```
A0A0L0BTL1    1  MNLSPMMYRLNHAQGQKLPMAAEQQQQQSQQHPQQQQQPTTTPQTTPPTPTQSQQQQIIP
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1   61  SHILLQQQLDAASANTNSSNSNTTISNSHSITHHSNNEPVFLNNFTEQESNTPLALRQAS
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  121  NASSTSNSALLHTIHQHHSHTHQAPPTPPNHSRLIHSTAANNLVITNAAAAAALVAASAA
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  181  AAAANQVDVLSSNSSVEENLKLLKTAIKSEPLPHDLNSASNSNHNNHNNVNNLSSNKEDL
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  241  ISLAACSALANVVIPASSANAVASSSTSVSSTMSSAKANVLKNVSTSALSNAITQVALAA
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  301  GAGNGGSVSASGGVVSAAAGSVAGAGGSGGEALTSSNGSMVFVPSKRARMELREEWISTP
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  361  SPGSVPSTAPLSPSSASQNHMYGANMSNGYASPMSAGSYDPFSPNGKTGRDDLSPSSSLN
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  421  GFSTSDASDVKKIKKGPAPRLQEELCLVCGDRASGYHYNALTCEGCKGFFRRSVTKNAVY
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  481  CCKFGHACEMDMYMRRKCQECRLKKCLAVGMRPECVVPENQCAMKRREKKAQKEKDKIQT
4OZT_E           ............................................................
4OZR_E           ............................................................
consensus>50     ............................................................


A0A0L0BTL1  541  SVCATEIKKEILDLMTCEPPSHPTCPLLPEDILAKCQARNIPPLSYNQLAVIYKLIWYQD
4OZT_E        1  .......................................VKPISPEQEELIHRLVYFQN
4OZR_E        1  .......................................KPISPEQEELIHRLVYFQN
consensus>50     .......................................ikPiSp#QeelIhrL!y%Q#


A0A0L0BTL1  601  GYEQPSEEDLKRIMSSPDENESQHDVSFRHITEITILTVQLIVEFAKGLPAFTKIPQEDQ
4OZT_E       21  EYEQPSDEDLKRISNTPSEGEDQSDLNFRHITEITILTVQLIVEFAKRLPGFDKLLREDQ
4OZR_E       20  EYEQPSDED...............FRHITEITILTVQLIVEFAKRLPGFDKLLREDQ
consensus>50     eYEQPS#EDlkri...p.e.e.q.d..FRHITEITILTVQLIVEFAKrLPgFdKllrEDQ
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A0A0L0BTL1   661  ITLLKACSSEVMMLRMARRYDHNSDSIFFANNRSYTRDSYKMAGMADNIEDLLHFCRQMY
4OZT_E        81  IALLKACSSEVMMLRMARRYDVGSDSILFANNQPYTRDSYSLAGMGETVDDLLRFCRQMY
4OZR_E        62  IALLKACSSEVMMLRMARRYDVGSDSIL.........................ATVDDLLRFCRQMY
consensus>50     IaLLKACSSEVMMLRMARRYDvgSDSIlfann..ytrdsy.magm.dt!#DLLrFCRQMY


A0A0L0BTL1   721  SMKVDNVEYALLTAIVIFSDRPGLEEAELVEAIQSYYIDTLRIYILNRHCGDPMSLVFFA
4OZT_E       141  GMKVDNAEYALLTAIVIFSERPSLIEGWKVEKIQEIYLEALKVYVDNRR..KPASGTIFA
4OZR_E       104  GMKVDNAEYALLTAIVIFSERPSLIEGWKVEKIQEIYLEALKVYVDNRR..KPRSGTIFA
consensus>50     gMKVDNaEYALLTAIVIFS#RPsLiEgwkVEkIQeiYl#aLk!Y!dNRr..kP.SgtiFA


A0A0L0BTL1   781  KLLSILTELRTLGNQNAEMCFSLKLKNRKLPKFLEEIWDVHAIPPSVQSHIQATQAEKAA
4OZT_E       199  KLLSVLTELRTLGNLNSEMCFSLKLKNKKLPPFLAEIWDV....................
4OZR_E       162  KLLSVLTELRTLGNLNSEMCFSLKLKNKKLPPFLAE........................
consensus>50     KLLS!LTELRTLGNlNsEMCFSLKLKNkKLPpFLaEiwdv....................


A0A0L0BTL1   841  QEAQATTSAISAAATSSSSINTSMATSSSSSLSPSAASTPNGGAVDYVGTDMSMSLVQSD
4OZT_E            ............................................................
4OZR_E            ............................................................
consensus>50     ............................................................


A0A0L0BTL1   901  NA
4OZT_E            ..
4OZR_E            ..
consensus>50     ..
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