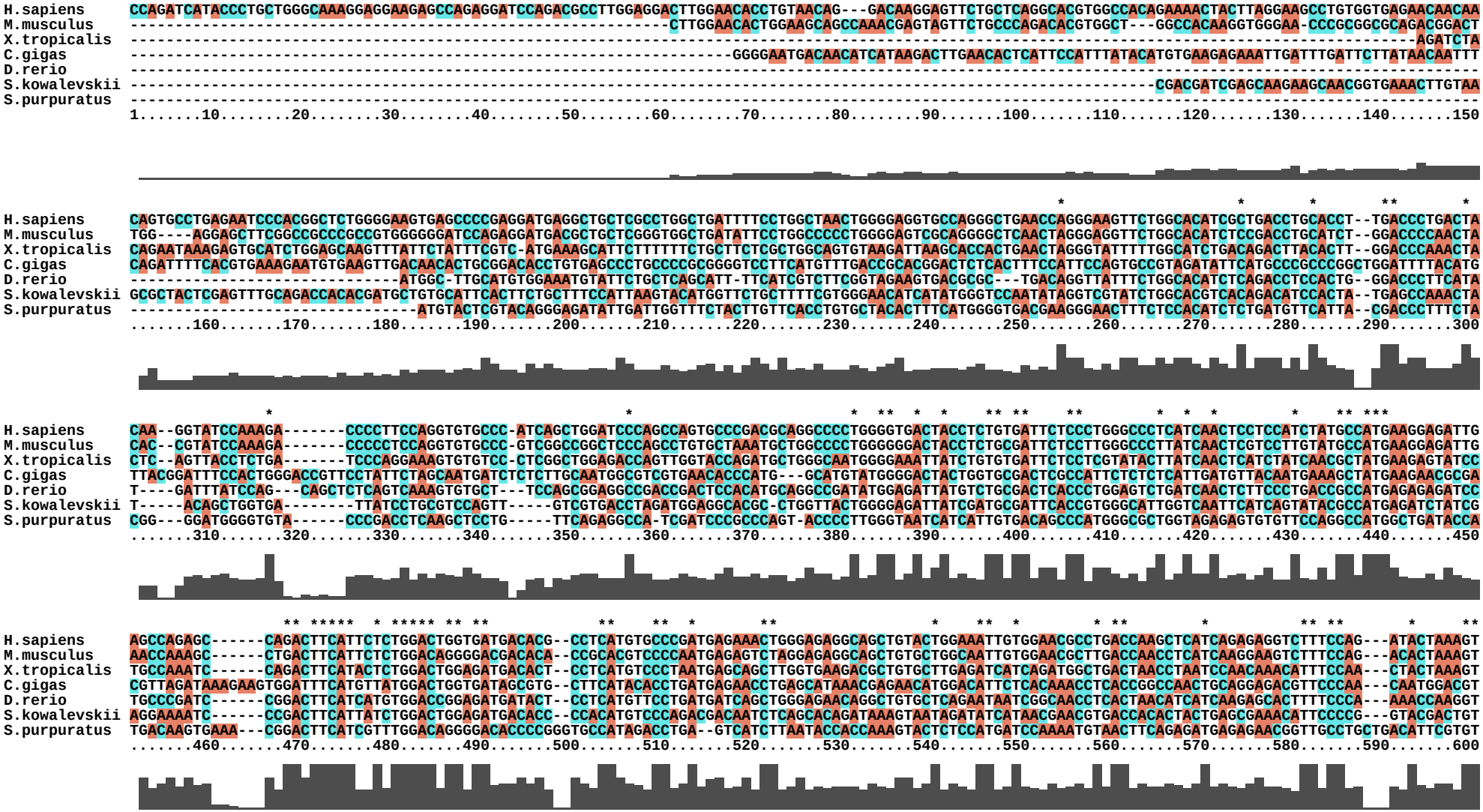
CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT



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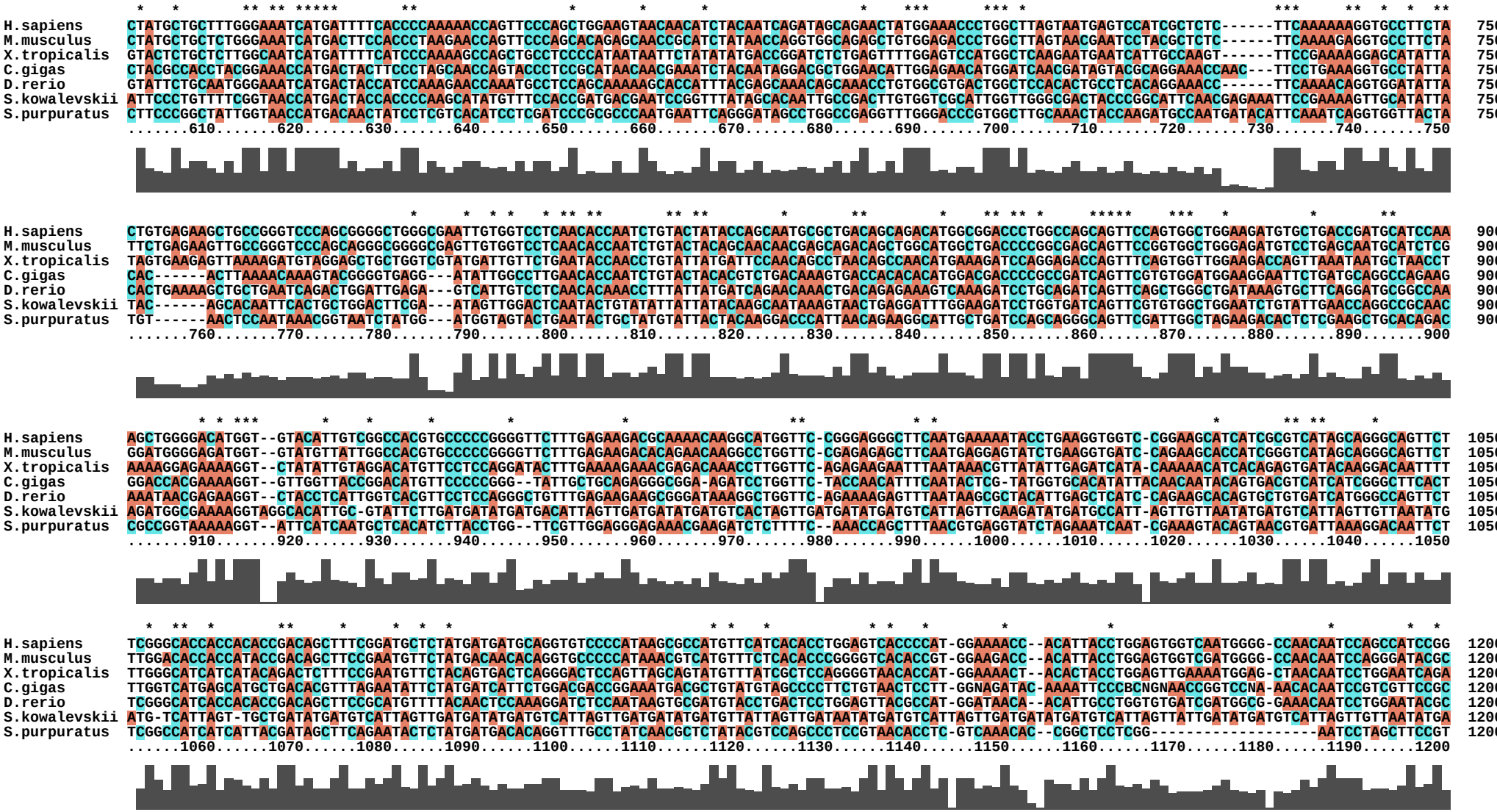
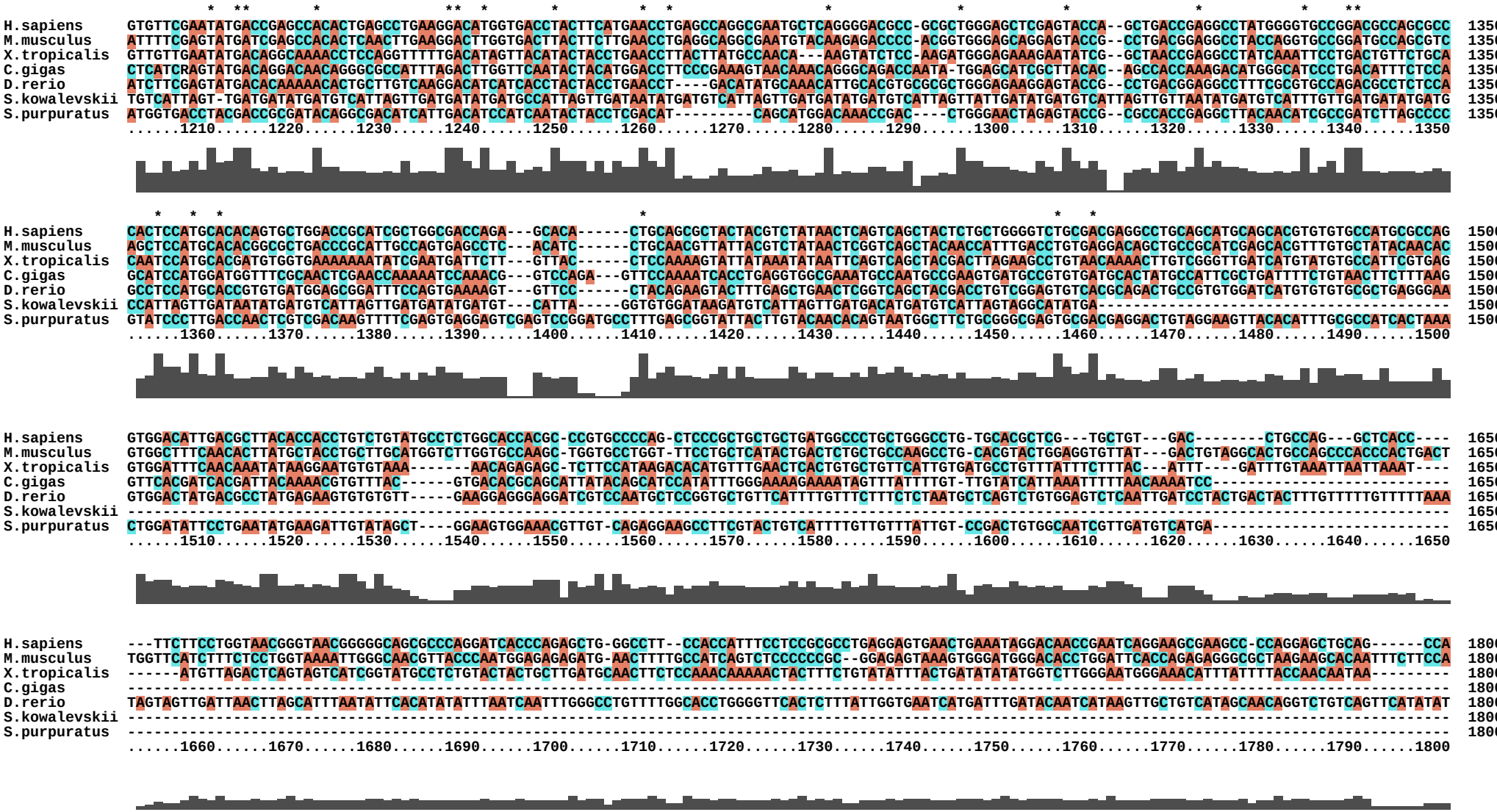


CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT



CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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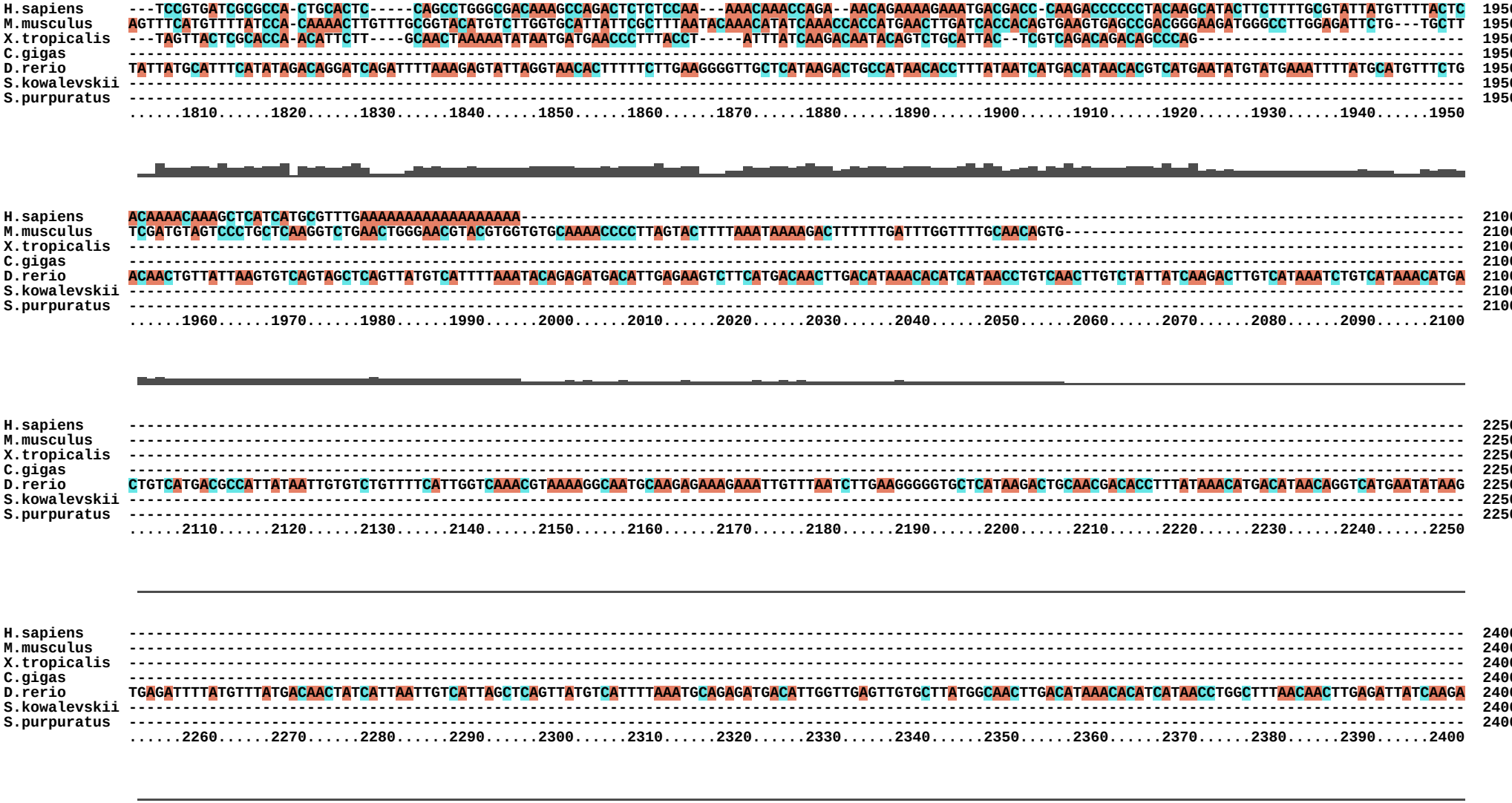
Page 4 of 5



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X.tropicalis  ------------------------------------------------------------------------------------------------------------------------  2550
C.gigas       ------------------------------------------------------------------------------------------------------------------------  2550
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S.kowalevskii ------------------------------------------------------------------------------------------------------------------------  2550
S.purpuratus  ------------------------------------------------------------------------------------------------------------------------  2550
              ......2410......2420......2430......2440......2450......2460......2470......2480......2490......2500......2510......2520......2530......2540......2550

H.sapiens     ------------------------------------------------------------------------------------------------------------------------  2700
M.musculus    ------------------------------------------------------------------------------------------------------------------------  2700
X.tropicalis  ------------------------------------------------------------------------------------------------------------------------  2700
C.gigas       ------------------------------------------------------------------------------------------------------------------------  2700
D.rerio       TGACATAACACGTGATAAATATGAATGAGATTTTATGCATGTTTATGACAACTGTTATTAATTGTCATTAGCTCAGTTATGCCATTTTAAATGCAGAGATGACATTGGTTGAGATTTGCTTCTGACAACTTGACATAAACACATCACAAC  2700
S.kowalevskii ------------------------------------------------------------------------------------------------------------------------  2700
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H.sapiens     ----------------------------------------------------------------------------------------------------------  2830
M.musculus    ----------------------------------------------------------------------------------------------------------  2830
X.tropicalis  ----------------------------------------------------------------------------------------------------------  2830
C.gigas       ----------------------------------------------------------------------------------------------------------  2830
D.rerio       CTGTCTTTGACAACTTGAGACTATAAAGACTTATCATAACTCTGTCATAAACATGACTGTCATGAGGCCATTATAATTGTGTCTTCTTTCATCTGTCCAATGCTAAAGCCAATAAAAGACAAAGAAATAG  2830
S.kowalevskii ----------------------------------------------------------------------------------------------------------  2830
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