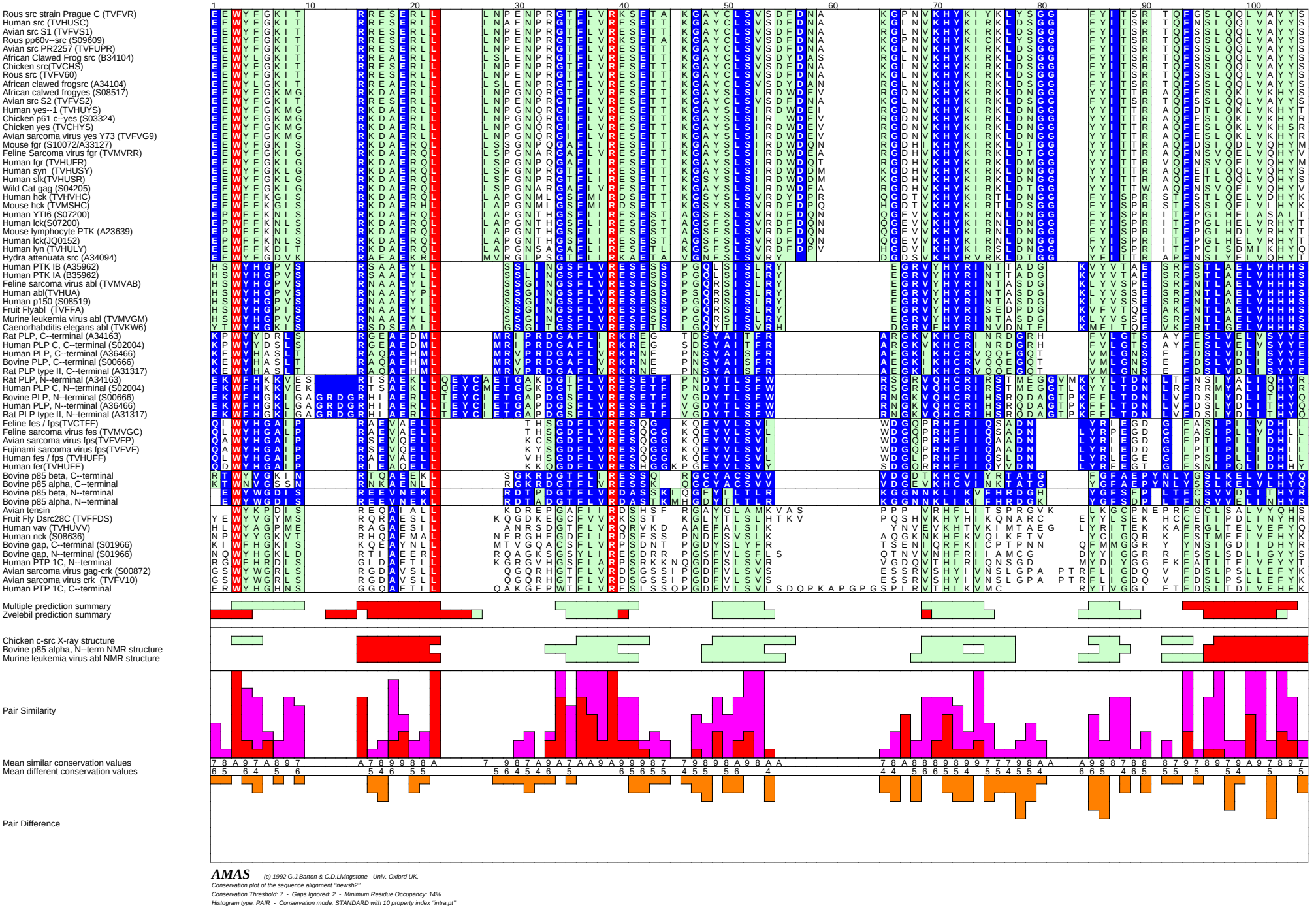

AMAS *(c) 1992 G.J.Barton & C.D.Livingstone - Univ. Oxford UK.*

Conservation plot of the sequence alignment "newsh2"

Conservation Threshold: 7 - Gaps Ignored: 2 - Minimum Residue Occupancy: 14%

Histogram type: PAIR - Conservation mode: STANDARD with 10 property index "intra.pt"