```
S-ADENOSYL-L-HOMOCYSTEINE (5C8S-SAH, SARS)
    S-ADENOSYLMETHIONINE (5C8T-SAM, SARS)
QHN73794
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SP|P0C6X7|R1AB_CVHSA
                            VIFTOTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLOFTSLEIPRRNVATLOAE 5904
TR | Q6UZF5 | Q6UZF5_CVHSA
                            VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEIPRRNVATLQAE 5904
TR Q6UZF1 Q6UZF1_CVHSA
                            VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEIPRRNVATLQAE 5904
TR Q6JH48 Q6JH48 CVHSA
                            VIFTOTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLOFTSLEIPRRNVATLOAE 5904
TR | Q692E6 | Q692E6_CVHSA
                            VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEIPRRNVATLQAE 5904
TR A0A0K1YZY7 A0A0K1YZY7 CVHSA VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEVPRRNVATLQAE 5904
SP POC6W2 R1AB BCHK3
                            VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEVPRRNVATLQAE 5898
SP POC6W6 R1AB_BCRP3
                            VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEVPRRNVATLQAE 5902
SP POC6V9 R1AB BC279
                            VIFTOTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLOFTSLEVPRRNVATLQAE 5910
TR | A0A0U1WH14 | A0A0U1WH14 CVHSA VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEVPRRNVATLQAE 5899
TR | A0A0U1WHG0 | A0A0U1WHG0_CVHSA VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEVPRRNVATLQAE 5899
TR A0A166ZL34 A0A166ZL34 9NIDO VIFTQTTETAHSCNVNRFNVAITRAKIGILCIMSDRDLYDKLQFTSLEVPRRNVATLQAE 5706
TR R9QTB2 R9QTB2_CVHSA
                            ______
TR | R9QTH2 | R9QTH2_CVHSA
                            ______
SP POC6U8 R1A CVHSA
TR Q6JH47 Q6JH47_CVHSA
                            _____
TR | Q692E5 | Q692E5_CVHSA
SP POC6F8 R1A BCHK3
TR AOAOK1ZON1 AOAOK1ZON1_CVHSA ------
SP | P0C6F5 | R1A_BC279
SP POC6T7 R1A BCRP3
QHN73794
                            NVTGLFKDCSKVITGLHPTQAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5987
SP|P0C6X7|R1AB CVHSA
                            NVTGLFKDCSKIITGLHPTQAPTHLSVDIKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5964
TR Q6UZF5 Q6UZF5 CVHSA
TR Q6UZF1 Q6UZF1 CVHSA
                            NVTGLFKDCSKIITGLHPTOAPTHLSVDIKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5964
                            NVTGLFKDCSKIITGLHPTQAPTHLSVDIKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5964
TR Q6JH48 Q6JH48 CVHSA
                            NVTGLFKDCSKIITGLHPTQAPTHLSVDIKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5964
TR Q692E6 Q692E6_CVHSA
                            NVTGLFKDCSKIITGLHPTQAPTHLSVDIKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5964
TR A0A0K1YZY7 A0A0K1YZY7 CVHSA NVTGLFKDCSKIITGLHPTQAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5964
SP POC6W2 R1AB BCHK3
                            NVTGLFKDCSKIITGLHPTQAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5958
SP POC6W6 R1AB_BCRP3
                            NVTGLFKDCSKIITGLHPTOAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5962
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TR A0A0U1WHI4 A0A0U1WHI4_CVHSA NVTGLFKDCSKIITGLHPTQAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5959
TR A0A0U1WHG0 A0A0U1WHG0_CVHSA NVTGLFKDCSKIITGLHPTQAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5959
TR A0A166ZL34 A0A166ZL34 9NIDO NVTGLFKDCSKIITGLHPTQAPTHLSVDTKFKTEGLCVDIPGIPKDMTYRRLISMMGFKM 5766
TR R9QTB2 R9QTB2 CVHSA
                           -----
TR R9QTH2 R9QTH2_CVHSA
                            _____
SP|P0C6U8|R1A CVHSA
TR Q6JH47 Q6JH47 CVHSA
TR Q692E5 Q692E5_CVHSA
SP POC6F8 R1A BCHK3
TR A0A0K1Z0N1 A0A0K1Z0N1 CVHSA ------
SP | P0C6F5 | R1A_BC279
SP POC6T7 R1A BCRP3
QHN73794
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATREAVGTNLPLQLGFSTGVNLVAVPT 6047
SP|P0C6X7|R1AB CVHSA
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6024
TR Q6UZF5 Q6UZF5 CVHSA
TR Q6UZF1 Q6UZF1 CVHSA
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6024
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6024
TR Q6JH48 Q6JH48 CVHSA
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6024
TR | Q692E6 | Q692E6 CVHSA
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6024
TR A0A0K1YZY7 A0A0K1YZY7 CVHSA NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6024
SP POC6W2 R1AB BCHK3
                            NYOVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLOLGFSTGVNLVAVPT 6018
SP POC6W6 R1AB_BCRP3
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAIPT 6022
SP | P0C6V9 | R1AB_BC279
                            NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6030
TR A0A0U1WH14 A0A0U1WH14 CVHSA NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6019
TR A0A0U1WHG0 A0A0U1WHG0 CVHSA NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 6019
TR A0A166ZL34 A0A166ZL34 9NIDO NYQVNGYPNMFITREEAIRHVRAWIGFDVEGCHATRDAVGTNLPLQLGFSTGVNLVAVPT 5826
TR R9QTB2 R9QTB2_CVHSA
                            ______
TR | R9QTH2 | R9QTH2_CVHSA
SP POC6U8 R1A CVHSA
TR Q6JH47 Q6JH47_CVHSA
TR Q692E5 Q692E5_CVHSA
SP POC6F8 R1A BCHK3
TR AOAOK1ZON1 AOAOK1ZON1_CVHSA ------
SP | P0C6F5 | R1A_BC279
SP|P0C6T7|R1A BCRP3
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GUANOSINE-P3-ADENOSINE-5',5'-TRIPHOSPHATE (5C8S-G3A, SARS)

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OHN73794
                              GYVDTPNNTDFSRVSAKPPPGDOFKHLIPLMYKGLPWNVVRIKIVOMLSDTLKNLSDRVV 6107
SP|P0C6X7|R1AB CVHSA
                              GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWNVVRIKIVQMLSDTLKGLSDRVV 6084
TR | Q6UZF5 | Q6UZF5_CVHSA
                              GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWNVVRIKIVQMLSDTLKGLSDRVV 6084
TR Q6UZF1 Q6UZF1 CVHSA
                              GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWNVVRIKIVQMLSDTLKGLSDRVV 6084
TR Q6JH48 Q6JH48_CVHSA
                              GYVDTENNTEFTRVNAKPPPGDOFKHLIPLMYKGLPWNVVRIKIVOMLSDTLKGLSDRVV 6084
TR | Q692E6 | Q692E6 CVHSA
                              GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWNVVRIKIVQMLSDTLKGLSDRVV 6084
TR A O A O K 1 Y Z Y 7 | A O A O K 1 Y Z Y 7 C V H S A G Y V D T ENTEFTR V N A K P P P G D O F K H L I P L M Y K G L P W N V R I K I V Q M L S D T L K G L S D R V V 6084
SP POC6W2 R1AB BCHK3
                              GYVDTENSTEFTRVNAKPPPGDOFKHLIPLMYKGLPWNVVRIKIVOMLSDTLKGLSDRVV 6078
SP POC6W6 R1AB_BCRP3
                              GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWNVVRIKIVQMLSDTLKGLSDRVV 6082
SP | P0C6V9 | R1AB BC279
                              GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWNVVRIKIVQMLSDTLKGLSDRVV 6090
TR A0A0U1WHI4 A0A0U1WHI4 CVHSA GYVDTENNTEFTRVNAKPPPGDOFKHLIPLMYKGLPWNVVRIKIVOMLSDTLKGLSDRVV 6079
TR A0A0U1WHG0 A0A0U1WHG0 CVHSA GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWSVVRIKIVQMLSDTLKGLSDRVV 6079
TR A0A166ZL34 A0A166ZL34 9NIDO GYVDTENNTEFTRVNAKPPPGDQFKHLIPLMYKGLPWSVVRIKIVQMLSDTLKGLSDRVV 5886
TR R9QTB2 R9QTB2 CVHSA
                              ______
TR R9QTH2 R9QTH2_CVHSA
SP POC6U8 R1A CVHSA
TR Q6JH47 Q6JH47_CVHSA
TR Q692E5 Q692E5_CVHSA
SP POC6F8 R1A BCHK3
TR | A0A0K120N1 | A0A0K120N1_CVHSA ------
SP | P0C6F5 | R1A_BC279
SP POC6T7 R1A BCRP3
OHN73794
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDRRATCFSTASDTYACWHHSIGFDYVYNPFMI 6167
SP|P0C6X7|R1AB CVHSA
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6144
TR Q6UZF5 Q6UZF5_CVHSA
TR Q6UZF1 Q6UZF1_CVHSA
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6144
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6144
TR Q6JH48 Q6JH48 CVHSA
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6144
TR Q692E6 Q692E6_CVHSA
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6144
TR A0A0K1YZY7 A0A0K1YZY7 CVHSA FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6144
SP POC6W2 R1AB BCHK3
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6138
SP | P0C6W6 | R1AB_BCRP3
SP | P0C6V9 | R1AB_BC279
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6142
                              FVLWAHGFELTSMKYFVKIGPERTCCLCDRRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6150
TR|A0A0U1WHI4|A0A0U1WHI4 CVHSA FVLWAHGFELTSMKYFVKIGPERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6139
TR A0A0U1WHG0 A0A0U1WHG0_CVHSA FVLWAHGFELTSMKYFVKIGSERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 6139
TR A0A166ZL34 A0A166ZL34 9NIDO FVLWAHGFELTSMKYFVKIGSERTCCLCDKRATCFSTSSDTYACWNHSVGFDYVYNPFMI 5946
TR R9QTB2 R9QTB2 CVHSA
TR R9QTH2 R9QTH2 CVHSA
SP POC6U8 R1A CVHSA
TR Q6JH47 Q6JH47_CVHSA
TR Q692E5 Q692E5_CVHSA
SP|P0C6F8|R1A BCHK3
TR|A0A0K1Z0N1|A0A0K1Z0N1 CVHSA ------
SP | P0C6F5 | R1A_BC279
SP|P0C6T7|R1A BCRP3
OHN73794
                              DVQQWGFTGNLQSNHDLYCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWTIEYPIIGDE 6227
SP|P0C6X7|R1AB_CVHSA
                              DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6204
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                              DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6204
TR | O6UZF1 | O6UZF1 CVHSA
                              DVOOWGFTGNLOSNHDOHCOVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6204
TR Q6JH48 Q6JH48_CVHSA
                              DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6204
TR | Q692E6 | Q692E6 CVHSA
                              DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6204
TR | A0A0K1YZY7 | A0A0K1YZY7 CVHSA DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6204
SP POC6W2 R1AB_BCHK3
                              DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6198
SP|P0C6W6|R1AB BCRP3
                              DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6202
SP POC6V9 R1AB BC279
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TR A0A0U1WHI4 A0A0U1WHI4_CVHSA DVQQWGLTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPIIGDE 6199
TR A0A0U1WHG0 A0A0U1WHG0 CVHSA DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPVIGDE 6199
TR A0A166ZL34 A0A166ZL34 9NIDO DVQQWGFTGNLQSNHDQHCQVHGNAHVASCDAIMTRCLAVHECFVKRVDWSVEYPVIGDE 6006
TR R9QTB2 R9QTB2_CVHSA
                              ______
TR R9QTH2 R9QTH2_CVHSA
SP POC6U8 R1A CVHSA
TR | Q6JH47 | Q6JH47_CVHSA
TR Q692E5 Q692E5 CVHSA
SP POC6F8 R1A BCHK3
TR A0A0K1Z0N1 A0A0K1Z0N1_CVHSA ------
SP | P0C6F5 | R1A BC279
SP POC6T7 R1A BCRP3
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LKINAACRKVQHMVVKAALLADKFPVLHDIGNPKAIKCVPQADVEWKFYDAQPCSDKAYK 6287 OHN73794 SP|P0C6X7|R1AB CVHSA LRVNSACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQAEVEWKFYDAQPCSDKAYK 6264 TR Q6UZF5 Q6UZF5 CVHSA LRVNSACRKVOHMVVKSALLADKFPVLHDIGNPKAIKCVPOAEVEWKFYDAOPCSDKAYK 6264 TR Q6UZF1 Q6UZF1_CVHSA LRVNSACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQAEVEWKFYDAQPCSDKAYK 6264 TR Q6JH48 Q6JH48 CVHSA LRVNSACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQAEVEWKFYDAQPCSDKAYK 6264 TR | Q692E6 | Q692E6 CVHSA LRVNSACRKVOHMVVKSALLADKFPVLHDIGNPKAIKCVPOAEVEWKFYDAOPCSDKAYK 6264 TR A0A0K1YZY7 A0A0K1YZY7 CVHSA LKINSACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQAEVEWKFYDAQPCSDKAYK 6264 SP POC6W2 R1AB BCHK3 LKINAACRKVQHMVVKSALLADKFTVLHDIGNPKAIRCVPQAEVDWKFYDAQPCSDKAYK 6258 SP POC6W6 R1AB BCRP3 LKINSACRKVOHMVVKSALLADKFPVLHDIGNPKAIKCVPOAEVEWKFYDAOPCSDKAYK 6262 SP | P0C6V9 | R1AB_BC279 LKINAACRKVQHMVVKSALLADKFSVLHDIGNPKAIKCVPQAEVDWKFYDAQPCSDKAYK 6270 TR A0A0U1WHI4 A0A0U1WHI4 CVHSA LKINAACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQADVEWKFYDAQPCSDKAYK 6259 TR A0A0U1WHG0 A0A0U1WHG0 CVHSA LKINAACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQADVEWKFYDVQPCSDKAYK 6259 TR A0A166ZL34 A0A166ZL34 9NIDO LKINAACRKVQHMVVKSALLADKFPVLHDIGNPKAIKCVPQADVEWKFYDVQPCSDKAYK 6066 TR R9QTB2 R9QTB2 CVHSA ______ TR R9QTH2 R9QTH2 CVHSA _____ SP POC6U8 R1A_CVHSA ______ _____ TR | Q6JH47 | Q6JH47_CVHSA TR | Q692E5 | Q692E5 CVHSA SP POC6F8 R1A BCHK3 TR A0A0K1Z0N1 A0A0K1Z0N1 CVHSA ------SP | P0C6F5 | R1A_BC279 SP POC6T7 R1A BCRP3 QHN73794 IEELFYSYATHSDKFTDGVCLFWNCNVDRYPANSIVCRFDTRVLSNLNLPGCDGGSLYVN 6347 SP|P0C6X7|R1AB CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6324 TR Q6UZF5 Q6UZF5 CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6324 TR Q6UZF1 Q6UZF1 CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6324 TR | Q6JH48 | Q6JH48_CVHSA TR | Q692E6 | Q692E6_CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6324 IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6324 TR | A0A0K1YZY7 | A0A0K1YZY7 CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6324 SP POC6W2 R1AB_BCHK3 IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6318 SP POC6W6 R1AB_BCRP3 IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6322 SP POC6V9 R1AB BC279 IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6330 TR A0A0U1WHI4 A0A0U1WHI4 CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6319 TR A0A0U1WHG0 A0A0U1WHG0 CVHSA IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6319 TR A0A166ZL34 A0A166ZL34 9NIDO IEELFYSYATHHDKFTDGVCLFWNCNVDRYPANAIVCRFDTRVLSNLNLPGCDGGSLYVN 6126 TR R9QTB2 R9QTB2_CVHSA TR R9QTH2 R9QTH2_CVHSA _____ ______ SP POC6U8 R1A CVHSA TR | Q6JH47 | Q6JH47_CVHSA TR | Q692E5 | Q692E5_CVHSA SP POC6F8 R1A BCHK3 TR A0A0K1Z0N1 A0A0K1Z0N1_CVHSA ------SP | P0C6F5 | R1A_BC279 SP POC6T7 R1A BCRP3 QHN73794 KHAFHTPAFDKSAFVNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6407 KHAFHTPAFDKSAFTNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6384 SP|P0C6X7|R1AB_CVHSA TR Q6UZF5 Q6UZF5_CVHSA KHAFHTPAFDKSAFTNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6384 TR Q6UZF1 Q6UZF1 CVHSA KHAFHTPAFDKSAFTNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6384 KHAFHTPAFDKSAFTNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6384 TR Q6JH48 Q6JH48_CVHSA TR | Q692E6 | Q692E6 CVHSA KHAFHTPAFDKSAFTNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6384 TR A0A0K1YZY7 A0A0K1YZY7_CVHSA KHAFHTPAFDKSAFTNLKOLPFFYYSDSPCESHGKOVVSDIDYVPLKSATCITRCNLGGA 6384 KHAFHTPAFDKSAFTHLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6378 SP | P0C6W2 | R1AB_BCHK3 SP POC6W6 R1AB_BCRP3 KHAFHTPAFDKSAFTNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6382 SP POC6V9 R1AB BC279 KHAFHTPAFDKSAFTYLKOLPFFYYSDSPCESHGKOVVSDIDYVPLKSATCITRCNLGGA 6390 TR | A0A0U1WH14 | A0A0U1WH14 CVHSA KHAFHTPAFDKSAFSNLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6379 TR A0A0U1WHG0 A0A0U1WHG0 CVHSA KHAFHTPAFDKSAFTHLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6379 TR A0A166ZL34 A0A166ZL34 9NIDO KHAFYTPAFDKSAFTHLKQLPFFYYSDSPCESHGKQVVSDIDYVPLKSATCITRCNLGGA 6186 TR R9QTB2 R9QTB2_CVHSA TR R9QTH2 R9QTH2 CVHSA SP POC6U8 R1A CVHSA TR Q6JH47 Q6JH47_CVHSA TR | Q692E5 | Q692E5 CVHSA SP POC6F8 R1A BCHK3 TR|A0A0K1Z0N1|A0A0K1Z0N1 CVHSA ------SP | P0C6F5 | R1A BC279

SP POC6T7 R1A BCRP3

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SP P0C6X7 R1AB_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6444
TR Q6UZF5 Q6UZF5_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6444
TR Q6UZF1 Q6UZF1_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6444
TR Q6JH48 Q6JH48_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6444
TR Q692E6 Q692E6_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6444
TR A0A0K1YZY7 A0A0K1YZY7_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6444
SP POC6W2 R1AB_BCHK3	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTKLQSLENVAYNVVNKGHF	6438
SP POC6W6 R1AB_BCRP3	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6442
SP POC6V9 R1AB_BC279	VCRRHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVVNKGHF	6450
TR A0A0U1WHI4 A0A0U1WHI4_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVINKGHF	6439
TR A0A0U1WHG0 A0A0U1WHG0_CVHSA	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVINKGHF	6439
TR A0A166ZL34 A0A166ZL34 9NIDO	VCRHHANEYRQYLDAYNMMISAGFSLWIYKQFDTYNLWNTFTRLQSLENVAYNVINKGHF	6246
TR R9QTB2 R9QTB2_CVHSA		
TR R9QTH2 R9QTH2_CVHSA		
SP POC6U8 R1A_CVHSA		
TR Q6JH47 Q6JH47_CVHSA		
TR Q692E5 Q692E5_CVHSA		
SP P0C6F8 R1A_BCHK3		
TR A0A0K1Z0N1 A0A0K1Z0N1_CVHSA		
SP P0C6F5 R1A_BC279		
SP P0C6T7 R1A_BCRP3		