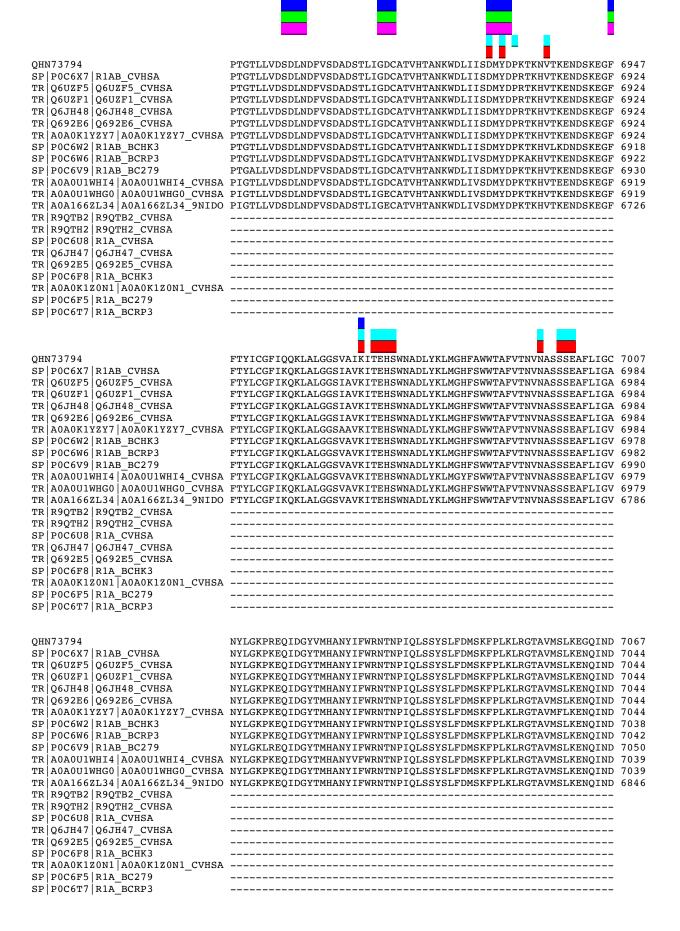
Ligand BS:

S-ADENOSYLMETHIONINE (5YN6-SAM, MERS)

S-ADENOSYL-L-HOMOCYSTEINE (5YN8-SAH, MERS)

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SINEFUNGIN (5YNB-SFG, MERS)
   P1-7-METHYLGUANOSINE-P3-ADENOSINE-5',5'-TRIPHOSPHATE (5YNF-GTA, MERS)
    7-METHYL-GUANOSINE-5'-TRIPHOSPHATE-5'-GUANOSINE (5YNI-GTG, MERS)
QHN73794
                              KVVKVTIDYTEISFMLWCKDGHVETFYPKLQSSQAWQPGVAMPNLYKMQRMLLEKCDLQN 6827
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6804
SP | P0C6X7 | R1AB_CVHSA
TR | Q6UZF5 | Q6UZF5_CVHSA
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6804
TR Q6UZF1 Q6UZF1 CVHSA
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6804
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6804
TR | Q6JH48 | Q6JH48_CVHSA
TR | Q692E6 | Q692E6 CVHSA
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6804
TR A0A0K1YZY7 A0A0K1YZY7 CVHSA KVVKVTIDYVEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6804
SP|P0C6W2|R1AB_BCHK3
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6798
SP POC6W6 R1AB_BCRP3
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6802
SP POC6V9 R1AB BC279
                              KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6810
TR A0A0UUWHI4 A0A0UUWHI4_CVHSA KVVKVTIDYAEISFMLWCKDGHVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6799
TR|A0A0U1WHG0|A0A0U1WHG0_CVHSA KVVKVTIDYAEISFMLWCKDGYVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6799
TR A0A166ZL34 A0A166ZL34 9NIDO KVVKVTIDYAEISFMLWCKDGYVETFYPKLQASQAWQPGVAMPNLYKMQRMLLEKCDLQN 6606
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
OHN73794
                              YGDSATLPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6887
                              YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6864
SP | P0C6X7 | R1AB_CVHSA
TR Q6UZF5 Q6UZF5_CVHSA
                              YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6864
TR Q6UZF1 Q6UZF1 CVHSA
                              YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6864
TR Q6JH48 Q6JH48 CVHSA
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TR | Q692E6 | Q692E6_CVHSA
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TR A0A0K1YZY7 A0A0K1YZY7 CVHSA YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6864
SP POC6W2 R1AB BCHK3
                              YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6858
SP | P0C6W6 | R1AB_BCRP3
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TR | A0A0U1WH14 | A0A0U1WH14_CVHSA YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6859
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TR A0A166ZL34 A0A166ZL34 9NIDO YGENAVIPKGIMMNVAKYTQLCQYLNTLTLAVPYNMRVIHFGAGSDKGVAPGTAVLRQWL 6666
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 POC6T7 R1A_BCRP3
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OHI	N73794	MILSLLSKGRLIIRENNRVVISSDVLVNN	7096
_	POC6X7 R1AB CVHSA	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7073
	Q6UZF5 Q6UZF5_CVHSA	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7073
TR	Q6UZF1 Q6UZF1_CVHSA	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7073
TR	Q6JH48 Q6JH48_CVHSA	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7073
TR	Q692E6 Q692E6 CVHSA	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7073
TR	A0A0K1YZY7 A0A0K1YZY7_CVHSA	MIYSLLEKGRLIIRENNTVVVSSDVLVNH	7073
SP	POC6W2 R1AB_BCHK3	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7067
SP	POC6W6 R1AB BCRP3	MIYSLLEKGRLIIRENNRVVVSSDILVNN	7071
SP	P0C6V9 R1AB BC279	MIYSLLENGRLIIRENNRVVVSSDILVNN	7079
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		MIYSLLEKGRLIVRENNRVIVSSDVLVNN	7068
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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	P0C6T7 R1A_BCRP3		