## Aligned using an external algorithm

## Consensus

- **1.** B.1.1.7
- **2.** B.1.351
- 3. B.1.427
- **4.** B.1.429
- **5.** P.1

1

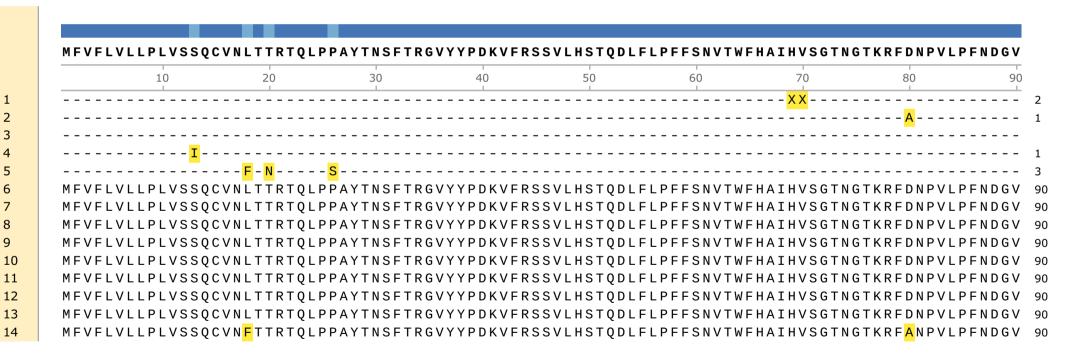
10

11

12

13

- **6.** Ref YP 009724390.1 surface glycoprotein [Severe acute respiratory syndrome coronavirus 2]
- 7. Ref OHD43416.1 surface glycoprotein [Severe acute respiratory syndrome coronavirus 2]
- 8. mRNA-1273 vaccine translated
- 9. BNT-162b2 vaccine translated
- 0. Ad26.COV2-S vaccine translated
- 1. NVX-CoV2373 vaccine translated
- 2. Sputnik V alleged unmodified YP 009724390.1
- 3. AZD1222 alleged unmodified YP 009724390.1
- **4.** AZD2816 modified YP 009724390.1 variants from B.1.351

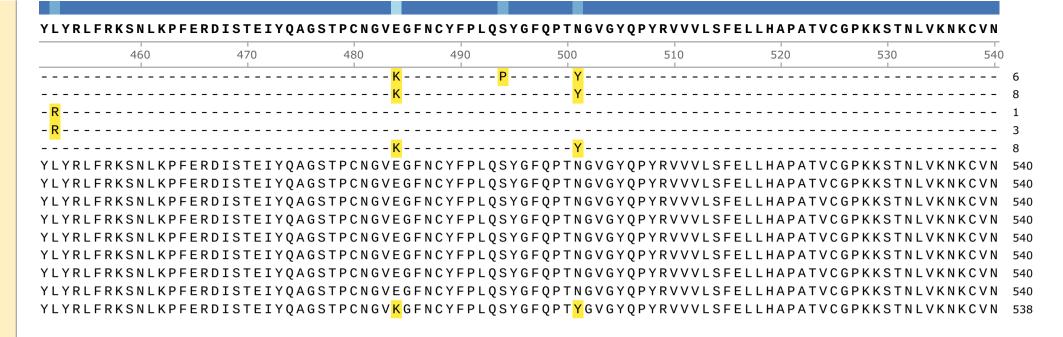


| 100          | 110        | 120        | 130         | 140              | 150                   | 160         | 170         | 1     |
|--------------|------------|------------|-------------|------------------|-----------------------|-------------|-------------|-------|
|              |            |            |             | <mark>X</mark> - |                       |             |             |       |
|              |            |            |             |                  |                       |             |             |       |
|              |            |            |             |                  |                       |             |             |       |
|              |            |            |             |                  |                       |             |             |       |
| STEKSNIIRGWI |            |            |             |                  |                       |             |             |       |
| STEKSNIIRGWI |            | •          | -           |                  |                       |             |             |       |
| STEKSNIIRGWI | -          |            | -           |                  |                       |             | _           |       |
| STEKSNIIRGWI | -          |            |             |                  |                       |             | =           |       |
| STEKSNIIRGWI | FGTTLDSKTQ | SLLIVNNATN | VVIKVCEFQF  | CNDPFLGVYY       | / H K N N K S W M E S | SEFRVYSSANN | ICTFEYVSQPI | FLMDL |
| STEKSNIIRGWI | FGTTLDSKTQ | SLLIVNNATN |             | CNDPFLGVYY       | / H K N N K S W M E S | SEFRVYSSANN | ICTFEYVSQPI | FLMDL |
| STEKSNIIRGWI | FGTTLDSKTQ | SLLIVNNATN |             | CNDPFLGVYY       | / H K N N K S W M E S | SEFRVYSSANN | ICTFEYVSQPI | FLMDL |
| STEKSNIIRGWI | FGTTLDSKTQ | SLLIVNNATN |             | CNDPFLGVYY       | / H K N N K S W M E S | SEFRVYSSANN | ICTFEYVSQPI | FLMDL |
| STEKSNITEGHT | ECTTLDSKTO | SILTVNNATA | IVVIKVCEFQF | CNDDELGVVV       | / H K N N K S W M E S | SEEDVVSSANA | ICTEEVVSOPI | ELMDI |

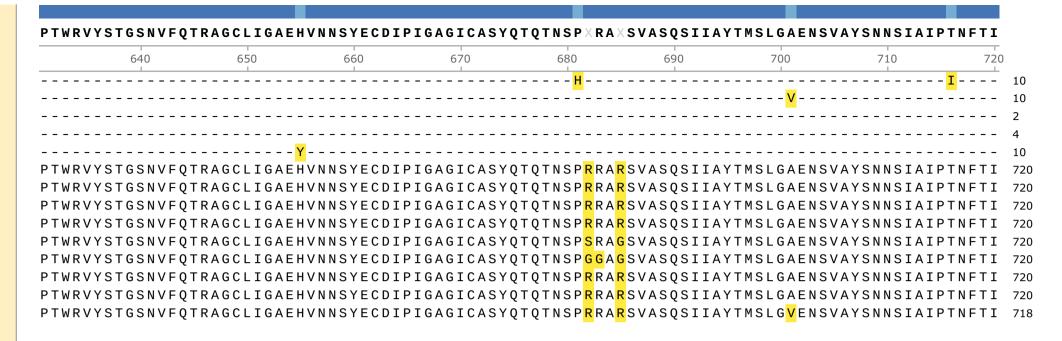
| 190            | 200         | 210        | 220       | 230         | 240        | 250        | 260       | 2     |
|----------------|-------------|------------|-----------|-------------|------------|------------|-----------|-------|
|                |             |            |           |             |            |            |           |       |
|                |             |            |           |             |            |            |           |       |
|                |             |            |           |             |            |            |           |       |
| <mark>S</mark> |             |            |           |             |            |            |           |       |
| NFKNLREFVFK    | NIDGYFKIYS  | KHTPINLVRD | LPQGFSALE | PLVDLPIGINI | TRFQTLLALH | RSYLTPGDSS | SGWTAGAAA | YYVGY |
| NFKNLREFVFK    | NIDGYFKIYS  | KHTPINLVRD | LPQGFSALE | PLVDLPIGINI | TRFQTLLALH | RSYLTPGDSS | SGWTAGAAA | YYVGY |
| NFKNLREFVFK    | NIDGYFKIYS  | KHTPINLVRD | LPQGFSALE | PLVDLPIGINI | TRFQTLLALH | RSYLTPGDSS | SGWTAGAAA | YYVGY |
| NFKNLREFVFK    | CNIDGYFKIYS | KHTPINLVRD | LPQGFSALE | PLVDLPIGINI | TRFQTLLALH | RSYLTPGDSS | SGWTAGAAA | YYVGY |
| NFKNLREFVFK    | CNIDGYFKIYS | KHTPINLVRD | LPQGFSALE | PLVDLPIGINI | TRFQTLLALH | RSYLTPGDSS | SGWTAGAAA | YYVGY |
| NFKNLREFVFK    | NIDGYFKIYS  | KHTPINLVRD | LPQGFSALE | PLVDLPIGINI | TRFQTLLALH | RSYLTPGDSS | SGWTAGAAA | YYVGY |
|                |             |            |           |             |            | RSYLTPGDSS | 001174044 |       |

|       | 280     | 290      | 30      |               | 310     |             | 320     | 330     |         | 340             | 350      | .30             |
|-------|---------|----------|---------|---------------|---------|-------------|---------|---------|---------|-----------------|----------|-----------------|
|       |         |          |         |               |         |             | 1       |         |         |                 |          |                 |
|       |         |          |         | . – – – – – – |         |             |         |         |         |                 |          |                 |
|       |         |          |         | . – – – – – – |         |             |         |         |         |                 |          |                 |
|       |         |          |         | . – – – – – – |         |             |         |         |         |                 |          |                 |
|       |         |          |         | . – – – – –   |         | . – – – – - |         |         |         |                 |          |                 |
| ·     | ·       |          |         |               | ·       | ·           |         |         |         |                 |          |                 |
|       |         |          |         |               |         | -           | -       |         |         |                 | RFASVYAV |                 |
| TFLLK | NENGTIT | DAVDCAL  | OPLSETK | CCTLKSFT      | VEKGIY  | 'QTSNFF     | ₹VQPTES | IVRFPN: | ĹTNLCPF | - G E V F N A T | RFASVYAV | √NRKRISN        |
| TFLLK | NENGTIT | DAVDCAL  | DPLSETK | CTLKSFT       | ·VEKGIY | 'QTSNFI     | RVQPTES | IVRFPN: | [TNLCPF | -GEVFNAT        | RFASVYAV | <b>VNRKRISN</b> |
| TFLLK | NENGTIT | DAVDCALI | DPLSETK | CTLKSFT       | VEKGIY  | 'QTSNFI     | RVQPTES | IVRFPN: | ITNLCPI | - G E V F N A T | RFASVYAV | NRKRISN         |
| TFLLK | NENGTIT | DAVDCALI | DPLSETK | CTLKSFT       | VEKGIY  | OTSNF1      | RVOPTES | IVRFPN: | ITNLCPI | - G E V F N A T | RFASVYAV | NNRKRISN        |
|       |         |          |         |               |         | _           | -       |         |         |                 | RFASVYAV |                 |
|       |         |          |         |               |         | _           | =       |         |         |                 | RFASVYAV |                 |
|       |         |          |         |               |         | -           | -       |         |         |                 | RFASVYAV |                 |

| 370         | 380         | 390                  | 400        | 410         | 420              | 430        | 440        | 45    |
|-------------|-------------|----------------------|------------|-------------|------------------|------------|------------|-------|
|             |             |                      |            |             |                  |            |            |       |
|             |             |                      |            |             | - N              |            |            |       |
|             |             |                      |            |             |                  |            |            |       |
|             |             |                      |            |             |                  |            |            |       |
|             |             |                      |            |             | - <mark>T</mark> |            |            |       |
| DYSVLYNSASF | STFKCYGVSP1 | <b>「KLNDLCFTNV</b>   | 'YADSFVIRG | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
| DYSVLYNSASF | STFKCYGVSP1 | <b>「KLNDLCFTNV</b>   | YADSFVIRGE | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
| DYSVLYNSASF | STFKCYGVSP1 | <b>FKLNDLCFTNV</b>   | YADSFVIRGE | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
| DYSVLYNSASF | STFKCYGVSP1 | <b>FKLNDLCFTNV</b>   | YADSFVIRGE | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
| DYSVLYNSASF | STFKCYGVSP1 | rk L N D L C F T N V | YADSFVIRGE | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
| DYSVLYNSASF | STFKCYGVSP1 | rk L N D L C F T N V | YADSFVIRGE | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
| DYSVLYNSASF | STFKCYGVSP1 | rk L N D L C F T N V | YADSFVIRGE | DEVRQIAPGQT | GKIADYNYKI       | PDDFTGCVIA | WNSNNLDSKV | GGNYN |
|             |             |                      |            | DEVRQIAPGQT |                  |            |            |       |
|             |             |                      |            | DEVRQIAPGQT |                  |            |            |       |



| 550          | 560         | 570            | 580         | 590        | 600        | 610                                       | 620       | 63                 |
|--------------|-------------|----------------|-------------|------------|------------|---|-----------|--------------------|
|              |             | <mark>D</mark> |             |            |            | <mark>G</mark> -                          |           |                    |
|              |             |                |             |            |            | <mark>G</mark> -                          |           |                    |
|              |             |                |             |            |            | <mark>G</mark> -                          |           |                    |
|              |             |                |             |            |            | <mark>G</mark> -                          |           |                    |
| NGLTGTGVLTES | SNKKFLPFQQF |                | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | <mark>G</mark> -<br>S N Q V A V L Y Q D V | NCTEVPVAI | HADQL <sup>-</sup> |
| NGLTGTGVLTES |             |                | =           |            |            |   |           | _                  |
| NGLTGTGVLTES | NKKFLPFQQF  | GRDIADTTDA     | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | SNQVAVLYQDV                               | NCTEVPVAI | HADQL              |
| NGLTGTGVLTES | NKKFLPFQQF  | GRDIADTTDA     | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | SNQVAVLYQDV                               | NCTEVPVAI | HADQL              |
| NGLTGTGVLTES | NKKFLPFQQF  | GRDIADTTDA     | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | SNQVAVLYQDV                               | NCTEVPVAI | HADQL              |
| IGLTGTGVLTES | NKKFLPFQQF  | GRDIADTTDA     | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | SNQVAVLYQDV                               | NCTEVPVAI | HADQL              |
| IGLTGTGVLTES | NKKFLPFQQF  | GRDIADTTDA     | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | SNQVAVLYQDV                               | NCTEVPVAI | HADQL              |
| NGLTGTGVLTES | NKKFLPFQQF  | GRDIADTTDA     | AVRDPQTLEIL | DITPCSFGGV | SVITPGTNTS | SNQVAVLYQDV                               | NCTEVPVAI | HADQL              |
| NGLTGTGVLTES | NKKFLPFOOF  | GRDIADTTDA     | AVRDPOTLEIL | DITPCSFGGV | SVITPGINIS | SNOVAVLYO <mark>G</mark> V                | NCTEVPVAT | HADOL.             |



| 730          | 740         | 750        | 760         | 770        | 780        | 790                     | 800              | 81    |
|--------------|-------------|------------|-------------|------------|------------|-------------------------|------------------|-------|
|              |             |            |             |            |            |                         |                  |       |
|              |             |            |             |            |            |                         |                  |       |
|              |             |            |             |            |            |                         |                  |       |
|              |             |            |             |            |            |                         |                  |       |
|              |             | STECSNILLO |             |            |            |                         | <br>DFGGFNFSQILF |       |
|              |             | _          | =           | =          | = = =      | =                       | DFGGFNFSQ1LF     |       |
|              |             | -          | •           | -          |            | -                       | DFGGFNFSQILF     |       |
|              |             | -          | •           | -          |            | -                       | DFGGFNFSQILF     |       |
|              |             | _          | =           | =          | = = =      | =                       | DFGGFNFSQILF     |       |
| TEILPVSMTKTS | SVDCTMYICGD | STECSNLLLQ | YGSFCTQLNR  | ALTGIAVEQD | KNTQEVFAQ\ | /KQIYKTPPIK             | DFGGFNFSQILF     | DPS   |
| TEILPVSMTKTS | SVDCTMYICGD | STECSNLLLQ | YGSFCTQLNR  | ALTGIAVEQD | KNTQEVFAQ\ | /KQIYKTPPIK             | DFGGFNFSQILF     | DPS   |
| TEILPVSMTKTS | SVDCTMYICGD | STECSNLLLQ | YGSFCTQLNR  | ALTGIAVEQD | KNTQEVFAQ\ | /KQIYKTPPIK             | DFGGFNFSQILF     | DPS   |
|              |             | OTEOCNILLO | WOOFOTOL NE |            | KNTOEVEAGY | /// O T V // T D D T // | DFGGFNFSQILF     | D D C |

| 90     | 890       | 880      | 870          | 860       | 850          | 840                 | 830        | 820         |
|--------|-----------|----------|--------------|-----------|--------------|---------------------|------------|-------------|
|        |           |          |              |           |              |                     |            |             |
|        |           |          |              |           |              |                     |            |             |
|        |           |          |              |           |              |                     |            |             |
|        |           |          |              |           |              |                     |            |             |
|        |           |          |              |           |              | (QYGDCLGDI <i>A</i> | VTLADAGFIK | KRSFIEDLLFN |
| _      |           |          | <del>-</del> |           | <del>-</del> | =                   |            | KRSFIEDLLFN |
| QIPFAM | WTFGAGAAL | AGTITSGV | TDEMIAQYTS   | NGLTVLPPL | ARDLICAQKF   | QYGDCLGDI <i>A</i>  | VTLADAGFI  | KRSFIEDLLFN |
| QIPFAM | WTFGAGAAL | AGTITSGV | TDEMIAQYTS   | NGLTVLPPL | ARDLICAQKF   | QYGDCLGDI <i>A</i>  | VTLADAGFI  | KRSFIEDLLFN |
| QIPFAM | WTFGAGAAL | AGTITSGV | TDEMIAQYTS   | NGLTVLPPL | ARDLICAQKF   | QYGDCLGDI <i>A</i>  | VTLADAGFI  | KRSFIEDLLFN |
| QIPFAM | WTFGAGAAL | AGTITSGV | TDEMIAQYTS   | NGLTVLPPL | ARDLICAQKF   | QYGDCLGDI <i>A</i>  | VTLADAGFI  | KRSFIEDLLFN |
| QIPFAN | WTFGAGAAL | AGTITSGV | TDEMIAQYTS   | NGLTVLPPL | ARDLICAQKF   | QYGDCLGDI <i>A</i>  | VTLADAGFI  | KRSFIEDLLFN |
| QIPFAN | WTFGAGAAL | AGTITSGV | TDEMIAQYTS   | NGLTVLPPL | ARDLICAQKF   | QYGDCLGDI <i>A</i>  | VTLADAGFI  | KRSFIEDLLFN |
| -      |           |          | TDEMIAQYTS   |           | -            | •                   |            |             |

| 910          | 920          | 930       | 940         | 950         | 960          | 970         | 980                 | 9                    |
|--------------|--------------|-----------|-------------|-------------|--------------|-------------|---------------------|----------------------|
|              |              |           |             |             |              |             | <mark>A</mark>      |                      |
|              |              |           |             |             |              |             |                     |                      |
|              |              |           |             |             |              |             |                     |                      |
|              |              |           |             |             |              |             |                     |                      |
| /RFNGIGVTQNV | 'LYENOKLIAN  | OFNSAIGKI | DDSLSSTASAI |             |              |             |                     |                      |
| 'RFNGIGVTQNV |              |           |             |             |              |             |                     |                      |
| 'RFNGIGVTQNV |              |           |             |             |              |             |                     |                      |
| /RFNGIGVTQNV | 'LYENQKLIAN  | QFNSAIGKI | QDSLSSTASAI | LGKLQDVVNQN | NAQALNTLVK   | QLSSNFGAISS | SVLNDILSRL          | D <mark>PP</mark> EA |
| /RFNGIGVTQNV | 'LYENQKLIAN  | QFNSAIGKI | QDSLSSTASAI | LGKLQDVVNQN | NAQALNTLVK   | QLSSNFGAISS | SVLNDILSRL          | D <mark>PP</mark> EA |
| 'RFNGIGVTQNV | 'LYENQKLIAN  | QFNSAIGKI | QDSLSSTASAI | LGKLQDVVNQN | NAQALNTLVK   | QLSSNFGAISS | SVLNDILSRL          | D P P E A            |
| / DENCTONY   | /I VENOKITAN | OENSATCKT | 1021221201  | CKIODYVNOV  | JAOAL NTI VK | QLSSNFGAISS | I O D I T O N I V S | DKVEV                |

| 108    | 1070                                  | 1060       | 1050         | 1040       | 1030     | 1020        | 1010                | 1000                       |
|--------|---------------------------------------|------------|--------------|------------|----------|-------------|---------------------|----------------------------|
|        |                                       |            |              |            |          |             |                     |                            |
|        |                                       |            |              |            |          |             |                     |                            |
|        |                                       |            |              |            |          |             |                     |                            |
|        |                                       |            |              |            |          |             |                     |                            |
|        |                                       |            |              |            |          |             |                     |                            |
|        |                                       |            |              |            |          |             |                     | DRLITGRLQSL                |
|        | =                                     |            | =            | =          |          |             |                     | DRLITGRLQSL<br>DRLITGRLQSL |
|        | · · · · · · · · · · · · · · · · · · · |            | <del>-</del> | =          |          |             |                     | <del>-</del>               |
|        | =                                     |            | =            | =          |          |             |                     | DRLITGRLQSL                |
| FTTAPA | VTYVPAQEKNI                           | SAPHGVVFLH | KGYHLMSFPQS  | GQSKRVDFCG | TKMSECVL | AAEIRASANLA | )                   | DRLITGRLQSL                |
| FTTAPA | VTYVPAQEKNI                           | SAPHGVVFLH | KGYHLMSFPQS  | GQSKRVDFCG | TKMSECVL | AAEIRASANLA | )TYVTQQLIR <i>A</i> | DRLITGRLQSL                |
| FTTAPA | VTYVPAQEKNI                           | SAPHGVVFLH | KGYHLMSFPQS  | GQSKRVDFCG | TKMSECVL | AAEIRASANLA | TYVTQQLIRA          | DRLITGRLQSL                |
| FTTAPA | VTYVPAQEKNI                           | SAPHGVVFLH | KGYHLMSFPQS  | GQSKRVDFCG | TKMSECVL | AAEIRASANLA | TYVTQQLIRA          | DRLITGRLQSL                |
|        | =                                     |            | =            | =          |          |             |                     | DRLITGRLQSL                |
|        | _                                     |            | -            | -          |          |             |                     | DRLITGRLQSL                |

| 1090   | 1100   | 1110        | 1120       | 1130                  | 1140         | 1150       | 1160        | 1170  |
|--|--|-------------|------------|-----------------------|--------------|------------|-------------|-------|
|  |  |             |            |                       |              |            |             |       |
|  |  |             |            |                       |              |            |             |       |
|  |  |             |            |                       |              |            |             |       |
|  |  |             |            |                       |              |            |             |       |
| DGKAHFPREGV  | FVSNGTHWFVT                                    | QRNFYEPQII  |            |                       |              |            |             |       |
| DGKAHFPREGV  |  | -           |            |                       | =            |            |             |       |
| DGKAHFPREGV  | F V S N G T H W F V T                          | QRNFYEPQII  | TTDNTFVSGN | NCDVVIGIVNN           | ITVYDPLQPEI  | DSFKEELDKY | 'FKNHTSPDVI | LGDIS |
| DGKAHFPREGV  | F V S N G T H W F V T                          | QRNFYEPQII  | TTDNTFVSGN | NCDVVIGIVNN           | ITVYDPLQPEI  | DSFKEELDKY | 'FKNHTSPDVI | LGDIS |
|  | FVSNGTHWFVT                                    | QRNFYEPQII  | TTDNTFVSGN | NCDVVIGIVNN           | ITVYDPLQPEI  | DSFKEELDKY | 'FKNHTSPDVI | LGDIS |
| DGKAHFPREGV  |  |             |            | 1 C D \               | ITVVDDI ODEI | DSEKEELDKY | 'FKNHTSPDVI | LGDIS |
|  |  | TQRNFYEPQII | TTDNTFVSGN | A C D A A T G T A W V | 1 1 V 1 D    |            |             |       |
| DGKAHFPREGV  | FVSNGTHWFVT                                    | -           |            |                       | -            |            |             |       |
| IDGKAHFPREGV<br>IDGKAHFPREGV<br>IDGKAHFPREGV<br>IDGKAHFPREGV | F V S N G T H W F V T<br>F V S N G T H W F V T | QRNFYEPQII  | TTDNTFVSGN | NCDVVIGIVNN           | ITVYDPLQPEI  | DSFKEELDKY | 'FKNHTSPDVI | LGDIS |

| 1260   | 1250      | 1240        | 1230         | 1220        | 1210       | 1200          | 1190            | 1180          |
|--------|-----------|-------------|--------------|-------------|------------|---------------|-----------------|---------------|
|        |           |             |              |             |            |               |                 |               |
|        |           |             |              |             |            |               |                 |               |
|        |           |             |              |             |            |               |                 |               |
|        |           |             |              |             |            |               |                 |               |
|        |           |             |              |             |            |               |                 |               |
|        |           |             |              |             |            |               | . – – – – – – . |               |
|        |           |             |              |             | •          | •             |                 | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | [AIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | CAIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | CAIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | CAIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | CAIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | CAIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KFDEDD | GCCSCGSCC | CCMTSCCSCLK | CAIVMVTIMLO  | /IWLGFIAGL: | YEQYIKWPW  | ESLIDLQELGK   | RLNEVAKNLNI     | NASVVNIQKEIDI |
| KEDEDD | GCCSCGSCC | COMTSCOSOLK | TATVMVTTMI ( | TWIGETAGI ' | YEOYTKWPWY | ESITDI OFI GK | PINEVAKNINI     | NASVVNTOKETDI |

GINASVVNIQKEIDRLNEVAKNLNESLIDLQELGKYEQYIKWPWYIWLGFIAGLIAIVMVTIMLCCMTSCCSCLKGCCSCGSCCKFDEDD

## SEPVLKGVKLHYT---

| 1270          |      |
|---------------|------|
|               | 13   |
|               | 10   |
|               | 2    |
|               | 4    |
|               | 11   |
| SEPVLKGVKLHYT | 1273 |
| SEPVLKGVKLHYT | 1271 |

**Consensus Threshold:** >50%

Compare to: the consensus

Amino acids that don't match the reference are marked with yellow highlighting.

**Created:** 28 Jun 2021

Last Modified: 28 Jun 2021

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