

Clustal Omega

Tools (//Tools/) > Multiple Sequence Alignment (//Tools/msa) > Clustal Omega

Results for job clustalo-l20180817-120426-0368-49877271-p2m

CLUSTAL O(1.2.4) multiple sequence alignment			
18RS-1924-6	-----TGTTTGGTGTTTACACTAGACTGATGAGTTGCGAACGGGTGAGTAACGCGTAGG	54	
18RS-1924-7	CGCTGATGTTTGGTGTTTACACTAGACTGATGAGTTGCGAACGGGTGAGTAACGCGTAGG	60	

18RS-1924-6	TAACCTGCCTCATAGCGGGGATAAATTGGAACGATAGCTAATACCGCATAAGAGTA	114	
18RS-1924-7	TAACCTGCCTCATAGCGGGGATAAATTGGAACGATAGCTAATACCGCATAAGAGTA	120	

18RS-1924-6	ATTAACACATGTTAGTTATTTAAAAGGAGCAATTGCTTCACTGTGAGATGGACCTGCGTT	174	
18RS-1924-7	ATTAACACATGTTAGTTATTTAAAAGGAGCAATTGCTTCACTGTGAGATGGACCTGCGTT	180	

18RS-1924-6	GTATTAGCTAGTTGGTGAGGTAAGGCTCACCAAGGCGACGATACATAGCCGACCTGAGA	234	
18RS-1924-7	GTATTAGCTAGTTGGTGAGGTAAGGCTCACCAAGGCGACGATACATAGCCGACCTGAGA	240	

18RS-1924-6	GGGTGATCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTAG	294	
18RS-1924-7	GGGTGATCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTAG	300	

18RS-1924-6	GGAATCTTCGGCAATGGACGGAAGTCTGACCAGCAACGCCGCGTGAGTGAAGAAGGTTT	354	
18RS-1924-7	GGAATCTTCGGCAATGGACGGAAGTCTGACCAGCAACGCCGCGTGAGTGAAGAAGGTTT	360	

18RS-1924-6	TCGGATCGTAAAGCTCTGTTGTTAGAGAAGAAGCTGGTAGGAGTGGAAAACTACCAAG	414	
18RS-1924-7	TCGGATCGTAAAGCTCTGTTGTTAGAGAAGAAGCTGGTAGGAGTGGAAAACTACCAAG	420	

18RS-1924-6	TGACGGTAACCTAACAGAAAGGACGGCTAACTACGTGCGCAGCAGCCGCGTAATACGTA	474	
18RS-1924-7	TGACGGTAACCTAACAGAAAGGACGGCTAACTACGTGCGCAGCAGCCGCGTAATACGTA	480	

18RS-1924-6	GGTCCCAGAGCGTTGTCCGGATTATTGGGCGTAAAGCGAGCGCAGGCGGTTCTTTAAGTC	534	
18RS-1924-7	GGTCCCAGAGCGTTGTCCGGATTATTGGGCGTAAAGCGAGCGCAGGCGGTTCTTTAAGTC	540	

18RS-1924-6	TGAAGTTAAAGGCAGTGGCTTAACCATTTGTACGCTTTGGAACTGGAGGACTTGAGTGCA	594	
18RS-1924-7	TGAAGTTAAAGGCAGTGGCTTAACCATTTGTACGCTTTGGAACTGGAGGACTTGAGTGCA	600	

18RS-1924-6	GAAGGGGAGAGTGAATTCCATGTGTAGCGGTGAAATGCGTAGATATATGGAGGAACACC	654	
18RS-1924-7	GAAGGGGAGAGTGAATTCCATGTGTAGCGGTGAAATGCGTAGATATATGGAGGAACACC	660	

18RS-1924-6	GGTGGCGAAAGCGGCTCTCTGGTCTGTAACGTGACGCTGAGGCTCGAAAGCTGGGGAGCA	714	
18RS-1924-7	GGTGGCGAAAGCGGCTCTCTGGTCTGTAACGTGACGCTGAGGCTCGAAAGCTGGGGAGCA	720	

18RS-1924-6	AACAGGATTAGATACCTGGTAGTCCACGCCGTAAACGATGAGTGCTAGGTGTTAGGCC	774	
18RS-1924-7	AACAGGATTAGATACCTGGTAGTCCACGCCGTAAACGATGAGTGCTAGGTGTTAGGCC	780	

18RS-1924-6	TTTCCGGGGCTTAGTGCCGAGCTAACGCATTAAAGC---	810	
18RS-1924-7	TTTCCGGGGCTTAGTGCCGAGCTAACGCATTAAAGCACTC	820	
