

Exercise 5 – Analysing and aligning newly discovered proteins

Objectives:

- to apply what has been learned today

Anonymous Test Proteins:

below, we provide 20 randomly chosen proteins. All have been derived from DNA on the teeth of ancient skeletons found in a german monastery (same as for the previous exercises). None of the proteins have been analyzed in detail before ... Please select arbitrarily one of the proteins below, and analyze it like we did in exercises #1 through #3.

Optionally you can also study some protein sequence related to SARS-CoV-2, which are listed at the bottom below.

Questions:

- what protein family does your protein belong to?
- which domain(s), if any, does the protein contain?
- from which organism is it, likely?
- what function might it have?
- is it complete?
- how can it be best aligned to other members of its family?

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>NODE_4178_length_1047_cov_6.240688_S6
SATSAAMKLIAPSWPSRYVSPRRRQGAAMAEWWSAQLVDRDGRIRGELPDIRGGSLEW
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STLKLEAPTSRLRSQLGWYQYEAGIVVTDVAMTLRQLGESQLALTPSPQTLRTPLTWD
PDKTWGTLYSELDAIGYGGIWCANGWWRAAPYVAPMERPLAATYGGDPADYRCRTTYG
DEADWTDVPNRVLLYTRATSEAPALTSEVWITDPANPWHPDVGPHTRCEAVEATSQEV
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RGGKMKIIIIAGIGNIGAGLAGRLLNEGHDIVLVDRDIDRLEYNEETQDVMTVKGTCAAME
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LSMIIDPGMYAAKDIEAILTYPGFLHREFTKGMTDVVEAELHPESELGKPVSAIQEIT
GSGALVCVVKRGDTAITPGRDFILREKDRIYVTAEEDLSTLLRLFGKKKETVEKVMIVG
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>NODE_60099_length_1027_cov_6.267770_S6
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QRNRADLAAENLGVNFIMLSPPNFWWFQQSNLEMSAKLKDAPGLTLVYQKNSIRIYAVNA
KFKDAELTRMRASGPASNQLPVPQCPKDSADGKAAATAGETTQVEYDPTGEQTTVTTPK
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IAAGIAGVLIGIPVLRRLRGDYLAIVTLAFGEIIRNIMNLLYVSVDQGRLRMAFNDGALPG
EQVIAGPKGAVGIEKIAFTFTMGFILVMITLFFVVLNLINSRSGRAIMAIRDSRIA AESVGI
NVTKYKMAFVISSVLAMAGALFGLNYSTVSAGKFKFDMSILVLVFFVLGGIGNIRGSV

>NODE_87482_length_1095_cov_5.276712_S6
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DPIDGTANYARGFGESAVSIALAKDDEPVLGVVRNPYARETYCAIKGRGAFLNGTPIHVS
GRSKENAMICLSWSAYDKSRSDCFRISQDLYAVCEDIRRTGSAAYELCLLARGSVDMHF
EIRLAPWDYAAGGLIIIEAGGRTGSLEGR LDMRRQCLVMAANSEKNFAFLKGVVSENLSL
RRRLAPVHV

>NODE_107984_length_1345_cov_80.271378_S6
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LADGFGFESLFQHCWRIRETVLDLFEQTTGGRVIFSICKVGGLNKDIDNETLNKIVKTLR
GIEKEIREYTSVFINDTSVKNRLTGVGVLSREDAEALCTVGP MARASGLRQDMRLAGEGK
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SCTER

>NODE_123020_length_4291_cov_7.623631_S6
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FGAALAFLLIVQVARTMISNKLQNL FASKLYDEFLDTVDEPLTRFFIPAYNR TYLRLNAF
MAKGSVEKAMEAFDQLLAMRSTRAQRDDLLFKAFQFYMQQEDFKGAKAVLDEM QSYGRHE
KRVEECVQAYEIFGNNSYAYIDEMEA AFDEAPYALKVSYALMLAAQYTSKKDGEAAEKWQ
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>NODE_182329_length_1939_cov_4.566271_S6
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GRNRPSE RALLPSRSRHLRSRVGRTPHRTRNALLYRAYRTGELHDCRPGSNGARVRGKHR
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>NODE_212586_length_1033_cov_30.919651_S6
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QWGAQHCPKIIVTEKGRERKKFGWQSDAAGTEYRK FLEQFLPALRSALQGMGYPDEKVYY
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RVGGGIYTKAADVGADLVGKVEAGIPEDDPRNPAVIADNVGD NVGDVAGMGADLFESYVG
SVVSAVTLGLVAYNQEGAVFPLLIAALGIGASIIIGSFFVKGDEKSSPHKALKFGSYASSV
LVAVGSLALS YKFFGNLNAGMAIVFGLVVG LLIGLVTEIYTSDDYKFVKKIADQSETGAA
TTVISGIAVGMQ

>NODE_264747_length_1361_cov_29.963263_S6
GICQGGHSSRQPYHRLWLHRTGGYMIRLLLRRELSALFFLLIIFLIAGIVNPAFLTLNN
VFLSINSSVVYAVVAMGIAFVIITGEIDVSVGAIVGISATVVGSMIRDGQWPWLLALLAGI
GIGMLIGLINGFGVVTLRIPSIIMTLGTSSIIRGLMYVYTDGKWVENVPFEFKQLSQQKF
LDSFTYFYLAAILLFMLLVHLIMMRSKRGKYAAVGDNAAGANLLGIPVARTKLTAFVICG
VLSALGGVIFVSRVGFVTPIAGVGYEMKVIAACVIGGISLSSGGVGNILGACIGAAFMASI
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TV

>NODE_301074_length_916_cov_4.279476_S6
VVVGTMARSAELPLIIQIGATFNSIFGNFLGFCIPLIIIGFVVSIGIAELGDGAGKTLGLT
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TALLLSFIFGIGIAVTHSTSLKNGFSEVQHIIIEKLVAGVLIPLPLHVYGIFANMTYAGT
VMDIMSVFIRVFIIILLHVAVILIQYTIAGTVVGRNPIKLIIRMLPAYFTAIGTQSSAA
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FILMLGITMVAAPG

>NODE_313178_length_2508_cov_7.222488_S6
MLNKYGADATRWYLLHVSPAWSPTKFDEGGLQELASKFFGTLRNVYNFFVLYGNLDKIDV
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RRRFYTPGMSADKESVFATTFEVLEGVARLIAPIAPFISDEMYSKLTGEETVHIAYYPKT
NAALIDEKVEKRM DIVRSVCNLGRGIREKKGLKVRQPLSEILVDGKYKDLISDMIPLIMD
ELNVKQVVFADDELGEYMNFEKPNFKVAGPALGKKINTFAGVLAKEDA EKFTKLEKDGF
VTCKMDGEDFKIEKEFVDIGINAKQGF AVAMENNVFVIIDTNLSQELIDEGIAREVISKI
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>NODE_338494_length_1128_cov_14.833333_S6
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PHPAVPAGLALLPARDHRGGPSRRSERDRHLRAYVRAYNEVDLRGGRRRHFFERVEQLHV
AQDHPRRRQVPDDADARVQPRGRVRHRLRRPHARRPHRVAARDGGLPRPAALLRQRNHGI
SQVNTELSHLTDPTCFADNRLPAHSDHLWYATEAEVASGRSSFQVCLDGVWKLHYATNPS
QAVEGFEVPSYDVSEWDDIAVPAHLQLHGYDKPQYANIQYPWDGHEQLEPGQVPSRYNPT
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>NODE_377851_length_1918_cov_6.185089_S6
LRALARLDEAHRAARTHLPLETGRKDRIMTMSRRAFLSTCSGLGAAALAGCAPASGTD
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ALIGAGGADMGLTYQDYIANSLSANPLPYTAVAAVVQHNTSGIMSR AEDGIVRPKDMEG
HSYATWGLPIEQATVKQVVEEDGGDFSKVALVPYEVDDEVMGLQAGLFDTVWVYEWVAVQ
NAKLQEYPVNYFAFADISPQFDFYTPVIAANDAF AAADPELVRAFLRACEQGYELAATSP
ERAAEILCGAVPELDPALIAAAQASISPQYTADASRWGVIDRSRWTRFYEWLNDTGLVEN
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DRKKIALVMDHFPNKKDIKSAEQCKKCRTF AKRLDIENYDVGEMGIEHALLPEKGLVAS
GEAIIIGADSHTCTY GALGAFSTGVGSTDVTA AIATGKTWFKVPQAVRFVLRGALKPYVCG
KDVLHIIIGMIGVDGALYKSMEFTGDGVRSLTIDDRLTIANMAIEAGAKNGIFPVDSVTE
EYMAGRVT RPYKVCEADEDAEY EKYTNIDLSSIEPTVSFPHLPENTKAISECPDIEIDQV
IIGSCTNGRMQDMQAADILRGK HMAKVG RGVIPATMTVYKECIRLG YINDFIDAGCIV
STPTCGPCLGGYMGILADGERCVSTTNRNFVGRMGASGSEVYLAGPAVAAAASGIAGKIAD
PRKTL

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RLYELTNKIAKPAVSFGGKYRIIDFPLSNCANSNINIVGVLTQYESVFLNSYVTADARWG
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KMLEEHKASYADASIAVIEVPMKEASRFGIMNADATGRILEFEEKPEKPKSNLASMGIIYI
FNWVKVLRMLVSDQKNDLSSHDFGKDII PKMLDENKILHAYKFSGYWKDVGTVDVSFEAN
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ERENEIKAFQPQYEWTIHADVTGKKKFEAVLSKVDGKKPKLNNEEDSHVILERCKEGDF
IVGKRTKRAKKKQARIPFTTSTLQQEASTKLNFGARRTMSIAQKLYEGIDLGGQQEGLIS
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YLANEPACQRAERNFFDPYLQVRARLALLESNGHITDKIELIVLGGTWSDDYDPSYQIWF
SELFRLNDGDGEAERICAEERAFYRSCGLIAEADTLAEQTRDLQRCVTAGALSYNQAI
RLYASEAWVRARARQTATFGELEEQQRINESAHHRTVGLCVETRPDLVDDASAQLMRHLG
CTKVQMGIIQSLDQILDACGRHIRVEQIARAFSVLRHLHGFKILAHMMVNLVGSTPEHRL
DYGRLVGDPRFLPDEIKLYPCVLVESAAALRLYDQGIWRPYTEDELDDVLAADVAATPAY
VRISMIRDISSGDIVAGNKKTNLRQMVDARTEAAESAIAEIRSREIATGDVSACDVRD
CISYTTAVSEERFLQWITDAGSIAGFLRLSLPHGRSTAMIREVHIYGRVAELGSI EAGGA
QHLGLGSALVETACKQASAAGCSAINVISSVGTTRAYYRKLGFIDGLYQRRVLGT
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>NODE_1102966_length_2142_cov_5.032213_S6
WLRVAVPAVSRCEYLTPLLRVAVCVRCQFVTLPPPLASKADRKRDSRYSRERACELPACFLG
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LVRLVLRHSPFDLMAVTSTELSGRRLDEAYPAFAGQCDLAFSPHDADDLQSCDVVFLAVP
HTAALTFAFALIARGATVIDLSADFRLLKDPAYEEWYRVPHTPELLARAAFGFLPELFGE
ELAALAQRSSAGEVVLVACAGCYPTATSLAAAPVLRAGLSPAGLVVVDVAVSGVTGAGRKA
TERTHFCFANEGVEAYGVGAHRHTPEIEQILGLEGRLI FTPLAPYNRGLLSTVTMPVTR
GAFDQAEALAEYRSFFKDAPFVTVLPEGRQPRTVSVAGTNYAHVSACYNERAGAVVATCA
IDNIGKAAGQAVQCANIVCGLPETCGLDAVALPI
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SARS-CoV-2 related proteins:

```
>pdb|6YLA|A Chain A, SARS-CoV-2 RBD
ETGPNITNLCPFGEVFNATRFASVYAWNKRKISNCVADYSVLYNSASFSTFKCYGVSPTKLNDLCFTNVY
ADSFVIRGDEVQRQIAPGQTGKIADYNYKL PDDFTGCVIAWNSNNLDSKVGGNYNLYRLFRKSNLKPFER
DISTEIIYQAGSTPCNGVEGFNCYFPLQSYGFQPTNGVGYQPYRVVVL SFELLHAPATVCGPKKSTNKH
HHH
```

Or go to this URL:

<https://tinyurl.com/2f8h5vwf>

(https://www.ncbi.nlm.nih.gov/protein/?term=Severe+acute+respiratory+syndrome+coronavirus+2%5Borganism%5D+AND+protein_structure_direct%5Bfilt%5D)

click any SARS-CoV-2 related protein name and then click “FASTA” button at the top of the new page and use them to repeat exercise 1-3