

CLUSTAL O(1.2.4) multiple sequence alignment

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A0A086BTX7|A0A086BTX7_PSEAI      -MSLELQEQCSSLKPYNSTFGIDVRARLLAHARGEADVREALALARQSLPLLVIGGSNLL  59
W1MJW5|W1MJW5_PSEAI              -MSLELQEHCSLKPYNSTFGIDVRARLLAHARDEADVREALALARERGLPLLVIGGSNLL  59
A0A0H2ZD69|A0A0H2ZD69_PSEAB     -MSLELQEHCSLKPYNSTFGIDVRARLLAHAQDEADVREALALARERGLPLLVIGGSNLL  59
Q9HZM7|MURB_PSEAE                 -MSLELQEHCSLKPYNSTFGIDVRARLLAHARDEADVREALALARERGLPLLVIGGSNLL  59
4JAY                               SMSLELQEHCSLKPYNSTFGIDVRARLLAHARDEADVREALALARERGLPLLVIGGSNLL  60
A0A0E1AVK6|A0A0E1AVK6_PSEAI     -MSLELQEHCSLKPYNSTFGIDVRARLLAHARDEADVREALALARERGLPLLVIGGSNLL  59
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A0A086BTX7|A0A086BTX7_PSEAI      LTRDVEALVLRMASQGRRLSEADDMVLVEAEAGETWDPFVQWSLEQGLAGLENLSLIPG  119
W1MJW5|W1MJW5_PSEAI              LTRDVEALVLRMASQGRRIVSDAADSVLVEAEAGEAWDPFVQWSLERGLAGLENLSLIPG  119
A0A0H2ZD69|A0A0H2ZD69_PSEAB     LTRDVEALVLRMASQGRRIVSDAADSVLVEAEAGEAWDPFVQWSLERGLAGLENLSLIPG  119
Q9HZM7|MURB_PSEAE                 LTRDVEALVLRMASQGRRIVSDAADSVLVEAEAGEAWDPFVQWSLERGLAGLENLSLIPG  119
4JAY                               LTRDVEALVLRMASQGRRIVSDAADSVLVEAEAGEAWDPFVQWSLERGLAGLENLSLIPG  120
A0A0E1AVK6|A0A0E1AVK6_PSEAI     LTRDVEALVLRMASQGRRIVSDAADSVLVEAEAGEAWDPFVQWSLERGLAGLENLSLIPG  119
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A0A086BTX7|A0A086BTX7_PSEAI      TVGAAPMQNIGAYGVELKDVFDLSLTALDRQDGSLEFDRQACRFGYRDSLQEPDRWLI  179
W1MJW5|W1MJW5_PSEAI              TVGAAPMQNIGAYGVELKDVFDLSLTALDRQDGTLEFDRQACRFGYRDSLQEPDRWLI  179
A0A0H2ZD69|A0A0H2ZD69_PSEAB     TVGAAPMQNIGAYGVELKDVFDLSLTALDRQDGTLEFDRQACRFGYRDSLQEPDRWLI  179
Q9HZM7|MURB_PSEAE                 TVGAAPMQNIGAYGVELKDVFDLSLTALDRQDGTLEFDRQACRFGYRDSLQEPDRWLI  179
4JAY                               TVGAAPMQNIGAYGVELKDVFDLSLTALDRQDGTLEFDRQACRFGYRDSLQEPDRWLI  180
A0A0E1AVK6|A0A0E1AVK6_PSEAI     TVGAAPMQNIGAYGVELKDVFDLSLTALDRQDGTLEFDRQACRFGYRDSLQEPDRWLI  179
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A0A086BTX7|A0A086BTX7_PSEAI      LRVRLRLRRRSLHLDYGPVQRLEEEGIASPTARDVSRVICAIRREKLPDPAVLGNAGS  239
W1MJW5|W1MJW5_PSEAI              LRVRLRLRRRERLHLDYGPVQRLEEEGIASPTARDVSRVICAIRREKLPDPAVLGNAGS  239
A0A0H2ZD69|A0A0H2ZD69_PSEAB     LRVRLRLRRRERLHLDYGPVQRLEEEGIASPTARDVSRVICAIRREKLPDPAVLGNAGS  239
Q9HZM7|MURB_PSEAE                 LRVRLRLRRRERLHLDYGPVQRLEEEGIASPTARDVSRVICAIRREKLPDPAVLGNAGS  239
4JAY                               LRVRLRLRRRERLHLDYGPVQRLEEEGIASPTARDVSRVICAIRREKLPDPAVLGNAGS  240
A0A0E1AVK6|A0A0E1AVK6_PSEAI     LRVRLRLRRRERLHLDYGPVQRLEEEGIASPTARDVSRVICAIRREKLPDPAVLGNAGS  239
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A0A086BTX7|A0A086BTX7_PSEAI      FFKNPLVSATQAAMLRRTPDLVAYPQADGQVKLAAGWLIDKGGWKGFRDGPVGVHAQQA  299
W1MJW5|W1MJW5_PSEAI              FFKNPLVDAAQAERLRQAFPDLVGYPPQADGRLKLAAGWLIDKGGWKGFRDGPVGVHAQQA  299
A0A0H2ZD69|A0A0H2ZD69_PSEAB     FFKNPLVDAAQAERLRQAFPDLVGYPPQADGRLKLAAGWLIDKGGWKGFRDGPVGVHAQQA  299
Q9HZM7|MURB_PSEAE                 FFKNPLVDATQAERLRQAFPDLVGYPPQADGRLKLAAGWLIDKGGWKGFRDGPVGVHAQQA  299
4JAY                               FFKNPLVDATQAERLRQAFPDLVGYPPQADGRLKLAAGWLIDKGGWKGFRDGPVGVHAQQA  300
A0A0E1AVK6|A0A0E1AVK6_PSEAI     FFKNPLVDAAQAERLRQAFPDLVGYPPQADGRLKLAAGWLIDKGGWKGFRDGPVGVHAQQA  299
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A0A086BTX7|A0A086BTX7_PSEAI      LVLVNHGGASGAQVQALAERIQADVRFVLELEPEPNLY  339
W1MJW5|W1MJW5_PSEAI              LVLVNHGGATGAQVQALAERIQEDVRRRFVLELEPEPNLY  339
A0A0H2ZD69|A0A0H2ZD69_PSEAB     LVLVNHGGATGAQVQALAERIQEDVRRRFVLELEPEPNLY  339
Q9HZM7|MURB_PSEAE                 LVLVNHGGATGAQVQALAERIQEDVRRRFVLELEPEPNLY  339
4JAY                               LVLVNHGGATGAQVQALAERIQEDVRRRFVLELEPEPNLY  340
A0A0E1AVK6|A0A0E1AVK6_PSEAI     LVLVNHGGATGAQVQALAERIQEDVRRRFVLELEPEPNLY  339
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