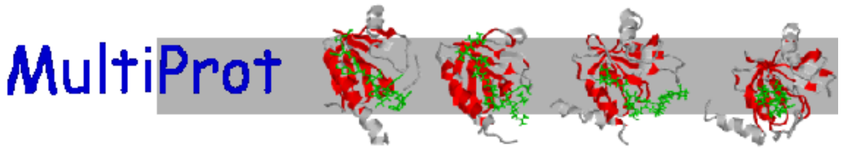

Number of aligned molecules : 2

Alignment Size	RMSD	Molecules	PDB alignment
86	1.54	alpha omega	aligned.pdb

Alignment syntaxis: *ChainID.AminoAcidCode.ResidueNumber*

alpha	omega
A M 1	A M 1
A K 2	A K 2
A G 3	A G 3
A M 4	A M 4
A L 5	A L 5
A T 6	A T 6
A P 8	A V 9
A V 9	A T 10
A T 10	A I 11
A I 11	A L 12
A L 12	A N 13
A N 13	A W 14
A S 15	A S 15
A W 16	A W 16
A P 17	A P 17
A R 18	A R 18
A E 19	A E 19
A D 20	A D 20
A I 21	A I 21
A T 22	A T 22
A H 23	A H 23
A E 24	A E 24
A E 25	A E 25
A Q 26	A Q 26
A T 27	A T 27
A K 28	A K 28
A Q 29	A Q 29
A L 30	A L 30
A A 31	A A 31
A L 32	A L 32
A A 33	A A 33
A I 34	A I 34
A R 35	A R 35
A D 36	A D 36
A E 37	A E 37
A V 38	A V 38
A L 39	A L 39
A D 40	A D 40
A L 41	A L 41
A E 42	A E 42
A A 43	A A 43
A A 44	A A 44
A G 45	A G 45
A I 46	A I 46
A K 47	A K 47
A I 48	A I 48
A I 49	A I 49
A Q 50	A Q 50
A I 51	A I 51
A D 52	A D 52
A E 53	A E 53
A A 55	A A 54
A L 56	A A 55
A R 57	A L 56
A E 58	A L 60
A L 60	A L 62
A L 62	A K 64
A R 63	A S 65
A D 66	A D 66
A H 68	A A 69
A K 70	A K 70
A Y 71	A Y 71
A L 72	A L 72
A D 73	A D 73
A W 74	A W 74
A A 75	A A 75
A I 76	A I 76
A P 77	A P 77
A A 78	A A 78
A F 79	A F 79
A R 80	A R 80
A L 81	A L 81
A V 82	A V 82
A H 83	A H 83
A S 84	A S 84
A A 85	A A 85
A V 86	A V 86
A K 87	A K 87
A P 88	A P 88
A T 89	A T 89
A T 90	A T 90
A Q 91	A Q 91
A I 92	A I 92
A H 93	A H 93
A T 94	A H 95
A M 96	A C 97