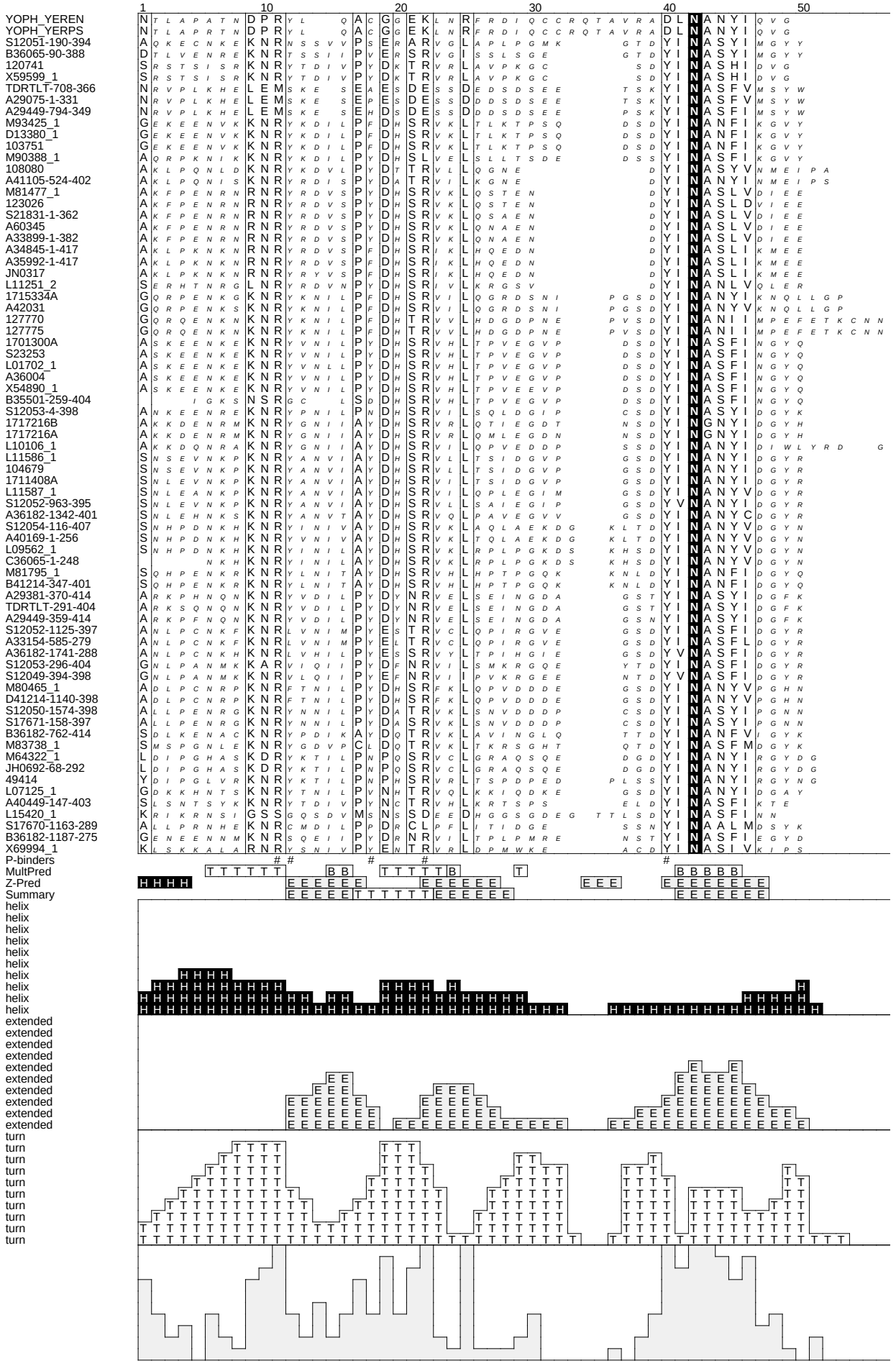
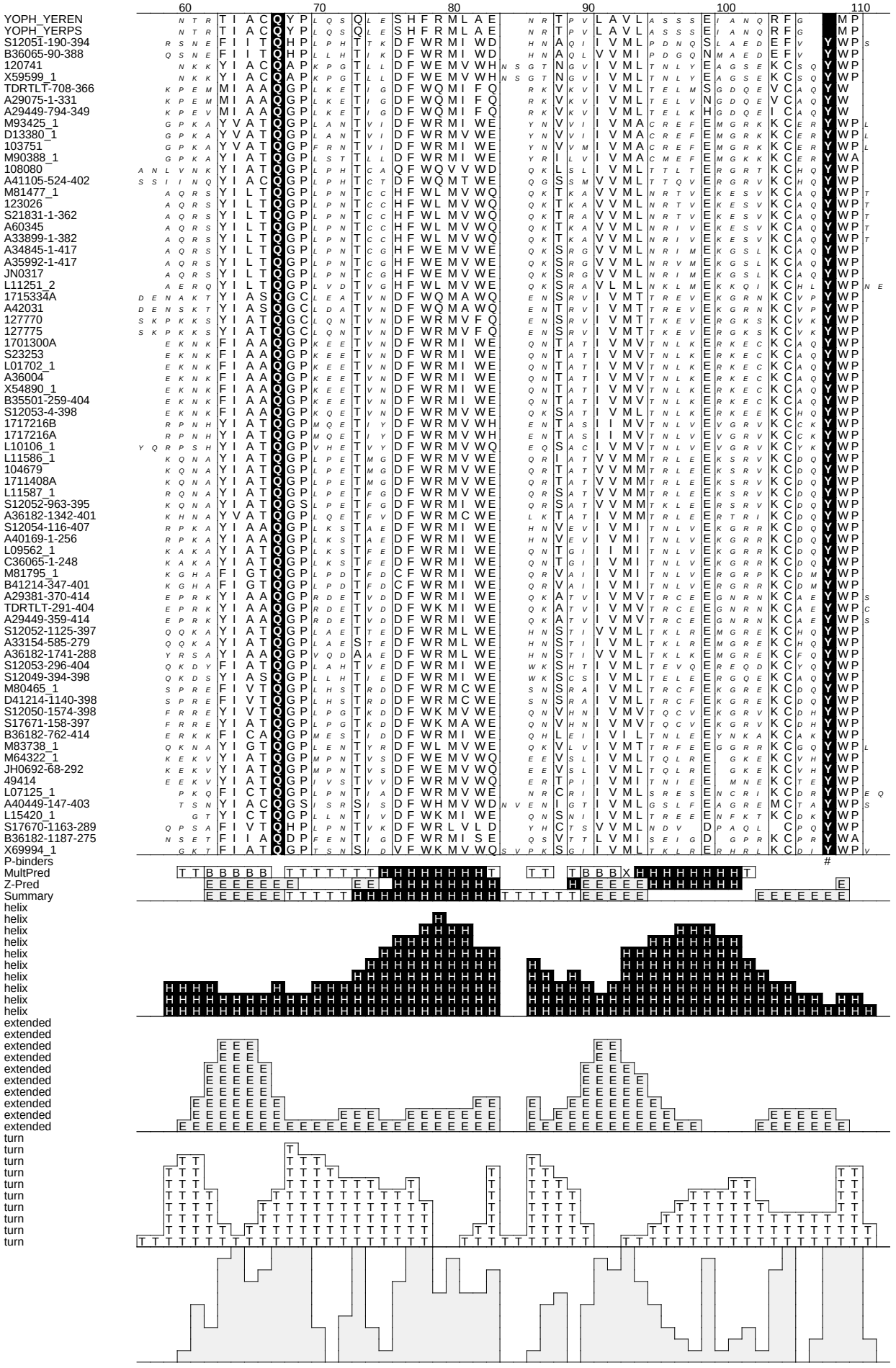


AMAS *(c) 1992 G.J.Barton & C.D.Livingstone - Univ. Oxford UK.*

Conservation plot of the sequence alignment "6.amas"

Conservation Threshold: 6 - Gaps Ignored: 4 - Minimum Residue Occupancy: 10%

Histogram type: AVERAGE - Conservation mode: STANDARD with 10 property index "extra.pt"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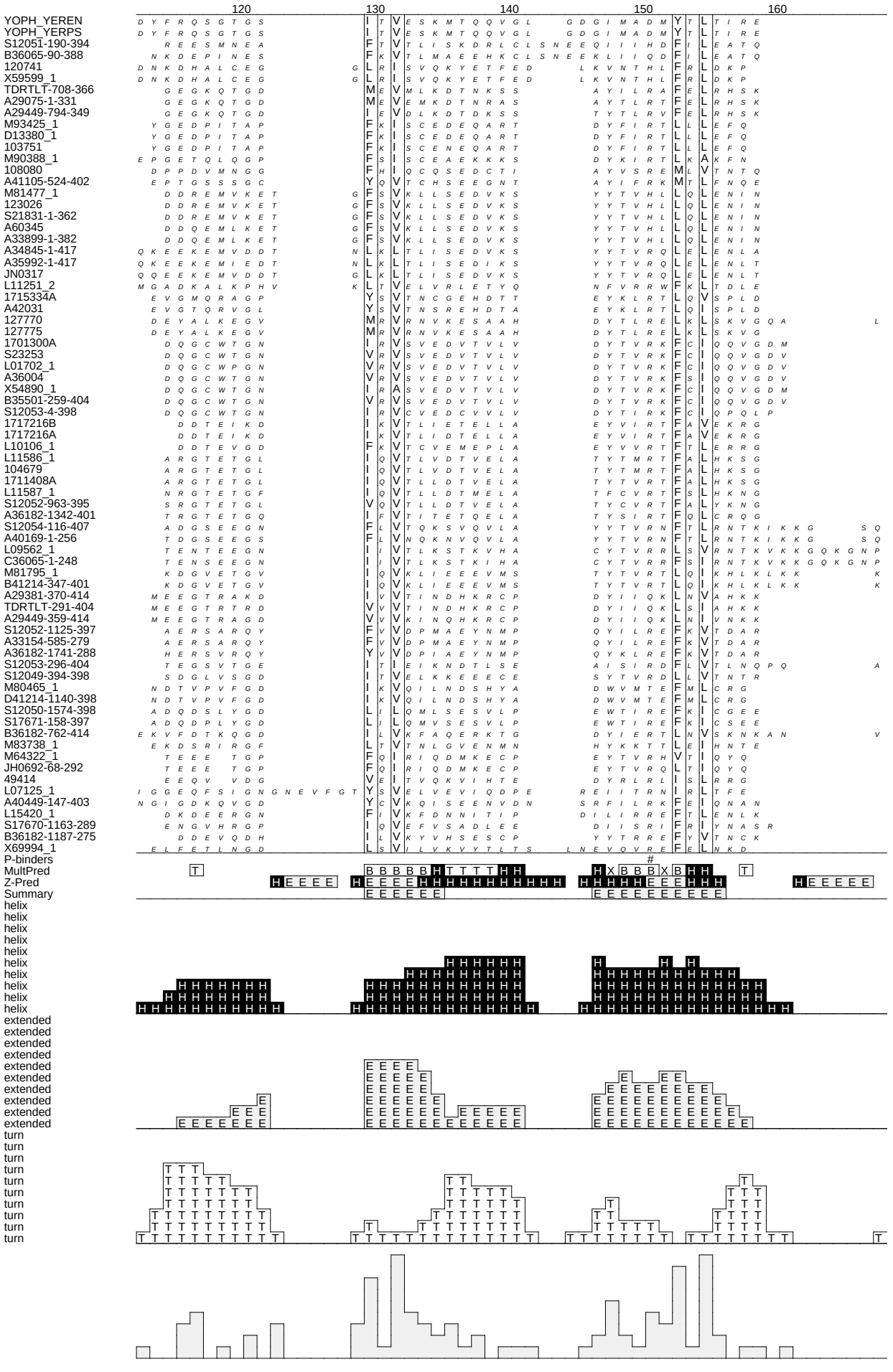
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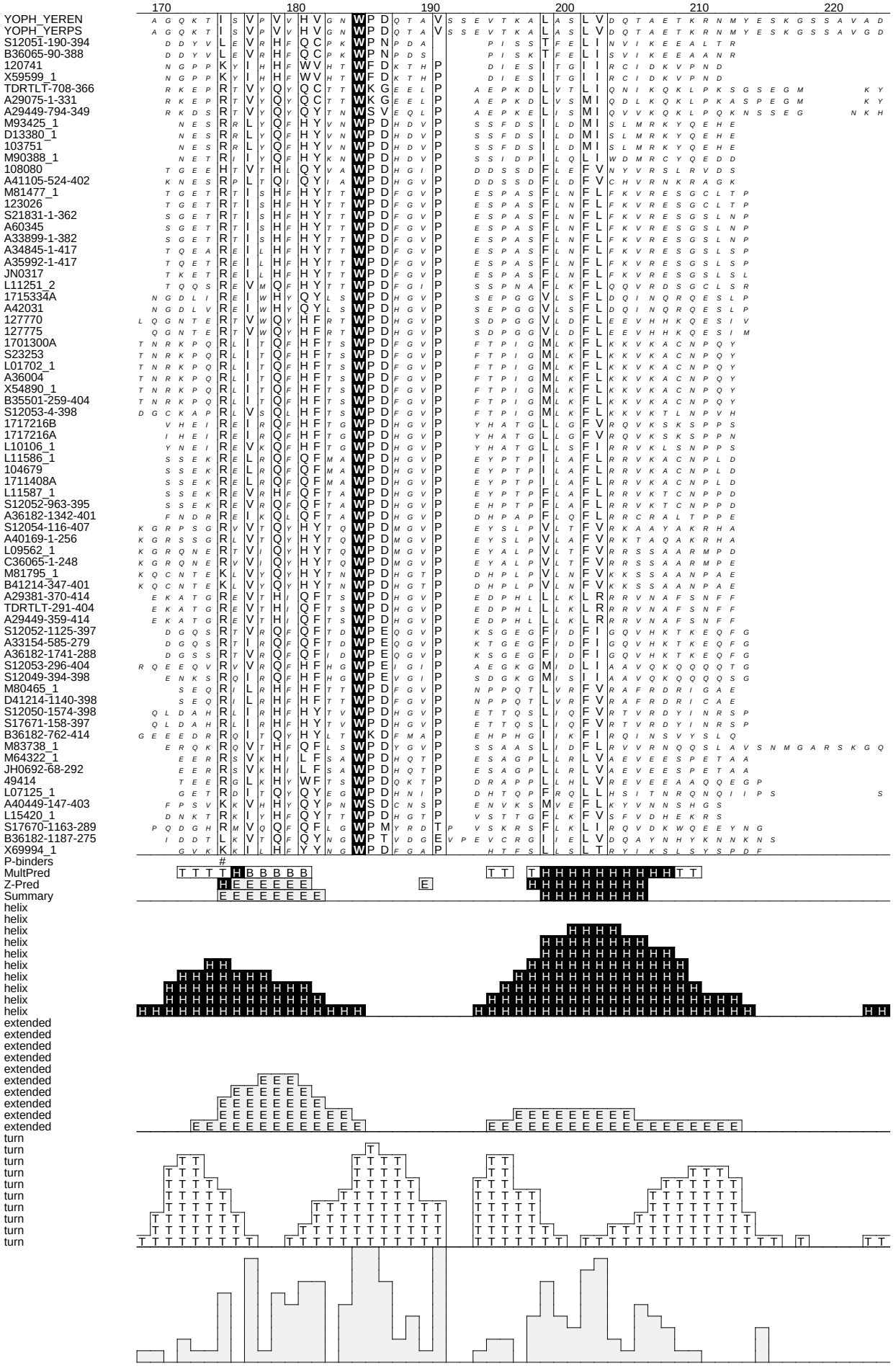
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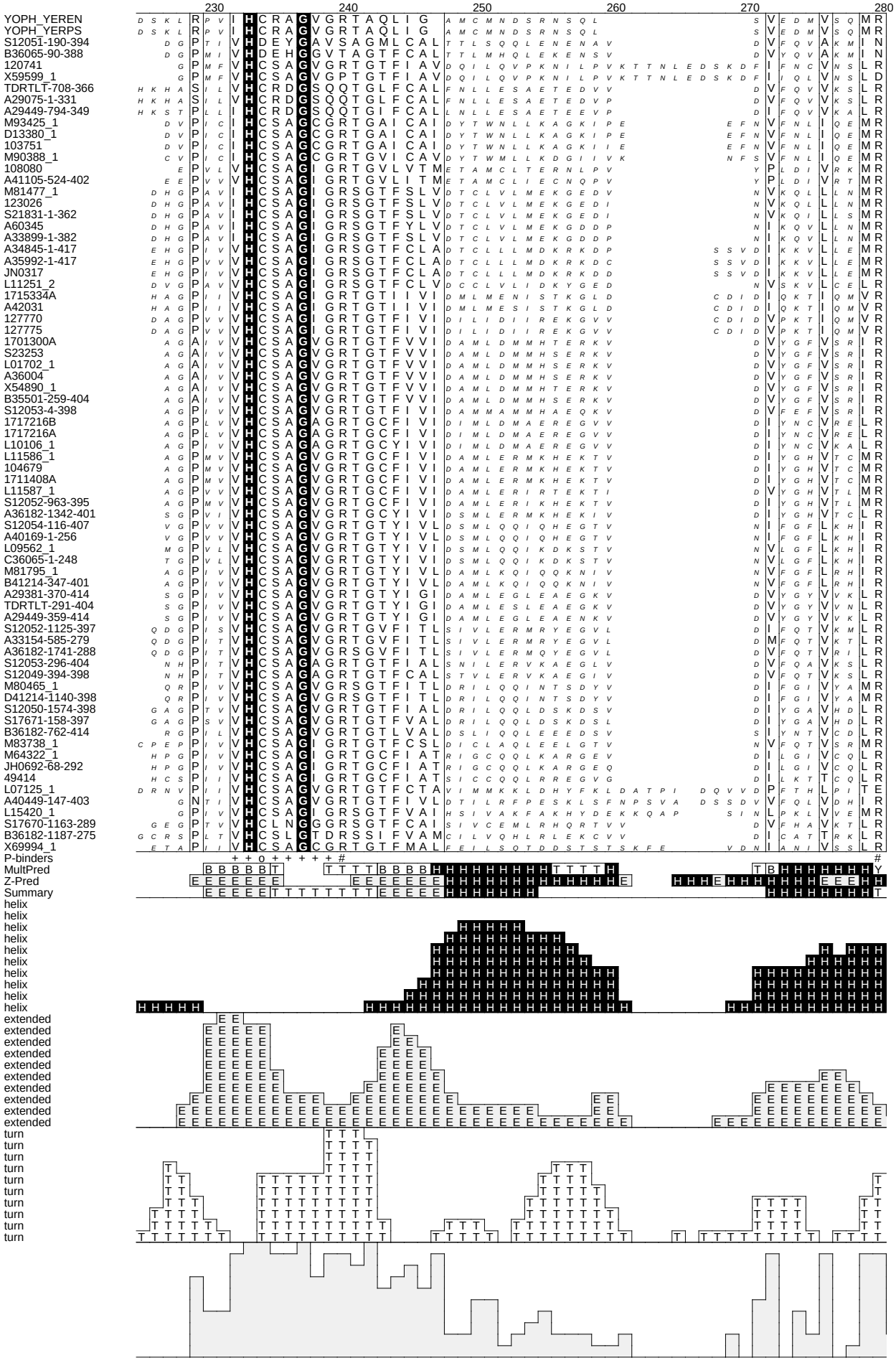
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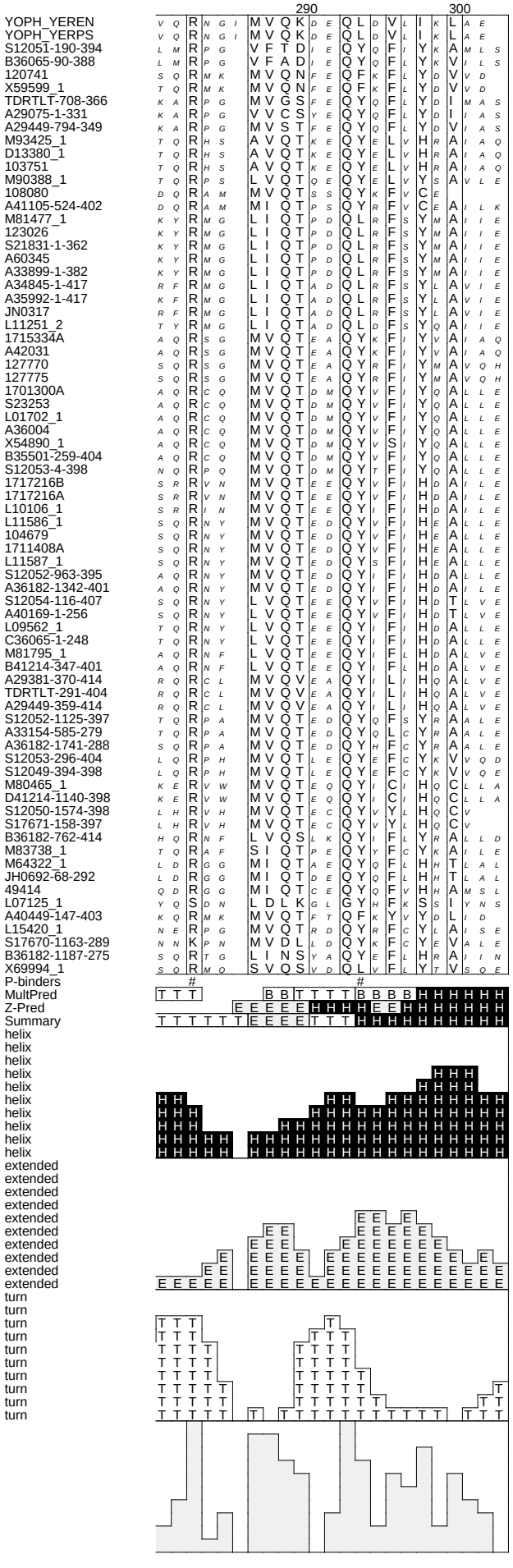
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