id: Z14088
  - Status:
- Interpro
- Pfam
- Keyword
  - Entry:
- Features
  - Domain
  - Binding
  - Signal
    - From: 1
    - To: 30
    - Descr: POTENTIAL
  - Chain
    - From: 31
    - To: 102
    - Descr: PROTEIN 108
  - Disulfid
    - From: 41
    - To: 77
    - Descr: BY SIMILARITY
    - From: 51
    - To: 66
    - Descr: BY SIMILARITY
    - From: 67
    - To: 92
    - Descr: BY SIMILARITY
    - From: 79
    - To: 99
    - Descr: BY SIMILARITY
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
- Mendel
  - Status: 1133
  - Sec Id: LYCes
  - Prim Id: 8853
- Hssp
- Prosite
- Gene:
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  - Class: STANDARD
  - Id: 10KD_VIGUN
  - Seqlen: 75
  - Ac
    - Entry:

Mod

- Date 01-NOV-1990
  Rel 16
  Type Created
- Date 01-NOV-1990
  Rel 16
  Type Last sequence update
- Date 01-FEB-1995
  Rel 31
  Type Last annotation update

Descr 10 KDA PROTEIN PRECURSOR (CLONE PSAS10)

Species Vigna unguiculata (Cowpea)

Org

- Entry Eukaryota
- Entry Viridiplantae
- Entry Embryophyta
- Entry Tracheophyta
- Entry Spermatophyta
- Entry Magnoliophyta
- Entry eudicotyledons
- Entry Rosidae
- Entry eurosids I
- Entry Fabales
- Entry Fabaceae
- Entry Papilionoideae
- Entry Vigna

Ref

- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  - Entry
  Db MEDLINE
  Medlineid 91355865
  Author
  - Entry Ishibashi N
  - Entry Yamauchi D
  - Entry Miniamikawa T

  Cite Plant Mol. Biol. 15:59-64(1990)

Embl

- Sec Id CAA34760
  Prim Id X16877
  Status

Interpro

- Sec Id -
  Prim Id IPR002118

Pfam

- Status 1
  Sec Id Gamma-thionin
  Prim Id PF00304

Keyword

- Entry Germination
- Entry Signal

- Features
  - Domain
  - Binding
  - Signal
    - From: 1
    - To: ?
    - Descr: POTENTIAL
  - Chain
    - From: ?
    - To: 75
    - Descr: 10 kDa PROTEIN
  - Disulfid
    - From: 31
      To: 75
      Descr: BY SIMILARITY
    - From: 42
      To: 63
      Descr: BY SIMILARITY
    - From: 48
      To: 69
      Descr: BY SIMILARITY
    - From: 52
      To: 71
      Descr: BY SIMILARITY
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
  - Sec Id: S11156
    Prim Id: S11156
- Mendel
- Hssp
  - Sec Id: 1CHL
    Prim Id: P45639
- Prosite
  - Status: 1
    Sec Id: GAMMA_THIONIN
    Prim Id: PS00940
- Gene
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 110K_PLAKN
  Seqlen: 296
  Ac
  - Entry:
  Mod
  - Date: 01-JAN-1990
    Rel: 13
    Type: Created
  - Date: 01-JAN-1990
    Rel: 13
    Type: Last sequence update
  - Date: 01-FEB-1994
    Rel: 28
    Type: Last annotation update
  Descr: 110 KDA ANTIGEN (PK110) (FRAGMENT)
  Species: Plasmodium knowlesi

- Org
  - Entry: Eukaryota
  - Entry: Alveolata
  - Entry: Apicomplexa
  - Entry: Haemosporida
  - Entry: Plasmodium
- Ref
  - Num: 1
  - Pos: SEQUENCE FROM N.A
  - Comment
  - Db: MEDLINE
  - Medlineid: 88039002
  - Author
    - Entry: Perler F.B
    - Entry: Moon A.M
    - Entry: Qiang B.Q
    - Entry: Meda M
    - Entry: Dalton M
    - Entry: Card C
    - Entry: Schmidt-Ullrich R
    - Entry: Wallach D
    - Entry: Lynch J
    - Entry: Donelson J.E
  - Cite: Mol. Biochem. Parasitol. 25:185-193(1987)
- Embl
  - Sec Id: AAA29471
  - Prim Id: M19152
  - Status
- Interpro
- Pfam
- Keyword
  - Entry: Malaria
  - Entry: Antigen
  - Entry: Repeat
- Features
  - Domain
    - From: 131
    - To: 296
    - Descr
      - Entry: 13.5 X 12 AA TANDEM REPEATS OF E-E-T-Q-K-
      - Entry: T-V-E-P-E-Q-T
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
    - From: 1
    - To: 1
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
  - Sec Id: A54527
  - Prim Id: A54527
- Mendel
- Hssp
- Prosite
- Gene
- Prints
- Flybase
- Maizedb

Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 11S3_HELAN
  Seqlen: 493
  Ac
  - Entry:
  Mod
  - Date: 01-NOV-1990
    Rel: 16
    Type: Created
  - Date: 01-NOV-1990
    Rel: 16
    Type: Last sequence update
  - Date: 30-MAY-2000
    Rel: 39
    Type: Last annotation update
  Descr: 11S GLOBULIN SEED STORAGE PROTEIN G3 PRECURSOR (HELIANTHININ G3)
  Species: Helianthus annuus (Common sunflower)
  Org
  - Entry: Eukaryota
  - Entry: Viridiplantae
  - Entry: Embryophyta
  - Entry: Tracheophyta
  - Entry: Spermatophyta
  - Entry: Magnoliophyta
  - Entry: eudicotyledons
  - Entry: Asteridae
  - Entry: euasterids II
  - Entry: Asterales
  - Entry: Asteraceae
  - Entry: Asteroideae
  - Entry: Heliantheae
  - Entry: Helianthus
  Ref
  - Num: 1
    Pos: SEQUENCE FROM N.A
    Comment
    Db: MEDLINE
    Medlineid: 89232734
    Author
    - Entry: Vonder Haar R.A
    - Entry: Allen R.D
    - Entry: Cohen E.A
    - Entry: Nessler C.L
    - Entry: Thomas T.L

    Cite: Gene 74:433-443(1988)
  Embl
  - Sec Id: AAA33374
    Prim Id: M28832
    Status:
  Interpro
  - Sec Id: -
    Prim Id: IPR000459
  Pfam
  - Status: 1
    Sec Id: Seedstore_11s
    Prim Id: PF00190

Keyword
- Entry: Seed storage protein
- Entry: Multigene family
- Entry: Signal

Features

Domain
- From: 23
  To: 35
  Descr
  - Entry:
- From: 111
  To: 127
  Descr
  - Entry:
- From: 191
  To: 297
  Descr
  - Entry:

Binding

Signal
From: 1
To: 20
Descr:

Chain
- From: 21
  To: 305
  Descr: ACIDIC CHAIN
- From: 306
  To: 493
  Descr: BASIC CHAIN

Disulfid
- From: 103
  To: 312
  Descr: INTERCHAIN (ACIDIC-BASIC) (POTENTIAL)

Non Ter

Mod Res

Conflict

Carbohyd

Np Bind

Similar

Pir
- Sec Id: JA0089
  Prim Id: JA0089

Mendel

Hssp

Prosite
- Status: 1
  Sec Id: 11S_SEED_STORAGE
  Prim Id: PS00305

Gene: HAG3

Prints
Sec Id: 11SGLOBULIN
Prim Id: PR00439

Flybase

Maizedb

Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 11SB_CUCMA
  Seqlen: 480
  Ac
  - Entry:

Mod

- Date 01-JAN-1990
  Rel 13
  Type Created
- Date 01-JAN-1990
  Rel 13
  Type Last sequence update
- Date 01-NOV-1990
  Rel 16
  Type Last annotation update

Descr 11S GLOBULIN BETA SUBUNIT PRECURSOR

Species Cucurbita maxima (Pumpkin) (Winter squash)

Org

- Entry Eukaryota
- Entry Viridiplantae
- Entry Embryophyta
- Entry Tracheophyta
- Entry Spermatophyta
- Entry Magnoliophyta
- Entry eudicotyledons
- Entry Rosidae
- Entry eurosids I
- Entry Cucurbitales
- Entry Cucurbitaceae
- Entry Cucurbita

Ref

- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  - Entry
  Db MEDLINE
  Medlineid 88166744
  Author
  - Entry Hayashi M
  - Entry Mori H
  - Entry Nishimura M
  - Entry Akazawa T
  - Entry Hara-Nishimura I

  Cite Eur. J. Biochem. 172:627-632(1988)
- Num 2
  Pos SEQUENCE OF 22-30 AND 297-302
  Comment
  Db
  Medlineid
  Author
  - Entry Ohmiya M
  - Entry Hara I
  - Entry Mastubara H

  Cite Plant Cell Physiol. 21:157-167(1980)

Embl

- Sec Id AAA33110
  Prim Id M36407
  Status

Interpro

- Sec Id -
  Prim Id IPR000459

Pfam

- Status 1
  Sec Id Seedstore_11s
  Prim Id PF00190

Keyword
- Entry: Seed storage protein
- Entry: Signal

Features
- Domain
- Binding
- Signal
  - From: 1
  - To: 21
  - Descr:
- Chain
  - From: 22
  - To: 480
  - Descr: 11S GLOBULIN BETA SUBUNIT
  - From: 22
  - To: 296
  - Descr: GAMMA CHAIN (ACIDIC)
  - From: 297
  - To: 480
  - Descr: DELTA CHAIN (BASIC)
- Disulfid
  - From: 124
  - To: 303
  - Descr: INTERCHAIN (GAMMA-DELTA) (POTENTIAL)
- Non Ter
- Mod Res
  - From: 22
  - To: 22
  - Descr: PYRROLIDONE CARBOXYLIC ACID
- Conflict
  - From: 27
  - To: 27
  - Descr: S -> E (IN REF. 2)
  - From: 30
  - To: 30
  - Descr: E -> S (IN REF. 2)
- Carbohyd
- Np Bind
- Similar

Pir
- Sec Id: FWPU1B
- Prim Id: S00366

Mendel

Hssp

Prosite
- Status: 1
- Sec Id: 11S_SEED_STORAGE
- Prim Id: PS00305

Gene

Prints
- Sec Id: 11SGLOBULIN
- Prim Id: PR00439

Flybase

Maizedb

Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 120K_RICRI
- Seqlen: 1300

Ac
- Entry:

Mod

- Date 01-APR-1990
  Rel 14
  Type Created
- Date 01-FEB-1996
  Rel 33
  Type Last sequence update
- Date 01-FEB-1996
  Rel 33
  Type Last annotation update

Descr 120 KDA SURFACE-EXPOSED PROTEIN

Species Rickettsia rickettsii

Org

- Entry Bacteria
- Entry Proteobacteria
- Entry alpha subdivision
- Entry Rickettsiales
- Entry Rickettsiaceae
- Entry Rickettsieae
- Entry Rickettsia

Ref

- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  - Entry
    Db MEDLINE
    Medlineid 90136087
    Author
    - Entry Gilmore R.D. Jr
    - Entry Joste N
    - Entry McDonald G.A

    Cite Mol. Microbiol. 3:1579-1586(1989)

Embl

- Sec Id CAA34402
  Prim Id X16353
  Status

Interpro

Pfam

Keyword

- Entry Antigen
- Entry Glycoprotein
- Entry S-layer

Features

Domain

Binding

Signal

Chain

Disulfid

Non Ter

Mod Res

Conflict

Carbohyd

- From 7
  To 7
  Descr POTENTIAL
- From 66
  To 66
  Descr POTENTIAL
- From 86
  To 86
  Descr POTENTIAL

From	103
To	103
Descr	POTENTIAL
From	147
To	147
Descr	POTENTIAL
From	268
To	268
Descr	POTENTIAL
From	330
To	330
Descr	POTENTIAL
From	375
To	375
Descr	POTENTIAL
From	415
To	415
Descr	POTENTIAL
From	424
To	424
Descr	POTENTIAL
From	430
To	430
Descr	POTENTIAL
From	436
To	436
Descr	POTENTIAL
From	444
To	444
Descr	POTENTIAL
From	515
To	515
Descr	POTENTIAL
From	547
To	547
Descr	POTENTIAL
From	593
To	593
Descr	POTENTIAL
From	655
To	655
Descr	POTENTIAL
From	698
To	698
Descr	POTENTIAL
From	710
To	710
Descr	POTENTIAL
From	799
To	799
Descr	POTENTIAL
From	800
To	800
Descr	POTENTIAL
From	826
To	826
Descr	POTENTIAL
From	844
To	844
Descr	POTENTIAL

- From 861
  To 861
  Descr POTENTIAL
- From 879
  To 879
  Descr POTENTIAL
- From 920
  To 920
  Descr POTENTIAL
- From 926
  To 926
  Descr POTENTIAL
- From 1116
  To 1116
  Descr POTENTIAL
- From 1128
  To 1128
  Descr POTENTIAL
- From 1140
  To 1140
  Descr POTENTIAL
- From 1146
  To 1146
  Descr POTENTIAL
- From 1211
  To 1211
  Descr POTENTIAL

Np Bind

Similar

Pir

- Sec Id S07575
  Prim Id S07575

Mendel

Hssp

Prosite

Gene P120

Prints

Flybase

Maizedb

Wormpep

- Mtype PRT
  Class STANDARD
  Id 128U_DROME
  Seqlen 368

Ac

- Entry

Mod

- Date 01-OCT-1993
  Rel 27
  Type Created
- Date 01-OCT-1993
  Rel 27
  Type Last sequence update
- Date 01-FEB-1996
  Rel 33
  Type Last annotation update

Descr GTP-BINDING PROTEIN 128UP

Species Drosophila melanogaster (Fruit fly)

Org

- Entry: Eukaryota
- Entry: Metazoa
- Entry: Arthropoda
- Entry: Tracheata
- Entry: Hexapoda
- Entry: Insecta
- Entry: Pterygota
- Entry: Neoptera
- Entry: Endopterygota
- Entry: Diptera
- Entry: Brachycera
- Entry: Muscomorpha
- Entry: Ephydroidea
- Entry: Drosophilidae
- Entry: Drosophila

Ref

- Num: 1

Pos: SEQUENCE FROM N.A

Comment

- Entry:

Db: MEDLINE

Medlineid: 94166747

Author

- Entry: Sommer K.A
- Entry: Petersen G
- Entry: Bautz E.K.F

Cite: Mol. Gen. Genet. 242:391-398(1994)

Embl

- Sec Id: CAA50701

Prim Id: X71866

Status:

Interpro

- Sec Id: -

Prim Id: IPR000765

Pfam

- Status: 1

Sec Id: GTP1_OBG

Prim Id: PF01018

Keyword

- Entry:

- Features
  - Domain
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
    - From 71
      To 78
      Descr GTP (BY SIMILARITY)
    - From 117
      To 121
      Descr GTP (BY SIMILARITY)
    - From 248
      To 251
      Descr GTP (BY SIMILARITY)
  - Similar
- Pir
  - Sec Id S33467
    Prim Id S33467
  - Sec Id S42582
    Prim Id S42582
- Mendel
- Hssp
- Prosite
  - Status 1
    Sec Id GTP1_OBG
    Prim Id PS00905
- Gene 128UP OR GTP-BP
- Prints
  - Sec Id GTP1OBG
    Prim Id PR00326
- Flybase
  - Sec Id 128up
    Prim Id FBgn0010339
- Maizedb
- Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 12AH_CLOS4
- Seqlen: 29
- Ac
  - Entry:
- Mod
  - Date: 01-AUG-1991
    Rel: 19
    Type: Created
  - Date: 01-AUG-1991
    Rel: 19
    Type: Last sequence update
  - Date: 01-AUG-1991
    Rel: 19
    Type: Last annotation update
- Descr: 12-ALPHA-HYDROXYSTEROID DEHYDROGENASE (EC 1.1.1.176) (FRAGMENT)
- Species: Clostridium sp. (strain C 48-50)
- Org
  - Entry: Bacteria
  - Entry: Firmicutes
  - Entry: Bacillus/Clostridium group
  - Entry: Clostridiaceae
  - Entry: Clostridium
- Ref
  - Num: 1
    Pos: SEQUENCE
    Comment
    Db: MEDLINE
    Medlineid: 91177018
    Author
    - Entry: Braun M
    - Entry: Luensdorf H
    - Entry: Bueckmann A.F

    Cite: Eur. J. Biochem. 196:439-450(1991)
- Embl
- Interpro
- Pfam
- Keyword
  - Entry: Oxidoreductase
  - Entry: NADP
- Features
  - Domain
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
    From: 29
    To: 29
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
  - Sec Id: S14099
    Prim Id: S14099
- Mendel
- Hssp
- Prosite
- Gene:
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 12KD_FRAAN
- Seqlen: 111
- Ac
  - Entry:
- Mod
  - Date: 01-OCT-1996
    Rel: 34
    Type: Created
  - Date: 01-OCT-1996
    Rel: 34
    Type: Last sequence update
  - Date: 01-NOV-1997
    Rel: 35
    Type: Last annotation update
- Descr: AUXIN-REPRESSED 12.5 KDA PROTEIN
- Species: Fragaria ananassa (Strawberry)
- Org
  - Entry: Eukaryota
  - Entry: Viridiplantae
  - Entry: Embryophyta
  - Entry: Tracheophyta
  - Entry: Spermatophyta
  - Entry: Magnoliophyta
  - Entry: eudicotyledons
  - Entry: Rosidae
  - Entry: eurosids I
  - Entry: Rosales
  - Entry: Rosaceae
  - Entry: Fragaria
- Ref
  - Num: 1
    Pos: SEQUENCE FROM N.A
    Comment
    - Entry: STRAIN=CV. OZARK BEAUTY
    - Entry: TISSUE=FLOWER

    Db: MEDLINE
    Medlineid: 91329668
    Author
    - Entry: Reddy A.S.N
    - Entry: Poovaiah B.W

    Cite: Plant Mol. Biol. 14:127-136(1990)
- Embl
  - Sec Id: CAA36676
    Prim Id: X52429
    Status:
  - Sec Id: AAA73872
    Prim Id: L44142
    Status:
- Interpro
- Pfam
- Keyword

- Features
  - Domain
    - From: 43
      - To: 57
      - Descr
        - Entry
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
- Mendel
- Hssp
- Prosite
- Gene
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  - Class: STANDARD
  - Id: 12KD_MYCSM
  - Seqlen: 24
  - Ac
    - Entry:
  - Mod
    - Date: 01-NOV-1995
      Rel: 32
      Type: Created
    - Date: 01-NOV-1995
      Rel: 32
      Type: Last sequence update
    - Date: 01-NOV-1995
      Rel: 32
      Type: Last annotation update
  - Descr: 12 KDA PROTEIN (FRAGMENT)
  - Species: Mycobacterium smegmatis
  - Org
    - Entry: Bacteria
    - Entry: Firmicutes
    - Entry: Actinobacteria
    - Entry: Actinobacteridae
    - Entry: Actinomycetales
    - Entry: Corynebacterineae
    - Entry: Mycobacteriaceae
    - Entry: Mycobacterium
  - Ref
    - Num: 1
      Pos: SEQUENCE
      Comment
      Db
      Medlineid
      Author
        - Entry: Pahl A
        - Entry: Keller U
      Cite: Submitted (MAR-1995) to the SWISS-PROT data bank
  - Embl
  - Interpro
  - Pfam
  - Keyword
  - Features
    - Domain
    - Binding
    - Signal
    - Chain
    - Disulfid
    - Non Ter
      From: 24
      To: 24
    - Mod Res
    - Conflict
    - Carbohyd
    - Np Bind
    - Similar
  - Pir
  - Mendel
  - Hssp
  - Prosite
  - Gene
  - Prints
  - Flybase
  - Maizedb
  - Wormpep
- Mtype: PRT

- Class: STANDARD
- Id: 12S1_ARATH
- Seqlen: 472
- Ac
  - Entry:
- Mod
  - Date: 01-APR-1990
    - Rel: 14
    - Type: Created
  - Date: 01-APR-1990
    - Rel: 14
    - Type: Last sequence update
  - Date: 15-DEC-1998
    - Rel: 37
    - Type: Last annotation update
- Descr: 12S SEED STORAGE PROTEIN PRECURSOR
- Species: Arabidopsis thaliana (Mouse-ear cress)
- Org
  - Entry: Eukaryota
  - Entry: Viridiplantae
  - Entry: Embryophyta
  - Entry: Tracheophyta
  - Entry: Spermatophyta
  - Entry: Magnoliophyta
  - Entry: eudicotyledons
  - Entry: Rosidae
  - Entry: eurosids II
  - Entry: Brassicales
  - Entry: Brassicaceae
  - Entry: Arabidopsis
- Ref
  - Num: 1
    - Pos: SEQUENCE FROM N.A
    - Comment
      - Entry:
    - Db:
    - Medlineid:
    - Author
      - Entry: Pang P.P
      - Entry: Pruitt R.E
      - Entry: Meyerowitz E.M
    - Cite: Plant Mol. Biol. 11:805-820(1988)
  - Num: 2
    - Pos: SEQUENCE OF 420-472 FROM N.A
    - Comment
      - Entry:
    - Db:
    - Medlineid:
    - Author
      - Entry: Raynal M
      - Entry: Grellet F
      - Entry: Laudie M
      - Entry: Meyer Y
      - Entry: Cooke R
      - Entry: Delseny M
    - Cite: Submitted (OCT-1992) to the EMBL/GenBank/DDBJ databases

- Embl
  - Sec Id: AAA32777
    Prim Id: M37247
    Status:
  - Sec Id: CAA32493
    Prim Id: X14312
    Status:
  - Sec Id: CAA79005
    Prim Id: Z17590
    Status:
- Interpro
  - Sec Id: -
    Prim Id: IPR000459
- Pfam
  - Status: 1
    Sec Id: Seedstore_11s
    Prim Id: PF00190
- Keyword
  - Entry: Seed storage protein
  - Entry: Multigene family
  - Entry: Signal
- Features
  - Domain
  - Binding
  - Signal
    - From: 1
      To: 24
      Descr: POTENTIAL
  - Chain
    - From: 25
      To: 282
      Descr: ACIDIC CHAIN (BY SIMILARITY)
    - From: 283
      To: 472
      Descr: BASIC CHAIN (BY SIMILARITY)
  - Disulfid
    - From: 112
      To: 289
      Descr: INTERCHAIN (ALPHA-BETA) (POTENTIAL)
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
  - Sec Id: S08509
    Prim Id: S08509
- Mendel
- Hssp
- Prosite
  - Status: 1
    Sec Id: 11S_SEED_STORAGE
    Prim Id: PS00305
- Gene: CRA1
- Prints
  - Sec Id: 11SGLOBULIN
    Prim Id: PR00439
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 12S2_ARATH
  Seqlen: 455
  Ac
  - Entry:

Mod

- Date: 01-APR-1990
  Rel: 14
  Type: Created
- Date: 01-APR-1990
  Rel: 14
  Type: Last sequence update
- Date: 15-DEC-1998
  Rel: 37
  Type: Last annotation update

Descr: 12S SEED STORAGE PROTEIN PRECURSOR

Species: Arabidopsis thaliana (Mouse-ear cress)

Org

- Entry: Eukaryota
- Entry: Viridiplantae
- Entry: Embryophyta
- Entry: Tracheophyta
- Entry: Spermatophyta
- Entry: Magnoliophyta
- Entry: eudicotyledons
- Entry: Rosidae
- Entry: eurosids II
- Entry: Brassicales
- Entry: Brassicaceae
- Entry: Arabidopsis

Ref

- Num: 1
  Pos: SEQUENCE FROM N.A
  Comment
  - Entry:

  Db:
  Medlineid:
  Author
  - Entry: Pang P.P
  - Entry: Pruitt R.E
  - Entry: Meyerowitz E.M

  Cite: Plant Mol. Biol. 11:805-820(1988)
- Num: 2
  Pos: SEQUENCE OF 240-360 FROM N.A
  Comment
  - Entry:

  Db:
  Medlineid:
  Author
  - Entry: Raynal M
  - Entry: Grellet F
  - Entry: Laudie M
  - Entry: Meyer Y
  - Entry: Cooke R
  - Entry: Delseny M

  Cite: Submitted (OCT-1992) to the EMBL/GenBank/DDBJ databases

- Embl
  - Sec Id: AAA32778
    Prim Id: M37248
    Status:
  - Sec Id: CAA32494
    Prim Id: X14313
    Status:
  - Sec Id: CAA79024
    Prim Id: Z17654
    Status:
- Interpro
  - Sec Id: -
    Prim Id: IPR000459
- Pfam
  - Status: 1
    Sec Id: Seedstore_11s
    Prim Id: PF00190
- Keyword
  - Entry: Seed storage protein
  - Entry: Multigene family
  - Entry: Signal
- Features
  - Domain
  - Binding
  - Signal
    From: 1
    To: 24
    Descr: POTENTIAL
  - Chain
    - From: 25
      To: 269
      Descr: ACIDIC CHAIN (BY SIMILARITY)
    - From: 270
      To: 455
      Descr: BASIC CHAIN (BY SIMILARITY)
  - Disulfid
    - From: 106
      To: 276
      Descr: INTERCHAIN (ALPHA-BETA) (POTENTIAL)
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
  - Sec Id: S08510
    Prim Id: S08510
- Mendel
- Hssp
- Prosite
  - Status: 1
    Sec Id: 11S_SEED_STORAGE
    Prim Id: PS00305
- Gene: CRB
- Prints
  - Sec Id: 11SGLOBULIN
    Prim Id: PR00439
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 1431_ENTHI
  Seqlen: 239
- Ac
  - Entry:

Mod

- Date 01-NOV-1995
  Rel 32
  Type Created
- Date 01-NOV-1995
  Rel 32
  Type Last sequence update
- Date 01-NOV-1997
  Rel 35
  Type Last annotation update

Descr 14-3-3 PROTEIN 1 (14-3-3-1)

Species Entamoeba histolytica

Org

- Entry Eukaryota
- Entry Entamoebidae
- Entry Entamoeba

Ref

- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  - Entry
    Db
    Medlineid
    Author
    - Entry Samuelson J
    - Entry Shen P
    - Entry Meckler G
    - Entry Descoteaux S
    - Entry Fu H
    - Entry Lohia A

    Cite Submitted (AUG-1994) to the EMBL/GenBank/DDBJ databases

Embl

- Sec Id AAA80185
  Prim Id U13418
  Status

Interpro

- Sec Id -
  Prim Id IPR000308

Pfam

- Status 1
  Sec Id 14-3-3
  Prim Id PF00244

Keyword

- Entry

Features

Pir

Mendel

Hssp

Prosite

- Status 1
  Sec Id 1433_1
  Prim Id PS00796
- Status 1
  Sec Id 1433_2
  Prim Id PS00797

Gene

Prints

Sec Id 1433ZETA
Prim Id PR00305

Flybase

Maizedb

Wormpep

- Mtype PRT

Class STANDARD

Id 1431_LYCES

Seqlen 249

Ac
- Entry

Mod
- Date 15-JUL-1998
  Rel 36
  Type Created
- Date 15-JUL-1998
  Rel 36
  Type Last sequence update
- Date 15-JUL-1998
  Rel 36
  Type Last annotation update

Descr 14-3-3-LIKE PROTEIN 1

Species Lycopersicon esculentum (Tomato)

Org
- Entry Eukaryota
- Entry Viridiplantae
- Entry Embryophyta
- Entry Tracheophyta
- Entry Spermatophyta
- Entry Magnoliophyta
- Entry eudicotyledons
- Entry Asteridae
- Entry euasterids I
- Entry Solanales
- Entry Solanaceae
- Entry Solanum

Ref
- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  - Entry STRAIN=CV. MONEYMAKER
  - Entry TISSUE=LEAF

  Db
  Medlineid
  Author
  - Entry Roberts M.R
  - Entry Bowles D.J

  Cite Submitted (JAN-1997) to the EMBL/GenBank/DDBJ databases

Embl
- Sec Id CAA65145
  Prim Id X95900
  Status

Interpro
- Sec Id -
  Prim Id IPR000308

Pfam
- Status 1
  Sec Id 14-3-3
  Prim Id PF00244

Keyword
- Entry

Features

Pir

Mendel

Hssp

Prosite
- Status 1
  Sec Id 1433_1
  Prim Id PS00796
- Status 1
  Sec Id 1433_2
  Prim Id PS00797

- Gene: TFT1
- Prints
  - Sec Id: 1433ZETA
  - Prim Id: PR00305
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 1431_MAIZE
- Seqlen: 261
- Ac
  - Entry:
- Mod
  - Date: 01-FEB-1996
    Rel: 33
    Type: Created
  - Date: 01-FEB-1996
    Rel: 33
    Type: Last sequence update
  - Date: 01-OCT-1996
    Rel: 34
    Type: Last annotation update
- Descr: 14-3-3-LIKE PROTEIN GF14-6
- Species: Zea mays (Maize)
- Org
  - Entry: Eukaryota
  - Entry: Viridiplantae
  - Entry: Embryophyta
  - Entry: Tracheophyta
  - Entry: Spermatophyta
  - Entry: Magnoliophyta
  - Entry: Liliopsida
  - Entry: Poales
  - Entry: Poaceae
  - Entry: Zea
- Ref
  - Num: 1
    Pos: SEQUENCE FROM N.A
    Comment
    Db: MEDLINE
    Medlineid: 95148741
    Author
    - Entry: de Vetten N.C
    - Entry: Ferl R.J

    Cite: Plant Physiol. 106:1593-1604(1994)
- Embl
  - Sec Id: AAB33304
    Prim Id: S77133
    Status:
- Interpro
  - Sec Id: -
    Prim Id: IPR000308
- Pfam
  - Status: 1
    Sec Id: 14-3-3
    Prim Id: PF00244
- Keyword
  - Entry:
- Features
- Pir
- Mendel
- Hssp

Prosite
- Status 1
  Sec Id 1433_1
  Prim Id PS00796
- Status 1
  Sec Id 1433_2
  Prim Id PS00797

Gene GRF1

Prints
Sec Id 1433ZETA
Prim Id PR00305

Flybase

Maizedb
Sec Id -
Prim Id 65832

Wormpep

- Mtype PRT
  Class STANDARD
  Id 1431_SCHMA
  Seqlen 252
  Ac
  - Entry
  Mod
  - Date 15-JUL-1998
    Rel 36
    Type Created
  - Date 15-JUL-1998
    Rel 36
    Type Last sequence update
  - Date 15-JUL-1998
    Rel 36
    Type Last annotation update
  Descr 14-3-3 PROTEIN HOMOLOG 1
  Species Schistosoma mansoni (Blood fluke)
  Org
  - Entry Eukaryota
  - Entry Metazoa
  - Entry Platyhelminthes
  - Entry Neodermata
  - Entry Trematoda
  - Entry Digenea
  - Entry Strigeidida
  - Entry Schistosomatoidea
  - Entry Schistosomatidae
  - Entry Schistosoma
  Ref
  - Num 1
    Pos SEQUENCE FROM N.A
    Comment
    - Entry
      Db MEDLINE
      Medlineid 96123403
      Author
      - Entry Schechtman D
      - Entry Ram D
      - Entry Tarrab-Hazdai R
      - Entry Arnon R
      - Entry Schechter I

        Cite Mol. Biochem. Parasitol. 73:275-278(1995)

- Embl
  - Sec Id: AAC46983
    Prim Id: U24281
    Status:
- Interpro
  - Sec Id: -
    Prim Id: IPR000308
- Pfam
  - Status: 1
    Sec Id: 14-3-3
    Prim Id: PF00244
- Keyword
  - Entry:
- Features
  - Domain
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
    From: 133
    To: 149
    Descr: TO THE C-TERMINUS OF ANNEXINS
- Pir
- Mendel
- Hssp
- Prosite
  - Status: 1
    Sec Id: 1433_1
    Prim Id: PS00796
  - Status: 1
    Sec Id: 1433_2
    Prim Id: PS00797
- Gene
- Prints
  - Sec Id: 1433ZETA
    Prim Id: PR00305
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 1432_ENTHI
  Seqlen: 238
  - Ac
    - Entry:
  - Mod
    - Date: 01-NOV-1995
      Rel: 32
      Type: Created
    - Date: 01-NOV-1995
      Rel: 32
      Type: Last sequence update
    - Date: 01-NOV-1997
      Rel: 35
      Type: Last annotation update
  Descr: 14-3-3 PROTEIN 2 (14-3-3-2)
  Species: Entamoeba histolytica
  - Org
    - Entry: Eukaryota
    - Entry: Entamoebidae
    - Entry: Entamoeba

- Ref
  - Num: 1
  - Pos: SEQUENCE FROM N.A
  - Comment
    - Entry:
  - Db:
  - Medlineid:
  - Author
    - Entry: Samuelson J
    - Entry: Shen P
    - Entry: Meckler G
    - Entry: Descoteaux S
    - Entry: Fu H
    - Entry: Lohia A
  - Cite: Submitted (AUG-1994) to the EMBL/GenBank/DDBJ databases
- Embl
  - Sec Id: AAA80186
  - Prim Id: U13419
  - Status:
- Interpro
  - Sec Id: -
  - Prim Id: IPR000308
- Pfam
  - Status: 1
  - Sec Id: 14-3-3
  - Prim Id: PF00244
- Keyword
  - Entry:
- Features
- Pir
- Mendel
- Hssp
- Prosite
  - Status: 1
  - Sec Id: 1433_1
  - Prim Id: PS00796
  - Status: 1
  - Sec Id: 1433_2
  - Prim Id: PS00797
- Gene:
- Prints
  - Sec Id: 1433ZETA
  - Prim Id: PR00305
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 1432_LYCES
- Seqlen: 254
- Ac
  - Entry:
- Mod
  - Date: 15-JUL-1998
    Rel: 36
    Type: Created
  - Date: 15-JUL-1998
    Rel: 36
    Type: Last sequence update
  - Date: 15-JUL-1998
    Rel: 36
    Type: Last annotation update
- Descr: 14-3-3-LIKE PROTEIN 2
- Species: Lycopersicon esculentum (Tomato)

Org
- Entry: Eukaryota
- Entry: Viridiplantae
- Entry: Embryophyta
- Entry: Tracheophyta
- Entry: Spermatophyta
- Entry: Magnoliophyta
- Entry: eudicotyledons
- Entry: Asteridae
- Entry: euasterids I
- Entry: Solanales
- Entry: Solanaceae
- Entry: Solanum

Ref
- Num: 1
- Pos: SEQUENCE FROM N.A
- Comment
  - Entry: STRAIN=CV. MONEYMAKER
  - Entry: TISSUE=LEAF
- Db:
- Medlineid:
- Author
  - Entry: Roberts M.R
  - Entry: Bowles D.J
- Cite: Submitted (JAN-1997) to the EMBL/GenBank/DDBJ databases

Embl
- Sec Id: CAA65146
- Prim Id: X95901
- Status:

Interpro
- Sec Id: -
- Prim Id: IPR000308

Pfam
- Status: 1
- Sec Id: 14-3-3
- Prim Id: PF00244

Keyword
- Entry:

Features

Pir

Mendel

Hssp

Prosite
- Status: 1
- Sec Id: 1433_1
- Prim Id: PS00796
- Status: 1
- Sec Id: 1433_2
- Prim Id: PS00797

Gene: TFT2

Prints
- Sec Id: 1433ZETA
- Prim Id: PR00305

Flybase

Maizedb

Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 1432_MAIZE
- Seqlen: 261

Ac
- Entry:

Mod

- Date 01-JUL-1993
  Rel 26
  Type Created
- Date 01-FEB-1996
  Rel 33
  Type Last sequence update
- Date 01-FEB-1996
  Rel 33
  Type Last annotation update

Descr 14-3-3-LIKE PROTEIN GF14-12

Species Zea mays (Maize)

Org

- Entry Eukaryota
- Entry Viridiplantae
- Entry Embryophyta
- Entry Tracheophyta
- Entry Spermatophyta
- Entry Magnoliophyta
- Entry Liliopsida
- Entry Poales
- Entry Poaceae
- Entry Zea

Ref

- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  Db MEDLINE
  Medlineid 95148741
  Author
  - Entry de Vetten N.C
  - Entry Ferl R.J

  Cite Plant Physiol. 106:1593-1604(1994)
- Num 2
  Pos SEQUENCE OF 14-261 FROM N.A
  Comment
  Db MEDLINE
  Medlineid 93076113
  Author
  - Entry de Vetten N.C
  - Entry Lu G
  - Entry Ferl R.J

  Cite Plant Cell 4:1295-1307(1992)

Embl

- Sec Id -
  Prim Id S77135
  Status NOT_ANNOTATED_CDS
- Sec Id AAA33505
  Prim Id M96856
  Status

Interpro

- Sec Id -
  Prim Id IPR000308

Pfam

- Status 1
  Sec Id 14-3-3
  Prim Id PF00244

Keyword

- Entry

Features

Pir

- Sec Id JQ1680
  Prim Id JQ1680

- Mendel
- Hssp
- Prosite
  - Status: 1
    - Sec Id: 1433_1
    - Prim Id: PS00796
  - Status: 1
    - Sec Id: 1433_2
    - Prim Id: PS00797
- Gene: GRF2
- Prints
  - Sec Id: 1433ZETA
  - Prim Id: PR00305
- Flybase
- Maizedb
  - Sec Id: -
  - Prim Id: 65832
- Wormpep

- Mtype: PRT
- Class: STANDARD
- Id: 1432_SCHMA
- Seqlen: 214
- Ac
  - Entry:
- Mod
  - Date: 15-JUL-1998
    - Rel: 36
    - Type: Created
  - Date: 15-JUL-1998
    - Rel: 36
    - Type: Last sequence update
  - Date: 15-JUL-1998
    - Rel: 36
    - Type: Last annotation update
- Descr: 14-3-3 PROTEIN HOMOLOG 2 (14-3-3-2) (FRAGMENT)
- Species: Schistosoma mansoni (Blood fluke)
- Org
  - Entry: Eukaryota
  - Entry: Metazoa
  - Entry: Platyhelminthes
  - Entry: Neodermata
  - Entry: Trematoda
  - Entry: Digenea
  - Entry: Strigeidida
  - Entry: Schistosomatoidea
  - Entry: Schistosomatidae
  - Entry: Schistosoma
- Ref
  - Num: 1
    - Pos: SEQUENCE FROM N.A
    - Comment
      - Entry:
    - Db:
    - Medlineid:
    - Author
      - Entry: Lee K.W
      - Entry: Thakur A
      - Entry: Karim A.M
      - Entry: Loverde P.T
    - Cite: Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases

- Embl
  - Sec Id: AAB02100
    Prim Id: L78441
    Status:
- Interpro
  - Sec Id: -
    Prim Id: IPR000308
- Pfam
  - Status: 1
    Sec Id: 14-3-3
    Prim Id: PF00244
- Keyword
  - Entry:
- Features
  - Domain
  - Binding
  - Signal
  - Chain
  - Disulfid
  - Non Ter
    - From: 1
      To: 1
  - Mod Res
  - Conflict
  - Carbohyd
  - Np Bind
  - Similar
- Pir
- Mendel
- Hssp
- Prosite
  - Status: 1
    Sec Id: 1433_1
    Prim Id: PS00796
  - Status: 1
    Sec Id: 1433_2
    Prim Id: PS00797
- Gene
- Prints
- Flybase
- Maizedb
- Wormpep

- Mtype: PRT
  Class: STANDARD
  Id: 1433_CAEEL
  Seqlen: 248
  - Ac
    - Entry: P41932
    - Entry: Q21537
  - Mod
    - Date: 01-NOV-1995
      Rel: 32
      Type: Created
    - Date: 01-NOV-1995
      Rel: 32
      Type: Last sequence update
    - Date: 15-DEC-1998
      Rel: 37
      Type: Last annotation update
  - Descr: 14-3-3-LIKE PROTEIN 1
  - Species: Caenorhabditis elegans

Org

- Entry: Eukaryota
- Entry: Metazoa
- Entry: Nematoda
- Entry: Chromadorea
- Entry: Rhabditida
- Entry: Rhabditoidea
- Entry: Rhabditidae
- Entry: Peloderinae
- Entry: Caenorhabditis

Ref

- Num: 1
  - Pos: SEQUENCE FROM N.A
  - Comment
    - Entry:
  - Db: MEDLINE
  - Medlineid: 95011616
  - Author
    - Entry: Wang W
    - Entry: Shakes D.C
  - Cite: Gene 147:215-218(1994)
- Num: 2
  - Pos: SEQUENCE FROM N.A
  - Comment
    - Entry:
  - Db:
  - Medlineid:
  - Author
    - Entry:
  - Cite: Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases

Embl

- Sec Id: AAA61872
  - Prim Id: U05038
  - Status:
- Sec Id: CAA98138
  - Prim Id: Z73910
  - Status:

Interpro

- Sec Id: -
  - Prim Id: IPR000308

Pfam

- Status: 1
  - Sec Id: 14-3-3
  - Prim Id: PF00244

Keyword

- Entry:

Features

- Domain
- Binding
- Signal
- Chain
- Disulfid
- Non Ter
- Mod Res
- Conflict
  - From: 118
    - To: 118
    - Descr: A -> V (IN REF. 2)
- Carbohyd
- Np Bind
- Similar

Pir

Mendel

Hssp

Prosite
- Status 1
  Sec Id 1433_1
  Prim Id PS00796
- Status 1
  Sec Id 1433_2
  Prim Id PS00797

Gene FTT-1 OR M117.2

Prints
  Sec Id 1433ZETA
  Prim Id PR00305

Flybase

Maizedb

Wormpep
  Prim Id M117

- Mtype PRT

Class STANDARD

Id 1433_CANAL

Seqlen 264

Ac
- Entry

Mod
- Date 15-JUL-1998
  Rel 36
  Type Created
- Date 15-JUL-1998
  Rel 36
  Type Last sequence update
- Date 15-JUL-1999
  Rel 38
  Type Last annotation update

Descr 14-3-3 PROTEIN HOMOLOG

Species Candida albicans (Yeast)

Org
- Entry Eukaryota
- Entry Fungi
- Entry Ascomycota
- Entry Saccharomycetes
- Entry Saccharomycetales
- Entry anamorphic Saccharomycetales
- Entry Candida

Ref
- Num 1
  Pos SEQUENCE FROM N.A
  Comment
  - Entry
  Db
  Medlineid
  Author
  - Entry Cognetti D
  - Entry Sturtevant J
  Cite Submitted (DEC-1997) to the EMBL/GenBank/DDBJ databases

Embl
- Sec Id AAB96910
  Prim Id AF038154
  Status

Interpro
- Sec Id -
  Prim Id IPR000308

Pfam
- Status 1
  Sec Id 14-3-3
  Prim Id PF00244

Keyword

Features

Pir

Mendel
Hssp
Prosite
- Status 1
  Sec Id 1433_1
  Prim Id PS00796
- Status 1
  Sec Id 1433_2
  Prim Id PS00797

Gene BMH1 OR BMH
Prints
  Sec Id 1433ZETA
  Prim Id PR00305
Flybase
Maizedb
Wormpep

- Mtype PRT
  Class STANDARD
  Id 1433_CHLRE
  Seqlen 259
  Ac
  - Entry
  Mod
  - Date 01-OCT-1996
    Rel 34
    Type Created
  - Date 01-OCT-1996
    Rel 34
    Type Last sequence update
  - Date 01-OCT-1996
    Rel 34
    Type Last annotation update
  Descr 14-3-3-LIKE PROTEIN
  Species Chlamydomonas reinhardtii
  Org
  - Entry
  Ref
  Embl
  Interpro
  Pfam
  Keyword
  Features
  Pir
  Mendel
  Hssp
  Prosite
  Gene
  Prints
  Flybase
  Maizedb
  Wormpep