02a-Large-Alignment

March 26, 2025

1 Test 02: Large Alignment

1.0.1 Overview

This notebook demonstrates iterative alignment of a large testcase.

For this demonstration, testcase 1a0cA_1ubpC from PREFAB is used as it contains 50 biological sequences.

Expected runtime: ~ 30 seconds or less

1.0.2 Context

This notebook is intended to test the following requirements of MAli:

Requirement 4.1 - Can align structural benchmarking test cases containing as many as 30 sequences. - The notebook demonstrates alignment of a testcase of 50 sequences.

Requirement 4.2 - Can align a structural benchmarking test case of at least 20 sequences within 60 seconds. - The testcase of 50 sequences is aligned in less than 10 seconds.

1.0.3 Installing Prerequisites

[1]: !pip install biopython

Requirement already satisfied: biopython in

c:\users\pdmoo\appdata\local\programs\python\python310\lib\site-packages (1.85)
Requirement already satisfied: numpy in

c:\users\pdmoo\appdata\local\programs\python\python310\lib\site-packages (from biopython) (1.26.2)

Imports

```
[2]: import os
  import shutil
  import subprocess
  import time
  from presentation_helper import PresentationHelper
```

MAli v1.31

```
[3]: ALIGNER_NAME = "MAli-v1.31"
ALIGNER_PATH = "MAli-v1.31/MAli.exe"
```

```
OUTPUT_FOLDER = "data/output"
```

```
[4]: # creating empty output folder
if os.path.exists(OUTPUT_FOLDER):
    shutil.rmtree(OUTPUT_FOLDER)
os.makedirs(OUTPUT_FOLDER)
```

Testcase

```
[5]: TESTCASE_NAME = "1a0cA_1ubpC"
INPUT_FILEPATH = f"data/input/{TESTCASE_NAME}"
OUTPUT_FILEPATH = f"data/output/{TESTCASE_NAME}"
```

Viewing Testcase

```
[6]: presenter = PresentationHelper()
```

```
[7]: presenter.present_unaligned_fasta(INPUT_FILEPATH)
```

Displaying Sequences from data/input/1a0cA_1ubpC:

>1a0cA

NKYFENVSKIKYEGPKSNNPYSFKFYNPEEVIDGKTMEEHLRFSIAYWHTFTADGTDQFGKATMQRPWNHYTDPMDIAKA RVEAAFEFFDKINAPYFCFHDRDIAPEGDTLRETNKNLDTIVAMIKDYLKTSKTKVLWGTANLFSNPRFVHGASTSCNAD VFAYSAAQVKKALEITKELGGENYVFWGGREGYETLLNTDMEFELDNFARFLHMAVDYAKEIGFEGQFLIEPKPKEPTKH QYDFDVANVLAFLRKYDLDKYFKVNIEANHATLAFHDFQHELRYARINGVLGSIDANTGDMLLGWDTDQFPTDIRMTTLA MYEVIKMGGFDKGGLNFDAKVRRASFEPEDLFLGHIAGMDAFAKGFKVAYKLVKDRVFDKFIEERYASYKDGIGADIVSG KADFRSLEKYALERSQIVNKSGRQELLESILNQYLFA

>13|1a0cA|gi|37526188

HRYFERVNRISYEGRQSNNPLAFRHYNPEEIILGKKMKDHLRFAVCYWHNFCWDGTDMFGSGAFERFWQKGGDALELAKL KADVAFEFFYKLNIPFYCFHDIDVAPEGCSLKEYIYNLGVMSDILADKQAETGVKLLWGTANCFTHPRYAAGASTNPDLN IFAYASAQVCQVMQMTKKLGGENYVLWGGREGYESLLNTDLRQEREQIGRFMQMVVDYKYKIGFQGTLLIEPKPQEPTKH QYDYDVATVYGFLKQFGLENEIKVNIEANHATLAGHSFQHEVATAIALGILGSIDANRGDAQLGWDTDQFPNSVEENSLV MYEILKAGGFTTGGLNFDAKVRRQSIDIDDLFYGHIGAIDTMALSLKSAVKILVDGKLDEYVAQRYSGWNSELGRDILEG KMTLDEVAHYAETLVQEPKHRSGQQELLENLINRYIY

>55|1a0cA|gi|139830

NEYWQGVDQIKYIGHQDKKSGLGFQYYNPDEVIGGKKMRDWLRFSVAYWHTFDQRLVDPFGDGTAQRPYDHITDPMDLAL AKVDAAFEFYHKLGVDYLCFHDRDLAPEGDTLRETNRNLDKVIDKIVDYQKQTGMKVLWNTSNMFTNPRFVAGAATSPDA DVFAYAAAQLKHSLEIGKRVGAENYVFWGGREGYESLWNTNMKLEQEHAAKFFHMAKDYANEIGFDAQMLLEPKPKEPST HQYDFDAATTIAFMKEYDLDKDFKLNLEGNHANLAGHTYQHEIRVAREANLLGSLDANQGDKLIGWDIDEFPSDLYEATA AMYEVVENGSIGPRGGLNFDAKPRRSSFAANDLFYGHIVGIDTFAAGLRVALKMKQDGFLEKLVADRYSSYQSGVGAEIE AGTADFKSLESYAIDKPQSELIAATSSDPLEEVKDTINHYI

>24|1a0cA|gi|3334499

YFDNISTIAYEGPASKNPLAFKFYNPEEKVGDKTMEEHLRFSVAYWHTFTGDGSDPFGAGNMIRPWNKYSGMDLAKARVE AAFEFFEKLNIPFFCFHDVDIAPEGETLKETYKNLDIIVDMIEEYMKTSKTKLLWNTANLFTHPRFVHGAATSCNADVFA YAAAKVKKGLEIAKRLGAENYVFWGGREGYETLLNTDMKLELDNLARFLHMAVDYAKEIGFDGQFLIEPKPKEPTKHQYD FDVATALAFLQTYGLKDYFKFNIEANHATLAGHTFEHELRVARIHGMLGSVDANQGDMLLGWDTDEFPTDLYSTTLAMYE ILKNGGLGRGGLNFDAKVRRGSFEPEDLFYAHIAGMDSFAVGLKVAHRLIEDRVFDEFIEERYKSYTEGIGREIVEGTAD FHKLEAHALQLGEIQNQSGRQERLKTLLNQYL

>28|1a0cA|gi|13474199

FFGDIQKIKYEGPDSTNPLAYRFYNPDEVVAGKRLEDHLRFAVAYWHSFAWPGGDPFGGQTFDRPWFPKAGGIDTMELAK LKADVAFEMFSLLGAPYFCFHDADVRPEGKDFSESAARLDEIADYFAGKMKQTGVKLLWGTANLFSHRRFMSGAATNPDP DVFAYAAATVKSCIDVTKRLKGENYVLWGGREGYETLLNTDLAREQEQAGRFLSLVVDYKHKIGFKGTILIEPKPQEPTK HQYDYDVATVYGFLKRFGLEKEVKLNIEQGHAILAGHSFEHELALANALGVFGSIDMNRNDYQSGWDTDQFPNNVPEMAL AYYQILQAGGFKTGGTNFDAKLRRQSLDPQDLLIGHIGGMDACARGLKAAARMVEDKALSAPLAERYAGWNSAEGKAMLS GKRTLEDIAERVVKKKIEPQPRSGRQELLENIVNRYV

>2|1a0cA|gi|32470780

KHGLSLTHAEADHRAKARDFVKQMIDFGGKYNAPAIIGSMQGKWGGDISRDQALAFLREALDELGPYAAQYNVPLFYEPL NRYETNLLRTVDEGVEFCKTLSTDNIKLLADLFHMNIEEADLAAAIRAGKGYVGHIHFVDSNRQ

>23|1a0cA|gi|34395196

EFFPGIPKIKYEGPSSKNPLAYKWYNAEEVILGKKMKDWMRFSVAFWHTFRGTGGDPFGAPTKSWPWEDGTNSLDMAKRR MRAHFEFMEKLGVDRWCFHDRDIAPDGKTLTETNKNLDEIVELAKKLQEETNIKPLWGTAQLFMHPRYMHGAATSPEVKV YAYGAAQVKKALEVTHYLGGENYVFWGGREGYQTLLNTDMKRELDHLASFLQAAVDYKKKIGFNGTLLIEPKPQEPTKHQ YDWDVATTFSFLQKYGLTGEFKINVECNHATLSGHSCHHELETARINGLLGNIDANTGDPQIGWDTDQFLTDISEATLVM SSVVKNGGLAPGGFNFDAKLRRESTDVEDMFLAHISGMDTLARGLRNVVKLIEDGSLDELVRKRYQSFDSEIGALIEAGK GDFETLEKKVLEWGEPIVPSGKQELAEILFQAAL

>3|1a0cA|gi|32034802

SNYFDKIEKVKYEGADSTNPFAFKHYNPNEVILGKTMAEHLRLAVCYWHTFCWTGNDMFGVGSLDRSWQKTGDLLEGAKQ KAEIAFEFFQKLGIPYYCFHDVDIAPEGNSYKEYVHNFHTMVDILEKKQAETGVKLLWGTANCFTNPRYMSGASTNPNPE VFSWAASQVFNAMNATKRLGGENYVLWGGREGYETLLNTDLKREREQIGRFMQMVVEHKHKIGFNGTLLIEPKPQEPTKH QYDYDVATVYGFLKQFGLEKEIKVNIEANHATLAGHTFQHEIATAAALDIFGSIDANRGDPQLGWDTDQFPNSVEENTLV MYEILKAGGFTTGGFNFDAKIRRQSTDPYDLFHGHIGAIDVLALSLKRAAKMIEDKTLQGIVDQRYAGWNGDLGQQILAG KASLEDLAKIVESKALDPKPVSGQQEYLENLVNNYIY

>29|1a0cA|gi|22984909

YFEHIPEIRYEGPQSDNPLAYRHYDKSKKVLGKTLEEHLRIAVCYWHTFVWPGVDIFGQGTFRRPWQQAGDAMERARQKA DSAFEFFSKLGTPYYTFHDTDVSPEGSSLKEYSENFLRITDYLARKQEDTGIKLLWGTANLFSHPRYAAGAATSPDPEVF AFAATQVRHALDATQRLGGDNYVLWGGREGYDTLLNTDLMRERDQLARFLHMVVDHAHKIGFKGSLLIEPKPQEPTKHQY DYDVATVHGFLLQHGLEKEIRVNIEANHATLAGHSFHHEIATAYALGIFGSVDANRGDPQNGWDTDQFPNSVEELTLAFY EILKHGGFTTGGMNFDSKVRRQSVDPEDLFYGHIGAIDNLALAVERAAVLIENDRLDQFKRQRYAGWDADFGRKILSGDY SLSTLATDALARGLNPQHASGHQELMENIVNQAIYS

>39|1a0cA|gi|6136455

NHFESANKVLYEGKDSKNPLAFKYYNPEEVVGGKTMKDQLRFSVAYWHQFTADGTDPFGAATMQRSWDRYDGMDLAKARV EAAFQLFETLNVPFFAFHDRDIAPEGSTLQETNKNLDVIVTMIKEYMQTSNVKLLWNTANMFTNPRFVHGAATSCNADVF RYAAAQVKKGLETAKELGAENYVFWGGREGYETLLNTNLQLELDNLARFMHMAVDYATEIGYTGQFLIEPKPKEPTTHQY DTDAATTISFLRQYGLDKYFKLNLEANHATLAGHTFEHELRVARVQGLLGSVDANQGDPLLGWDTDEFPTDLYSTTLAMY EILQNGGLGSGGLNFDAKVRRGSFEQDDLLYAHVAGMDAFARGLKVAHKLVEDRVFENVINERYSSFKEGIGLEIVEGKA NFHTLEQYAFKNPNIANKSGRQERLKSILNQYI

>53|1a0cA|gi|6136456

YFENVPKVQYEGKNAKSKYAFRHYNPEEIIMGKPMKDHLRFSVAFWHTMTEDGSDPFGDGTYQRNWEGSTPMETAKNRVD AFFEILEKLGAEYFCFHDVDIAPQGDSLKEFLENIDVMTDYIKGKMDKTGVKLLWNTANMFTHPTFVNGAATTNNADVYS MAAAQVKKGLDVSKKLNGENYVFWGGREGYENLLNTDMNFELDNLARFYQMVIDYAQKIDYHPQFLIEPKPKEPTKHQYD YDAATAMAFIQKYNLEDSFKLNLEANHATLAGHTFEHELNVAKNYNALGSLDANQGDLLLGWDTDEFPTDIYTATLAMYE VLDFGGIAPGGLNFDAKVRRTSFAMDDLILAHIAGMDTYARGLRAAAKMKDDNFFEQIIANRYESFSSGIGKQIVENKED LESLTNYALSLNGVENKSGHIEHLKSLLNDYL

>22|1a0cA|gi|15615320

FFNDVEKVQYEGPRSTNPYAFKYYNPEEIVAGKTMAEHLRFSIAYWHTFVGDGSDPFGVGTMQRPWDQYSGMDLAKARVE AAFEFFEKLNVPYFCFHDVDVAPEGETLADTYKNLDEIVLMIKDYMKTSKTKLLWNTANMFTHPRWIHGAATSPNADVYA YAAAKVKKGLEIGKELGAENYVFWGGREGYETLLNTNMKLELDNLARFFHMAVDYAKEIGFDAQFLIEPKPKEPTKHQYD FDVATGIAFLKTHGLDEHFKFNIEANHATLAGHTFEHELHLARIHDMLGSVDANQGDTLLGWDTDEFPTDLYTTTLAMYE ILKNDGLGKGGLNFDAKVRRGSFEANDLFHAHIAGMDSFAIGLKVANQLLEDRVLEDVIDNRYKSYQSGIGQKIANNDTN LKELEAYALSLGEIKHSSGQQERIKATLNQYL

>43|1a0cA|gi|21233522

EYFPGIGKIGFEGRDSDNPLAFKVYDANKQVAGKTMAEHLRFAVAYWHSFCGNGADPFGPGTRAYPWDVGNTALARAEAK SDAAFEFFTKLGVPYYCFHDIDLAPDADDIGEYENNLKHMVGIAKQRQADTGVKLLWGTANLFSHPRYMNGASTNPDFNV VARAAVQVKAAIDATVELGGENYVFWGGREGYACLHNTQMKREQDNMARFLTLARDYGRAIGFKGNFLIEPKPMEPMKHQ YDFDSATVIGFLRQHGLDQDFKLNIEANHATLSGHSFEHDLQVASDAGLLGSIDANRGNPQNGWDTDQFPTDLYDTVGAM LVVLRQGGLAPGGLNFDAKVRRESSDPQDLFLAHIGGMDAFARGLEVADALLTSSPLETWRAQRYASFDSGAGADFANGT STLADLATYAAGKGEPTQLSGRQEAYENLINQYL

>8|1a0cA|gi|16766946

YFDQLDRVRYEGPQSTNPLAFRHYNPDELVLGKRMEDHLRFAACYWHTFCWNGADMFGVGAFNRPWQQPGEALELAKRKA DVAFEFFHKLNVPFYCFHDVDVSPEGASLKEYKNNFAQMVDVLAAKQEQSGVKLLWGTANCFTNPRYGAGAATNPDPEVF SWAATQVVTAMNATHKLGGENYVLWGGREGYETLLNTDLRQEREQIGRFMQMVVEHKHKMGFQGTLLIEPKPQEPTKHQY DYDVATVYGFLKQFGLEKEIKVNIEANHATLAGHSFHHEIATAIALGIFGSVDANRGDAQLGWDTDQFPISVEENALVMY EILKAGGFTTGGLNFDAKVRRQSTDKYDLFYGHIGAMDTMALSLKIAARMVEDGELDKRVAKRYAGWNGELGQQILKGQL SLGELAQYAEQHNLAPVHQSGHQELLENLVNRYLF

>54|1a0cA|gi|139832

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>42|1a0cA|gi|32472084

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>30|1a0cA|gi|15890502

FFGDIAKIKYEGPDSTNPLAFRHYNPDEIVGGKRMEDHLRFAVAYWHTFTWPGGDPFGGQTFQRPWFEDTMQAAKLKADV AFEFFSLLGSPFYCFHDADVRPEGKNFAENTKNLNEIVDYFAQKQADTGVKLLWGTANLFSNRRFMSGAATNPDPDVFAF SAATVKTCMDATKTLGGANYVLWGGREGYETLLNTDLSRELDQLGRFLNLVVEYKYKIGFEGTILIEPKPQEPTKHQYDY DVATVYAFLQKNGLEKEVKVNIEQGHAILAGHSFEHELAMANAFGIFGSIDMNRNDYQSGWDTDQFPNNVPEMALAYYHV LAGGGFKNGGTNFDSKLRRQSLDPQDLLIGHIGGMDCCARGLKAAAKMIEDGALSKPLSERYAKWDSPEAQKMLRGELKL EEIAALVERDDINPEPKPGRQEYLENVVNRYV

>18|1a0cA|gi|1175034

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>31|1a0cA|gi|28870180

YFPTVDKVIYEGPDSDSPLAFRHYDADKRVLGKPMREHLRMAACYWHTFVWPGADMFGVGTFKRPWQRAGDPMELAIGKA EAAFEFFSKLGIDYYSFHDTDVAPEGSSIKEYRNNFAQMVDRLERHQEQSGIKLLWGTANCFSNPRFAAGAASNPDPEVF AYAGAQVFSAMNATQRLKGSNYVLWGGREGYETLLNTDLKREREQLGRFMRMVVEHKHKIGFKGDLLIEPKPQEPTKHQY DYDSATVFGFLHQYGLQDEIKVNIEANHATLAGHSFHHEIATAVSLGIFGSIDANRGDPQNGWDTDQFPNSVEEMTLATY EILKAGGFTHGGYNFDSKVRRQSLDDVDLFHGHVAAMDVLALSLERAAAMVQNDKLQQFKDQRYAGWQQPFGQSVLSGGF SLASLAEHAFANELNPQAVSGRQELLEGVVNRFIYT

>59|1a0cA|gi|23026381

EYFPGVGKIEFEGPDSDNPFAFKYYDENRVVAGKTLKEHFKFSVCYWHTFCGTGADPFGPGPLQFPWNKADTADARARLK MDAAFEFITKLGAPYYCFHDIDLVEEGASRQETQRRVQTIVEYAKEKQAASGVKLLWGTANLFSNPRYMNGASTNPDFNV VAYAGAQLKDALDATIALDGENYVFWGGREGYMSLLNTDMGREQEHLARFLTMARDYARGQGFTGNFFIEPKPMEPSKHQ YDFDSETVIGFLRHHGLDKDFKLNIETNHATLANHTMDHEMTVAANAGMLGSIDANRGDYQNAWDTDQFPYNINETVEMM MVVLRAGGFQGGGVNFDAKRRNSTDLEDTFHGHIGGMDTFARALIIADRIISESKIESMRKERYASFDSGAGKQFETGA LDLAALAEIGNAIGEPALTSGKQEYYENIINQYI

>51|1a0cA|gi|17433696

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>37|1a0cA|gi|29375173

YFPKIEKIKYEGTNTKNMYAFRHYNSEEIIMGKTMKEHLRFAVAYWHTMTQDGSDPFGKAVNKRSWLGESPMETAKKRVI AFFEILEKLDVEYFCFHDIDIAPEGNSLKEFFSNIDEITDLIKEKMDETGIKLLWNTANMFSNPRYVNGAASTNNANVYA IAAAQVKKGLDVSKKLGGENYVFWGGREGYETLLNTDMKFEQDNIARLFKMAIFYGEKIGHKPQFLIEPKPKEPSKHQYD FDAATTMAFILKYGLEKDFKLNLEANHATLAGHTFEHELNVARNYGALGSIDANQGDVLLGWDTDEFPTNVYDVTLAMYE ILENGGIEPGGINFDSKVRRSSFEMEDLLLAHIAGMDTFARGLKSAMKLKEDRFFEDLKEQRYSSFKKGIGAKIISGKEN LESLTNYALKNDEPIIESSHIEYVKNILNDYLY

>35|1a0cA|gi|23100574

YFNEIGTIQYEGAKSDNPFAFKFYNPTESINGQTMEEHLRFGVAYWHTFTEDLSDPFGNGTAIRPWDKFNGMDLAKARVE AAFEFFEKLQIPYFCFHDVDIAPEGSSLKESNENLDTVVAMIKEYMKDSKTKLLWNTVNNFTHPRFVHGAASSNNADVFA YAAAKVKKGLEVGKELGAENYVFWGGREGYETLLNTNMKLELDNLARFFHMAKDYANEIDFNAQFLIEPKPKEPTSHQYD FDVASGYAFLQNYDLQDTFKFNIEANHATLAGHTFEHELHYARIHNMLGSVDANQGHPLLGWDTDEFPSDIYTTTLAMYE ILKNDGLGKGGLNFDAKVRRGSFTAEDLFHAHIAGMDSFAIGLKVAQRLMDDRVLENVVEDRYRSFNQGIGRDIVSGKTD FHQLEKHALTLSEITQPSGQLEVIKNKINQYLLT

>48|1a0cA|gi|7161893

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>36|1a0cA|gi|6136454

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>1ubpC

MKINRQQYAESYGPTVGDEVRLADTDLWIEVEKDYTTYGDEVNFGGGKVLREGMGENGTYTRTENVLDLLLTNALILDYT GIYKADIGVKDGYIVGIGKGGNPDIMDGVTPNMIVGTATEVIAAEGKIVTAGGIDTHVHFINPDQVDVALANGITTLFGG GTGPAEGSKATTVTPGPWNIEKMLKSTEGLPINVGILGKGHGSSIAPIMEQIDAGAAGLXIHEDWGATPASIDRSLTVAD EADVQVAIHSDTLNEAGFLEDTLRAINGRVIHSFHVEGAGGGHAPDIMAMAGHPNVLPSSTNPTRPFTVNTIDEHLDMLM VCHHLKQNIPEDVAFADSRIRPETIAAEDILHDLGIISMMSTDALAMGRAGEMVLRTWQTADKMKKQRGPLAEEKNGSDN FRLKRYVSKYTINPAIAQGIAHEVGSIEEGKFADLVLWEPKFFGVKADRVIKGGIIAYAQIGDPSASIPTPQPVMGRRMY GTVGDLIHDTNITFMSKSSIQQGVPAKLGLKRRIGTVKNCRNIGKKDMKWNDVTTDIDINPETYEVKVDGEVLTCEPVKE LPMAQRYFLF

>36|1ubpC|gi|22297549

RIDRQTYAETYGPTVGDRLRLADTDLIIEIEHDYTHYGDEVKFGGGKVIRDGMGQSPIANAEGAVDVVITNAVILDWWGV VKADVGIKDGKIYKIGKAGNPYTQEGVDIIIGPGTEAIAGEGMILTAGGIDAHIHFICPQQIATAIAAGITTMIGGGTGP ATGTNATTCTPGPWNIYRMLQAADAFPVNLGFLGKGNSSQPQGLIEQVQAGVVGLKLHEDWGSTPNAIDTCLSVAEDYDI QVAIHTDTLNESGFVEDTIAAFKNRTIHAYHTEGAGGGHAPDIIKLCGQANVLPSSTNPTRPYTVNTLDEHLDMLMVCHH LDPSIPEDVAFAESRIRRETIAAEDILHDLGAFSIISSDSQAMGRVGESIIRTWQTAHKMKVQRGHLRDPQRPGVDHDNF RARRYVAKYTINPAITHGIAEYVGSVEVGKIADLCLWKPAFFGVKPELVIKGGVIAYAQMGDANASIPTPQPVHMQPMFA SYGGCRTTTSVTFMSQAGIANNIPEQLGLQKTVLPVGHIRQLRKADLKLNDYLPHIEVDPETYEVRADGELLTCEPATVL PLAQRYFLF

>12|1ubpC|gi|27462198

KISRKEYVSMYGPTTGDKVRLGDTDLILEVEHDCTTYGEEIKFGGGKTIRDGMAQTNSPSSHELDLVLTNALIVDYTGIY KADIGIKNGKIHGIGKAGNKDMQDGVCNNLCVGPATEALAAEGLIVTAGGIDTHIHFISPQQIPTAFASGITTMIGGGTG PADGTNATTITPGRWNLKTMLRASEEYAMNLGYLGKGNVSYEPSLVDQLEAGAIGFKIHEDWGSTPAAIYHCLNVADKYD VQVAIHTDTLNEAGCVEDTLQAIAGRTIHTFHTEGAGGGHAPDVIKMSGEFNILPASTNPTIPFTVNTEAEHMDMLMVCH HLDKNIKEDVQFADSRIRPQTIAAEDKLHDMGIFSITSSDSQAMGRVGEVITRTWQTADKNKKEFGRLPEEKGDNDNFRI KRYISKYTINPAIAHGISEYVGSVEVGKFADLVLWSPAFFGIKPNMIIKGGFIALSQMGDANASIPTPQPVYYREMFGHH GKAKFDTNITFVSQVAYDNGIKEELGLQRVVLPVKNCRNITKKDLKFNDVTAHIEVNPETYKVKVDGKEVTSKAADKISL AQLYNLF

>86|1ubpC|gi|33519971

LNKKIYAEMYGPTVGDSVRLADTDLWVKIEKDFTIYGEEVKFGGGKVIRDGMGQGQMNNKECVDLVLTNAIIIDYWGVIK ADIGINNGRIVGIGKSGNPDIQPGVTIYIGPGTEVISAEGKIVTAGGIDVHVHFICPQQIEEALSSGMTTLIGGGTGPTT GSNATTCTPGVWFISRMLQAADALPINIGFTGKGSSALPESLIEQIHAGAIGLKVHEDWGATPATIDCCLNVADNFDVQV SIHTDTLNESGFLETTISAIKNRTIHAYHVEGAGGGHSPDIIRICELNNVLPSSTNPTLPYTINTVDEHLDMMMVCHNLN CNLPEDIAFAESRIRRETIAAEDILHDLGALSMISSDSQAMGRIGETILRTWQTAHKMKLQRGSLLGDDQYNDNTRIKRY IAKYTINPAITHGIAHEVGSVEIGKLADLVLWSPVFFGVKPELIVKGGMIVSSVMGDPNASIPTPQPVYYRLMFGSYDLA KCATRMTFISQSSYDVGVVENLRLNSLIGISKRCRNIKKDHMINNNLKPVMEVDPQTYEVRANGELLVCEPVSVVPMSQR YFLF

>6|1ubpC|gi|19310948

KISRHAYADTYGPTTGDRVRLADTDLWIKVERDLTTYGDEVKFGGGKVIRDGMGQSQLSRDQVMDLAITNALIVDHWGII KADVGIKDGRIAAIGKAGNPDTQPNVNINIWRPATEVIDGEGKIFTAGGIDSHVHFICPQLVEEAANLGGDYPDWLAGTG PATGTNATTCTPGSWNIGRMLQAADAFPINMGFLGKGNASLPQSLEEQVRAGVLGLKLHEDWGTTPAAIDNCLSVAERFD VQVTIHTDTLNESGFVEDTIAAFKDRTIHTYHTEGAGGGHAPDIIKACGEANVLPSSTNPTTPFTANTVDEHLDMLMVCH HLDPSIPEDVAFAESRIRRETISAEDALHDMGALSMHGSDSQAMGRVGEVILRTWQSASVMKRDRGTLPEDKGDHDNFRI KRYIAKYTHNPAITHGIAHEIGSIEVGKLADLVLWKPAFFGVKPSLVLKGGVIVTAPMGDPNASIPTPQPVHYRPMFGSFGGSRTASCVSFVSQAGLDEGIGEKLGLQKRLVAVKNIRGLRKSDLIHNNALPRIEVDPQNYQVRADGQLLWFEPSKVLPMAQRYFLF

>7|1ubpC|gi|23000615

LDRGSYAAMFGPTVGDRVRLADTELIIEVEQDYTTYGEEVKFGGGKVIRDGMGQSQHSRQEGAVDTVITNALIVDHWGIV KADVGIKDGRICAIGKAGNPDVQPNVDIIIGPGTEAIAGEGRILTAGGIDAHIHWICPQQIEDALHSGITTMLGGGTGPA EGTNATTCTPGPWHIARMLQAAEGLPMNMGFFGKGNASRPQGLEEQLRAGACGLKLHEDWGTTPSAIDVCLSVAEKWDVQ VAIHTDTLNESGFVENTTAAFKGRTIHAFHTEGAGGGHAPDIIKLCGEANVLPSSTNPTRPFTRNTLDEHLDMLMVCHHL DSRIPEDVAFAESRIRRETIAAEDILHDLGAFSMIASDSQAMGRVGEVIIRTWQTADKMKRQRGALPEERGQNDNQRVKR YIAKYTINPAITHGIAHYVGSVAVGKLADLVLWKPMFFGVKPDLVLKCGTIASAAMGDPNASIPTPQPVHYRPMFGAFGG ALTHSAVNFVSQAGLDGEIAQQFGLRKTLLPVVGCRTIGKADMVHNSATPHMEVDPETYEVRADGRLLTCEPATVLPLAQ RYFLF

>16|1ubpC|gi|586164

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>2|1ubpC|gi|15673051

MTLLIKNGRLVDPKSNRDEVLDILIENDKIVKIGLNLSVENAEIVDARGLVVAPGLIDVHVHFREPGQTHKETIHSGAKS

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>42|1ubpC|gi|22976226

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>19|1ubpC|gi|23027998

KISRRAYADMYGPTVGDKVRLADTALWIQVEKDFTIYGEEVKFGGGKVIRDGMGQSQATSDKTPDTVITNALILDHWGIV KADVAIKNGRISAIGKAGNPDIQPGVTIIVGPCTEVIAGEGQILTAGAIDSHIHFICPQQIDEALMSGTTTMIGGGTGPA TGTNATTCTPGKWHIGKMLQAGESFAMNLGFLGKGNASLPGGLNEQLEAGALGLKLHEDWGTTPASIDNCLTVAENYDVQ VAIHTDTLNESGFVEDTLAAFKGRTIHTYHTEGAGGGHAPDIIKACGSSNVLPSSTNPTRPYTVNTVDEHLDMLMVCHHL DPNIPEDVAFADSRIRKETIAAEDILHDLGAFSMIASDSQAMGRVGEVICRTWQTAHKMKVQRGLLPEDEGTGADNFRAK RYIAKYTINPALAHGVAHEVGSIEVGKLADIILWKPAFFGVKPSLIIKGGAIAAAPMGDPNASIPTPQPVHYRKMFGAYG QACKQTSVSFVSQAAIDASISDYFQLERRLVAVKNCRSVTKADMVHNAYQPHMEVDPETYEVRADGQLLTCEPAEELPMA ORYFLF

>25|1ubpC|gi|13474128

RISRAAYAQMYGPTVGDKVRLADTELFIEVEKDLTIHGEEVKFGGGKVIRDGMGQSQVSRAQGAVDTVITNALVVDAGAG IFKADIGLKDGRIAAIGKAGNPDTQDGVTIIIGPGTEIIAGEGKILTAGGFDAHIHFICPQQIEEALMSGITTMLGGGTG PAHGTLATTCTPGPWHMARMIQSFDAFPMNIGLSGKGNASLPAALEEMVLGGACSLKLHEDWGTTPAAIDCCLSVADDYD VQVMIHTDTLNESGFVENTVAAIKGRTIHAFHTEGAGGGHAPDIIKVCGLPNVIPSSTNPTRPYTVNTLAEHLDMLMVCH HLSPSIPEDIAFAESRIRKETIAAEDILHDIGAFSIISSDSQAMGRVGEVAIRTWQTADKMKRQRGALPQETGDNDNFRV RRYIAKYTINPAIAHGLSKDIGSIAVGKRADLVLWNPAFFGVKPDMVLVGGMIAAAPMGDPNASIPTPQPMHYRPMFGAY GKARTNSSVTFVSKAALESGLHGRLGVEKQFVAVENTRGGIGKHSMVLNDATPHVEVDPETYEVRADGELLTCEPATVLP MAQRYFLF

>43|1ubpC|gi|16080717

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>80|1ubpC|gi|33864500

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>39|1ubpC|gi|15800666

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>53|1ubpC|gi|15220459

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>85|1ubpC|gi|17402589

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>44|1ubpC|gi|32265907

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>72|1ubpC|gi|25990713

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>50|1ubpC|gi|22960876

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>3|1ubpC|gi|23020534

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TGPADGTNATTCTPGAFNIRKMLEAAEDFPVNLGFLGKGNASFETPLIEQIEAGAIGLKLHEDWGTTPKAIDTCLKVADL
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CHHLDSKVKEDVAFADSRIRPETIAAEDILHDMGVFSMMSSDSQAMGRVGEVIIRTWQTAHKMKLQRGALPGEKSGCDNI
RAKRYLAKYTINPAITHGISQYVGSLEKGKIADLVLWKPAMFGVKPEMIIKGGFIIAGRMGDANASIPTPQPVIYKNMFG
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PMAQRYFLF

>81|1ubpC|gi|465008

IHRKEYANKYGPTTGDKIRLGDTNLLAEIEKDYALYGDECVFGGGKVIRDGMGQSCGHPPAISLDTVITNAVIIDYTGII KADIGIKDGLIASIGKAGNPDIMNGVFSNMIIGANTEVIAGEGLIVTAGAIDCHVHYICPQLVYEAISSGITTLVGGGTG PAAGTRATTCTPSPTQMRLMLQSTDDLPLNFGFTGKGSSSKPDELHEIIKAGAMGLKLHEDWGSTPAAIDNCLTIAEHHD IQINIHTDTLNEAGFVEHSIAAFKGRTIHTYHSEGAGGGHAPDIIKVCGIKNVLPSSTNPTRPLTSNTIDEHLDMLMVCH HLDREIPEDLAFAHSRIRKKTIAAEDVLNDIGAISIISSDSQAMGRVGEVISRTWQTADKMKAQTGPLKCDSSDNDNFRI RRYIAKYTINPAIANGFSQYVGSVEVGKLADLVMWKPSFFGTKPEMVIKGGMVAWADIGDPNASIPTPEPVKMRPMYGTL GKAGGALSIAFVSKAALDQRVNVLYGLNKRVEAVSNVRKLTKLDMKLNDALPEITVDPESYTVKADGKLLCVSEATTVPL SRNYFLF

>83|1ubpC|gi|26989564

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YFLF

>21|1ubpC|gi|15612817

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>13|1ubpC|gi|10336584

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>70|1ubpC|gi|12313641

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Performing Alignment

```
[8]: ALIGNMENT_COMMAND = f"{ALIGNER_PATH} -input {INPUT_FILEPATH} -output

→{OUTPUT_FILEPATH}"

print(f"CLI command to be run: '{ALIGNMENT_COMMAND}'")
```

CLI command to be run: 'MAli-v1.31/MAli.exe -input data/input/1a0cA_1ubpC -output data/output/1a0cA_1ubpC'

```
[9]: start_time = time.perf_counter()
    subprocess.run(ALIGNMENT_COMMAND)
    end_time = time.perf_counter()

    time_in_milliseconds = (end_time - start_time) * 1000
    time_in_milliseconds_rounded = round(time_in_milliseconds, 0)
    time_in_seconds = time_in_milliseconds_rounded / 1000
```

```
print(f"Performed alignment of {TESTCASE_NAME} in {time_in_seconds} seconds")
```

Performed alignment of 1a0cA_1ubpC in 5.278 seconds

Viewing Alignment Produced by MAli

```
[10]: ALIGNMENT_FILEPATH = OUTPUT_FILEPATH + ".faa" presenter.present_interleaved_aligned_fasta(ALIGNMENT_FILEPATH)
```

Displaying interleaved alignment from 'data/output/1a0cA_1ubpC.faa:

```
1a0cA
              -----N
13|1a0cA|gi|375 -----
55|1a0cA|gi|139 -----NEYWQGVDQIKYIGH
24|1a0cA|gi|333 -----
28|1a0cA|gi|134 ------FGDIQKIKYEGPDSTNPLAYRFYNPDEVVAGKR
2|1a0cA|gi|3247 -----
23|1a0cA|gi|343 -----EF-FP-GIPKIKYE-GPSSKNPL-A------
3|1a0cA|gi|3203 -----S
29|1a0cA|gi|229 -----
39|1a0cA|gi|613 -----N-HFESANKVLYEGKDSKNPLAFKYYNPEEVVGG
53|1a0cA|gi|613 -----
22|1a0cA|gi|156 -----FF-NDVEKVQYEGPRSTNPYAFKYYNPEEIVAGKT
43|1a0cA|gi|212 -----EYF-PGIGKIGFEGRDSDNPLAFKVYDANKQVAGKT
8|1a0cA|gi|1676 -----YFDQLDRVR
54|1a0cA|gi|139 -----YFDINKVNYEGPKSNNAFSFKYYNPE
42|1a0cA|gi|324 -----F-PDVPVIQYEGPQSDNPLAFRWYNPDEVIEGKT
30|1a0cA|gi|158 -----FFGDIAKIKYEGPDSTNPLAFRHYNPDEIVGGKR
18|1a0cA|gi|117 -----EFFPEI-PKVQFEGKESTNPLAFKFYDPEEIIDGKPLKD
31|1a0cA|gi|288 -----YFPTVDKV-IYEGPDSDSPLAFRHYDADKRVLGKPMREHLR
59|1a0cA|gi|230 -----
51|1a0cA|gi|174 ------
37|1a0cA|gi|293 -----
35|1a0cA|gi|231 -----YFN-EI-GTIQYEGAKSDNPFAFKFYNPTESINGQTMEE
48|1a0cA|gi|716 -EYFPQIQKIK-FEGKDSKNPLAFHYY-DAEKEVMGKKMKDWLRFAMAWWHTLCAEGADQ
36|1a0cA|gi|613 -----
         -----MKI-NRQQYAESYGPTVGD-EVR-----LAD
1ubpC
36|1ubpC|gi|222 R-----ID-RQTYAETYGPTVGDR-LRLADTDLII------
12|1ubpC|gi|274 -----KI-SRKEYVSMYGPTTGD-KVRLGDTDLI----------
86|1ubpC|gi|335 -----
6|1ubpC|gi|1931 -----KI-SRHAYADTYGPTTGD-RVRLADTDLW------
7|1ubpC|gi|2300 -----LD-RGSYAAMFGPTVGDR-VRLADTELII-----
16|1ubpC|gi|586 ------
2|1ubpC|gi|1567 ------MTLLIK-NGRLVDPKSNRDEVLDILIENDKIVKIGLNLS
42|1ubpC|gi|229 ------
19|1ubpC|gi|230 -----KIS-RRAYADMYGPTVGDK-VRLADTALWIQVEKDFTIYGEEVKFGGGKVIR
25|1ubpC|gi|134 R-----IS-RAAYAQMYGPTVGDK-VRLADTELFI------
43|1ubpC|gi|160 -----KMSREEYAELFGPTTGDKIRLGD
```

80 1ubpC gi 338	RID-RQAYAETYGPTTGDR-IRLADTELIL
39 1ubpC gi 158	I-SRQAYADMFGPTTGD-KIRLADTELW
53 1ubpC gi 152	IS-REKYANMYGPTTGDK-LRLGDTNLYA
85 1ubpC gi 174	VD-HEKYASMYGPTTGDK-IRLGDTDLFA
44 1ubpC gi 322	
72 1ubpC gi 259	KI-SRQTYVEMMGPTTGD-RVRLADTELFIEIEKDFTTYGEEVKFGGGKVI
50 1 ubpC gi 229	
3 1ubpC gi 2302	
81 1ubpC gi 465	
83 1ubpC gi 269	
21 1ubpC gi 156	KL-TRAQHASLYGPTVGD-KVRLADTDLL
	KISRQAYADMFGPTTGDC-VRLADTELFL
	IS-RATYADMFGPTTGDK-VRLADTDLII
1 0	
1a0cA	KYFENVSKIKYEGPKSNNPYSFKFYNPEEVIDGKTMEEHLRFSIA-YWHTFTADGTDQ
13 1a0cA gi 375	
24 1a0cA gi 333	QDKKSGLGFQYYNPDEVIGGKKMRDWLRFSVAYWHTFDQRLVDPF-GDGTAQRPYDHI
28 1a0cA gi 134	LEDHLRFAVAYWHSFAWPGGDPFGGQTFDRPWFPKAGGIDTMELAKLKADVAFEMFSLLG
2 1a0cA gi 3247	
23 1a0cA gi 343	
3 1a0cA gi 3203	NYFDKIEKVKYEGADSTNPFAFKHYNPNEVILGKTMAEHLRLAVC-YWHTFCWTGNDM
29 1a0cA gi 229	
	KTMKDQLRFSVAYWHQFTADGTDPFGAATMQRSWDRYDGMDLAKA-RVEAAFQLFETL
53 1a0cA gi 613	YFEN-VPKVQYEGKNAK
	MAEHLRFSIAYWHTFVGDGSDPFGVGTMQRPWDQYSGMD-LAKAR-VEAAFEFFEKLN
_	MAEHLRFAVAYWHSFCGNGADPFGPGTRAYPWDVGNTALARAEAK-SDAAFEFFTKLG
	YEGPQSTNPLAFRHYNPDELVLGKRMEDHLRFAACYWHTFCWNGA-DMFGVGAFNRPW
-	EKLGNHSMSELLRFSVAYWHTFTADLSDPFGVGVAERDWDSLDEM-EKAKARVEAIFE
_	MKDHMRFSIVYWHTFRGTGADPFGPGTAVRPWDNGSESVENAQKR-AVVAFELFTKLQ
•	MEDHLRFAVAYWHTFTWPGGDPFGGQTFQRPWFEDTMQAAKLKADVAFEFFSLLG
	HLKFSVAFWHTFVNEGRDPFGDPTADRPWNRYTDPMDKAFARVDA-LFEFCEKLNIEY
	MAACYWHTFVWPGADMFGVGTFKRPWQRAGDPMELAIGKAEAAFE-FFSKLGIDYYSF
51 1a0cA g1 200	
37 1a0cA g1 174	EFFSNIGKIQYQGPKSTDPLSFKYYNPE-EVINGKTMREHL
0	HLRFGVAYWHTFTEDLSDPFGNGTAIRPWDKFN-GMDLAKARVEA-AFEFFEKLQIPY
	FGGGTKSFPWNEGTDAIEIAKQKVDAGFEIMQKLGIPYYCFHDVD-LVSEGNSIEEYE
	FFRNIGMIEYEGADSENPYAFKYYNPDEFVGG-KTMKEHLRFAVA
1ubpC	TDLWIEVEKDYTTYGDEVNFGGGKVLREGMGENGTYTRTENVLDL-LLTNALILDYTG
	LEVEHDCTT
	LNKK-IYAEMYGPTVGD
/ 1ubpC gi 2300	EVEQDYTTYGEEVK-FGGGKVIRDGMG
16 1ubpC gi 586	MSRKQ
2 1ubpC gi 1567	VENAEIVDARGLVVAPGLID
42 1ubpC gi 229	

19 1ubpC gi 230	DGMGQ
25 1ubpC gi 134	DGMGQ
	TDLWIEVEKDFTVYGEEMIFGGGKTIRDGMGQNGRITGKDGALDL-VITNVVLLDYTG
80 1ubpC gi 338	
39 1ubpC gi 158	
53 1ubpC gi 152	
85 1ubpC gi 174	
44 1ubpC gi 322	
	RDGMGQSQRNHADVMDTVITNAVIVDHWGIVKADVGLKDGKIAAI-GKAGNPDIQPNV
50 1ubpC gi 229	
3 1ubpC gi 2302	KISGKD
81 1ubpC gi 465	IHRKEYANKYGPTTGDKIRLGDTNLLAEIEKDYALYGDECV-FGGGKVIRDGMG
	LE-IEKDYTVYGDEC
	QVEQDFTIYGEEVK-FGGGKVIRDGMG
70 1ubpC gi 123	
1a0cA	FGKATMQRPWNHYTDPMDIAKARV
13 1a0cA g1 375	HRYFERVNRISYEGRQSNNPLAFRHYNPEEIILGKKMKDHLR-FAVCYWH
55 1a0cA g1 139	TDPMDLALAKVDAAFEFYHKLGVDYLCFHDRDLAPEGDTLRETNRNLDKV-IDKIVDY
	YFDNISTIAYEGPASKNPLAFKFYNPEEK-VGDKTME
28 1a0cA g1 134	APYFCFHDADVRPEGKDFSESAARLDEIADYFAGKMKQTGVKLLWGTANL-FSHRRFM
2 1a0cA g1 3247	
•	
	FGVGSLDRSWQKTGDLLEGAKQKAEIAFEFFQKLGIPYYCFHDVDIAPEG-NSYKEYV
_	YFEHIPEIRYEGPQSDNPLAYRHYDKSKKVLGKT-LEEHLRI
_	NVPFFAFHDRDIAPEGSTLQETNKNLDVIVTMIKEYMQTSNVKLLWNTAN-MFTNPRF
_	SKYAFRHYNPEEIIMGKPMKDHLRFSVAFWHTMTEDGSDPFGDGTYQRNW-EGSTPMETA
•	VPYFCFHDVDVAPEGETLADTYKNLDEIVLMIKDYMKTSKTKLLWNTANM-F
	VPYYCFHDIDLAPDADDIGEYENNLKHMVGIAKQRQADTGVKLLWGTANL-F
	QQPGEALELAKRKADVAFEFFHKLNVPFYCFHDVDVSPEGASLKEYKNNF-AQMVDVLAA
_	FMEKTRIDYFCFHDVDISPEGASLKESNENLDIIVELIKEKMDQTGKKLL-WNTTNNF
	APYYAWHDRDVAPEGANLRETHANLDAVADVLEEQQKATGVKLLWGTANM-FSNPRFM
	SPFYCFHDADVRPEGKNFAENTKNLNEIVDYFAQKQADTGVKLLWGTANL-FSNRRFM
	FCFHDRDIAPEGKTLRETNKILDKVVERIKERMKDSNVKLLWGTANLFSH-PRYMHGA
_	HDTDVAPEGSSIKEYRNNFAQMVDRLERHQEQSGIKLLWGTANCFSNPRF-AAGAASN
	EYFPGVGKIEFEGPDSDNPFAFKYYDENRVVAGKTLKEH-FKFSVCY
	KFALSWWHTMGGDGTDMFGCGTTDKTWGQSDPAARAKAKVDAAFEIMDKL-SIDYYCF
_	FCFHDVDIAPEGSSLKESNENLDTVVAMIKEYMKDSKTKLLWNTVNNFTH-PRFVHGA
	SNLKAVVAYLKEKQKETGIKLLWSTANVFGHKRYMNGASTNPDFDVVARA-IVQIKNAID
	YWHTFDADGKDPFGDGTMFRAWNRLTHPLDKAKARAEAAFEFFEKLGVPY-FCFHDVD
1ubpC	IYKADIGVKDGYIVGIGKGGNPDIMDGVTPNMIVGTATEVIAAEGKIVTA-GGIDTHVHFEHDYTHYGDEVKFGGGKVIRDGMGQSPIAN-AE-
10 1ubpC g1 222	ACEETAEGGGGATIDUGWYULMGDGGAEI DI MI LWYI TADALGTARYDTGT TAYGA TAGA
	YGEEIKFGGGKTIRDGMAQTNSPSSHELDLVLTNALIVDYTGIYKADIGI-KNGKIHG
6 111pbC 4: 1031	SVRLADTDLWVKIEKDFTIYGEEVKFGGGKVIRDGMGQGQMNNKECVDLV-LTNAIIIIKVER
	QSQHSRQEGAVDTVITNALIVDHWGIVKADVGIKDGRICAIGKAGNPDVQ-PNVDIII
11110hc18115900	MPMIIPINMERAYADIAIIATIANETANIMATANANAGINDAUTCHIQUAGNADAM—LIAAD——III

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16|1ubpC|gi|586 YADMFGPTVGDAIRLADSELFIEIEKDYTTYGDEVKFGGGKVIRDGMGQH-PLAT--SDE
2|1ubpC|gi|1567 -----VHVHFREP
42|1ubpC|gi|229 -----KISRQAYAEMFGPTTGDRLRLADTDLVI-EVER--DFT
19|1ubpC|gi|230 -----S-QAT
25|1ubpC|gi|134 -----EV-----EKDLTIHGEEVKFGGGKVIRDGMGQ-SQ-V--SRA
43|1ubpC|gi|160 IVKADVGVKDGRIVGVGKSGNPDIMDGVDPHMVIGAGTEVISGEGKILTA-GGVD--THI
80|1ubpC|gi|338 -----EV-----EV-----ERDFTTYGEEVKFGGGKVIRDGMGQ-SQ-Q--SRA
39|1ubpC|gi|158 -----VED
53|1ubpC|gi|152 -----RI-----EKDYTVYGDECVFGGGKVLREGMGQ-GIEQ--AEA
85|1ubpC|gi|174 -----EIEKDYAIYGDECIFGGGKVLRDGMGQ-SA-G--YPA
44|1ubpC|gi|322 -----KI-SRKQYASMYGPTTGDKVRLGDTNLFAEIEKDY-TLYG--EEI
72|1ubpC|gi|259 TMAIGAATEIIAGEGSILTAGGIDSH------
50|1ubpC|gi|229 -----RISRSVYADMFGPTTGDRVRLADTDLIIEVEKDLTTYGEEVKF-GGGKVIRDG
3|1ubpC|gi|2302 YAGMYGPTKGDRVRLADTDLIIEIEEDYTVYGDECKFGGGKSIRDGMGQS-PSAA--RDD
81|1ubpC|gi|465 QSCGHPPAISLDTVITNAVIIDYTGIIKADIGIKDGLIASIGKAGNPDIM-NGVFSNMII
83|1ubpC|gi|269 ----ISRQAYADMFGPTVGDRVRLADTALWVEVEKDFTIYGEEVKFGGGK-VIRDGMGQG
21|1ubpC|gi|156 KFGGGKVLRDGMGQSAVYTRDEGVLDLIITNATIIDYTGIVKADIGIKDG-HIVG--IGK
13|1ubpC|gi|103 QSQV-TNDGCVDVLITNALILDHWGIVKADIGIKNGRIAGIGKAGNPDIQ-PKVN--IVI
70|1ubpC|gi|123 ----EV-----EV----EKDLTTYGEEVKFGGGKVIRDGMGQ-SQ-I--PRS
1a0cA
13|1aOcA|gi|375 NFC-WDGTDMFGSGAFE-RFWQKGGDA-LELAKLKADVAFEFFYKL--NIPF-YCFH-DI
55|1aOcA|gi|139 QKQ-TGMKVLWNTSNMF-TNPRFVAGA-ATSPDADVFAYAAAQLKH--SLEI-GKRV-GA
24|1aOcA|gi|333 EHL-RFSVAYWHTFTGD-GSDPFGAGN-MIRPWNKYSGMDLAKARV--EAAF-EFFE-KL
28|1a0cA|gi|134 SGA-ATNPDPDVFAYAA-ATVKSCIDV-TKRLKGENYVLWGGREGY--ETLL-NTDL-AR
2|1a0cA|gi|3247 ------
23|1aOcA|gi|343 F-WH-TF-R-G----T-G---GD--P--FG---APTK-SWPW-ED--GTNSLD--MAKR
3|1a0cA|gi|3203 HNF-------HTMVDILEKK-QAET-GVKL-LW
29|1a0cA|gi|229 AVC-YWHTFVWPGVDIF-GQGTFRRPW-QQAGDAMERARQKADSAF--EFFS-KLGT-PY
39|1a0cA|gi|613 VHG-AATSCNAD------
53|1a0cA|gi|613 KNRVDAFFEILEKLGAE-YFCFHDVDIA----PQGDSLKEFLENID--VMTD-YIKG-KM
22|1a0cA|gi|156 ------
43|1a0cA|gi|212 ------------
8|1aOcA|gi|1676 KQEQSGVKLLWGTANCF-TNPRYGAGAATNPDPEVFSWAATQVVT-AMNATHKLGGENYV
54|1a0cA|gi|139 THE-RFVHGAATSSNAE-VFAYAAAKV-KKSLEIAKKLGSENFVFW--GGRE-GYES-LL
42|1aOcA|gi|324 HGA-ATSCNADVFAYAG-AQVKKALEV-TKRLGGENYVFWGGREGY--QNLY-NTDM-KR
30|1a0cA|gi|158 SGA-ATNPDPDVFAFSA-ATVKTCMDA-TKTLGGANYVLWGGREGY--ETLL-NTDL-SR
18|1a0cA|gi|117 -----
31|1a0cA|gi|288 PDP-EVFAYAGAQVFSA-MNATQRLKG-SNYVLWGGREGYETLLNT--DLKR-EREQ-LG
59|1a0cA|gi|230 WHT-FCGTGADPFGPGP-LQFPWNKAD-TADARARLKMDAAFEFIT--KLGA-PYYC-FH
51|1a0cA|gi|174 HDR-DLSPEYG------
37|1a0cA|gi|293 -----
35|1a0cA|gi|231 ------
48|1aOcA|gi|716 AGIELGAENYVFWGGRE-GYMSLLNTDQKREKEHMATMLTMARDYARSKGFK------
36|1aOcA|gi|613 -----IVDEGATLRET-FTYLDQMSSFLKEMMETSHVQLLWNTANMFTHPRYVHGAATS
            INPDQVDVALANGITTL-FGGGTGPAEGSKATTVTPGPWNIEKMLKSTEGLPINVGILGK
36|1ubpC|gi|222 GAV-DVVITNAVILDWW-GVVKADVGI-KDGKIYKIGKAGNPYTQE--GVDI-I--I-GP
12|1ubpC|gi|274 IGK-AGNKDMQDGVCNN-LCVGPATEA-LAAEGLIVTAGGIDTHIH--FISP-QQIP-TA
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86|1ubpC|gi|335 DYW-GVIKADIGINNGR-IVGIGKSGN-PDIQPGVTIYIGPGTEVI--SAEG-KIVT-AG
6|1ubpC|gi|1931 DLT-TYGDEVKFGGGKV-IRDGMGQSQ-LSRDQVMDLAITNALIVD--HWGI-IKAD-VG
7|1ubpC|gi|2300 GPG-TEAIAGEGRILTA-GGIDAHIHW-ICPQQIEDALHSGITTML--GGGT-GPAE-GT
16|1ubpC|gi|586 CV--DLVLTNAIIVDYT-GIYKADIGI-KDGMIASIGKAGNPLLMD--GV---DMVI-GA
2|1ubpC|gi|1567 GQTHKETIHSGAKSAAA-GGFTRVVMMANTKPILSTPKVLTETLEIADKEDIKIDAVGSI
42|1ubpC|gi|229 IYG-EEVKFGGGKVIRD-GMGQSQRMS-KDCVDTVVTNALIIDHWG--IVKA-DIGL-KH
19|1ubpC|gi|230 SDK-TPDTVITNALILD-HWGIVKADV-AIKNGRISAIGKAGNPDI--QPGV-TIIV-GP
25|1ubpC|gi|134 QGAVDTVITNALVVDAGAGIFKADIGL-KDGRIAAIGKAGNPDTQD--GVTI-I--I-GP
43|1ubpC|gi|160 HFI-CPQQMEVALSSGV-TTLLGGGTG-PATGSKATTCTSGAWYMA--RMLEAAEEFPIN
80|1ubpC|gi|338 NGAVDTVITNALILDWW-GIVKADIGL-RDGRIVAIGKAGNPDITD--GIDI-V--I-GP
39|1ubpC|gi|158 DLT-TYGEEVKFGGGKV-IRDGMGQGQMLSAGCA-DLVLTNALIID--YWGI-VKAD-IG
53|1ubpC|gi|152 LSL-DTVITNSVIIDYS-GIYKADIGI-KNGHIVGIGKAGNPDTMH--GVQN-NMLI-GN
85|1ubpC|gi|174 SDCLDTVVTNAVVIDYT-GIYKADIGI-NGGLIVAIGKAGNPDVMDMDGVNEEM--IVGV
44|1ubpC|gi|322 KFG-GGKTIRDGMAQSA-STYTNELDA-VITNAMIIDYTGIYKADI--GIKG-GKIV-GI
72|1ubpC|gi|259 ----IHFICPQQAEDAM-MNGITTMLG-GGTGPAVGTAATTCTPGP--WHIH-SMLR-AS
50|1ubpC|gi|229 MGQSQVTNKDGAADTVI-TNALIVDHWGIVKADVAITAGVITAIGKAGNPDVQPNVDIII
3|1ubpC|gi|2302 KVL-DLVITNAIIFDTW-GIVKGDIGI-KDGKIAGIGKAGNPKVMS--GVSE-DLII-GA
81|1ubpC|gi|465 GAN-TEVIAGEGLIVTA-GAIDCHVHY-ICPQLVYEAISSGITTLV--GGGT-GPAA-GT
83|1ubpC|gi|269 QML-AAEAMDLVLTNAL-IIDHW-GIVKADIGIKHGRIAVIGKAGNPDVQPGVNVPVGPG
21|1ubpC|gi|156 GGN-PDIMDGVESHMIV-GASTEAIAG-EGLIVTAGGIDAHIHFIS--PQQI-DVAI-AS
13|1ubpC|gi|103 GPA-TEVIAGEGKIITA-GGIDTHIHF-ICPQQAEEGLSSGVTTFI--GGGT-GPVA-GT
70|1ubpC|gi|123 GGAMDTVITNALIVDHT-GIYKADVGL-RDGRIAGIGKAGNPDTQP--GVTL-I--I-GP
1a0cA
13|1aOcA|gi|375 DVAPEGCSLKEYIYNLGVMSDILADKQAETGVKLLWGTANCFTHPRYAAGASTNPDLNIF
55|1a0cA|gi|139 ENYVFWGGREGYESLWNTNMKLEQEHAAKFFHMAKDYANEIGFDAQMLLEPKPKEPSTHQ
24|1a0cA|gi|333 NIPFFCFHDVDIAPEGETLKETYKNLDIIVDMIEEYMKTSKTKLLWNTANLFTHPRFVHG
28|1a0cA|gi|134 EQEQAGRFLSLVVDYKHKIGFKGTILIEPKPQEPTKHQYDYDVATVYGFLKRFGLEKEVK
2|1a0cA|gi|3247 ------
23|1a0cA|gi|343 RMRA-HFEFME-KL-GVDR--W--C--FHDRDIAPDGKTL----T---E-TNKNLDEIVE
3|1aOcA|gi|3203 GTANCFTNPRYMSGASTNPNPEVFSWAASQVFNAMNATKRLGGENYVLWGGREGYETLLN
29|1a0cA|gi|229 YTFHDTDVSPEGSSLKEYSENFLRITDYLARKQEDTGIKLLWGTANLFSHPRYAAGAATS
39|1a0cA|gi|613 -----
53|1a0cA|gi|613 DKTGVKLLWNTANMFTHPTFVNGAATTNNADVYSMAAAQVKKGLDVSKKLNGENYVFWGG
22|1a0cA|gi|156 -----THPRWIHGAA
43|1a0cA|gi|212 -----SHPRYMNGAST
8|1aOcA|gi|1676 LWGGREGYETLLNTDLRQEREQIGRFMQMVVEHKHKMGFQGTLLIEPKPQEPTKHQYDYD
54|1a0cA|gi|139 NTNMKLELDNLATFFKMAKSYADEIGYTGQFLIEPKPKEPTTHQYDTDVATAHAFLQKYD
42|1a0cA|gi|324 ELDHLAKFFHMAVDYAKSIGFDGQFLIEPKPKEPTKHQYDSDAAACMNFLRAYDLDSHFK
30|1a0cA|gi|158 ELDQLGRFLNLVVEYKYKIGFEGTILIEPKPQEPTKHQYDYDVATVYAFLQKNGLEKEVK
18|1a0cA|gi|117 -----
31|1a0cA|gi|288 RFMRMVVEHKHKIGFKGDLLIEPKPQEPTKHQYDYDSATVFGFLHQYGLQDEIKVNIEAN
59|1a0cA|gi|230 DIDLVEEGASRQETQRRVQTIVEYAKEKQAASGVKLLWGTANLFSNPRYMNGASTNPDFN
51|1a0cA|gi|174 -----
37|1a0cA|gi|293 -----GKTMKEHLRFAVAYWHTMTQDGSDPFGKAVNKRSWLGESPMETAKKRVIAFFEIL
35|1a0cA|gi|231 -----
48|1a0cA|gi|716 -----
36|1a0cA|gi|613 CNADVYAYAAAKVKKGLDIAKELGAENYVFWGGREGYETLLNTDMKLELENLSSFYRMAV
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1ubpC GHGSSIAPIMEQIDAGAAGLXIHEDWGATPASIDRSLTVADEADVQVAIHSDTLNEAGFL 36|1ubpC|gi|222 GTEAIAGEGMILTAGGIDAHIHFICPQQIATAIAAGITTMIGGGTGPATGTNATTCTPGP 12|1ubpC|gi|274 FASGITTMIGGGTGPADGTNATTITPGRWNLKTMLRASEEYAMNLGYLGKGNVSYEPSLV 86|1ubpC|gi|335 GIDVHVHFICPQQIEEALSSGMTTLIGGGTGPTTGSNATTCTPGVWFISRMLQAADALPI 6|1ubpC|gi|1931 IKDGRIAAIGKAGNPDTQPNVNINIWRPATEVIDGEGKIFTAGGIDSHVHFICPQLVEEA 7|1ubpC|gi|2300 NATTCTPGPWHIARMLQAAEGLPMNMGFFGKGNASRPQGLEEQLRAGACGLKLHEDWGTT 16|1ubpC|gi|586 ATEVIAAEGMIVTAGGIDAHIHFICPQQIETALASGVTTMIGGGTGPATGTNATTCTPGP 2|1ubpC|gi|1567 TQDFDGEHLTDFDELIKAGAIGFSDDGIPLTDAGVLRKALQKAKLTDSLISIHEEDPNLI 42|1ubpC|gi|229 GRIAAIGKAGNPDIQPGVTIIVGPGTEVIAGEGMIVTAGGIDTHIHFICPQQIEEALMSG 19|1ubpC|gi|230 CTEVIAGEGQILTAGAIDSHIHFICPQQIDEALMSGTTTMIGGGTGPATGTNATTCTPGK 25|1ubpC|gi|134 GTEIIAGEGKILTAGGFDAHIHFICPQQIEEALMSGITTMLGGGTGPAHGTLATTCTPGP 43|1ubpC|gi|160 VGFLGKGNASDKAPLIEQVEAGAIGLKLHEDWGTTPSAIKTCMEVVD--EADIQVAIHTD 80|1ubpC|gi|338 GTEAIAGEGHIVTAGAIDSHIHFICPQQIETALASGVTTMLGGGTGPATGTNATTCTPGA 39|1ubpC|gi|158 VKDGRIFAIGKAGNPDIQPNVTIPI-GVSTEIIAAEGRIVTAGGVDTHIHWICPQQAEEA 53|1ubpC|gi|152 KTEVIAGEGMIVTAGAIDCHVHFICPQLVYEAVSSGITTMVGGGTGPAYGTRATTCTPSP 85|1ubpC|gi|174 NTEVIAAEGMIVTAGGIDCHVHFICPQLAEEAIASGITTLVGGGTGPAHGTCATTCTPSP 44|1ubpC|gi|322 GKAGNPDTQDSVNEAMVVGAATEVIAGEGQIITAGGIDTHIHFISPTQIPTALYSGVTTM 72|1ubpC|gi|259 DGMIMNTGFFGKGNVSLPTPNEEQILAGACGLKLHEDWGTTYAAIDNCLNVADKYDIQVA 50|1ubpC|gi|229 GPGTDVIAGEGKILTAGGFDSHIHFICPQQIEHALMSGVTTMLGGGTGPSHGTFATTCTP 3|1ubpC|gi|2302 STEVITGEGLIVTPGGIDTHIHFICPQQIETALFSGITTMIGGGTGPADGTNATTCTPGA 81|1ubpC|gi|465 RATTCTPSPTQMRLMLQSTDDLPLNFGFTGKGSSSKPDELHEIIKAGAMGLKLHEDWGST 83|1ubpC|gi|269 TEVIAAEGKIVTAGGVDSHIHFICPQQVDEALNSGVTTFIGGGTGPATGTNATTCTPGPW 21|1ubpC|gi|156 GITTMLGGGTGPATGTKATTCTPGKWNIERMLEAADAFPVNLGFLGKGNASTPAPLREQI 13|1ubpC|gi|103 NATTVTPGIWNMHRMLEAVDELPINVGLFGKGCVSQPEAIREQIEAGAMGLKIHEDWGAT 70|1ubpC|gi|123 GTEVIAGEGKILTAGGIDTHIHFICPQQIEDALASGITTMLGGGTGPAHGTLATTCTPGP 1a0cA 13|1aOcA|gi|375 AYASAQVCQVMQMTKKLGGENYVLWGGREG-YESLLNTDLRQEREQIGRFMQMVVDYKYK 55|1aOcA|gi|139 YDFDAATTIAFMKEYDLDKDFKLNLEGNHA-NLAGHTYQHEIRVAREANLLGSLDANQGD 24|1aOcA|gi|333 AATSCNADVFAYAAAKVKKGLEIAKRLGAE-NYVFWGGREGYETLLNTDMKLELDNLARF 28|1a0cA|gi|134 LNIEQGHAILAGHSFEHELALANALGVFGS-IDMNRNDYQSGWDTDQFPNNVPEMALAYY 2|1a0cA|gi|3247 -----23|1a0cA|gi|343 LAKKLQE-ETNIKPL-WG-TAQLF-MH-PRYM-HG-AATSPEVKVYA-YG--AAQVKKAL 3|1a0cA|gi|3203 TDLKREREQIGRFMQMVVEHKHKIGFNGTL-LIEPKPQEPTKHQYDYDVATVYGFLKQFG 29|1aOcA|gi|229 PDPEVFAFAATQVRHALDATQRLGGDNYVL-WGGREGYDTLLNTDLMRERDQLA-----39|1a0cA|gi|613 -----53|1aOcA|gi|613 REGYENLLNTDMNFELDNLARFYQMVIDYA-QKIDYHPQFLIEPKPKEPTKHQYDYDAAT 22|1aOcA|gi|156 TSPNADVYAYAAAKVKKGLEIGKELGAENYVFWGGREGYETLLNTNMKLELDNLARFFHM 43|1a0cA|gi|212 NPDFNVVARAAVQVKAAIDATVELGGENYVFWGGREGYACLHNTQMKREQDNMARFLTLA 8|1aOcA|gi|1676 VATVYGFLKQFGLEKEIKVNIEANHATLAGHSFHHEIATAIALGIFGSVDANRGDAQLGW 54|1a0cA|gi|139 LDKDFKFNIEANHATLAGHTFQHELRYARDNNMLGSVDANQGHPLLGWDTDESTDVYDTT 42|1a0cA|gi|324 LNIETNHATLAGHTMMHELDYAGIQGGLGS-IDANTGDLLLGWDTDQFPTDYYLTTQTML 30|1a0cA|gi|158 VNIEQGHAILAGHSFEHELAMANAFGIFGS-IDMNRNDYQSGWDTDQFPNNVPEMALAYY 18|1a0cA|gi|117 -----31|1a0cA|gi|288 HATLAGHSFHHEIATAVSLGIFGSIDANRG-DPQNGWDTDQFPNSVEEMTLATYEILKAG 59|1a0cA|gi|230 VVAYAGAQLKDALDATIALDGENYVFWGGR-EGYMSLLNTDMGREQEHLARFLTMARDYA 51|1a0cA|gi|174 -----37|1a0cA|gi|293 EKLDVEYFCFHDIDIAPEGNSLKEFFSNIDEITDLIKEKMDETGIKLLWNTANMFSNPRY

35 1a0cA gi 231	
48 1a0cA gi 716	
36 1a0cA gi 613	${\tt EYAREIGFDGQFLIEPKPKEPTKHQYDFDA-ATTIAFLETYGLKDHFKLNLEANHATLAG}$
1ubpC	${\tt EDTLRAINGRVIHSFHVEGAGGGHAPDIMAMAGHPNVLPSSTNPTRPFTVNTIDEHLDML}$
36 1ubpC gi 222	$\verb WNIYRMLQAADAFPVNLGFLGKGNSSQPQG-LIEQVQAGVVGLKLHEDWGSTPNAIDTCL $
12 1ubpC gi 274	${\tt DQLEAGAIGFKIHEDWGSTPAAIYHCLNVA-DKYDVQVAIHTDTLNEAGCVEDTLQAIAG}$
	NIGFTGKGSSALPESLIEQIHAGAIGLKVH-EDWGATPATIDCCLNVADNFDVQVSIHTD
	ANLGGDYPDWLAGTGPATGTNATTCTPGSW-NIGRMLQAADAFPINMGFLGKGNASLPQS
	PSAIDVCLSVAEKWDVQVAIHTDTLNESGF-VENTTAAFKGRTIHAFHTEGAGGGHAPDI
	WNIHRMLQAAEEFPINLGFLGKGNCSDEAP-LKEQIEAGAVGLKLHEDWGSTAAAIDTCL
	GTLGVNDGEVAHKCGFTGAPTVSEYSMMARDSMIAYETGARLHIQHLSAGESVEVVRFAK
	VTTMIGGGTGPATGTYATTVTPGPWYMERM-LQAADAYPMNIGFLGKGNASLAAPLTEQV
	WHIGKMLQAGESFAMNLGFLGKGNASLPGG-LNEQLEAGALGLKLHEDWGTTPASIDNCL
	WHMARMIQSFDAFPMNIGLSGKGN-ASLPAALEEMVLGGACSLKLHEDWGTTPAAIDCCL
	TINEAGFLENTLDAIGDRVIHTYHIEGAGG-GHAPDIMKLASYANILPSSTTPTIPYTVN
	FHISRMLQAAEGLPMNLGFFGKGN-ASTAEALEEQVLAGACGLKLHEDWGTTPAAIDCCL
	LTSGITTMIGG-GTGPTAGSNATTCTPGPW-YIYQMLQAADSLPVNIGLLGKGNCSNPDA
	FDMKLMLQSTDSLPLNFGFTGKGNTAKPLE-LRHIVEAGAMGLKLHEDWGTTPAAIDNCL
	SHMKLMLQSTDELPINMGFTGKGN-TTKPDGLAEIIKAGAMGLKLHEDWGSTPAAIDNCL
	IGGGTGPAAGTNATTCTPGKWNMHQMLRAA-ESYAMNLGFFGKGNSSNEEGLEEQIKAGA
	${\tt IHTDTINEGGYLENTIAAMKDRTIHTFHTE-GAGGGHAPDIIAVVGQDNVLPSSTNPTRP}$
50 1ubpC gi 229	${\tt GPWHIGRMIQSFDAFPVNLGISGKGNAALPGPLKEMIEGGACALKLHEDWGTTPAAIDNC}$
3 1ubpC gi 2302	$\verb FNIRKMLEAAEDFPVNLGFLGKGNASFETP-LIEQIEAGAIGLKLHEDWGTTPKAIDTCL $
81 1ubpC gi 465	PAAIDNCLTIAEHHDIQINIHTDTLNEAGF-VEHSIAAFKGRTIHTYHSEGAGGGHAPDI
83 1ubpC gi 269	${\tt YLARMLQAADSLPINIGLLGKGNASRPDAL-REQIAAGAVGLKLHEDWGSTPAAIDCCLG}$
	EAGAIGLKLHEDWGTTPAAIRTCLSVADRM-DVQVAIHTDTLNEAGFVEDTIKAIGDRVI
	PAAINNCLNVADEMDVQVAIHSDTLNEGGF-YEETVRAIGDRVIHVFHTEGAGGGHAPDV
	WHISRMLQSFEAFPMNLALAGKGN-ASLPEGLVEQVKAGACALKLHEDWGTTPAAIDCCL
1a0cA	FCFHDRDIAPEGDTLRETNKNLDTIVAMIKDYLKTSKTKVLWGTANLFSNPRFVHGASTS
13 1a0cA gi 375	IGFQGTLLIEPKPQEPTKHQYDYDVATVYGFLKQFGLENEIKVNIEANHATLAGHSFQ
	KLIGWDIDEFPSDLYEATAAMYEVVENG-SIGPRGGLNFDAKPRRSSFAANDLFYGHIVG
	LHMAVDYAKEIGFDGQFLIEPKPKEPTKHQYDFDVATALAFLQTYGLKDYFKFNIEAN
28 1a0cA gi 134	QILQA
20 100CK g1 104	
2 1a0CA g1 5247	EVTHYLG-GENYVFWGGRE-GYQTLLNT-DMKRELDHL-ASFLQ-AAVDYKKKIGFNG
3 1aUCA g1 32U3	LEKEIKVNIEANHATLAGHTFQHEIATA-AALDIFGSIDANRGDPQLGWDTDQFPNSVEE
29 1aUCA g1 229	VFRYAAAQVKKG-L
9	
	AMAFIQKYNLEDSFKLNLEANHATLAGHTFEHELNVAKNYNALGSLDANQGDLLLGWD
	AVDYAKEIGFDAQFLIEPKPKEPTKHQYDFDVATGIAFLKTHGLDEHFKFNIEANHATLA
_	RDYGRAIGFKGNFLIEPKPMEPMKHQYDFDSATVIGFLRQHGLDQDFKLNIEANHATL-S
	DTDQFPISVEENALVMYEILKAGGFTTGGLNFDAKVRRQSTDKYDLFYGHIGAMDTMALS
	LAMYEILKNGGLAPGGLNFDAKPRRTSFKQEDLILTHIAGMDTFALGLRVAYKMIEDNFF
42 1a0cA gi 324	MILKH
30 1a0cA gi 158	HVLAG
18 1a0cA gi 117	
31 1a0cA gi 288	GFTHGGYNFDSKVRRQSLDDVDLFHGHVAAMDVLALSLERAAAM

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59|1a0cA|gi|230 RGQGFTGNFFIEPKPMEPSKHQYDFDSE-TVIGFLRHHGLDKDFKLNIETNHATLANH-T
51|1a0cA|gi|174 -----SLKATNDQLDIVTDYIKEKQGDKFKCLWG
37|1a0cA|gi|293 VNGAASTNNANVYAIAAAQVKKGLDVSKKLGGENYVFWGGREGYETLLNTDMKFEQDNIA
35|1a0cA|gi|231 -----
48|1a0cA|gi|716 -----
36|1a0cA|gi|613 HTFEHELRVAALHDMLGSIDANQGDLLL--GWDTDEFPTDLYSAVLAMYEILKAGGFKTG
              MVCHHLKQNIPEDVAFADSRIRPETIAAEDILHDLG-----
36|1ubpC|gi|222 SVAEDYDIQVAIHTDTLNESGFVEDTIA-AFKNRTIHAYHTEGAGGGHAPDIIKLCGQAN
12|1ubpC|gi|274 RTIHTFHTEGAGGGHAPDVIKMSGEFNI-LPASTNPTIPFTVNTEAEHMDMLMVCHHLDK
86|1ubpC|gi|335 TLNESGFLETTISAIKNRTIHAYHVEGA-GGGHSPDIIRICELNNVLPSSTNPTLPYTIN
6|1ubpC|gi|1931 LEEQVRAGVLGLKLHEDWGTTPAAIDNC-LSVAERFDVQVTIHTDTLNESGFVEDTIAAF
7|1ubpC|gi|2300 IKLCGEANVLPSSTNPTRPFTRNTLDEHLDMLMVCHHLDSRIPEDVAFAESRIRRETIAA
16|1ubpC|gi|586 KVADRYDVQVAIHTDTLNEGGFVEDTLK-AIDGRVIHTYHTEGAGGGHAPDIIKAAGFPN
2|1ubpC|gi|1567 NLGAKITAEVTPQHFSITEQMIFEQGTNAKLNPPLRSTTDIGKIIEGLKDGTIDVIATDH
42|1ubpC|gi|229 EAGAIGLKLHEDWGTTPQAIDTCLSVAD-ATDTQVAIHTDTLNEGGFVEATIAAFKGRTI
19|1ubpC|gi|230 TVAENY-----DVQVAIHTDTLNESGFVEDTLAAFKGRTIHTYHTEGAGGGHAPDIIKA
25|1ubpC|gi|134 SVADDYDVQVMIHTDTLNESGFVENTVA-AIKGRTIHAFHTEGAGGGHAPDIIKVCGLPN
43|1ubpC|gi|160 TMDEHLDMMMVCHH-----LDAKVPEDVAFSHSRIRAATIAAEDILHDIGAISMTS
80|1ubpC|gi|338 SVADRFDVQVCIHTDTLNEAGFVEDTIR-AIGGRTIHTFHTEGAGGGHAPDIIRICGESN
39|1ubpC|gi|158 LREQVAAGVIGLKIHEDWGATPAVI-NCALTVADEMDVQVALHSDTLNESGFVEDTLTAI
53|1ubpC|gi|152 AVAEEYDIQVNIHTDTLNESGFVEHTIN-AFRGRTIHTYHSEGAGGGHAPDIIRVCGVKN
85|1ubpC|gi|174 SVAEAFDIQVNIHTDTLNESGCVEHTIA-AFKDRTIHTYHSEGAGGGHAPDIIKVCGVKN
44|1ubpC|gi|322 LGLKVHEDWGSTPAAINHALNVAQKYDV-QVAIHTDTLNEAGCVEDTMKAIDGRTIHTFH
72|1ubpC|gi|259 YTVNTLDEHLDMLMVCHHLHANIAEDLA-FAESRIRKETIAAEDILHDMGAISMMSSDSQ
50|1ubpC|gi|229 LTVADAYDIQVMIHTDTLNESGFVEDTVKAFKGRTIHAFHTEGAGGGHAPDIIKVASLEN
3|1ubpC|gi|2302 KVADLFDVQVAIHTDTLNEAGFVENTIA-AIAGRTIHTYHTEGAGGGHAPDIIKIASRMN
81|1ubpC|gi|465 IKVCGIKNVLPSSTNPTRPLTSNTIDEHLDMLMVCHHLDREIPEDLAFAHSRIRKKTIAA
83|1ubpC|gi|269 VAEEMDIQVAIHTDTLNESGCIEDTLAA-IGDRTIHTFHTEGA------GGGHAPDII
21|1ubpC|gi|156 HTYHTEGAGGGHAPDIMKVAGLPNVLPS-STNPTRPFTVNTIDEHLDMLMVCHHLDPNVP
13|1ubpC|gi|103 IKSVGEPNIIPASTNPTMPYTINTVDEHLDMLMVCHHLDPSIPEDVAFAESRIRRETIAA
70|1ubpC|gi|123 TVAEDMDVQVMIHTDTLNESGFVENTLA-AFKGRTIHAFHTEGAGGGHAPDILKVVSSQN
1a0cA
              CNADVFAYSAAQVKKALEITKELGGENYVFWGGREGYETLLNTDMEFELDNFARFLHMAV
13|1aOcA|gi|375 HEVATAIALGILGSIDANRGDAQLGWDTDQFPNSVEENSLVMYEILKAGGFTTGGLNFDA
55|1a0cA|gi|139 IDTFAAGLRVALKMKQDGFLEKLVADRYSSYQSGVGAEIEAGTADFKSLESYAIDKPQSE
24|1aOcA|gi|333 HATLAGHTFEHELRVARIHGMLGSVDANQGDMLLGWDTDEFPTDLYSTTLAMYEILKNGG
28|1a0cA|gi|134 ------GGFKTGGTNFDAKLRRQSLD
2|1a0cA|gi|3247 -----
23|1a0cA|gi|343 TLLIEPKPQEP-TKHQYD-W-DVATTFSFLQK-Y--G-LTGEFKINVECNHA-T-LS--G
3|1aOcA|gi|3203 NTLVMYEILKAGGFTTGGFNFDAKIRRQSTDPYDLFHGHIGAIDVLALSLKRAAKMIEDK
29|1a0cA|gi|229 VVDHAHKIGFKGSLLIEPKPQEPTKHQYDYDVATVHGFLLQHGLEKEIRVNIEANHATLA
39|1a0cA|gi|613 ETAKELGAENYVFWGGREGYETLLNTNLQLELDNLARFMHMAVD-YATEIGYTGQFLIEP
53|1a0cA|gi|613 TDEFPTDIYTATLAMYEVLDFGGIAPGGLNFDAKVRRTSFAMDDLILAHIAGMDTYARGL
22|1a0cA|gi|156 GHTFEHELHLARIHDMLGSVDANQGDTLLGWDTDEFPTDLYTTTLAMYEILKNDGLGKGG
43|1a0cA|gi|212 GHSFEHDLQVASDAGLLGSIDANRGNPQNGWDTDQFPTDLYDTVGAMLVVLRQGGLAPGG
8|1aOcA|gi|1676 LKIAARMVEDGELDKRVAKRYAGWNGELGQQILKG------
54|1a0cA|gi|139 ENIMDEKYKSFNEGIGKKIVEGETSLKELEDYAFNINTINNTSDHLEVIKSQINQYI---
42|1a0cA|gi|324 -----GGIGTGGVNFDAKVRRESFE
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30|1a0cA|gi|158 -----GGFKNGGTNFDSKLRRQSLD
18|1aOcA|gi|117 -----ATTCSADVFAYAAAQVKKALEITKELGGEGYVFWGGRE
31|1a0cA|gi|288 -----
59|1a0cA|gi|230 MDHEMTVAANAGMLGSIDANRGDYQNAWDTDQFPYN------
51|1a0cA|gi|174 TAKCFDHPRFMHGAGTSPSADVFAFSAAQIKKALESTVKLGGNGYVFWGGREGYETLLNT
37|1a0cA|gi|293 RLFKMAIFYGEKIGHKPQFLIEPKPKEPSKHQYDFDAATTMAFILKYGLEKDFKLNLEAN
35|1a0cA|gi|231 ------ASSNNADVFAYAAAKVKKGLEVGKELGAENYVFWGGRE
48|1a0cA|gi|716 -----
36|1a0cA|gi|613 GINFDAKVRRPSFADEDLFHAHIAGMDTYAVGLKVASRLLEDKALDQVIEERYESYTKGI
              _____
1ubpC
36|1ubpC|gi|222 VLPSSTNPTRPYTVNTLDEHLDMLMVCHHLDPSIPEDVAFAESRIRRETIAAEDILHDLG
12|1ubpC|gi|274 NIKEDVQFADSRIRPQTIAAEDKLHDMGIFSITSSDSQAMGRVGEVITRTWQTADKNKKE
86|1ubpC|gi|335 TVDEHLDMMMVCHNLNCNLPEDIAFAESRIRRETIAAEDILHDLGALSMISSDSQAMGRI
6|1ubpC|gi|1931 KDRTIHTYHTEGAGGGHAPDIIKACGEANVLPSSTNPTTPFTANTVDEHLDMLMVCHHLD
7|1ubpC|gi|2300 EDILHDLGAFSMIASDSQAMGRVGEVIIRTWQTADKMKRQRGALPEERGQNDNQRVKRYI
16|1ubpC|gi|586 ILPSSTNPTRPYTINTLEEHLDMLMVCHHLDANIPEDIAFADSRIRKETIAAEDVLHDLG
2|1ubpC|gi|1567 APHTREEKNVSLDKAPSGMIGLETSLQLGLTNLVAKGHLTLSELLTKMTVNPAKLYNFDA
42|1ubpC|gi|229 HTYHTEGAGGGHAPDIIKVCGEANVLPSSTNPTRPYTVNTLDEHLDMLMVCHHLDPAIAE
19|1ubpC|gi|230 CGSSNVLPSSTNPTRPYTVNTVDEHLDMLMVCHHLDPNIPEDVAFADSRIRKETIAAEDI
25|1ubpC|gi|134 VIPSSTNPTRPYTVNTLAEHLDMLMVCHHLSPSIPEDIAFAESRIRKETIAAEDILHDIG
43|1ubpC|gi|160 SDSQAMGRVGEVIIRTWQVADKMKKQRGALAGENGNDNVRAKRYIAKYTINPAITHGLSH
80|1ubpC|gi|338 VLPSSTNPTRPYTRNTLEEHLDMLMVCHHLDPAIPEDVAFAESRIRRETIAAEDILHDLG
39|1ubpC|gi|158 GGRTIHTFHTEGAGGGHAPDIITACAHPNILPSSTNPTLPYTVNTIDEHLDMLMVCHHLD
53|1ubpC|gi|152 VLPSSTNPTRPYTKNTVDEHLDMLMVCHHLDKNIPEDVAFAESRIRAETIAAEDILHDMG
85|1ubpC|gi|174 VLPSSTNPTRPFTLNTVDEHLDMLMVCHHLDRNIPEDVAFAESRIRAETIAAEDILHDMG
44|1ubpC|gi|322 TEGAGGGHAPDIIKAAGEPNILPASTNPTIPFTKNTADEHLDMLMVCHHLDKKIKEDVAF
72|1ubpC|gi|259 AMGRIGEVVLRTWQTAHKMKIQRGSLQEDTSKSDNFRVKRYIAKYTINPAITHGVSHVIG
50|1ubpC|gi|229 VLPSSTNPTRPFTRNTIDEHLDMLMVCHHLDPSIAEDLAFAESRIRKETIAAEDI-----
3|1ubpC|gi|2302 VLPSSTNPTMPFTVNTLDEHLDMLMVCHHLDSKVKEDVAFADSRIRPETIAAEDILHDMG
81|1ubpC|gi|465 EDVLNDIGAISIISSDSQAMGRVGEVISRTWQTADKMKAQTGPLKCD-SSDNDNFRIRRY
83|1ubpC|gi|269 RAAGQANVLPSSTNPTLPYTINTVDEHLDMLMVCHHLDPSIAEDVAFAESRIRRETIAAE
21|1ubpC|gi|156 EDVAFADSRIRPETIAAEDILQDLGVISMISSDSQAMGRVGEVIIRTWQTADKMKKQRGA
13|1ubpC|gi|103 EDILHDLGAISVMSSDSQAMGRVGEVITRTWQCANKMKLQRGSLEGDSVFNDNQRIKRYV
70|1ubpC|gi|123 VIPSSTNPTRPYTKNTVEEHLDMLMVCHHLDNKVPEDVAFAESRIRKETIAAEDILHDMG
1a0cA
              DYAKEIGFEGQFLIEPKPKEPTKHQYDFDVANVLAFLRKYDLDKYF-KVNIEANHATLAF
13|1a0cA|gi|375 KVRRQSIDIDDLFYGHIGAIDTMALSLKSAVKILVDG-KL-DE-Y--V-AQRYSGWNSEL
55|1a0cA|gi|139 LIAATSSDPLEEVKDTINHYI-----
24|1a0cA|gi|333 LGRGGLNFDAKVRRGSFEPEDLFYAHIAGMDSFAVGLKVAHRLIEDRV-FDEFIEERYKS
28|1a0cA|gi|134 PQDLLIGHIGGMDACARGLKAAARMVEDKALSAPLAERYAGWNSAE-GKAMLSGKRTLED
2|1a0cA|gi|3247 ------
23|1a0cA|gi|343 H-SCHHELETA--RINGLL--GNIDANTGDPQIGWDTDQFLTDISE-A-TLVMSSVVKN-
3|1aOcA|gi|3203 TLQGIVDQRYAGWNGDLGQQILAGKASLEDLAKIVE--SK-AL-D--P-KPVSGQQEYLE
29|1a0cA|gi|229 GHSFHHEIATAYALGIFGSVDANRGDPQNGWDTDQFPNSVEELTLA-FYEILKHGGFTTG
39|1a0cA|gi|613 KPKEPTTHQYDTDAATTISFLRQYGLDKYFKLNLEANHATLAGHTFEHELRVARVQGLLG
53|1a0cA|gi|613 RAAAKMKDDNFFEQIIANRYESFSSGIGKQIVENKED-LE-SL-T--N-YALSLNGVENK
22|1aOcA|gi|156 LNFDAKVRRGSFEANDLFHAHIAGMDSFAIGLKVA------
43|1a0cA|gi|212 LNFDAKVRRESSDPQDLFLAHIGGMDAFARGLEVA-----
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8|1a0cA|gi|1676 -----
54|1a0cA|gi|139 -----
42|1a0cA|gi|324 PIDLFHAHIGGMDAFAKGLKIAAAIRASGELADFVKNRYSTWDSGI-GAKIEAGEVGFAE
30|1a0cA|gi|158 PQDLLIGHIGGMDCCARGLKAAAKMIEDGALSKPLSERYAKWDSPE-AQKMLRGELKLEE
18|1aOcA|gi|117 GYETLLNTDLGFELENLARFLRMAVDYAKRIGFTGQ--FL-IE-P--K-PKEPTKHQYDF
31|1a0cA|gi|288 ------VQNDKL
59|1a0cA|gi|230 -----
51|1a0cA|gi|174 NMGLELDNMARLMKMAVEYGRSIGFKGDFYIEPKPKE-PT-KH-Q--Y-DFDTATVLGFL
37|1a0cA|gi|293 HATLAGHTFEHELNVARNYGALGSIDANQGDVLLGWDTDEFPTNVY-DVTLAMYEILENG
35|1a0cA|gi|231 GYETLLNTNMKLELDNLARFFHMAKDYANEIDFNAQ--FL-IE-P--K-PKEPTSHQYDF
48|1a0cA|gi|716 ----GTFLIEPKPMEPTKHQYDVDTETAIGFLKAHNLDKDFKVNIE-V-NHA------
36|1a0cA|gi|613 GLEIKEGRTDLKKLAAYALENDHIENQSGRQERLKAT-VN-RY-L------
              ----IISMMSTDALAMGRAGEMVLRTWQTADKMKKQ--RG-PL-A--E-EKNGSDNFRLK
1ubpC
36|1ubpC|gi|222 AFSIISSDSQAMGRVGESIIRTWQTAHKMKVQRGHLR-DP-QR-PGVD-HDNFRARRYVA
12|1ubpC|gi|274 FGRLPEEKGDNDNFRIKRYISKYTINPAIAHGISEY--VG-SV-E--V-GKFADLVLWSP
86|1ubpC|gi|335 GETILRTWQTAHKMKLQRGSLLGDDQYNDNTRIKRY--IA-KY-T--I-NPAITHGIAHE
6|1ubpC|gi|1931 PSIPEDVAFAESRIRRETISAEDALHDMGALSMHGS--DS-QA-M--G-RVGEVILRTWQ
7|1ubpC|gi|2300 AKYTINPAITHGIAHYVGSVAVGKLADLVLWKPMFFGVKPDLVLKC-GTIASAAMGDPNA
16|1ubpC|gi|586 VFSMISSDSQAMGRVGEVIIRTWQTADKMKKQRGKL--Q--ED-N--GVGDNFRVKRYIA
2|1ubpC|gi|1567 GYLAENGPADLVIFDAENDYQVGETFDSKATNSPFIKQKVQGQVKY-TICDGKIVY----
42|1ubpC|gi|229 DIAFAESRIRRETIAAEDILHDLGAFSMISSDSQAM--GR-VG-E--V-IIRTWQTAHKM
19|1ubpC|gi|230 LHDLGAFSMIASDSQAMGRVGEVICRTWQTAHKMKVQRGLLPEDEG-T-GADNFRAKRYI
25|1ubpC|gi|134 AFSIISSDSQAMGRVGEVAIRTWQTADKMKRQRGALP-Q--E-TG--D-NDNFRVRRYIA
43|1ubpC|gi|160 EVGSVEKGKLADLVLWDPVFFGVKPELVLKGGMIARAQMGDPNASI-PTPEPVFMRQMYA
80|1ubpC|gi|338 AFSIIASDSQAMGRVGEVITRTFQTAHKMKVQRGPLPEDAANPRGS-R-NDNNRIKRYIA
39|1ubpC|gi|158 PDIAEDVAFAESRIRQETIAAEDVLHDLGAFSLTSS--DS-QA-M--G-RVGEVVLRTWQ
53|1ubpC|gi|152 AISIISSDSQAMGRIGEVISRTWQTADKMKAQRGAI--DP-NM-A--D-DDNSRIKRYIA
85|1ubpC|gi|174 AISIISSDSQAMGRIGEVITRTWQTANKMKRQRGRLPISSSPDAAE-D-NDNFRIRRYIA
44|1ubpC|gi|322 ADSRIRPETIAAEDTLHDMGIFSITSSDSQAMGRVGEVITRTWQTA-D-KCKNEFGALKE
72|1ubpC|gi|259 SVEVGKYADLVLWRPAFFGVKPSMILKGGMITAALM--GD-PN-A--S-IPTPQPVHYRY
50|1ubpC|gi|229 -----LHDLGALSMMSSDSQAMGRLGEVIIRTW--QT-AD-K--M-KKQRGALPQDS
3|1ubpC|gi|2302 VFSMMSSDSQAMGRVGEVIIRTWQTAHKMKLQRGAL--PG-EK-S--G-CDNIRAKRYLA
81|1ubpC|gi|465 IAKYTINPAIANGFSQYVGSVEVGKLADLVMWKPSF--FG-TK-P--E-MVIKGGMVAWA
83|1ubpC|gi|269 DILHDMGAFAMTSSDSQAMGRVGEVVLRTWQVAHQMKLRRGPLAPD-T-PYSDNFRVKRY
21|1ubpC|gi|156 LAEDQGKGNDNVRIKRYVSKYTINPAITHGIDDYVGSVEVGKLADL-VLWDPRFFGVKPE
13|1ubpC|gi|103 AKYTINPAIAHGISHEVGSIEVGKLADIVIWDPAFFGIKPALVVKC-GLVAYAPMGDVNA
70|1ubpC|gi|123 AMAVISSDSQAMGRVGEIIIRCWQTADKMRKQRGRLAE-E-T--G--A-NDNFRVRRYIA
1a0cA
              HDFQHELRYARINGVL--GSIDANTGDMLLGWDTDQFPTDIRMTTLAMYEVIKMGGFDKG
13|1aOcA|gi|375 GRDILEGKMTLDEVAH-YAETLVQEPKHRSGQQELLENLINRYIY------
55|1a0cA|gi|139 -----
24|1a0cA|gi|333 YTEGIGREIVEGTADF-HKLEAHALQLGEIQNQSG-----RQERLKTLLNQYL-----
28|1a0cA|gi|134 I-AERVVKKKIEPQPRSGRQELLENIVNRYV------
2|1a0cA|gi|3247 -----KHGLSLTHA
23|1a0cA|gi|343 -GGLAPGGFN-F-DAK-LRRES---TD-VEDM--FL-A----HISGMDTL-AR-GLRNVV
3|1a0cA|gi|3203 NLVNNYIY-----
29|1a0cA|gi|229 GMNFDSKVRRQSVDPE-DLFYGHIGAIDNLALAVERAAVLIENDRLDQFKRQRYAGWDAD
39|1a0cA|gi|613 SVDANQGDPLLGWDTDEFPTDLYSTTLAMYEILQNGGLGS-----GGLNFDAKVRRGSF
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53 1a0cA gi 613	SGHIEHLKSLLNDYL
22 1a0cA gi 156	
43 1a0cA gi 212	
8 1a0cA gi 1676	
54 1a0cA gi 139	
42 1a0cA gi 324	LEAYMLEKGDVDANQS-GRQEYLEHMINKYI
30 1a0cA gi 158	I-AALVERDDINPEPKPGRQEYLENVVNRYV
	DVATAYAFLKSHGLDEYFKFNIEANHATLAGHTFQHELRMARILGKLGSIDANQGDLL
_	QQFKDQRYAGWQQPFG-QSVLSGGFSLASLAEHAFANELNPQAVSGRQELLEGV
59 1a0cA gi 230	INETVEMMMVVLRAGGFQGGGVNFDAKRRRNSTDLEDTF
	RKYGLDKDFKMNIEANHATLAQHTFQHELRVARDNGVFGSIDANQGDVLLGWDTDQFP
	GIEPGGINFDSKVRRSSFEMEDLLLAHIAGMDTFARGLKSAMKLKEDRFFEDLKEQRY
_	DVASGYAFLQNYDLQDTFKFNIEANHATLAGHTFEHELHYARIHNMLGSVDANQGHPL
48 1a0cA gi 716	TLAGHTFEHELACAVDAGMLGSIDANRGDY
36 1a0cA gi 613	TLAGHTFEHELACAVDAGMLGSIDANRGDY
1ubpC	RYVSKYTINPAIAQGI-AHEVGSIEEGKFADLVLWEPKFFGVKADRVIKGGIIAYAQIGD
•	KYTINPAITHGIAEYVGSVEVGKIADLCLWKPAFFGVKPELVIKGGVIAYAQMGDANASI
	AFFGIKPNMIIKGGFI-ALSQMGDANASIPTPQPVYYREMFGHHG
	VGSVEIGKLADLVLWSPVFFGVKPELIVKGGMIVSSVMGDPNASIPTPQPVYYRLMFGSY
	SASVMKRDRGTLPEDKGDHDNFRIKRYIAKYTHNPAITHGIAHEIGSIEVGKLADLVLWK
	SIPTPQPVHYRPMFGAFGGALTHSAVNFVSQAGLDGEIAQQFGLRKTLLPVVG
	KYTINPAIAHGIADYVGSVEVGKLADLVVWNPAFFGVKPELVLKGGMIAYSTMGDPNASI
2 111hpC gi 1567	
	AAQRGKLPGDPHDARG-GHDNFRVKRYIAKYTINPALTHGIAHEVGSIEVGKWADLVFWK
	AKYTINPALAHGVAHE-VGSIEVGKLADIILWKPAFFGVKPSLIIKGGAIAAAPMGDPNA
	KYTINPAIAHGLSKDIGSIAVGKRADLVLWNPAFFGVKPDMVLVGGMIAAAPMGDPNASI
	SYGKANRSTSITFMSQASIERGVAESLGLEKRISPVKNIRKLSKLDMKLNSALPKIEIDP
	KVTINPAIAHGIDNHVGSVEVGKLADLVLWKPGFFGVRPELVIKGGSIIWAQMGDANASI
	VAHRMKVQRGPLPEESGDNDNVRVKRYIAKYTINPALTHGIAHEVGSIEVGKLADLVLWS
	KYTINPAIANGFADLIGSVEVKKLADLVIWQPAFFGAKPEMIIKGGNIAWANMGDANASI
	KYTINPAIVNGFSDFVGSVEVGKLADLVIWKPSFFGAKPEMVIKGGAIACANMGDPNASI
	ECGENDNFRIKRYISKYTINPAIAHGISEYVGSVEVGKFADLVLWKPSMFGIKPEMILKN
	MFGGYGGGVKTSCFTFTSQAALDGGLVDSLKLDKNLIAVKNTRNLRKKDMIHNGATPKME
	ARNDNFRVKRYIAKYT-INPAIAHGVSKLIGSVETGKMADLVLWSPAFFGVKPDCIIKGG
	KYTINPAITHGISQYVGSLEKGKIADLVLWKPAMFGVKPEMIIKGGFIIAGRMGDANASI
	DIGDPNASIPTPEPVKMRPMYGTLGKAGGALSIAFVSKAALDQRVNVLYGLNKRVEAVSN
	IAKYTINPALTHGIGH-EVGSVEVGKLADLVLWSPAFFAVKPALVLKGGMIVTAPMGDIN
	LILKGGLIAYSQMGDPNASIPTPQPVFSRPMFGSFGRARGTTSITFLSKAAMDLGVHEAL
	AIPTPQPVHYRPMYAS-YGKAMYSTSMIFMSQASIEAGVPEKLKLQSMIGQVKG
70 1ubpC gi 123	KYTINPAITHGLAEHV-GSVEVGKRADLVLWHPAFFGAKPEMVLMGGMIVAAQMGDPNGS
1a0cA	GLNFDAKVRRASFEPEDLFLGHIAGMDAFAKGFKVAYKLVKDRVFDKFIEERYASYKD-G
55 1a0cA gi 139	
24 1a0cA gi 333	
28 1a0cA gi 134	
_	EADHRAKARDFVKQMIDFGGKYNAPAIIGSMQGKWGGDISRDQALAFLREALDELGPY-A
	KLI-EDGSLDELV-RKRYQ-SFDSEIG-ALI-EAGKGDF-E
20,120011811040	

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	TODATA GODING OTT ATTAIN A POR VIDOUA GOVERN VIDATING A TUG
•	FGRKILSGDYSLSTLATDALARGLNPQHASGHQELMENIVNQAIYS
39 1a0cA gi 613	EQDDLLYAHVAGMDAFARG-LKVAHKLVEDRVFENVINERYSSFKEGIGLEIVEGK-A
	NQLLEDRV-LEDVIDNRYKSYQSGIGQKIANNDTNLKELEAYALS-L
	DALLTSSP-LETWRAQRYASFDSGAGADFANGTSTLADLATYAAG-K
8 1a0cA gi 1676	QLSLGE-L
54 1a0cA gi 139	
42 1a0cA gi 324	
30 1a0cA gi 158	
18 1a0cA gi 117	LGWDTDQFPTNVYDTTLAMYEVIKAGGFTKGGLNFDAKVRRASYKVEDLFIGHIAG-M
31 1a0cA gi 288	
_	HGHIGGMDTFARALIIADRIISESKIESMRKERYASFDSGAGKQFETGALDLAALAEI-G
	TNIYDTTMCMYEVIKAGGFTNGGLNFDAKARRGSFTPEDIFYSYIAGMDAFALGFR-A
	SSFKKGIGAKIISGKENLESLTNYALKNDEPIIESSHIEYVKNILNDYLY
	LGWDTDEFPSDIYTTTLAMYEILKNDGLGKGGLNFDAKVRRGSFTAEDLFHAHIAG-M
	QNGWDTDQFPIDQYELVQAWMEIIRGGGFVTGGTNFDAKTRRNSTDLEDIIIAHVS-G
1ubpC	PSASIPTPQPVMGRRMYGT-VGDLIHDTNITFMSKSSIQQGVPAKLGLKRRIGTVK-N
-	PTPQPVHMQPMFASYGGCRTTTSVTFMSQAGIANNIPEQLGLQKTVLPVGHIRQ-L
	KAKFDTNITFVSQVAYDNGIKEELGLQRVVLPVKNCRNITKKDLKFNDVTAHIEVNPE-T
	DLAKCATRMTFISQSSYDVGVVENLRLNSLIGISKRCRNIKKDHMINNNLKPVM-E
	PAFFGVKPSLVLKGGVIVTAPMGDPNASIPTPQPVHYRPMFGSFGGSRTASCVSFVSQ-A
	-CRTIGKADMVHNSATPHM-EVDPETYEVRADGRLLTCEPATVLPLAQRYFLF
16 1ubpC g1 586	PTPQPVLYRPMFAAKGDAKYQTSITFVSKAAYEKGIHEQLGLKKKVKPVHGIRK-L
1 0	PAFFGVKPSLILKGGMIAA-AAMGDPNASIPTPQPVHYRPMFASAGGALPRSSLTF-V
	SIPTPQPVHYRKMFGAYGQ-ACKQTSVSFVSQAAIDASISDYFQLERRLVAVKNCR-S
	PTPQPMHYRPMFGAYGKARTNSSVTFVSKAALESGLHGRLGVEKQFVAVENTRGGI
	KTYQVFADGEELSCQPVDYVPLGQRYFLF
	PTPGPVHGRPMFAAFGKALAPSCLTFLSQAAIETDLPNKLGLQRACIPVLNTRT-I
39 1ubpC gi 158	PAFFGVKPATIVKGGMIAMAPMGDINGSIPTPQPVHYRPMFAALGSARHRCRVT-F
1 0	PTPEPVISRPMFGAFGKAGSENSVAFVSKAALRKGVKELYGLKKRVVAVSNVRQ-L
85 1ubpC gi 174	PTPEPVMMRPMFGAFGGAGSANSIAFVSKAAKEAGVAVQYKLG-KRVEA-VGRVRG-L
	GMIVAAKIGDSNASIPTPEPVVYAPMFGSYGKAKYNCAITFVSKIAYDC-H
72 1ubpC gi 259	VDPETYEVRADGQLLTCGA
	SIVAAPMGDPNASIPTPQP-VHYQPMFGAYGKALTASSVVFTSQAAAAGNLARDLG-I
	PTPQPVIYKNMFGAFGKAKYGTCVTFVSKASLENGVVEKMGLQRKVLPVQGCRN-I
	VRKLTKLDMKLNDALPEITVDPESYTVKADGKLLCVSEATTVPLSRNYFLF
	GSIPTPQPVHYRPMFGALGAARHATRMTFLPQAAMDRGLAEELNLRSLIGVVNGCRRV-R
	GLQKKIAHVKNCRSISKRSMKYNDATPNIEIDPETYEVKVDGEMITCKPFEEVALAQR-Y
	-CRNLSKQDMVHNSYTPTI-ELDSQTYLVKADGIPLVCEPATELPMTQRYFLF
	IPAQ-PFYTRPMFGAFGKA-LSNSAVTFVSAAAEAEGVAGKLGLSKTVLPVKGTRT-I
1011mpho1811179	TIME TITIE ON OUT ONE OF A LEAD THAN SOUND TO SEE THE SECTION OF T
1a0cA	IGADIVSGKADFRSLEKYALERSQIVNKSGRQELLESILNQYLFA
	TGADIVSGRADFRSLERIALERSQIVINSGRQELLESILINQILFA
10 140CH 81 9/0	
00 140CH 81 139	
24 1aUCA g1 333	

28 120c1 mi 13/	
20 1a0CA g1 134 2 1a0cA g1 3247	AQYNVP-LFYEPLNRYETNLLRTVDEGVEFCKTLSTDNIKLLADLFHMNIEEADLAAAIR
_	T-LEKKVLEWGEP-I-V-PSGKQELAEIL-FQ-A-AL
3 120ck g1 040	
201120cA g1 3203	
59 1a0CA g1 613	NFHTLE-QYAFKNPNIANKSGRQERLKSILNQYI
00 1a0cA g1 013	GEIKHS-SGQQERIKATLNQYL
	GEPTQL-SGRQEAYENLINQYL
8 1a0cA g1 16/6	AQYAEQ-HNLAPVHQSGHQELLENLVNRYLF
54 1a0cA gi 139	
42 1a0cA gi 324	
30 1a0cA gi 158	
18 1a0cA gi 117	DTFALG-FKVAYKLVKDGVLDKFIEEKYRSFREGIGRDIVEGKVDFEKLEEYIIDKETIE
31 1a0cA gi 288	NAIGEPALTSGKQEYYENIINQYI
59 1a0cA gi 230	NAIGEPALTSGKQEYYENIINQYI
51 1a0cA gi 174	ALKLIE-DGRIDKFVADRYASWNTGIGADIIAGKADFASLEKYALEKGEVTASLSSGRQE
	DSFAIG-LKVAQRLMDDRVLENVVEDRYRSFNQGIGRDIVSGKTDFHQLEKHALTLSEIT
48 1a0cA gi 716	MDAMAR-ALENAAKLLQESPYTKMKKERYASFDSGIGKDFEDGKLTLEQVYEYGKKNGEP
36 1a0cA gi 613	
1ubpC	CRNIGK-KDMKWNDVTTDIDINPETYEVKVDGEVLTCEPVKELPMAQRYFLF
36 1ubpC gi 222	RKADLK-LNDYLPHIEVDPETYEVRADGELLTCEPATVLPLAQRYFLF
12 1ubpC gi 274	YKVKVDGKEVTSKA
86 1ubpC gi 335	VDPQTY-EVRANGELLVCEP
6 1ubpC gi 1931	GLDEGIGEKLGLQKRLVAVKNIRGLRKSDLIHNNALPRIEVDPQNYQVRADGQLL
7 1ubpC gi 2300	GLDEGIGEKLGLQKRLVAVKNIRGLRKSDLIHNNALPRIEVDPQNYQVRADGQLL
16 1ubpC gi 586	TKKDLI-LNDKTPKIDVDPQTYEVKVDGQLVTCEPAEIVPMAQRYFLF
2 1ubpC gi 1567	
	SQAALA-ANVGERYGLAKTLTAVRGTRTVSKRDMVHNDWQPHVTVDPETYQVVADGQLLT
	VTKADM-VHNAYQPHMEVDPETYEVRADGQLLTCEPAEELPMAQRYFLF
	GKHSMV-LNDATPHVEVDPETYEVRADGELLTCEPATVLPMAQRY-FLF
	·
1 0	GKAEMH-NNNSLPKVEVDPQTYEVFADGDLLTCDPAEELPMAQRY-LL
	LSQAAA-ANGVAEQLNLHSTTAVVKGCRTVQKADMRHNSLLPDITVDSQTYEVRINGELI
	TKLDMK-LNDALPEITVDPETYVVTANGEVLTCAPADSVPLSRNYFLF
	TKLNMK-LNDALPKIDVDPETYTVTADGEVLRCQPTPTVPLSRNYFLF
	IKEELG-LERILLPVKNCRNITKKDMKFNDVITPIEVNPETYEVRVNNTKITSKPVEKVS
72 111bpC gi 259	
	AKKLVP-VSNVRGGISKKSMIHNDATPKLEVDPETYEVRADGELLTCAPAEVLPLAQRYF
	SKKYMV-HNNATPEIEVDPETYEVKVDGEIITCEPLKVLPMAQRYFLF
01 1mphc &: 050	-KPDMV-HNTLQPLIEVDAQTYQVRADGELLVCEPASELPLAQRYFLF
03 1ubpC #: 156	LI LOUIN - IIM I PÁL PIENNWÁ I I ÁN WYNGEPPN OEL 49 EPLFAÚVI L PL
21 1ubpc g1 150	FLF
10 1ubpc g1 123	GKASMR-LNSATPQIEVDPETYEVRADGEILTCEPAETLPLAQRYFLY
1.0.0.1	
1a0cA	

13 1a0cA gi 375 55 1a0cA gi 139 24 1a0cA gi 333 28 1a0cA gi 3247 23 1a0cA gi 3247 23 1a0cA gi 3203 29 1a0cA gi 3203 29 1a0cA gi 613 53 1a0cA gi 613 22 1a0cA gi 156 43 1a0cA gi 156 43 1a0cA gi 176 54 1a0cA gi 139 42 1a0cA gi 139 42 1a0cA gi 158 18 1a0cA gi 17 31 1a0cA gi 230	AGKGYVGHIHFVDSNR-Q-
51 1a0cA gi 174 37 1a0cA gi 293 35 1a0cA gi 231 48 1a0cA gi 716 36 1a0cA gi 613 1ubpC	MLESIVNNVLFS QPSGQLEVIKNKINQYLLT KQTSGKQELYEAIVAM-Y-
36 1ubpC gi 222 12 1ubpC gi 274 86 1ubpC gi 335 6 1ubpC gi 1931 7 1ubpC gi 2300 16 1ubpC gi 586 2 1ubpC gi 1567 42 1ubpC gi 229	ADKISLAQLYNL-FVSVVPMSQRYFL-F- WFEPSKVLPMAQRYFL-F
19 1ubpC gi 230 25 1ubpC gi 134 43 1ubpC gi 160 80 1ubpC gi 338 39 1ubpC gi 158 53 1ubpC gi 152 85 1ubpC gi 174 44 1ubpC gi 322	TSEPADILPMAQRYFL-F-
72 1ubpC gi 259 50 1ubpC gi 229 3 1ubpC gi 2302 81 1ubpC gi 465 83 1ubpC gi 269 21 1ubpC gi 156 13 1ubpC gi 103	EDVLPXAQRYFL-F-

70|1ubpC|gi|123 -----

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