protein segfasta dictionary function

October 11, 2020

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[35]: x = """>gi|27544883|ref|NP_775396.1| putative ribosomal protein S3
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     NQNKRFISFGKYITMKRRRNLLRKIYVSNPVFKFSYFSAFITLFSLSREKVFYKKNILKRIKKYKVIKCNVIYFYIKIIK
     WIYKIFLAKQYYKNKNISYFLLKNKNKFIQYKLKYLSKFLLLKNLYLKRVWSKIIKNFIKRHLIFLRKYELLYSLNQLKF
     NKLTLLNKLSLLLNKILGKKIEYNIINLKSIIFNSDLFTQAITLKFQKRKSFNYKKNILSILGRVNFPYFKDVVALAYNE
     STNYILNKYKDGKILSYINNHQNLDNFINKIHNATNKNIHKEIFNSIQYKNIEGIRIETNGRLTKRYRADRAVHYRKWKG
     GLQKTSLNSTLFRGNVNPNISYSIVNNTRRVGSFAIRGWISGK
     >gi|27544884|ref|NP_775397.1| NADH dehydrogenase subunit 2 (mitochondrion)
      → [Lecanicillium muscarium]
     MIIISILSLLLSNAVNLRRDVSILYNRIAILILLYCIVHDYTSLTVVTKGIGLHGGLLLINNLTQIFHIFVFIVTIFILT
     LTSFYPRKVWVSEYSSIKDLEPVDPLFNKFIYYNTKIINKMGEHLKIIEYPLILLFIVRPGAIFLMSSNDLITIFLSIEL
     QSYGLYILSTVYRNSELSTTGGLIYFLLGGLSSCFILLGTALLYAKSGRTSLESLYIITSISDIHSSTNDLWYSPNYISL
     SLVIFTVGFLFKISAAPFHFWSPDVYDAIPTIVTTFVALIAKISILILLLQLVYYTNADITMNNWTFILLISSLFTLVIG
     TVVGLTQFRIKRLFAYSTISQIGFMLLALSISSIESTQAPIFYLTQYIIINLNAFIILLAITYSLYFYTNYNKEHKDLLD
     KTNSPIQLITQLKGYFFINPVLALNLTITIFSFSQIPPLLVCFCKQIVLSASLNQNLYFITLIAILTNFIERIYYLTIIK
     KIFFNKSDYKINTLLCNFKLKINIYNNNNSINSVEYNYKNIFLSIPISFIISILTLTILFFLFINKKWLIMITIFVQLIF
     >gi|27544885|ref|NP_775398.1| NADH dehydrogenase subunit 3 (mitochondrion)⊔
      MSGTTFLFIFVCVIAILFLALNFILAPHNPYQEKYSIFECGFHSFLGQNRTQFGFKFFIFSLVYLLLDLDLEILVIYPYG
     LSSYENGIYGLIIVLLFIGIITAGFVFELGKGALKIDSRQSYNYFNVQSTKNFINTFFENK"""
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[37]: protein_dict(x)
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[37]: {'gi|27544883|ref|NP_775396.1| putative ribosomal protein S3 (mitochondrion) [Lecanicillium muscarium]': 'MEKNIKLKNILLSKVPQKITLNNKIADIRKTKYLPSFSKEWKDTIYSYNKN

IMKNIPSHHLNINKIIQSYFNLFFSTNKNNQNKRFISFGKYITMKRRRNLLRKIYVSNPVFKFSYFSAFITLFSLSREKV FYKKNILKRIKKYKVIKCNVIYFYIKIIKWIYKIFLAKQYYKNKNISYFLLKNKNKFIQYKLKYLSKFLLLKNLYLKRVW SKIIKNFIKRHLIFLRKYELLYSLNQLKFNKLTLLNKLSLLLNKILGKKIEYNIINLKSIIFNSDLFTQAITLKFQKRKS FNYKKNILSILGRVNFPYFKDVVALAYNESTNYILNKYKDGKILSYINNHQNLDNFINKIHNATNKNIHKEIFNSIQYKN IEGIRIETNGRLTKRYRADRAVHYRKWKGGLQKTSLNSTLFRGNVNPNISYSIVNNTRRVGSFAIRGWISGK',

'gi|27544884|ref|NP_775397.1| NADH dehydrogenase subunit 2 (mitochondrion) [Lecanicillium muscarium]': 'MIIISILSLLLSNAVNLRRDVSILYNRIAILILLYCIVHDYTSLTVVTKGI GLHGGLLLINNLTQIFHIFVFIVTIFILTLTSFYPRKVWVSEYSSIKDLEPVDPLFNKFIYYNTKIINKMGEHLKIIEYP LILLFIVRPGAIFLMSSNDLITIFLSIELQSYGLYILSTVYRNSELSTTGGLIYFLLGGLSSCFILLGTALLYAKSGRTS LESLYIITSISDIHSSTNDLWYSPNYISLSLVIFTVGFLFKISAAPFHFWSPDVYDAIPTIVTTFVALIAKISILILLLQ LVYYTNADITMNNWTFILLISSLFTLVIGTVVGLTQFRIKRLFAYSTISQIGFMLLALSISSIESTQAPIFYLTQYIIIN LNAFIILLAITYSLYFYTNYNKEHKDLLDKTNSPIQLITQLKGYFFINPVLALNLTITIFSFSQIPPLLVCFCKQIVLSA SLNQNLYFITLIAILTNFIERIYYLTIIKKIFFNKSDYKINTLLCNFKLKINIYNNNNSINSVEYNYKNIFLSIPISFII SILTLTILFFLFINKKWLIMITIFVQLIFNS',

'gi|27544885|ref|NP_775398.1| NADH dehydrogenase subunit 3 (mitochondrion) [Lecanicillium muscarium]': 'MSGTTFLFIFVCVIAILFLALNFILAPHNPYQEKYSIFECGFHSFLGQNRT QFGFKFFIFSLVYLLLDLDLEILVIYPYGLSSYENGIYGLIIVLLFIGIITAGFVFELGKGALKIDSRQSYNYFNVQSTK NFINTFFENK'}

[38]: fastaSeqs = """>CCDS2.2|Hs109|chr1

MSKGILQVHPPICDCPGCRISSPVNRGRLADKRTVALPAARNLKKERTPSFSASDGDSDG SGPTCGRRPGLKQEDGPHIRIMKRRVHTHWDVNISFREASCSQDGNLPTLISSVHRSRHL VMPEHQSRCEFQRGSLEIGLRPAGDLLGKRLGRSPRISSDCFSEKRARSESPQEALLLPR ELGPSMAPEDHYRRLVSALSEASTFEDPQRLYHLGLPSHGEDPPWHDPPHHLPSHDLLRV RQEVAAAALRGPSGLEAHLPSSTAGQRRKQGLAQHREGAAPAAAPSFSERELPQPPPLLS PQNAPHVALGPHLRPPFLGVPSALCQTPGYGFLPPAQAEMFAWQQELLRKQNLARLELPA DLLRQKELESARPQLLAPETALRPNDGAEELQRRGALLVLNHGAAPLLALPPQGPPGSGP PTPSRDSARRAPRKGGPGPASARPSESKEMTGARLWAQDGSEDEPPKDSDGEDPETAAVG CRGPTPGQAPAGGAGAEGKGLFPGSTLPLGFPYAVSPYFHTGAVGGLSMDGEEAPAPEDV TKWTVDDVCSFVGGLSGCGEYTRVFREQGIDGETLPLLTEEHLLTNMGLKLGPALKIRAQ VARRLGRVFYVASFPVALPLQPPTLRAPERELGTGEQPLSPTTATSPYGGGHALAGQTSP KQENGTLALLPGAPDPSQPLC

>CCDS3.1|Hs109|chr1

>CCDS4.1|Hs109|chr1

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MGNSHCVPQAPRRLRASFSRKPSLKGNREDSARMSAGLPGPEAARSGDAAANKLFHYIPG
TDILDLENQRENLEQPFLSVFKKGRRRVPVRNLGKVVHYAKVQLRFQHSQDVSDCYLELF
PAHLYFQAHGSEGLTFQGLLPLTELSVCPLEGSREHAFQITGPLPAPLLVLCPSRAELDR
WLYHLEKQTALLGGPRRCHSAPPQRRLTRLRTASGHEPGGSAVCASRVKLQHLPAQEQWD
RLLVLYPTSLAIFSEELDGLCFKGELPLRAVHINLEEKEKQIRSFLIEGPLINTIRVVCA
SYEDYGHWLLCLRAVTHREGAPPLPGAESFPGSQVMGSGRGSLSSGGQTSWDSGCLAPPS
TRTSHSLPESSVPSTVGCSSQHTPDQANSDRASIGRRRTELRRSGSSRSPGSKARAEGRG
PVTPLHLDLTQLHRLSLESSPDAPDHTSETSHSPLYADPYTPPATSHRRVTDVRGLEEFL
SAMQSARGPTPSSPLPSVPVSVPASDPRSCSSGPAGPYLLSKKGALQSRAAQRHRGSAKD
GGPQPPDAPQLVSSAREGSPEPWLPLTDGRSPRRSRDPGYDHLWDETLSSSHQKCPQLGG
PEASGGLVQWI

>CCDS5.1|Hs109|chr1

MAADTPGKPSASPMAGAPASASRTPDKPRSAAEHRKSSKPVMEKRRARINESLAQLKTL ILDALRKESSRHSKLEKADILEMTVRHLRSLRRVQVTAALSADPAVLGKYRAGFHECLAE VNRFLAGCEGVPADVRSRLLGHLAACLRQLGPSRRPASLSPAAPAEAPAPEVYAGRPLLP SLGGPFPLLAPPLLPGLTRALPAAPRAGPQGPGGPWRPWLR

>CCDS6.1|Hs109|chr1

MGWDLTVKMLAGNEFQVSLSSSMSVSELKAQITQKIGVHAFQQRLAVHPSGVALQDRVPL ASQGLGPGSTVLLVVDKCDEPLSILVRNNKGRSSTYEVRLTQTVAHLKQQVSGLEGVQDD LFWLTFEGKPLEDQLPLGEYGLKPLSTVFMNLRLRGGGTEPGGRS

>CCDS7.2|Hs109|chr1

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>CCDS8.1|Hs109|chr1

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[39]: protein_dict(fastaSeqs)

[39]: {'CCDS2.2|Hs109|chr1': 'MSKGILQVHPPICDCPGCRISSPVNRGRLADKRTVALPAARNLKKERTPSFSASDG DSDGSGPTCGRRPGLKQEDGPHIRIMKRRVHTHWDVNISFREASCSQDGNLPTLISSVHRSRHLVMPEHQSRCEFQRGSL EIGLRPAGDLLGKRLGRSPRISSDCFSEKRARSESPQEALLLPRELGPSMAPEDHYRRLVSALSEASTFEDPQRLYHLGL PSHGEDPPWHDPPHHLPSHDLLRVRQEVAAAALRGPSGLEAHLPSSTAGQRRKQGLAQHREGAAPAAAPSFSERELPQPP PLLSPQNAPHVALGPHLRPPFLGVPSALCQTPGYGFLPPAQAEMFAWQQELLRKQNLARLELPADLLRQKELESARPQLL APETALRPNDGAEELQRRGALLVLNHGAAPLLALPPQGPPGSGPPTPSRDSARRAPRKGGPGPASARPSESKEMTGARLW AQDGSEDEPPKDSDGEDPETAAVGCRGPTPGQAPAGGAGAEGKGLFPGSTLPLGFPYAVSPYFHTGAVGGLSMDGEEAPA PEDVTKWTVDDVCSFVGGLSGCGEYTRVFREQGIDGETLPLLTEEHLLTNMGLKLGPALKIRAQVARRLGRVFYVASFPV ALPLQPPTLRAPERELGTGEQPLSPTTATSPYGGGHALAGQTSPKQENGTLALLPGAPDPSQPLC',

'CCDS3.1|Hs109|chr1': 'MAAAGSRKRRLAELTVDEFLASGFDSESESESENSPQAETREAREAARSPDKPGGS PSASRRKGRASEHKDQLSRLKDRDPEFYKFLQENDQSLLNFSDSDSSEEEEGPFHSLPDVLEEASEEEDGAEEGEDGDRV

PRGLKGKKNSVPVTVAMVERWKQAAKQRLTPKLFHEVVQAFRAAVATTRGDQESAEANKFQVTDSAAFNALVTFCIRDLI GCLQKLLFGKVAKDSSRMLQPSSSPLWGKLRVDIKAYLGSAIQLVSCLSETTVLAAVLRHISVLVPCFLTFPKQCRMLLK RMVIVWSTGEESLRVLAFLVLSRVCRHKKDTFLGPVLKQMYITYVRNCKFTSPGALPFISFMQWTLTELLALEPGVAYQH AFLYIRQLAIHLRNAMTTRKKETYQSVYNWQYVHCLFLWCRVLSTAGPSEALQPLVYPLAQVIIGCIKLIPTARFYPLRM HCIRALTLLSGSSGAFIPVLPFILEMFQQVDFNRKPGRMSSKPINFSVILKLSNVNLQEKAYRDGLVEQLYDLTLEYLHS QAHCIGFPELVLPVVLQLKSFLRECKVANYCRQVQQLLGKVQENSAYICSRRQRVSFGVSEQQAVEAWEKLTREEGTPLT LYYSHWRKLRDREIQLEISGKERLEDLNFPEIKRRKMADRKDEDRKQFKDLFDLNSSEEDDTEGFSERGILRPLSTRHGV EDDEEDEEGEEDSSNSEDGDPDAEAGLAPGELQQLAQGPEDELEDLQLSEDD',

 $\label{thm:cods.1} $$ \cds.1 + s.109 + c.1': \mbox{\sc Maadtpgkpsaspmagapasasrtpdkprsaaehrksskpvmekrrarineslag lktlildalrkessrhsklekadilemtvrhlrslrrvqvtaalsadpavlgkyragfheclaevnrflagcegvpadvr srllghlaaclrqlgpsrrpaslspaapaeapapevyagrpllpslggpfpllappllpgltralpaapragpqgpggpw rpwlr',$

'CCDS6.1|Hs109|chr1': 'MGWDLTVKMLAGNEFQVSLSSSMSVSELKAQITQKIGVHAFQQRLAVHPSGVALQD RVPLASQGLGPGSTVLLVVDKCDEPLSILVRNNKGRSSTYEVRLTQTVAHLKQQVSGLEGVQDDLFWLTFEGKPLEDQLP LGEYGLKPLSTVFMNLRLRGGGTEPGGRS',

'CCDS7.2|Hs109|chr1': 'MALRHLALLAGLLVGVASKSMENTAQLPECCVDVVGVNASCPGASLCGPGCYRRWN ADGSASCVRCGNGTLPAYNGSECRSFAGPGAPFPMNRSSGTPGRPHPGAPRVAASLFLGTFFISSGLILSVAGFFYLKRS SKLPRACYRRNKAPALQPGEAAAMIPPPQSSVRKPRYVRRERPLDRATDPAAFPGEARISNV',

'CCDS8.1|Hs109|chr1': 'MGSSQEEGLRCQPSQPDHDADGHCGPDLEGAERASATPGPPGLLNSHRPADSDDTN AAGPSAALLEGLLLGGGKPSPHSTRPGPFFYIGGSNGATIISSYCKSKGWQRIHDSRRDDYTLKWCEVKSRDSYGSFREG EQLLYQLPNNKLLTTKIGLLSTLRGRARAMSKASKVPGGVQARLEKDAAAPALEDLPWTSPGYLRPQRVLRMEEFFPETY RLDLKHEREAFFTLFDETQIWICKPTASNQGKGIFLLRNQEEVAALQAKTRSMEDDPIHHKTPFRGPQARVVQRYIQNPL LVDGRKFDVRSYLLIACTTPYMIFFGHGYARLTLSLYDPHSSDLGGHLTNQFMQKKSPLYMLLKEHTVWSMEHLNRYISD TFWKARGLAKDWVFTTLKVRPLCPPVWE'}

[]: