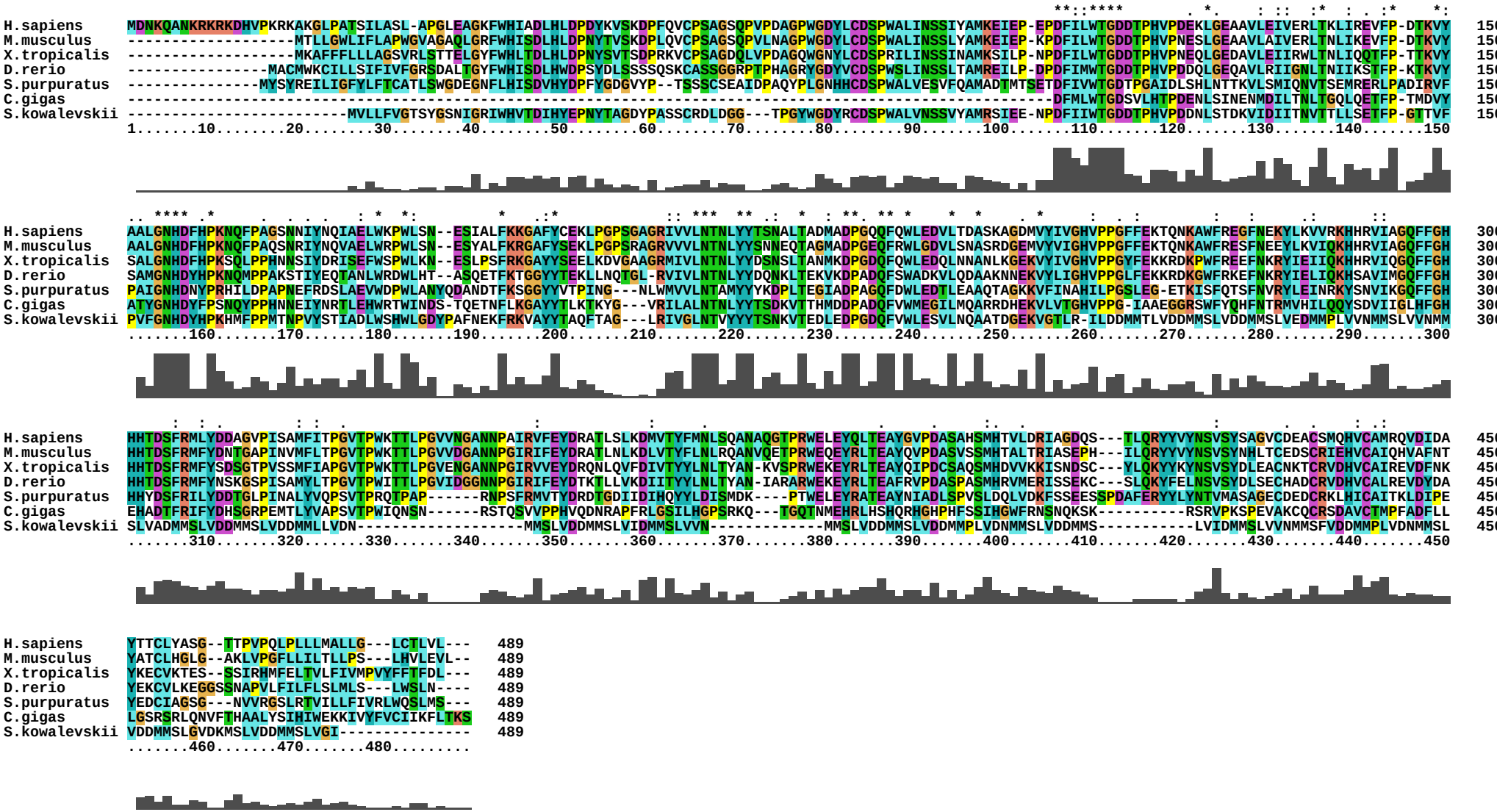
CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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