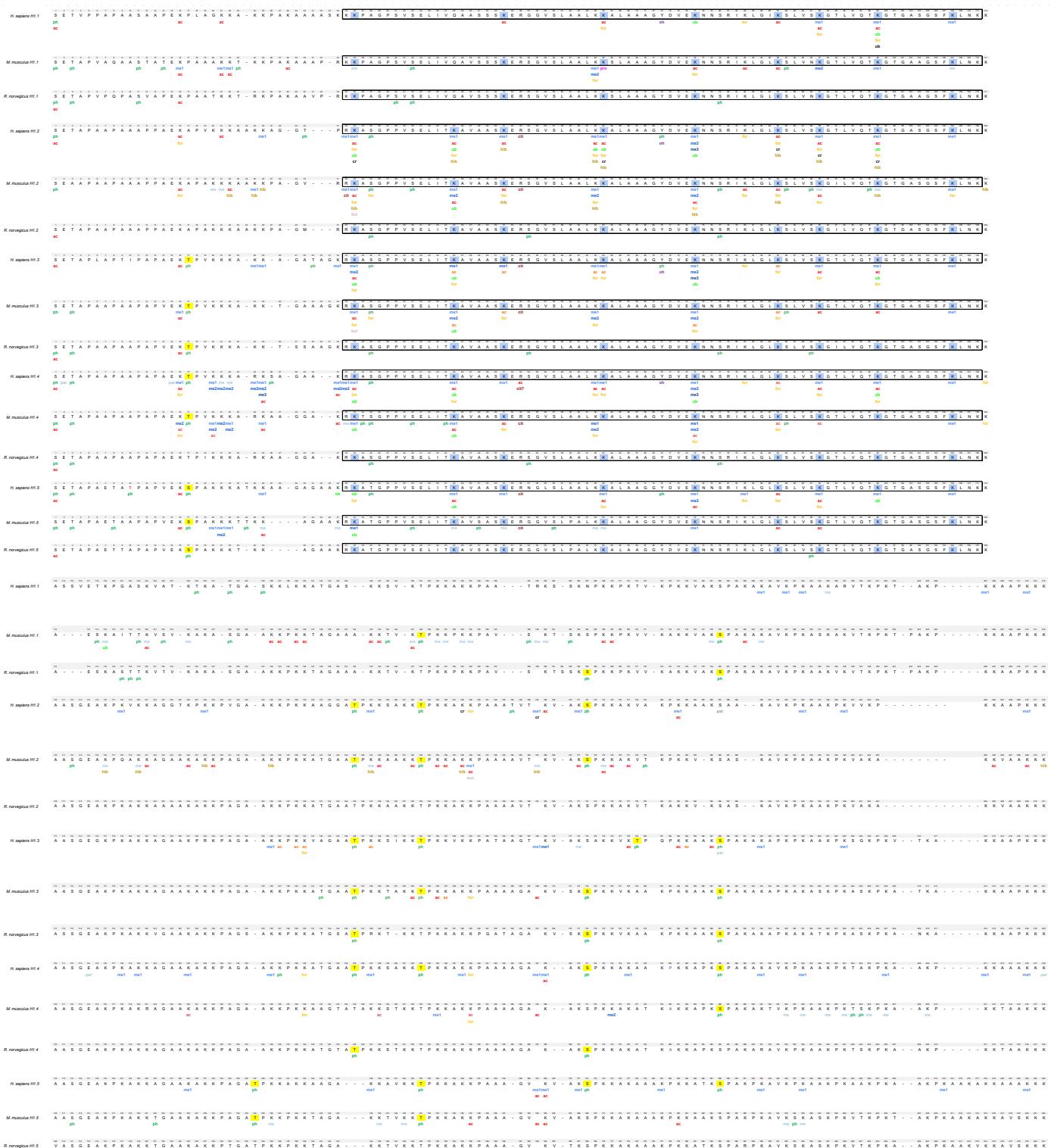
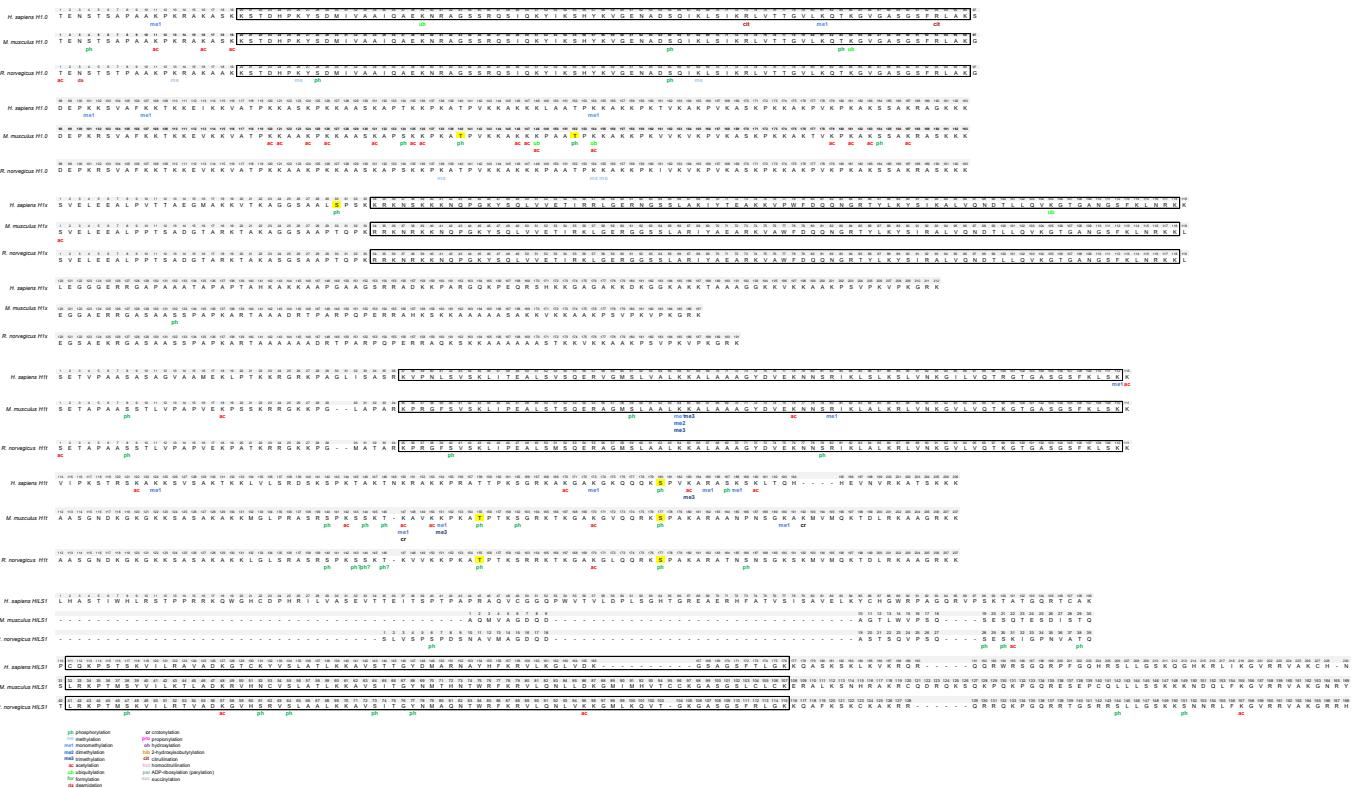


A



B



**Figure S2. Post-translational modifications in mammalian H1 subtypes.** A, Sequence alignment H1.1-H1.5 subtypes. B, Sequence of H1.0, H1x, H1t and HILS1. The residues within the box correspond to the globular domain. Data were extracted from references 29-30, 34-47, 55-58, 61-62, 64-65, 70, 75, 77, 83, 95, 96, 99, and 101.