

Key to Gene Annotations

- **General organization:** The medial acid-base dyads (ABDs) are “parsed” into “strings that initiate with a dyad. N-term domain is denoted the head, C-term the tail.
- **Highlights:** Yellow, tyrosine; Green, cysteine; Pink, VPV; Gray, ABD in predicted head and tail domains.
- **Colored font:** Red, glycine residues in ABD domains; Bold-faced red, last G residue in head and first G residue in tail; Blue, alanine residues in ABD domains; Green, repeated string domains.
- **Underscores:** Acid-base triads or tetrads.
- **Commentary:** Notes on distinctive gene features (e.g. orthologues, localization patterns) in green font at top of some pages. Predicted homology domains (e.g. PDZ, coiled-coil) are given in Supplement Table 2.
- **Secondary structure:** PSIPRED predictions for a subset of proteins. Yellow, amino acids predicted in β -strand; pink, amino acids predicted in α -helix; no highlight, amino acids predicted in disordered (random coil) domain.

Articulin

Orthologue of AM931979.1 and AM931980.1

MMYPGAYGAPLVGGTIGGGYGGYPTGAVETIGAGYGGYGGYSTGAVETI
GYGGYGGY**G**ATPIMSAPMVTAAPQVQTV

ERVVEVPQLQVQEIVRQVPRVMTQEVVXN**VPV**PQIQTV
EKVVEVPQVQTFETIIPVPQVQVQEVRQVPRVQMQEVRQMPVPQVQTV
EKVIEVPQTQVM
EKV**VPV**PQVVTQEVRQVPRPQTVELTRQ**VPV**PQVQTV
EKTMQVPQVQMV
ERTMPVPQVTVQEVRQVPR**A**MQVEVVR**A****VPV**PQ**A**TVTKRVEVPQVQTV
ERT**VPV**PQLSVQEVV
REVKVVPQEVRP**VPV**PQIQTV
EKVVEVPQVQVV
EKVMPVPQVQVQEIVRQVPR**A**IPVEVVVQ**A**PVPQV**A**TVTKRVEVPQIQTV
EKV**VPV**PQVSIQEVRQVPRVMVQEVVKQVPRVMVQEVVKQ**VPV**PQVQAV
ERVVEVPEVQVV
EKIVEVPQVQTV
EKIVQ

APAPVMAAPMIETFAPAPIMTSSIPMTTMAAPTY**G**AGFGYPAPMVGSISAALP
MTTMAAPTYGAGLGYPAPMVGSISAALPMTTMAAPTYGAGFGYPTTIF"

Karlodinium veneficum AM931979.1

Articulin

Orthologue of AM931978.1 and AM931980.1

MYPGAYGAPLVGGTIGGGYGGYSTGAVETIGAGYGGYGGYSTGAVETIGYGGYG
GYGATPIMSAPMVTAAPQVQTV

ERVVEVPQLQVQEIVRQVPRVMTQEVRNVPVPQIQTV
EKVVEVPQVQTFETIIPVPQVQVQEVRQVPRVQMQEVRQMPVPQVQTV
EKVIEVPQTQVM
EKIVPVPQVVTQEVRQVPRPQTVELTRQVPVPQVQTV
EKTMQVPQVQMV
ERTMPVPQVTVQEVRQVPRAMPVEVVKAVPVPQATVTKRVEVPQVQTV
ERTVPVPQLSVQEVV
REVKVVPQEVRPVPVPQVQTV
EKVVEVPQVQVV
EKVMPVPQVQVQEIVRQVPRAPVEVVVQAPVPQVVTVTKRVEVPQVQTV
EKVVPVPQVSVQEVRQVPRVMVQEVVKQVPVPQVQAV
ERVVEVPEVQVV
EKIVEVPQVQTI
EKIVQ

APAPVMAAPMIETFAPAPIMTSSIPMTTMAAPMVGSISAALPMTTMAAPTYGAGFGYP
APMVGSISAALPMTTMAAPTYGAGFGYPTTIF

Articulin

Orthologue of AM931978.1 and AM931979.1

MYPGTIGAPLMGGYGATYGSPVETFGAPLVGGYGI**G**APVFETFASAPIMSAP
MVTAAPQVQTV

ERVVEVPQLQVQEIVRQVPRVMTQEVRN**VPV**PQIQTV
EKVVEVPQVQTFETIIPVPQVQVQEVRQVPRVQMQEVRQMPVPQVQTV
EKVIEVPQTQVM
EKI**VPV**PQVVTQEVRQVPRPQTVELTRQ**VPV**PQVQTV
EKTMQVPQMCMV
ERTMPVPQVTVQEVRQVPR**AMP**VEVV**KAVPV**PQ**TA**TVTKRVEVPQVQTV
ERT**VPV**PQLSVQEVV
REVVKVVPQEVRP**VPV**PQVQTV
EKVVEVPQVQVV
EKVMPVPQVQVQEIVRQVPR**AI**PVEVV**QA**PVPQVVTVTKRVEVPQVQTV
EKV**VPV**PQVSVQEVRQVPRVMVQEVLKQVPRVMVQEVVKQ**VPV**PQV**QA**V
ERVVEVPQVQVV
EKIVEVPQVQTV
EKIVQ

APAPVMAAPMIETFAPAPIMTSSIPMTTMAAPMFETFAPAPIMASSIPMTTMAAPM
V**G**SISAALPMTTMAPTYGAGFGSPTYGAGFGYPTTIF

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 1 M Y P G T I G A P L M G G Y G A T Y G S P V V E T F G A P L V G G Y G I G A P V F E T F A S A P I M S A P
51 S A P M V T A A P Q V Q T V E R V V E V P Q L Q V Q E I V R Q V P R V M T Q E V V R N V P V P Q I Q
101 T V E K V V E V P Q V Q T F E T I I P V P Q V Q V Q E V V R Q V P R V Q M Q E V V R Q M P V P Q V Q
151 T V E K V I E V P Q T Q V M E K I V P V P Q V V T Q E V V R Q V P R P Q T V E L T R Q V P V P Q V Q
201 T V E K T M Q V P Q M Q M V E R T M P V P Q V T V Q E V V R Q V P R A M P V E V V K A V P V P Q T A
251 T V T K R V E V P Q V Q T V E R T V P V P Q L S V Q E V V R E V V K V V P Q E V V R P V P V P Q V Q
301 T V E K V V E V P Q V Q V V E K V M P V P Q V Q V Q E V I R Q V P R A I P V E V V V Q A P V P Q V V
351 T V T K R V E V P Q V Q T V E K V V P V P Q V S V Q E V V R Q V P R V M V Q E V L K Q V P R V M V Q
401 E V V K Q V P V P Q V Q A V E R V V E V P E V Q V V E K I V E V P Q V Q T V E K I V Q A P A P V M A
451 A P M I E T F A P A P I M T S S I P M T T M A A P M F E T F A P A P I M A S S I P M T T M A A P M V
501 G S I S A A L P M T T M A A P T Y G A G F G S P T Y G A G F G Y P T T I F

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MAYSNRQSPSSPTAREPSVQRQTPVATRQDPPQASPVSSSASRKVEYHTVI

REQPQIETV
ERQVDVPTVEFQ
ERVVEMPQVITH
ERIVEVPEVSQTDIIKQVPRVEY
REVT
KEIALPVYHGI
DKVVEVPQQLV
KEKITEVPQVQYVELIRQVPKPQLQYI
DKHV
EKPVIELQ
ERPVEVPVTT
REEIVEVPRYEHMELV
KEVP
REEIQQVQKQVIRPEVQLV
ERPVEVVLTELH
DRITEVPEVEY
REIIRQVPKVEIRYI
DKEVPITKVEYI
EKVVEVPQIIE
ERIVEVPEVEV
REVIKQVPVAVTEYV
EKEVPKHIMEYY
EKIVDVPTVLNHETVIEVPQVLSMDLQTRVPRIDYQSVH
KEVPKYQIMAQ
ERVVDVPEVIQH
ERPRHVTEVQTVDAVTQVSKPYTEYV
EKQVPRFETQAVEMVVEVPQVL
REEFIQEVPPQVQI

AEAIKQVHQPKFEQAVKQVQVQMEYVERVVQVESKMDREKEYIMQPVIQKETV
YQPIVQRTIFTAPQEPVAVAPPPPPQEPRVVSPPPPPPQEPTVAPLVVTTVTAPPRTT
ATTTITAPPQYMTQAAPQYAAATTITAPPQYMTAPAATQSLGSPQYMTAPAEPAT
RMVQPMVFDALDTNHDGKLTREEFNRGVAQQAAPTAPGAFAPPARTTTVVAP
GIVATGPQPMVAKLPPTTVRAQSVTIGQPSVTSIAPRPATQVVTTAAGAPMSVFD
ALDKNHDGKLSREEFAQLRVG

6

Kryptoperidinium_MMETSP0120_c13725
_g1_i1_g45335

Articulin

...MAMMVAPSVGTTQWYS

REIPQVQVV
ERVVEVPQVQV
KENIVEVPTVL^YQ
ERVVEVPKIQDTILTKQVPRVEVQEVT
KEVVRPVNTIT
ERVVEVPMTLTTEVVHE^{VPV}VQQVDVMKQVPKPTV
REVQKQV^YKPTPQ^Y
RERIVEVPHVMFSEQIVEVPQVQIEELI
KEVPRHEMQVVQ
KEV
EKPV^YQLV
ERVVEVPMTTV
KE^CILEVPQVVNVDVVKQVPRHEVQVVQ
REVPKTEIQ^YV
EKIVEIPQIV^YE
ERIVE^{VPV}VET
REVVKQVPKTEVQYI
DKRVPKKIVQ^YV
EKIVEVPQII^YE
ER^AVEVPQVVNVE^AITSVPRPQTQMVR
KEIPKHIFQ^AQ
EKIVEVPLTLNH
ERPVE^{VPV}TQ^YVDVTVQVPS^AEVQII
DKQ^{VPV}VQIE^A
REQVVEVPQ^AF^YE
ERLVDVPQVQMA^AEVVRQVLVPQVQEI

AKQIPKIVKTEVRQRVVQVPSTLIQETAVEVPKMMMGEVVTQKASAQMQRIV
QNWTQYQQPISREEIVQ^GSKEAVVGGVHEASVIGTRSYAVPPDQIERSERVV
TESRLSQEAPVLTTVNMPPDPQPVYRAAPPQMMYAPPVAQAAPPVITTTTT
TAAPVYLHPTHAAPPTTAYGGAGSGVPMEPVAVAHP



Kryptoperidinium MMETSP0120_c12023
_g1_i1_g37885

Articulin

MTQEELVHVPKIIPQ

ERVVQRHVEQVVEVPITMTQEEVHVPVWVQQARTHHIHVEEIVEVPVPMT
QEELVHVPKVITQQRDHHIHVEEIVDVMVEQRVEEIVHVPVIQTQ
ERIVQNPIEHVVEIPKPQVI
EKTIEVPKIQIE
EKIIKVPKVVTVIDTTVQNQVQTIEIVKPKIIQKIVQRKKPIIQDEIRQVPKIEKQ
MVPIQKVV
EREVQMPQVQIIDEIVEVPVQKQVQVPMVMKVQKVVDIPQVEIVEQDVHVPV
QKRRHVPMV

SVAQKFVDIPQVEFVD

*Kryptoperidinium_MMETSP0120_c320_
g2_i1_g1076*

MGSRAGPMIQEVVREVPGREVITHCTVIQYVPEIV

EKEKVVEIPQFNLYLEEEVVQRLTH

EKIVEVEQEQLHFELLTQESRVEYQEVA

KEIQKPVFEP

REVIVEVPHVLH

KEVLVEVPQVQIIDLLKQVP

KEHRQRKVVHV

EKPVVQY

RERRVEVPAVTV

KEQLVEVPVTQELEVVRQIAGPPQVQEV

EKQVN

KEIFEINETTQEVPIETM

EKVVEFPQVEI

REVVKQVPRIEVQYI

DKEVQKIEYQYV

ERIVEVPHVVYE

ERIIEVPEIEV

REVVRQIPKPMVQYV

EKRIPKQELRYL

ERTVQQPLLLQIEQPVEVPRVEIIETVAQVPRPVPQYV

DKQVQKVVIEP

REIVEEVPVLVQQEQ

AVEIADVQIMDTVTQVSAPRTEYRDKPVPVAVETKVQERIQQVPYVLT KDRLVE

VPQVQYVDIVTEEPSYDIREVLKEVPRYTVEYRHTIEEVSQQIAGPGGAPSPR

LSA

*Kryptoperidinium*_MMETSP0120_c1965
_g1_i1_g5573

...TAPMAMMVAPSVGTSQFYS

REIPQVQVV

ERVVEVPQVQV

KEHVIEVPTVLYQ

ERVVEVPKIQDTILTKQVPRVEVQEVT

KEVVRPVNTIS

ERVVEVPLTLTTEVLHEVPVQQVDVVKQVPKATV

REVQKQVYKPTPQYR

...QVEYV

DRVVEVPQVEYV

DRVVEVPQVEYVDRPIQVPRVEYI

DRPIEVPKVQRI

ERQEVPQVDWVL

EKVMEP

TTMQQVVRPPTYHRAPPVQVAVAPPRTTAMLPPQHYVMA**G**TAPV

...IVEVPQVQVQE VVKPVPKVRVQEVV

REVPK**Y**ETKIV

ERIVDVPQVE**Y**I

DRVVEVPEVR**Y**V

EKLVEVPRVS**Y**QDVVRHVPKIEVQE VVRTVPKVEIQT

RERVVEIPEVQ**Y**V

ERVVEVPEVQIQEVVRHV**Y**PKIEIQ

ERIRHV**Y**PK**Y**ETQLV

ERLVEVPEVQ**Y**V

DRIVEVPQID**Y**QEVV**Y**KPVPKIEVQE VIRQV**Y**PK**Y**ETQIV

EKLVEV**Y**PKIQ**Y**T

ERVVEVPEVQ**Y**V

EKIVE

*Kryptoperidinium*_MMETSP0120_c11415
_g1_i2_g35718

...NWSVLELVLRARMVCMSCALRWLAAEVRRLR

REV

DKMVKQKTEEIVHVPMGQQEQKQIVPIRKAVG

REVQTPQVPIFNETVDVPVQKQVVELD

EREVLQILIHDEVDVPVQKQVVELD

EREVLQILIHDE

...DASLVARALRCIEEAAPRGSMSPDRSRTIDRQKHEDLLKTATHVEHKG
KGGGGP

DKVVLPPGKSRHVHTAVKQIHVPV
ERTVRVPVKKHV
ERQTETKVVNGVKMVPVKKYRTV
KETVLET
KEEVVEGF
REEWKKVKVPT
KEVVKKQVPRTVTRQVPYVEYVP
KEVV
REVRVP
RDVIK

EKHGIRVDKHLGTKVMTIEEDHHYEMRPVKVKTGDVRVMETGYQHGGKSQ
HGRSQFNP

MSQPAASAQVPMEPLA**G**SPMPPQHAEQHFSSMPNAAEEFQL
EPEYY

ERVVEVPRIHVEN
RERIIEVPQPQIV
DRIVEIPQVQEVIKQIP**G**EIETHIIT
KEVPKIEVKQV
ERIVEVPQIE**Y**Q
DR**Y**VEVEEV
REIVRRVPRIEVHEIPI
ERIVRVPKKIIQEIEQPI**Y**RPVPHLVQQRV
EREPI

AKTQVQTLEVVQQVPVPMN**G**GSGMFMPMEPGPGMMQQP

Kryptoperidinium_MMETSP0120_c52233
 _g1_i1_g123926

MAYGGGPAYPMQFDGAQQRAAMGPGPSDAYQGNYPKPFAAHASAATTNF
 TSGGGGFQGGSSGYQQQSSGFQQQFGAQQAYGDYDPSVFESLPKELLQ
 GAVSFEVRDPTLVSERVVETQEHYFAAVPSSMFHSHDARDREAVRAVVKV
 ASGDGVAVA**G**YRPVYMM

EKVVEVPHVIT

RE**Y**

DR**Y**VPKPEVI

RERLVE**A**PKIEV

ERVQQLPPRVQ**Y**

KEQIVEVPEVVIE

ERIIHVPK

REVQ

ERLIEVPKV**T**YV

ERIE**Y**EDVIE**Y**

RE**VPV**

DKIVEVPEIE**Y**RV

KEVEHMVPQ**T**Y**I**QE**Y**FV

DR**Y**

KEMPVTQVQEVQRIEQ**VPV**

SV**G**QQPGMPPPGSSFLVPRGTYQPGHYGFDGSRQG

1 M A Y G G G P A Y P M Q F D **G A Q Q** R A A M G P G P S D A Y Q G N Y P K P F A A H A S A A T T N F T
 51 S G G G G F Q G Q S S G Y Q Q Q S S G F Q Q Q F G A Q Q A Y G D Y D P S V F E S L P K E L L Q G A
 101 **V S F E V R D P T L V S E R V V E T Q E H Y F A A V P S S M F H S H D A R D R E A V R A V V N K V A**
 151 S G D G V A V A G Y R P V Y M M E K V V E V P H V I T R E Y D R Y V P K P E V I R E R L V E A P K I
 201 E V E R V Q Q L P P R V Q Y K E Q I V E V P E V V I E E R I I H V P K R E V Q E R L I E V P K V T Y
 251 V E R I E Y E D V I E Y R E V P V D K I V E V P E I E Y R V K E V E H M V P Q T Y I Q E Y F V D R Y
 301 K E M P V T Q V Q E V Q R I E Q V P V S V G Q Q P G M P P P G S S F L V P R G T Y Q P G H Y G F D G
 351 S R Q G

...IEVQVQ

ERIVEVPQVQVV

EKTVEVPQVQIQEVVRHVHPKIHVQEVRHVHPKIEVQVQ

ERIVEVPQIQTV

EKMVEVPQIQVQEVRHVHPKVHIHEVVKHVHPKIEVQV

KERIVEVPQVEV

REKVVEVPRIEVQEVRHVHPKHIIET

REHIVEIPCVQIQ

ERIVEHPEIHVNEV

TRHVNKVVEVQEVRVKNVTKVDWQGANAAASLGTTFDGAATLPPGSPIHVASV

QTAMPGAPTMGWGGAPSSPVHAPTMRLPQQRVMMEPSGYS

Repeated strings in lower two-thirds of ABD domain. Homologous to string repeats in *Symbiodinium* MMETSP1367_c42055_g1_i1_g70685 (5.39).

...ELAARRPPDFDNWFDPRAFDDRPPGYAGSRPVTEPIAGSPMEPIACSSRQ
VVEPVACSSSPQPDASRSPSSPGSWHYGRNLGYARGELDI

RDSVV
ERSEQFR
REVP
KEVRKRQTVELITEVPKPEVQYV
EKLVEVPQVQYV
EKIVEVPQVQFV
EKIVEVPQVQYQEMVRHVPKIVTQEIVKHVPRVEIQTV
EKIVEVPQVQYV
EKIVEVPQVHIQEVVKHVPRVEVQEIVRHVVKIEVQVQ
ERIVEHRQVQFV
EKIVEVPQVMVQVV
DKHVQKIEVQEIVKHVVKIEVQVQ
ERIVEHPQIQVV
EKLVEVPQVMVQVV
DKHVSKIQVQEVVRHVVKIEVQVQ
ERIVEHRQVQFV
EKIVEVPQVMVQVV
DKHVQKIEVQEIVKHVVKIEVQVQ
ERIVEHPQIQVV
EKLVEVPQVMVQVV
DKHVSKIQVQEVVRHV

*Kryptoperidinium*_MMETSP0120_c72315
_g1_i1_g162326

...GGGAPAA**C**RALTPPAPLSLPMEALVGPLGGGVLPVAVMRVTLPAVVPA
HPPTPRHGAG**C**GGFFV**C**GTPRALTPPPRAPTPRASTVGTSGVALMPPPRVM
TRPLPSDSSSPIRRRSSGVASSPTRSDMSAPRSAKAVVRLSEPVAVVT**C**AAA
GACAMEP**GG**ESEATPVVT

RESQPA**S**QMLTD**Y**QTVI
REKPTIQEVVKEIEVPEFHV**Y**
ERVVEVPVTLIE
ERILEVTH
EREVE**A**VKQVPRVS**Y**QEL**A**IDVQKLVPTPV
ERIVEVPQVLE
ERLVEMPQPY**Y**VD**C**LKEVVQPQ**Y**Q**A**VQRNVFKPVVQLV
EKHVEVP**A**PTIKEELVEIPQHVEVELLRQV**A**RHEVQEV
EKRV
EKPEVNVRHLVKEVPQVEFQ
EKVVEIPRVEI
REVVRQVPKVEVV**Y**I
EKRVPK
REVQWV
EKIVEVPQIV**Y**E
ERIIIEVPEVEVHEVVRHVPVTTVQ**Y**I
DKPVPKFEMR**Y**T
ERQVDVP

TELTLTHEQAVEVPEVSSVEVVTQVARPVVQHVDKQVPRVSLQAEQRFVD**C**PVP
VR**C**EQPVEMPEVQYIDTVRQVLRPSYHIQDKPVASVEIQAVRIVEVPQVFLEER
VVEIPQVQIVEALKQELQPIVQEVIKEVPQYQIEYTERVVEVSSSVLQEINPESQLLQ
STATV**C****S**GRTLPPGAHGPM**L****C**QERVGSEMLPQRPVSWHPHYVQPPALPPPPP
LAPPHLGAPGGVREES**F****C**

...SHLGACLLAIATATPRPPPARARSALRCRTCD
SAATQQAMSTSAVSTGAY
RQEQPMHTGSQLSSSRFANVPPMEPTAMTSMMPHIQEVVREVPGREVI
THSTVIQYVPEIV

EKEKVVVEIPQFNLQYLEEEIVQRVTH
EKVVEIEQEQQFEILTQEARVEYQEVM
KEVQKPVFEP
REVIVEVPHVLH
KEVLVEVPQVQIFELIKQVP
KEHRQRKVVHV
EKPVVQF
RERYVEVPAVTV
KEQLVEVPVAQEMEVVRQVAGPQEVQQV
EKQVN
REVEINEATQEVPFVETM
EKVIEVPQVEV
REIVKQIPKIEVQYI
DKQVPKVEYQYV
EKIVEV

...SELACGASAGSQPMMPAQVMQYGGGGPPQEPTVMRPQTPGTTAI

REMPKVELV
EKVVEVPNVQVQ
EKFIDVPTTL^YQ
ERIVEVPQIQVAELIKQVPKVEVQE^VVKHVA^KPKVTEV^Y
ERVVEVPQVMSV
EKLVEVPQVQQVEVVRQIVRPQVQ^YV
EKQVPKPV^YEV
REKIVEVPQA^TIQEQLVEVPKIE^YVELV
KEVPRPEVQVVQKNV
DKPVVQ^CV
ERIVEVPQVTVQE^VIMEVPQVEIQEVVRQVPKVEVQMV
DKQVKK
REIQ^YI
EKIVEVPQIV^YYE
ERIVEVPKIE^Y
REIKQVPKPKQIQ^YI
DKKVPKHII^YV
EKIVEVPQVV^YYE
ER^AVDVPQVHHVE^AITEVPKPHVQQVQKPIPK^YYEIRA^AQ
ERIVEVPTILRVEQPIEVPQVLVAEVVTQVPRPEIQ^YI
DREIPVIRTEA
REQIVEVPQVQIEELI
KEVPR

MSPMAGGGLQQPVG TAPGASPASSRGFATPVGT SRSGMMMMHTSSLD **GQ**
SETLMEVPQH HFR EKIVEVPTI **CA**

KEVVK **A**VPRVEVHEVI
KEVPKVE **Y**QVT
ERIVEVPEVH **Y**V
EKIVEIPEVHM
REVVKTVPKVEVQEVIKNVPKVKT V FQ
EKIVEVPQVHQV
EKIIEVPQIHVQE VVKHVPKHMIQEIVKHVPRVELQTV
EKIVEVPQVSFV
EKVVEVPQIQVQE VVKHVPKVHVHEVV
REVPKIHTE **Y**
RERIVEIPQVHIQ
ERIVEVPRIEVQEIVKTVPKVEVQIV
DRMVEVPQVQ **Y**V
EKIVEVPQIHQV
ERVIEVPRVHIQEIVKQVPKVEIQEIVHEVPRWETRFV
EKIVEVP

Oxyrrhis marina AM931982

MSVSGKPQFSSSSWFVGQPQYVTAQAPQYMTTQAPQYITSQAPQYLTSQAPQ
 YVTSQAPQYVTSQAPQYVTSQAFEQVSYSTGVVIDTYQTGERNLVSE^{RT}GS^{QL}
 TND^GTIVNIRKTRQ^{VPV}MQVE

ERIVEVPKIEIQTV
 ERIVEIPQIQ^{YV}
 DKIVEVPQIQEVI
 KEVIKIQTQEQ
 REVPRTVIQTV
 EKVVVEVPQVQTV
 ERVVQVPQIQEVVRQVPRVEIQDV^{YY}
EREVQIPRLV^C

^GDGGSRSGRFL^{CL}RSWM^{CR}WR

1 M S V S G K P Q F S S S S W F V G Q P Q Y V T A Q A P Q Y M T T Q A P Q Y I T S Q A P Q Y L T S Q A
 51 P Q Y V T S Q A P Q Y V T S Q A P Q Y V T S Q A F E Q V S Y S T G V V I D T Y Q T G E R N L V S E R I
 101 V T G S Q L T N D G T I V N I R K T R Q V P V M Q V E E R I V E V P K I E I Q T V E R I V E I P Q I
 151 Q Y V D K I V E V P Q I Q E V I K E V I K I Q T Q E Q R E V P R T V I Q T V E K V V E V P Q V Q T V
 201 E R V V Q V P Q I Q E V V R Q V P R V E I Q D V Y Y E R E V Q I P R L V C G D G G S R S G R F L C L
 251 R S W M C R W R

Oxyrrhis marina AM931981

MSVSGQPTTQQYAGQPQFMGTQQFVGQPQFMGTQQFVGQPQYIASQAPQYVTTQAPQYVT
TQAPQYVTTQAPQYVTTQAPQYVTSQPAQFITSQAPQYITSQAVEQQQVSYSTGVVIDTYQT**G**

ERNLVS
ERVTGSQTLTND**G**TIVNIRKTRQ**VPV**MQVE
ERIVEVPKIEIQTV
ERIVEIPQIQ**Y**V
DKIVEVPQIQEVI
KEVVKIETQEVI
REVPRT VIQTV
EKVVEVPQVQTV
ERVVQVPQIQEVVRQVPRVEIQDVQVQ
REVQIPRLVVETVEQV
REIPVPQIV**VPV**EVIQEVVVQQ**AV**QQIV
DRPVQVDQ**PY**QV
ERVV
ERP**Y**TVVQEQEVVRQQIVDVPRIQQRQVTV
ERP**VYY**D

APVDQQYQEQVQY**G**QPQMVQSMAQPQMVQSTMVQQPMMQTVQQPMMQSIQYGAPVQYGA
PMQY GAPVSYGAPVQYGAPVQYGPPGSQGGSVGVRPQP

Articulin

MAAAGM**C**ASGGAC**C**GAGETTVTGERWEVVGKKGKGSYERVTSLKYYVGEHG
SISKEPNVLISGRSPVIGAI**V****C**ML**C**IPVLLGLGYLLIWSERPVGADDVLVPPVAT
EPD**C**VS**G**FDDW

EKSWT

KERQSY**CC**SHFQR**G****C**APLV

REVPKWVVDVN**VPV****N****VPV****K**

KE**VPV**PVTKVQTVQVK**APDP****Y****E**CHDT**ASVDAWSADHQRW****CCY****AASV****G****C**KP

IIV

DKTV**Y**KTVTKI

KE**VPV**PH**Y**IEPKKPKVHIVH

KEVP**Y**EVP**G**TPKIIPVKLPRK**VI****Y**

KDRVQVVPKLIK**VPE****AG**KVQIVHKPVPH**VH****G****HA****I****Y**VK**VPV**

KDS**Y****D****C**DE

GFENWFFGWSSVKKHF**CC**KTEQKG**C**PHTWHGSLHLHSHVHVEGEGHATGRIY
D**C**HAGFSNWQQGWSVSKKKW**CC**SREERG**C**EFY**C**EGDSKMWHAACKDW**CC**
SHFQQG**C**AATLSPLG**C**DAT**C**ELKGHSS**C**QERIHWTSHHFFSGRENS**C**ALAYS
KVQVE**C**DV**C**RAC**C**SIEAAG**C**TVQKVATSDAFD**C**HAAYNNFFRAWSPSKKKW**CC**TI
KRMGCEGESPPSVDPGVGMMWKHVQVNGYWTWQAVASGGGGFVKLPYD**C**QA
GLTHRSTGWSPGKKS**W****CC**QHKQIG**C**MDVSAGAAAGAAAAAAGGAAAGAAGA

Articulin

...TFHTFFPMKKRGGKKAPPPQEQPLAGDHPG**G**IPEVQII

EKVVEVPRVLKQPLEHLVEEIIHVPKVVVH

ERIQHRTVEQFIDIPVPQPQEEIVH**VPV**

KEY**Y**Q

DRHHHVEVEQFVDIP**VPV**QQEDVVQIPVELPQKRVIQQSVEQLIE**VPV**SITQE

EIVHVPRVVYQHRHHHAEEVEQVVDLHVPHHIEETVSVPKVIQQ

ERLVHHTVEQLVE**VPV**QIPQ

EKIVQVPKITVQ

ERQNVVHVEHVIEIPVPQTVEEIVQVPIVQTE

ERLVQNPVEITVEVPRPVIL

EKTIEVPKVQIEETIVKVPKIMQTVQHLLTVQDQIQKIEVVKPQIIQRTVRRKKPII

QENITEVPQITIEH**VPV**

EKVVKKPVDPVQIQ**Y**V

DKIVE**VPV**QKQRH**VPV**IQKVPKIVELPTLE**Y**VDHVVHVPITQHRH**VPV**VQTVK

KHVE**VPV**VKYEDEILHMPVQKHVHVPMVTKVPRTVEVEQVE**Y**VDDHH

VHIPVQTHRHTV

EKRMQRTVEVPQIE**Y**V

DKV**VPV**PVQKHVHVPQV**AAV**

ERHVEVPQIQ**Y**V

DKDLE**VPV**PVHHH**VPV**LSKVSK**AVE****VPV**

KETI

EKVIE**VPV**VNQVDIPQVQTV

EKIVEVPMVQEVQKVVE**VPV**V

GDTLPGQQRS**C**VDLPLVRREAPPEVRMEVVQGPPMPPEYVKGLVTAAPSS

AASHAASTLGGLATPPRTGSFMPPMMGSFAPPMGSGSLTPGSLWANSAGTP

TGSPISRRA

Articulin

MPIARTP**C**ESRGETE**EPC**STSAPVSSARRKVKRSASSRRARSKSRKRDGA
GAEGASE**G**EHIESAQVKDF**C**AKVEVAVQTDNVNLQDLRLMVPE**K**VVEVET
DERRQIIKYIPIERVVTV

EKRVPFV
KEVVRT**VPV**DRIV
EKIVE**VPV****E****G****F****G**
EKIVEVI
KE**VPV**
EKVVV
KEVV
KE**VPV**ERIV
EKIVEV
EKEVI
KE**VPV**
EKIV
EKVVI
KE**VPV**DKTL
EKEVI
KEVI
KE**VPV**
ERIV
EKIVEV
EKEVI
KEVPI
EKIV
EKEVI
KE**VPV**
EKIV
EKIVEVEVI
KE**VPV**QI

ANREVMNEMITDFAEMEPFEEVEEDAQLPEEVHQAVLEAQEIL**G**SGLEDLESF
EEQIVQRLRTETVDHERKRSVIREMVMEALTAWLKNATPETKKALMTWGQGS
KPKKPASWIGAALMKDDETF

Articulin

...

ERIMEVPEVFEDELVKQVPRIQYLEV
EKPIELPVFQAQQRIQELPTLLEQECLVEVPQVQCIELI
REVP
KEVVKTQKLV
ERPVKVVGPVETTLQEELVEVPVVEEVEIIKQVPVYELQQV
EKRVEVPAVELQ
ERIVEVPQTEVAENPVEVPEIEICVIRQVPKYEIHVV
DKEVVKHEIKYI
EKLVEVPHVIYE
ERIEVPQV
ERRELIRHVPVSRVQVV
DKKVPRHMLQVT
EKTVEVPTVITEEVPQETLEVVVVETITEVPKIQMLPVPVEVPKVVNLQIQ
ERLEEVHTTLE
ERVVEVPEPQVVEAFTQVVRPVVEYV
DK

...FGSSSQPALIKGWLRRARNSPRAQRRFSAMAPKLFHVLFALLVLDGLG**G**HK
RLEPNALVDSEEAIADQLVKEAEANV

ERESPAP

KESNSTKKVPVTPKKVPVEVNVS

KEV

KESSTPSK

KEA

KEDGDEDL

EKEILDMENE

T**G**NANKTKEVRSG**C**KAASMSVLAADVLSATLG

...CRKCGHKRGAVGAAVPAGSGAVPAYAAGPPVTMMAPATSVPVTA
 PAVV
 TTYTRAPVTAPAVAQVAVRPQVQLV

EKIVEVPQVEVQ
 EK^YV^YVDVPTTLY^YQ
 EK^VVEVPRVQVA^AEVVKQVPKIEVQEVI
 KEV^AKPVTEV^Y
 ERTVEVTQNL^YV
 EKLVEVPKVRVVDVVKQVPKPMKQV
 EKQVVKPI^YEVV
 EKLVEVPISTIEEQIVEVPKVE^YME^CI
 REVPRPELQKAQKQV
 EKPVMQ^CV
 EK^VVEVPQ^YTIQESIVE^VVPVTETHEIVKQVPSIEVQTV
 DKEVKK
 KEIH^YV
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 ERIIEVPKIE^Y
 REVIKQVPKPEVQ^YI
 DKKVPKHIE^YV
 EKIVEVPQVV^YE
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 EKKIPKIQLRAQ
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 KEERLVEVPQVQVA^AEFIKQVPKQQVQEIPKHIPKVEM^RCV
 EKIQN^VVPVKLMQEV^AVEVPQVLRHEVISQVSQQTEQRVVQ^AAEE^Y
 ERFVN
 RDEVVV

^GEQESQFGGAYEAKVVGVPISPSTASAEMYERGQVSHVKSTEVRRQGTASAAQ
 ASGLASG^CAGYGSAQRAGDLFTALDTNGDGVLSREEMAALRTGPSTQGMRAP
 VTMAAPMAYAGAVGAPAAMGQVTTYTAPPVYAHPMAMGPVTRAPGPPVTYGAA
^CGSGA^CVTYGQG^CQGC^CQGSSASMRGGDLFSALDTNGDGVLSREEMAALQPA
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...RGMGGMGMGYGPGGRGGKGGKGNLPVKVIT

REVPKIEVKQV
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DRLVEV
REV
REVVRRVPRIEV
REIPI
ERVIQVPKKVVQEVEQPVYRPVPHMVKQHV
EREIPVPKPYLQTLLEVVKQVSVP

TTEDGVVIPQSEVTPQAAVQPGQTSPPAQGASPGQPVL SAPVSRTYELPPVP
SGGTPVAPGGTVMYGSGVVSQTQVAGGDLFSKMDKDGSGAISREEF AQAQ
QVGIATMSSPGAAGPLYGPPQSVSMGSMVMGAGFGSTPQGC CAVAYGGAATP
PQAPSSASCIQSQRYPGQVGVVQSMGAMNPYATMSGAPGGAVAYASMPPQ
APCGSGCFTQTQRYASAGPAASGDLFSRLDRDGS GVISREEF EQALRSGVVG
APQSGSSQVATT

1 R G M G G M G M G Y G P G G R G G K G K G K G N L P V K V I T R E V P K I E V K Q V E K I V E I P N
51 I E Y Q D R L V E V R E V R E V V R R V P R I E V R E I P I E R V I Q V P K K V V Q E V E Q P V Y R
101 P V P H M V K Q H V E R E I P V P K P Y L Q T L E V V K Q V S V P T T E D G V V I P Q S E V T P Q A
151 A V Q P G Q T S P P A Q G A S P G Q P V L S A P V S R T Y E L P P V P S G G T P V A P G G T V M Y G
201 S G V V Q S Q T Q V A G G D L F S K M D K D G S G A I S R E E F A Q A Q Q V G I A T M S S P G A A G
251 P L Y G P P Q S V S M G S M V M G A G F G S T P Q G C A V A Y G G A A T P P Q A P S S A S C I Q S Q
301 R Y P G Q V G V V Q S M G A M N P Y A T M S G A P G G A V A Y A S M P P Q A P C G S G C F T Q T Q R
351 Y A S A G P A A S G D L F S R L D R D G S G V I S R E E F E Q A L R S G V V G A P Q S G S S Q V A T
401 T

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...PMVQKVQKEVEVPQIQYEDQVVEVPVTKQVHVPMVETVQKQVEVPQVQY
EDQVVHVPVAKHVHVPMVQTVQKQIEVPQIEYQDEIVEVPVQKQINVPMIQKV

ERSVDVPQIQYEDQVVQVPLAKQVHVPMIQEI
ERTVEVPQVEYVDNVVQVPV
EKQVHVPMIQKI
EKTVEIPQIEYVDNHVHIPVQKTRHVHVHVPV
ERPVEVNVIEL
EKVIDVPVVKQVEVPQVQTI
EKIVEVPFVQVV
EKVVEVPQV

GSTTQGSVREVDVETEPTRQEPAQVVQQVIAGTPYPVEHAAPEVIGASPVPT
QTEDAPVAEPIQTGA

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MIQKVQKIVDVPQIEIEEQVVEVPVAKHVQVPMIQKVQKMVEVPQVEYDDQV
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EKVQRSVEVPQVEFVDEVMQIPVAKQVNVPMSKV
EKVVEIPQVQYV
DKVVPIPVQKQVNVPMTQV
EKIVEIPQVEFVDTHIHIPVQKHRQVPVQVPV
EKPVEVNIETT
EKVIDVPVVKQI AVPQIQTI
EKIVEIPFVQVV
EKVVEVPQV

GSTTQGSVREEHVEGETKREEHPAQVVQQVLAGPSHPVEYVTETEEPPPQ
EPIQTAT

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...PRQMSGPVTKVVTAPPVTKVPSM**G**SPKRKLEYHTILREQATIEEV

EK**A**VEVPRVEVV

EKVEEVPI TLIHESLVEVPQLLQDEIIREVPLPQ**Y**QEVV

KEIELPQFE**A**

RERIEVPMSLIEET**Y**VEVPRVQRLEVIKQVPK**A**MARKMPKHV

EKPVIQV

RERQMPVTT**A**TV

KEEMVEVPKVEVVEMVRE**A**AVMEVQNV

EKLV

ERIEVELVT

KE**V**VPVVLQ**A**IHDNIVEVPQVE**Y**IEVV

KEIPKVEVR**Y**V

EKTVEVPKIE**Y**V

DRIVEVPQIT**Y**E

ERIVEVVQREVREIIKQVPKPVVQ**Y**V

DKKVPKH**Y**NY**Y**DNVQ

EKLAVLKQ**E**V**V**EVPEVH**A**VELLTELPKP**Q****Y**

EKVP

KEIPL**Y**QIEVV

ERIESTPVVL

KEER**A**V**V**DQVENLQVMRQVVVKPVVE**Y**V

EKTVPKVETQ**A**V

EKEVEVPMVLRQEVIVETPQLV**V****A**EVIREMPQEMTQQVV

KEVPKFQME**Y**VNKVLEVKP

ARYE**A**GMMSAPLTLGSSVASGSRYMEPGPVYQPSAGRAVSYGAQSMGSSMM
GLLGSATS

...DPSKATMSHWAAQFQAVSVEGEPQYLAGVGSEVQGAPQVMMQRREQV
VAGQVKQKITEIPTIQEIVEEQEVPEVQYINKIVEVPEIIQQQIEQVVEQVVHVP
QIVEQKRVQHRHVEQFVDVPVAEQAAPQEQVVEIPVPMQEEEEIVHVPKIITQ
TRQVQQPVEQVIEVPVPMMQEEVHVHPKIMTASRVQHQHVEELVEVPIPMT
QEEVVHVPKVMTATRQVQQPVEQVVEVPIHQRREEVVHVHPKVVTHTRTSH
QHVEQTVEVPVPMQEEHVHHPKILTQTRQIQQQQVEQSVEVPVPMTEEVMV
HVPKIMTQTRVRHEHVEQTVEVPIPHTQ

EKVVHVHPKVITQ
ERVTHQPVETT
KEVPIPMVQEQVVHVHPKVQQ
ERQHFFHVEEMVDIPIEQPVETQVHVPLVSKV
ERIVHNPVDLVVEVPRPVVV
EKSVQVPMIQVE
EKTVRVPKVLNTVVDTVVQHQQVQNVQIVRPTVVHKVVQRRKPIVQ
ERVQHVPKVMVQHVPVQKVI
EKQVHIPQIQYVDEFVDVPV

ARHRRVAMPAAQAPIVHKHVHVPMVQKVQKTVEVPQIQFEDQIVEVPVAKQVN
VPVVQTMQKQVEVPQMQYEDQIVHVPVQKQVHVPMVQKVQ

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MLASNC^CKN^CGNTFVSDSNF^CRRC^CGEKREVYGASAAPMAT^GSIRTAAPAATL
MQEPNTQVQIEEQIVEVPTILYQ

EKVVEVPKVQETILLKQVPRVEVQEVTKEVVRPI^YEVT
EKIVEVPHTLVQ
ERIEEVPKVEFVPLVKQVPKTQVREVQKQV^GVPH^YQP
RERIVEVPTTMVAEQMVEVPQVQI
KELL
KEVPRPEVKIVAKPV
EKPVNE^YV
ERIVEVPHTTIQE^CIIEVPKVEVQEIVKQVPRPEVQVV
DRQV
EKLEVQ^YV
EKLVEVPQII^YE
EKIVEVPTVEVREVIKQVSKPEVK^YI
EKRVPKKVIK^YV
EKIVEVPQVI^YE
ERPIEVELIHVEAVTQVPRPHVQFVK
KEVPKVVLQPQ
ERIVEVPLTLTQ
ERPVEVPQVHV^AELMVQVPKPQVEMV
DKEVPKIIVQ^AE
EKIVEVPQVF^YEE^CLKHVPQVQV^AEVI
KEVPH^YQKQVVQKHVPKIVETKVV
EKVVPVPV

NLVRE^TAVEVPQVVQHEVIRQKAS^GAMQQRIIQTGWQYQRSARKEEVVSGIAEAQQGG
IYDAPVASVRRVSLPTNEVEVSAVITEQSVSHEAPVTTVN^LKPGMFAQTQVAPA
PVTTYGTAIVGTMTGTAAAQSIFRAAQAC^PORTAWGTGDVVSASAAATSGAAAS
TMR^TTRA^EATA^CAN^CGNVLMPSDF^CRK^CGAARSVPGAPDPFLVTEPVAPGYG
TINTQSTLPTQPL

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...EQ

ERISHQHVEEVVEVPVPMTE

EKVVHVPKVVQQ

ERQHFFHVEETVDVPIEQHVEQVVHVPVVSRG

ERIVHNPVDLEVEVPRPIVV

EK**C**IEVPKIQVE

ERTVRVPKIVNTVV

DRVVRHSMETVEVVKPTVVHKVVQRKKPVVQ

ERVQHVPKVFIIQQVQV

EKVV

EKPVEVPQIQFVDEVVDVPMVKHRHVPTVVKQQ

KEVEVPQVE**Y**VD

Repeated strings in lower two-thirds of ABD domain. Homologous to string repeats in *Kryptoperidinium* MMETSP0120_c71290_g1_i1_G159348 (5.17).

...SDSPGQWFKENPLLQSWNASAQDALAGALP**G**QTSQLQTPRSSMSIPRAD
MSQEVLEVPQTLFREKVVVEVPTI**C**TQELVRAVPKVEVHEII

KEVPKIE**Y**QVT
ERVVEVPEVR**Y**V
EKIVEVPEVHTQEIVRTVPKIEVQEVEVKTVPKVQVQVQ
ERIEVPKV**C**HV
ERVVEVPQVQVQEVRVTVPRLEVQEVI
REVPKLEVRTV
EKLVEVPQV**A**QV
DKFVEVPQVQVQEVLRHVPKLEVHEVI
KEVPKIQME**Y**
RERIVEVPQLHLQ
ERLVEVPKIEVQEVRHVPKVEVRTV
DRVVEVPQVQ**Y**V
EKLVEVPKVEIQ
ERLHHVPKIHTQVV
EKLVEVPQVQFV
EKFVEVPQVQIQEVVRHVPKIETQEIVRTVPKVEIRTV
EKIVEVPQVQFV
EKIVEVPEVRIQEVRHVPKIEVQEVEVKHVPKIEVQLQ
ERIVEHPQVHQV
EKIVEVPQTVIQTV
DRHVPKIEVQEIVRHVPKIDVKIQ
ERIVEHPQIHQV
EKVVEVPQTVVQEVRHVPKIQVQEIVKHIPKIEVQVQ
EKIEVPQVQVV
EKIVEVPQKQVHEVIKHVPKLQIQEVVRHVPKVEVQVQ
ERIVEVPHIQTI
EKVVEVPKVQIQEVVRHVKKPVIET
RERIVEIPVVQVQ
ERVVEHPEIHVQEVI
KEIPKVMEVQEIVKNVQKVD

WT**G**ADVSAAMSGGSIEAPTLPPATPNAASTRYVPMSAAGDCTPSGRSVATVDIN
AATLTPSTLNTQLNKLPTLKASCELEAPVIRLPEEASRQGSEVPVFEVPRDRVTV
TSAELQSRPSGSPAVAAASLVAGFCVACGSSYPPGSSFCRQCGRQVDGPSQP
QPTALPQPGMAPGMPMHPPQTAYGAPCTGMGGRPSMPTVFGSAPTQYPHGL
RPGVDPFSFR

MPGMPGMPGMPGMPGMPGMPGMPGGPEMSGRYGMDLHMPGMPGLPGMP
GMPGMPGFGLDVGLPGMRGFGISGMGMGGFDGMGFGLYGNGVSGIGPG
GLAMD LGGYGGGGIQDILGEGGMELVNGGDLDAALGLLNGPGVDANGEIQ
LPPGVEPGTKEYNDFLQQLEAQGLIEI

REEIVEVPQVVVE
ERVVHVPGLKKQIQ
ERLIEVPKVDWV
ERVEYEDYVEY
REVPV
DKIIEVPEIEYKIKQVDQTVPQTYIQEHFV
DKF
REVPVTQVQET
ERVEHVPVMVP
KDWQPPNYE

ALGMKSSPANAPAQITVT

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EKPVEVTVIETLQKIVDVPVVKQVEVPQVQTI

EKIVEIPLVQTV

EKVVEVPMMGQTTQGTT

REVDIPLAPR

REEHPAQVVQQVIM

GEDHPVQVMAAQAASIAQPGISVTATPAAASASAAPAAIQEPVQTSA

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...SQVWNGFSDRRGLQIHAPHPASVVPPTWRPPWNVGSQTAAAGFHKAYS
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TVQGGIGPQEHVTQMTSNVSNVSRQYSAPQEAYAMRPAAMPQVREVEVT
QLATAAALEDSIKQCQVVIR

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REREKVVEVPTISLSENMVEVPVSLVH
EKVVEVPQVQVVEIMKQLPKVEYQ
ERVVQV
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REVAKVQSKQLAKHFARPVVQW
KESI
REVPQTTL
REQLQEVPLRTVEVL
REVPKPEVQRV
EKQVEVPTIQLQ
ERLVDVPCVEQ
REEPTEVAKVEI
REVVRQVPKVEVKYV
EKKVPKQVIQYV
EKTVEVPQVVYEEVIVEVPEVEI
KEMVRQVPVPVVQYI
DKPVPRRSLRAV
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KEFPKVLLEA
REVAVEVPVVL
REERVVEVPQVQSVPLVRQRLCPSMQAV
DKPVPYIETRVQ
ERQVQVPYVLQEEAIAEVPQQQLVELR
REELQPMAEVVLKQVPKYQVEYV
EKVVEVQ

SQVAQETSPSRESRSQA^{GG}TAVAVSTISTQKSASLHREIQGLGLGQFHESTCPK
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EKTDEE
EKVV
EKVIEE