Ligand BS:

SP Q0Q468 | NCAP BC279

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N~2~,N~2~-DIMETHYL-N~1~-(6-OXO-5,6-DIHYDROPHENANTHRIDIN-2-YL)GLYCINAMIDE (4KXJ-P34, OC43)
    ADENOSINE MONOPHOSPHATE (4LI4-AMP, OC43)
    URIDINE-5'-MONOPHOSPHATE (4LM7-U5P, OC43)
     GUANOSINE-5'-MONOPHOSPHATE (4LM9-5GP, OC43)
    CYTIDINE-5'-MONOPHOSPHATE (4LMC-C5P, OC43)
    6-chloranyl-7-(2-morpholin-4-ylethylamino)quinoline-5,8-dione (4LMT-CQD,0C43)
CLUSTAL O(1.2.4) multiple sequence alignment
                                MSDNGPO-NORNAPRITFGGPSDSTGSNONGERSGARSKORRPOGLPNNTASWFTALTOH 59
YP 009724397
SP | P59595 | NCAP_CVHSA
                                MSDNGPQSNQRSAPRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTASWFTALTQH 60
TR | Q6UZF2 | Q6UZF2 CVHSA
                                MSDNGPQSNQRSAPRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTASWFTALTQH 60
TR Q6JH41 Q6JH41_CVHSA
                               MSDNGPQSNQRSAPRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTASWFTALTQH 60
TR Q6UZE8 Q6UZE8_CVHSA
                                MSDNGPQSNQRSAPRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTASWFTALTQH 60
TR Q692D7 Q692D7_CVHSA
TR R9QTB4 R9QTB4_CVHSA
                                MSDNGPQSNQRSAPRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTASWFTALTQH 60
                                MSDNGPO-NORSAPRITFGGPSDSTDNNODGGRSGARPKORRPOGLPNNTASWFTALTOH 59
SP | Q3I5I7 | NCAP_BCRP3
                                MSDNGPQ-NQRSAPRITFGGPTDSTDNNQDGGRSGARPKQRRPQGLPNNTASWFTALTQH 59
TR A0A0K1YZZ7 A0A0K1YZZ7 CVHSA MSDNGPH-NQRSASRITFGGPTDSTDNNQNGGRNGARPKQRRPQGLPNNTASWFTALTQH 59
TR R9QTA2 R9QTA2 CVHSA
                                {\tt MSDNGPQQNQRSAPRITFGGPTDSADNNQDGGRSGARPKQRRPQGLPNNTASWFTALTQH~60}
SP Q3LZX4 NCAP BCHK3
                                MSDNGPQ-SQRSAPRITFGGPADSNDNNQDGGRSGARPKQRRPQGLPNNTASWFTALTQH 59
TR A0A0U1WHJ1 A0A0U1WHJ1 CVHSA MSDNGPQ-NQRSAPRITFGGPSDSTDNNQDGGRSGVRPKQRRPQGLPNNTASWFTALTQH 59
SP | Q0Q468 | NCAP_BC279
                               MSDNGPQ-NQRSAPRITFGGPSDSTDNNQDGGRSGARPKQRRPQGLPNNTASWFTALTQH 59
TR A0A166ZLE5 A0A166ZLE5_9NIDO MSDNGTQ-NQRSASRITFGGPSDSTDNNQDGGRSGARPKQRRPQGLPNNTASWFTALTQH 59
TR A0A0U1UZD6 A0A0U1UZD6 CVHSA MSDNGTQ-NQRSASRITFGGPSDSTDNNQDGGRSGARPKQRRPQGLPNNTASWFTALTQH 59
                                ***** : .** *******
                                                          ..**:* *.*.* ******
YP 009724397
                                GKEDLKFPRGQGVPINTNSSPDDQIGYYRRATRRIRGGDGKMKDLSPRWYFYYLGTGPEA 119
SP|P59595|NCAP CVHSA
                                GKEELRFPRGQGVPINTNSGPDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 120
TR | Q6UZF2 | Q6UZF2_CVHSA
                                GKEELRFPRGQGVPINTNSGPDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 120
TR | Q6JH41 | Q6JH41 CVHSA
                                GKEELRFPRGQGVPINTNSGPDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 120
TR Q6UZE8 Q6UZE8_CVHSA
                                GKEELRFPRGQGVPINTNSGPDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 120
TR Q692D7 Q692D7_CVHSA
                                GKEELRFPRGQGVPINTNSGPDDQIGFYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 120
TR R9QTB4 R9QTB4 CVHSA
                                GKEELRFPRGQGVPINTNSGKDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119
SP|Q3I5I7|NCAP_BCRP3
                                GKEELRFPRGQGVPINTNSGKDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119
TR A0A0K1YZZ7 A0A0K1YZZ7 CVHSA GKEELRFPQGQGVPINTNSGPDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119
TR R9QTA2 R9QTA2 CVHSA
                                GKEELRFPRGQGVPINTNSGKDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 120
SP | Q3LZX4 | NCAP_BCHK3
                                GKEELRFPRGQGVPINTNSGKDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119
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TR A0A0U1WHJ1 A0A0U1WHJ1_CVHSA GKEGLKFPQGQGVPINTNSGRDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119

TR|A0A166ZLE5|A0A166ZLE5_9NIDO GKEGLKFPQGQGVPINTNSGTDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119
TR|A0A0U1UZD6|A0A0U1UZD6_CVHSA GKEGLKFPQGQGVPINTNSGRDDQIGYYRRATRRVRGGDGKMKELSPRWYFYYLGTGPEA 119

GKEELRFPRGQGVPINTNSGKDDQIGYYRRATRRVRGGDGKMKKLSPRWYFYYLGTGPEA 119

	1 1	
	•	
YP_009724397	GLPYGANKDGIIWVATEGALNTPKDHIGTRNPANNAAIVLQLPQGTTLPKGFYAEGSRGG	
SP P59595 NCAP_CVHSA TR Q6UZF2 Q6UZF2 CVHSA	SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAATVLQLPQGTTLPKGFYAEGSRGG SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAATVLQLPQGTTLPKGFYAEGSRGG	
TR Q6JH41 Q6JH41_CVHSA	SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAATVLQLPQGTTLPKGFYAEGSRGG	
TR Q6UZE8 Q6UZE8_CVHSA	${\tt SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAATVLQLPQGTTLPKGFYAEGSRGG}$	
TR Q692D7 Q692D7_CVHSA TR R9QTB4 R9QTB4 CVHSA	SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAATVLQLPQGTTLPKGFYAEGSRGG SLPYGANKEGIIWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRGG	
SP Q31517 NCAP_BCRP3	SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRGG	
· · · · · · —	${\tt SLPYGANKEGIIWVATEGALNTPKDHIGTRNPNNNAATVLQLPQGTTLPKGFYAEGSRGG}$	
TR R9QTA2 R9QTA2_CVHSA SP O3LZX4 NCAP BCHK3	SLPYGANKEGIVWVAIEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRGG SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRGG	
1~ 1 _	SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRNG	
SP Q0Q468 NCAP_BC279	${\tt SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRGG}$	
· · · · · · · · · · · · · · · · · · ·	SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRSG SLPYGANKEGIVWVATEGALNTPKDHIGTRNPNNNAAIVLQLPQGTTLPKGFYAEGSRSG	
IK AOAOOTOZDO AOAOOTOZDO_CVIIDA	**************************************	1/)
VD 000724207	COACCDCCCDCDNCCDNCCDCCCCCCCCCDADADACNCCCDAATATITIDDINOLECUMCCUCO	220
YP_009724397 SP P59595 NCAP_CVHSA	SQASSRSSRSRNSSRNSTPGSSRGTSPARMAGNGGDAALALLLLDRLNQLESKMSGKGQ SQASSRSSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ	
TR Q6UZF2 Q6UZF2_CVHSA	SQASSRSSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ	
TR Q6JH41 Q6JH41_CVHSA	SQASSRSSRRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ	
TR Q6UZE8 Q6UZE8_CVHSA TR Q692D7 Q692D7 CVHSA	SQASSRSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ SQASSRSSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ	
TR R9QTB4 R9QTB4 CVHSA	SQASSRSSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ	
SP Q3I5I7 NCAP_BCRP3	SQASSRSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGRSQ	239
	SQASSRSSRRGNSRNSTPGSSRGTSPARIASGGGETALALLLLDRLNQLESKVSGKGQ	
TR R9QTA2 R9QTA2_CVHSA SP Q3LZX4 NCAP BCHK3	SQASSRSSRSRGNSRNSTPGSSRGNSPARMASGGGETALALLLLDRLNQLESKVSGKGQ SQSSSRSSSRSRGNSRNSTPGSSRGSSPARLASGGGETALALLLLDRLNQLESKVSGKGQ	
· · · · · · · · · · · · · · · · · · ·	SQASSRSSSRSRGNSRTSTPGSSRGNSPARVASGGGETALALLLLDRLNQLESKVSGKGQ	
SP Q0Q468 NCAP_BC279	${\tt SQASSRSSRSRGNSRNSTPGSSRGNSPARMASGSGETALALLLLDRLNQLESKVSGKGQ}$	239
	SQASSRSSRRGNSRTSTPGSSRGNSPARVASGGGETALALLLLDRLNQLESKVSGKGQ	
TR AUAUU1UZD6 AUAUU1UZD6_CVHSA	SQASSRSSRSRGNSRISTPGSSRGNSPARVASGGGETALALLLLDRLNQLESKVSGKGQ **:**********************************	239
VD 000724207		200
YP_009724397 SP P59595 NCAP_CVHSA	QQQGQTVTKKSAAEASKKPRQKRTATKAYNVTQAFGRRGPEQTQGNFGDQELIRQGTDYK QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK	
TR Q6UZF2 Q6UZF2_CVHSA	QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK	
TR Q6JH41 Q6JH41_CVHSA	$\tt QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK$	
TR Q6UZE8 Q6UZE8_CVHSA TR Q692D7 Q692D7_CVHSA	QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK	
TR R9QTB4 R9QTB4 CVHSA	QQQGQTVTKKSAAEASKKPRQKRTATKSYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK	
SP Q3I5I7 NCAP_BCRP3	QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQELIRQGTDYK	
	QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPDQTQGNFGDQDLIRQGTDYK	
TR R9QTA2 R9QTA2_CVHSA SP Q3LZX4 NCAP BCHK3	QQQGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQELIRQGTDYK QQPGQTVTKKSAAEASKKPRQKRTATKQYNVTQAFGRRGPEQTQGNFGDQELIRQGIDYK	
	QQQGQTVTKKSASEASKKPRQKRTATKQYNVTQAFGRRGPDQTQGNFGDQELIRQGTDYK	
SP Q0Q468 NCAP_BC279	QQQGQTVTKKSAAEASKKPRQKRTATKSYNVTQAFGRRGPEQTQGNFGDQDLIRQGTDYK	299
	QQQGQTVTKKSASEASKKPRQKRTATKQYNVTQAFGRRGPDQTQGNFGDQELIRQGTDYK	
TR AUAUU1UZD6 AUAUU1UZD6_CVHSA	QQQGQTVTKKSASEASKKPRQKRTATKQYNVTQAFGRRGPDQTQGNFGDQELIRQGTDYK ** **********************************	299
VD 000724207	HINDOTA OF A DCA CA FEGMENT CMENTING CORRUST THE NOVE DATE OF DATE OF THE CASE	250
YP_009724397 SP P59595 NCAP_CVHSA	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYTGAIKLDDKDPNFKDQVILLNKHIDA HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
TR Q6UZF2 Q6UZF2_CVHSA	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
TR Q6JH41 Q6JH41_CVHSA	${\tt HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA}$	
TR Q 6 U Z E 8 Q 6 U Z E 8 C V H S A	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
TR Q692D7 Q692D7_CVHSA TR R9QTB4 R9QTB4 CVHSA	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
SP Q31517 NCAP_BCRP3	${\tt HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA}$	359
· · · · · —	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
TR R9QTA2 R9QTA2_CVHSA SP Q3LZX4 NCAP BCHK3	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
SP Q0Q468 NCAP_BC279	${\tt YWPQIAQFAPSASAFFGMSRIGMEVTPLGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA}$	359
	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA	
IK AUAUUIUZDO AUAUUIUZDO_CVHSA	HWPQIAQFAPSASAFFGMSRIGMEVTPSGTWLTYHGAIKLDDKDPQFKDNVILLNKHIDA :************************************	309
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YP 009724397
                              YKTFPPTEPKKDKKKKADETOALPOROKKOOTVTLLPAADLDDFSKOLOOSMSSADSTOA 419
SP|P59595|NCAP CVHSA
                              YKTFPPTEPKKDKKKKTDEAQPLPQRQKKQPTVTLLPAADMDDFSRQLQNSMSGASADST 420
TR Q6UZF2 Q6UZF2 CVHSA
                              YKTFPPTEPKKDKKKKTDEAQPLPQRQKKQPTVTLLPAADMDDFSRQLQNSMSGASADST 420
TR Q6JH41 Q6JH41_CVHSA
                              YKTFPPTEPKKDKKKKTDEAQPLPQRQKKQPTVTLLPAADMDDFSRQLQNSMSGASADST 420
TR | Q6UZE8 | Q6UZE8 CVHSA
                              YKTFPPTEPKKDKKKKTDEAQPLPQRQKKQPTVTLLPAADMDDFSRQLQNSMSGASADST 420
TR | Q692D7 | Q692D7 CVHSA
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TR R9QTA2 R9QTA2_CVHSA
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TR A0A0U1WHJ1 A0A0U1WHJ1_CVHSA YKTFPPTEPKKDKKKKTDEAQPLPQRQKKQPTVTLLPAADMDDFSRQLQNSMSGASADST 419
                              YKAFPPTEPKKDKKKKTDEAQPLPQRK-KQPTVTLLPAADMDDFSRQLQNSMSGASADST 418
SP Q0Q468 NCAP_BC279
TR | A0A166ZLE5 | A0A166ZLE5 | 9NIDO YKTFPPTEPKKDKKKKTDEAQPLPQRK-KQPTVTLLPAADMDDFSRQLQNSMSGASADST 418
TR A0A0U1UZD6 A0A0U1UZD6_CVHSA YKTFPPTEPKKDKKKKTDEAQPLPQRK-KQPTVTLLPAADMDDFSRQLQNSMSGASADST 418
                              YP 009724397
SP|P59595|NCAP CVHSA
                              QA 422
TR Q6UZF2 Q6UZF2_CVHSA
                              QA 422
TR Q6JH41 Q6JH41 CVHSA
                              OA 422
TR | Q6UZE8 | Q6UZE8_CVHSA
                              QA 422
TR Q692D7 Q692D7_CVHSA
                              QA 422
TR R9QTB4 R9QTB4 CVHSA
                              OA 420
SP Q31517 NCAP_BCRP3
                              QA 421
TR A0A0K1YZZ7 A0A0K1YZZ7 CVHSA QA 421
TR R9QTA2 R9QTA2_CVHSA
                              OA 422
SP Q3LZX4 NCAP_BCHK3
                              QA 421
TR A0A0U1WHJ1 A0A0U1WHJ1 CVHSA QA 421
SP Q0Q468 NCAP BC279
                              QA 420
TR A0A166ZLE5 A0A166ZLE5 9NIDO QA 420
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TR A0A0U1UZD6 A0A0U1UZD6 CVHSA OA 420