

Sequence IDs and their corresponding colors:

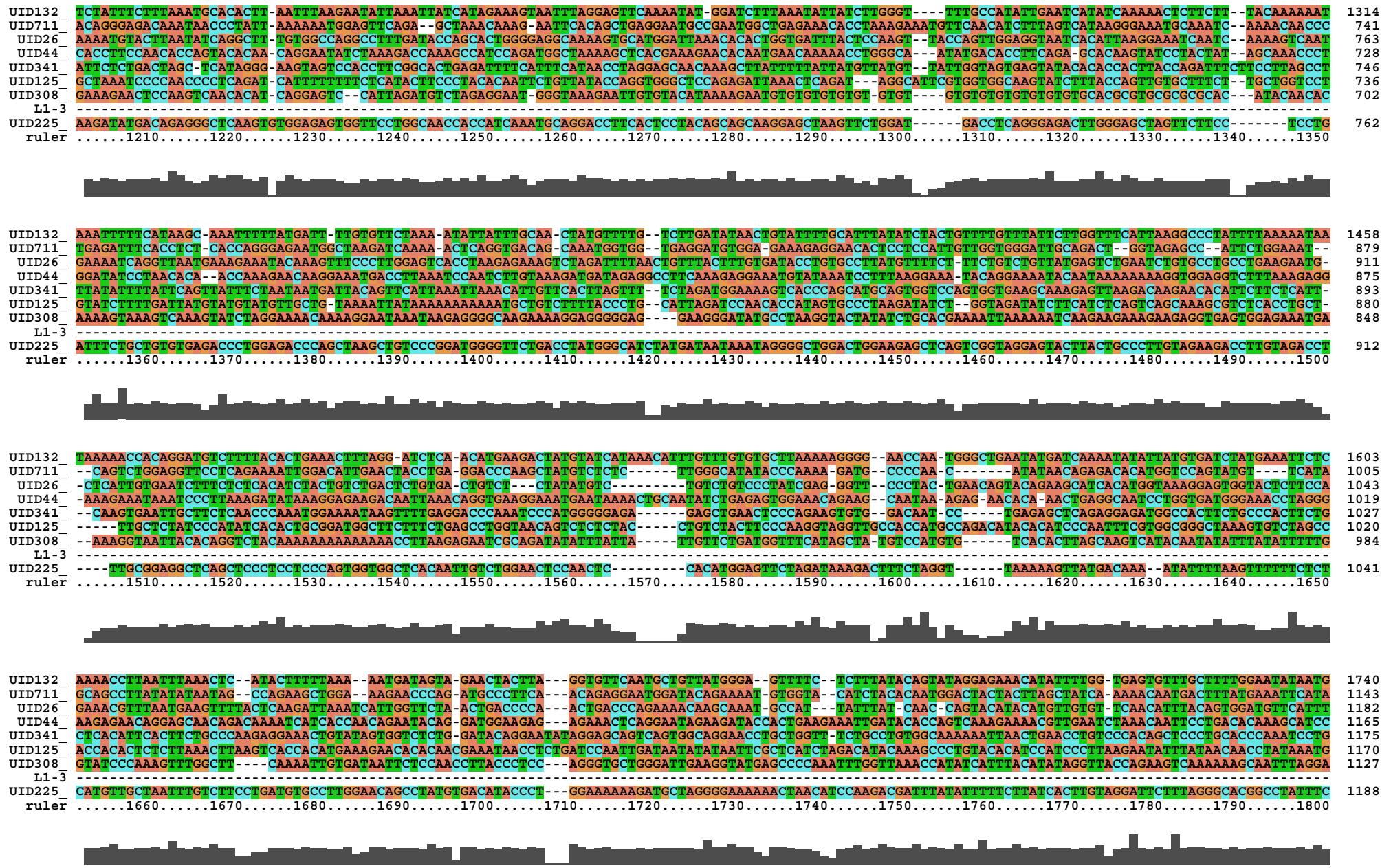
- Cluster 1 (Top): UID132 (red), UID711 (green), UID26 (blue), UID44 (orange), UID341 (yellow), UID125 (purple), UID308 (pink), L1-3 (grey).
- Cluster 2 (Second from Top): UID132 (red), UID711 (green), UID26 (blue), UID44 (orange), UID341 (yellow), UID125 (purple), UID308 (pink), L1-3 (grey).
- Cluster 3 (Third from Top): UID132 (red), UID711 (green), UID26 (blue), UID44 (orange), UID341 (yellow), UID125 (purple), UID308 (pink), L1-3 (grey).
- Cluster 4 (Bottom): UID132 (red), UID711 (green), UID26 (blue), UID44 (orange), UID341 (yellow), UID125 (purple), UID308 (pink), L1-3 (grey).

Sequence lengths (bp) indicated on the right side of the tree:

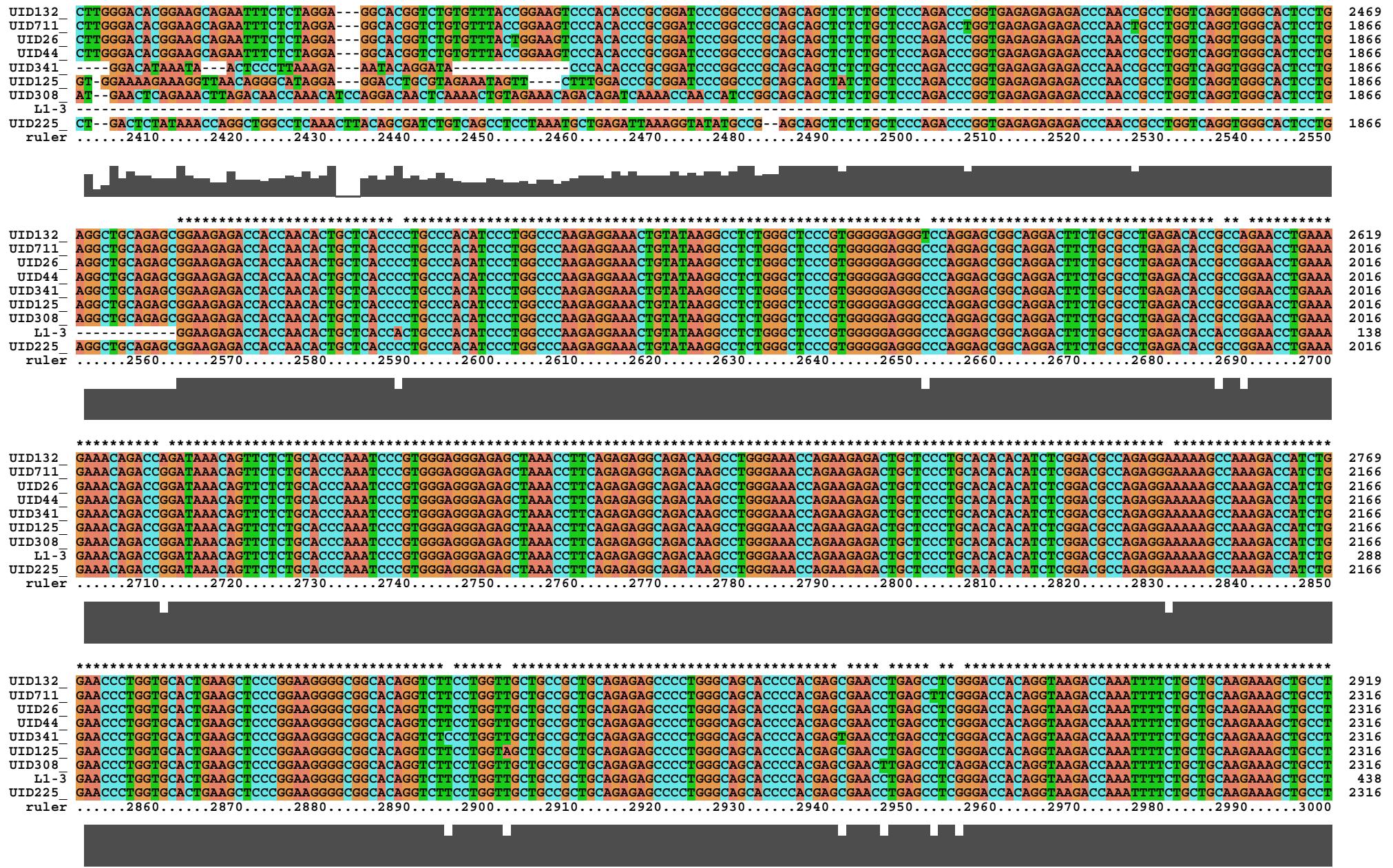
- Cluster 1: 741, 177, 194, 164, 198, 172, 137.
- Cluster 2: 886, 308, 339, 305, 338, 319, 277.
- Cluster 3: 1030, 455, 475, 445, 464, 456, 420.
- Cluster 4: 1172, 597, 618, 585, 600, 592, 564.
- Root: 625.

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Fig. S4A













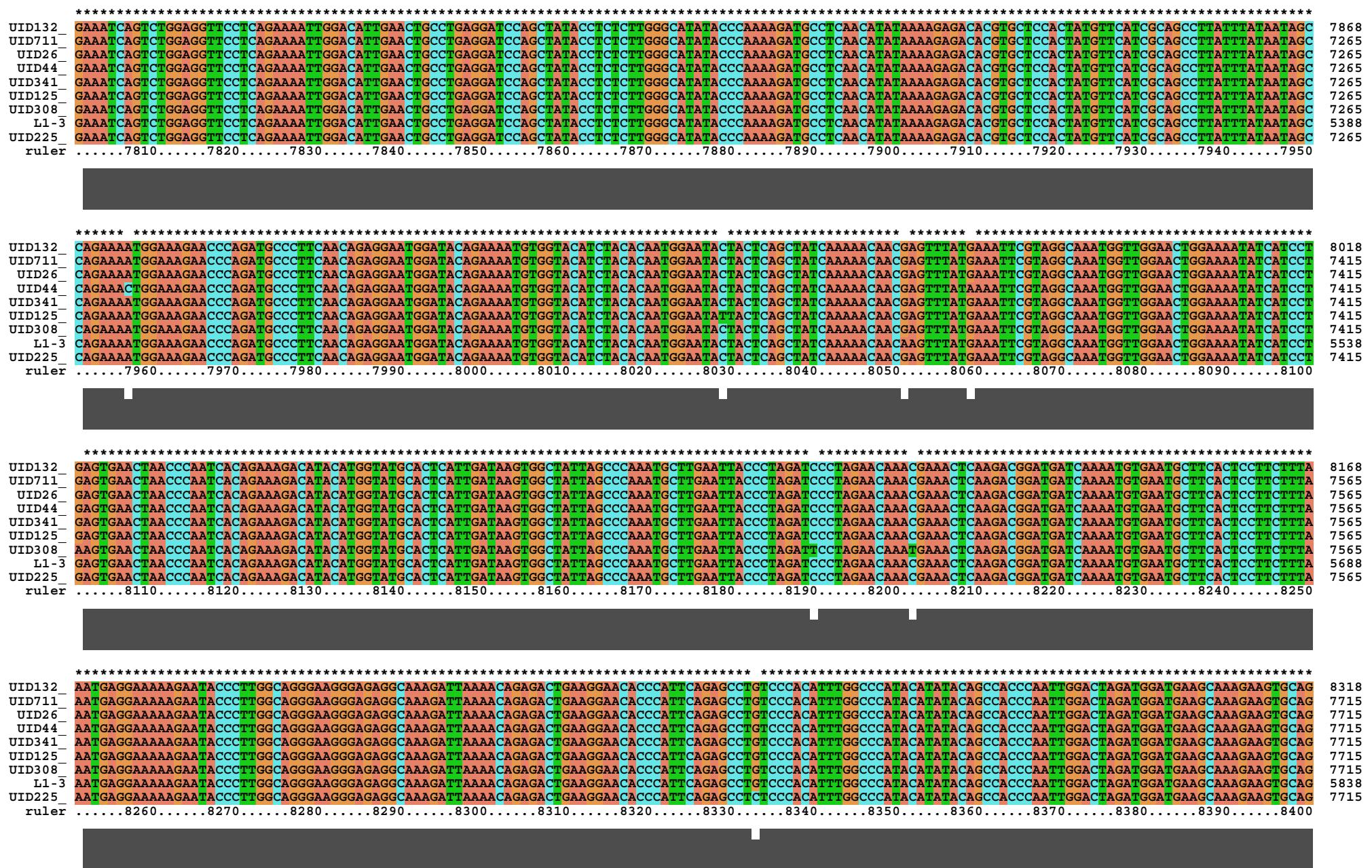




UID132	CATCAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	6068
UID711	CAICAAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
UID26	CAICAAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
UID44	CAICAAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
UID341	CAICAAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
UID125	CAICAAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
UID308	CAICAAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
L1-3	CATCAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	3588
UID225	CATCAAAACAAATCATCCACCATTGATCAAGTAGGCTTCATCCCAGGCATGCGAGGGATGCTTAAATACGGAAAACCATAACGTGATCCATTATAAACAACTGAAAGAACAGAACCCATGATCATTCTATTAGATGCTGAGAAAGC	5465
ruler	.....6010.....6020.....6030.....6040.....6050.....6060.....6070.....6080.....6090.....6100.....6110.....6120.....6130.....6140.....6150	
		
UID132	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	6218
UID711	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
UID26	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
UID44	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
UID341	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
UID125	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
UID308	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
L1-3	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	3738
UID225	ATTTGACAAAATTCAACACCCCCCTCATGATAAAAGCTCTGGAAAGAATAGGAATTCAAGGCCATACTAAACATAACGCAAACAGTTGCTAACATTAAACTGAGAAACCTTCCCACATAAA	5615
ruler	.....6160.....6170.....6180.....6190.....6200.....6210.....6220.....6230.....6240.....6250.....6260.....6270.....6280.....6290.....6300	
		
UID132	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	6368
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UID26	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	5765
UID44	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	5765
UID341	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	5765
UID125	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	5765
UID308	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	5765
L1-3	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	3888
UID225	ATCAGGGACTAGACAAGGCCTGCCACTCTCTCCCTACTTATTCAATATAGTTCTGAAGTTCTAGCCAGAGCAATCAGACAACAAAAGGGATCAAGGGATACAGATCGGAAAAAGAAGGGTCAAAATACACTATTGAGATGATAT	5765
ruler	.....6310.....6320.....6330.....6340.....6350.....6360.....6370.....6380.....6390.....6400.....6410.....6420.....6430.....6440.....6450	
		
UID132	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	6518
UID711	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
UID26	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
UID44	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
UID341	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
UID125	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
UID308	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
L1-3	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	4038
UID225	GATAGTATAATTCAAGTGAACCCAAAAAGTCTCCACCCAGAGAACCTACTAAAGCTGATAAAACAACTTCAAGCAAAGTGCTGGGTATAAAAATAACTCAAATAAATCAGTTGCCCTCTCTATAACAAAAGAGAAACAGCCGAGAAAGAAATTAG	5915
ruler	.....6460.....6470.....6480.....6490.....6500.....6510.....6520.....6530.....6540.....6550.....6560.....6570.....6580.....6590.....6600	
		









UID132 ATC - ACACATTGGTAGTTTAAAGTTAAATGAAAC - TTATGTCAACTGGATGATCTATTGTGCTGTTGGATAATGCTG - GTTGATATGACAAACT - AATCAATGCTGTTTAAATTATCTCCACAGTAATGGC 9048  
 UID711 CAAGAACATCAAGAGAGATGCCAATGCCATGCTTCACCTTCTTAAAGGGAAACAGAATACCCCTGGAGGGAAATAGGGAGGAAATTAGAACCTGAGGCTGAAGGAAACCCCCATTGAGGCTGCCTCACATGIGGCCA 8452  
 UID26 TCC - TTTAAAAATTAATTAAAGTACCTGAACAAC - CAAACTGTACACATTCTAGGT - ACTTGTTTTTTAAAGATTAAATTAAATTAAATTAGAGATGCTATTATATACATGAAGAGTCATAGAAGGCAGAAGG 8444  
 UID44 ATAT - TICAAGACAAATCC - AAATTGAAACAAATTA - CTATCIAAAAATCCAGCCCTATAGAAAATACTAGAGGAAACICCAA - TAIAAGGAAGATAACTACATCTAAGAAAACACAGGAAAATAAATAATTATATAA - CAGCTAA 8442  
 UID341 AAACTAAAGGAATTGAAAGAAATTATACAGGAATTAAAACGTAGAGAGAAACAAATAAAGAAAGCACAAAGGGAAATGTTTCAAGGCTCTCGGAATTGGTCAGAGTTCAAGTTGGAAACC 8450  
 UID125 GGATCTACTCACAAACACACTCAAAACTCAGGATGGC - AGTCTATCCAGAAAGAAACCAAGGCTCTCGGAATTGGTCAGAGTTCAAGTTGGAAACC 8456  
 UID308 ACAGAATCTGGTAGGGAGC - ATGGTTGGGATGCTCTGACTTGTGGTCTGAC - TTGGATAGTTTAAAGCACTGAATTCCAGCAACAGACTGAACCCACA - GAACAAAGCTTCTGTTCTCCACAGGGCACCAACACGGAAGCTAA 8446  
 L1-3 ..... 5860  
 UID225 TGCCCTTCTTAGCAAGGCCCTCACCACTGAGCTAAATC - CCCAACCCCTCAGAGCTTGAATCTTCATTTGGAGTATTGCACTAGACTTTAAGCTGATC - TGAAAGCCCTAAATCTGTAAGTCTCTGGATTAAATTAG 8453  
 ruler ..... 9010 ..... 9020 ..... 9030 ..... 9040 ..... 9050 ..... 9060 ..... 9070 ..... 9080 ..... 9090 ..... 9100 ..... 9110 ..... 9120 ..... 9130 ..... 9140 ..... 9150

UID132 CACCTCCCTTCACCAAGTTCTGCTTACCTT - 9079  
 UID711 TACGTATACTGCCAGCAAAGTAGATA - 8478  
 UID26 CATCAGATCTCCTAAACTCTAGAGTTATCCATG - 8476  
 UID44 AACAAAAATAGAAAAGGAGGGAGCTAGGACCGGG 8476  
 UID341 CAACTGTTCTGCACCCAAATCCCTG - 8476  
 UID125 TGCTTGTGAAAGGACATGT - 8476  
 UID308 CACCTGCACAGCCCCAACATTGTTCTC - 8476  
 L1-3 ..... 5860  
 UID225 AGATTTTGATCTTTGTTTTTC - 8476  
 ruler ..... 9160 ..... 9170 ..... 9180 .....