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|----------------|-----------|----------------|----------------|---------------------------------------------------------------------------------------|----|
| NTDB id 1159   | ABDM36    | RS04765        | WP 001166105.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLSSEKESKEVKEEKEEPL..... | 0  |
| NTDB id 1150   | A1552VC   | RS08580        | WP 001166105.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLSSEKESKEVKEEKEEPL..... | 0  |
| NTDB id 1395   | DSB67     | RS04445        | WP 010649155.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLSSEKESKEVKEEKEEPL..... | 0  |
| NTDB id 1362   | HI 1008   | AAC22669.1     |                | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLSSEKESKEVKEEKEEPL..... | 0  |
| NTDB id 1382   | A4U84     | RS07375        | WP 062924234.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLSSEKESKEVKEEKEEPL..... | 0  |
| NTDB id 595    | KW2       | RS08725        | WP 021037687.1 | .....MEKILDKLKEYWKIM.LLVVICALIAGGIFYIFTNSPKPAETLSVEN....LSSS...STKSSVSKFNSSSSEK       | 65 |
| NTDB id 521    | SMSK321   | RS04960        | WP 000443804.1 | .....MESIIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPHTPVKETNLQAE...VVAISKDSSTEKDVKKEEKEEPL        | 69 |
| NTDB id 493    | SM12261   | RS04300        | WP 000387351.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSVSEKEVVKKEEKEEPL       | 69 |
| NTDB id 263    | KZH43     | RS04195        | WP 000387330.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLVSEKEVNKEEKEEPL        | 69 |
| NTDB id 222    | SPD       | RS04535        | WP 000387330.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLVSEKEVNKEEKEEPL        | 69 |
| NTDB id 188    | SPR       | RS04295        | WP 000387330.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDLVSEKEVNKEEKEEPL        | 69 |
| NTDB id 153    | SP        | RS04730        | WP 000387344.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 69 |
| NTDB id 1069   | ABD1      | RS03060        | WP 043960841.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 1041   | HON27     | RS14740        | WP 005124713.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 105    | BSU 25590 | NP 390437.1    |                | .....MNWLNQHKKAIILAASAAVFTAIMIFLATG....KNKE....PVKQAVPT...ETENTVVVKQEANNDE            | 57 |
| NTDB id 36     | MW        | RS08230        | WP 001793467.1 | MVLLYQFLLRYKDFLTQWKLYIISAVVLMVLI GFIFWRQDDYTSRNFENKDNALKQSTSENSSLSKVEDVQVKDGDNSK      | 80 |
| NTDB id 20     | SA        | RS08005        | WP 001802022.1 | MVLLYQFLLRYKDFLTQWKLYIISAVVLMVLI GFIFWRQDDYTSRNFENKDTALKQSTSENSSLSKLEDVQVKDGDNSK      | 80 |
| NTDB id 1014   | ACIAD     | RS13855        | WP 004924483.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 104688 | R8661     | RS00060        | WP 002853093.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 1242   | Cj0011c   | YP 002343483.1 |                | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 1127   | NGFG      | RS10560        | WP 003703428.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 1126   | NGFG      | RS09375        | WP 003703428.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 1125   | NGFG      | RS07215        | WP 003703428.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| NTDB id 1108   | NGFG      | RS06525        | WP 003703428.1 | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL       | 0  |
| consensus      |           |                |                | .....MEALIEKIKKEYKLIYICTGLGLLVGGFFLLKPAPQTPVKETNLQAE...VVAISKDSSTEKEVVKKEEKEEPL.....  | 0  |

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| NTDB id 1159   | ABDM36    | RS04765        | WP 001166105.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 33  |
| NTDB id 1150   | A1552VC   | RS08580        | WP 001166105.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 33  |
| NTDB id 1395   | DSB67     | RS04445        | WP 010649155.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 25  |
| NTDB id 1362   | HI 1008   | AAC22669.1     |                | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 36  |
| NTDB id 1382   | A4U84     | RS07375        | WP 062924234.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 37  |
| NTDB id 595    | KW2       | RS08725        | WP 021037687.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 142 |
| NTDB id 521    | SMSK321   | RS04960        | WP 000443804.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 146 |
| NTDB id 493    | SM12261   | RS04300        | WP 000387351.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 146 |
| NTDB id 263    | KZH43     | RS04195        | WP 000387330.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 146 |
| NTDB id 222    | SPD       | RS04535        | WP 000387330.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 146 |
| NTDB id 188    | SPR       | RS04295        | WP 000387330.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 146 |
| NTDB id 153    | SP        | RS04730        | WP 000387344.1 | EQDLITVDVKGAVKSPGIYDLSPVGS.RVNDVAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAP.. | 146 |
| NTDB id 1069   | ABD1      | RS03060        | WP 043960841.1 | .....MQYSMRFWKNKYFVLLIMLCLGV.....ISPSVCAQSFQDNFQEWKVKQMQMYDQ                       | 48  |
| NTDB id 1041   | HON27     | RS14740        | WP 005124713.1 | .....MQYSMRFWKNKYFVLLIMLCLGV.....ISPSVCAQSFQDNFQEWKVKQMQMYDQ                       | 48  |
| NTDB id 105    | BSU 25590 | NP 390437.1    |                | SNETIVIDIKGAVQHGPVYEMRTGD.RVSQAIEKAGGTSEQADEAQVNLAELIQDGTIVVYIPKKEETAV.....Q       | 127 |
| NTDB id 36     | MW        | RS08230        | WP 001793467.1 | NKGPVYVDVKGAVKHPNVYKMTSKD.RVVDLLDKAQ..LLEDADVSQINLSEKLTQDKMIFIPH.....K.GQKNVEP     | 149 |
| NTDB id 20     | SA        | RS08005        | WP 001802022.1 | NKGPVYVDVKGAVKHPNVYKMTSKD.RVVDLLDKAQ..LLEDADVSQINLSEKLTQDKMIFIPH.....K.GQKNVEP     | 149 |
| NTDB id 1014   | ACIAD     | RS13855        | WP 004924483.1 | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 44  |
| NTDB id 104688 | R8661     | RS00060        | WP 002853093.1 | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 9   |
| NTDB id 1242   | Cj0011c   | YP 002343483.1 |                | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 9   |
| NTDB id 1127   | NGFG      | RS10560        | WP 003703428.1 | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 43  |
| NTDB id 1126   | NGFG      | RS09375        | WP 003703428.1 | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 43  |
| NTDB id 1125   | NGFG      | RS07215        | WP 003703428.1 | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 43  |
| NTDB id 1108   | NGFG      | RS06525        | WP 003703428.1 | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT                    | 43  |
| consensus      |           |                |                | .....MSIR.HFF...T....FFWAT....LL..TLFSMQIVFAQGFQDQYDAWKTQQKAHDT.....               | 43  |

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| NTDB id 1159   | ABDM36    | RS04765        | WP 001166105.1 | .....VEEGIVITVNINTASAEELATLLKGGIGLKKAAQIVDYREANGPFTHIDDLTNVKGIGEATV                | 94  |
| NTDB id 1150   | A1552VC   | RS08580        | WP 001166105.1 | .....VEEGIVITVNINTASAEELATLLKGGIGLKKAAQIVDYREANGPFTHIDDLTNVKGIGEATV                | 94  |
| NTDB id 1395   | DSB67     | RS04445        | WP 010649155.1 | .....KYEGIEITVNNSATAQEITALLIGIGEKKAAQDIVEYRNEHGPFKTAADLTQVKGIGEATV                 | 86  |
| NTDB id 1362   | HI 1008   | AAC22669.1     |                | QAEA.....QVAPAVVSDKLNINTATASEIQKSLTGIGAKKAAEIVQYREKHGNFXNAEQQLLEVQGIGKATL          | 103 |
| NTDB id 1382   | A4U84     | RS07375        | WP 062924234.1 | QQ.V.....AATQKVNANVVNINTATASEIQDKLVGIGAKKAAQIVVEYREKNGKFLNVEQLTEVSGIGKATL          | 103 |
| NTDB id 595    | KW2       | RS08725        | WP 021037687.1 | .....S.....S..VSNQISTTSGPKVNINKADLTTELQK.LTGIGQKKAQDIIDFRMKNNGDFKSIEDLGKVS         | 208 |
| NTDB id 521    | SMSK321   | RS04960        | WP 000443804.1 | .....SSTSCKDKKINLNKASLEELKQ.VKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGAKTI                | 206 |
| NTDB id 493    | SM12261   | RS04300        | WP 000387351.1 | .....SSTSKEKKVNLNKASLEELKQ.VKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGAKTI                 | 206 |
| NTDB id 263    | KZH43     | RS04195        | WP 000387330.1 | .....SSISKEKKVNLNKASLEELKQ.VKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTI                 | 206 |
| NTDB id 222    | SPD       | RS04535        | WP 000387330.1 | .....SSISKEKKVNLNKASLEELKQ.VKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTI                 | 206 |
| NTDB id 188    | SPR       | RS04295        | WP 000387330.1 | .....SSISKEKKVNLNKASLEELKQ.VKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTI                 | 206 |
| NTDB id 153    | SP        | RS04730        | WP 000387344.1 | .....SSTSKEKKVNLNKASLEELKQ.VKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTI                 | 206 |
| NTDB id 1069   | ABD1      | RS03060        | WP 043960841.1 | KLNISKSSSHSNSSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGIGEKKAAQIVVEYRQKNGSFKNIDEIKNVKGIGPAIF | 127 |
| NTDB id 1041   | HON27     | RS14740        | WP 005124713.1 | KLNITKSSHSNGSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGVGEKKAAQIVVEYRQKNGSFKNIDEIKNVKGIGPAIF  | 127 |
| NTDB id 105    | BSU 25590 | NP 390437.1    |                | QGGG.....GSV....QSDGGKGALVNINTATLEELQG.ISGVGPSKAAEIIAYREENGFRQTIEDITKVS            | 195 |
| NTDB id 36     | MW        | RS08230        | WP 001793467.1 | QIGVNSV.....HV....KNGNTNNTKVNLNTASVSELMS.VPGVGQAKANAIVEYRNQQGAFQEIDDLK             | 219 |
| NTDB id 20     | SA        | RS08005        | WP 001802022.1 | QIEVNSV.....HV....KNGNTNNTKVNLNTASVSELMS.VPGVGQAKANAIVEYRNQQGAFQEIDDLK             | 219 |
| NTDB id 1014   | ACIAD     | RS13855        | WP 004924483.1 | RLNSSASHHSLSKP...TQQSGTGKVNINSATAKELQQ.LHGVGEKKALAIVEYRQQYQKGFASVDDLQ              | 119 |
| NTDB id 104688 | R8661     | RS00060        | WP 002853093.1 | .....ALTAFLFGAVNINTATILKELKS.LNGIGEAKAKAILEYRKE.ANFTSIDDLKVKVGIGDKLF               | 69  |
| NTDB id 1242   | Cj0011c   | YP 002343483.1 |                | .....ALTAFLFGAVNINTATILKELKS.LNGIGEAKAKAILEYRKE.ANFTSIDDLKVKVGIGDKLF               | 69  |
| NTDB id 1127   | NGFG      | RS10560        | WP 003703428.1 | .....FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKVKVGIGPAVL                | 104 |
| NTDB id 1126   | NGFG      | RS09375        | WP 003703428.1 | .....FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKVKVGIGPAVL                | 104 |
| NTDB id 1125   | NGFG      | RS07215        | WP 003703428.1 | .....FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKVKVGIGPAVL                | 104 |
| NTDB id 1108   | NGFG      | RS06525        | WP 003703428.1 | .....FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKVKVGIGPAVL                | 104 |
| consensus      |           |                |                | ***!! !** !* * !!* **!!*!!*!!* ** !***** *!***!! ***                               |     |

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| NTDB id 1159   | ABDM36    | RS04765        | WP 001166105.1 | RNNAARILL.....              | 103 |
| NTDB id 1150   | A1552VC   | RS08580        | WP 001166105.1 | RNNAARILL.....              | 103 |
| NTDB id 1395   | DSB67     | RS04445        | WP 010649155.1 | RKNEDRILL.....              | 95  |
| NTDB id 1362   | HI 1008   | AAC22669.1     |                | EKNRDRRIIF.....             | 112 |
| NTDB id 1382   | A4U84     | RS07375        | WP 062924234.1 | EKNRDCIVIE.....             | 113 |
| NTDB id 595    | KW2       | RS08725        | WP 021037687.1 | EKLKDEISID.....             | 218 |
| NTDB id 521    | SMSK321   | RS04960        | WP 000443804.1 | EKLKDYVTVD.....             | 216 |
| NTDB id 493    | SM12261   | RS04300        | WP 000387351.1 | EKLKDYVTVD.....             | 216 |
| NTDB id 263    | KZH43     | RS04195        | WP 000387330.1 | EKLKDYVTVD.....             | 216 |
| NTDB id 222    | SPD       | RS04535        | WP 000387330.1 | EKLKDYVTVD.....             | 216 |
| NTDB id 188    | SPR       | RS04295        | WP 000387330.1 | EKLKDYVTVD.....             | 216 |
| NTDB id 153    | SP        | RS04730        | WP 000387344.1 | EKLKDYVTVD.....             | 216 |
| NTDB id 1069   | ABD1      | RS03060        | WP 043960841.1 | EKNKSRIAL.....              | 136 |
| NTDB id 1041   | HON27     | RS14740        | WP 005124713.1 | EKNKSRIAL.....              | 136 |
| NTDB id 105    | BSU 25590 | NP 390437.1    |                | EKLKSSITVK.....             | 205 |
| NTDB id 36     | MW        | RS08230        | WP 001793467.1 | DKLKSIFTI.....              | 228 |
| NTDB id 20     | SA        | RS08005        | WP 001802022.1 | DKLKSIFTI.....              | 228 |
| NTDB id 1014   | ACIAD     | RS13855        | WP 004924483.1 | EKNRDRRIAI.....             | 128 |
| NTDB id 104688 | R8661     | RS00060        | WP 002853093.1 | EKLKNDITIE.....             | 79  |
| NTDB id 1242   | Cj0011c   | YP 002343483.1 |                | EKLKNDITIE.....             | 79  |
| NTDB id 1127   | NGFG      | RS10560        | WP 003703428.1 | AKLKDQASVGAPAPKGPAPVLPVAVKK | 131 |
| NTDB id 1126   | NGFG      | RS09375        | WP 003703428.1 | AKLKDQASVGAPAPKGPAPVLPVAVKK | 131 |
| NTDB id 1125   | NGFG      | RS07215        | WP 003703428.1 | AKLKDQASVGAPAPKGPAPVLPVAVKK | 131 |
| NTDB id 1108   | NGFG      | RS06525        | WP 003703428.1 | AKLKDQASVGAPAPKGPAPVLPVAVKK | 131 |
| consensus      |           |                |                | ***** **                    |     |

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  $\geq 50\%$  conserved