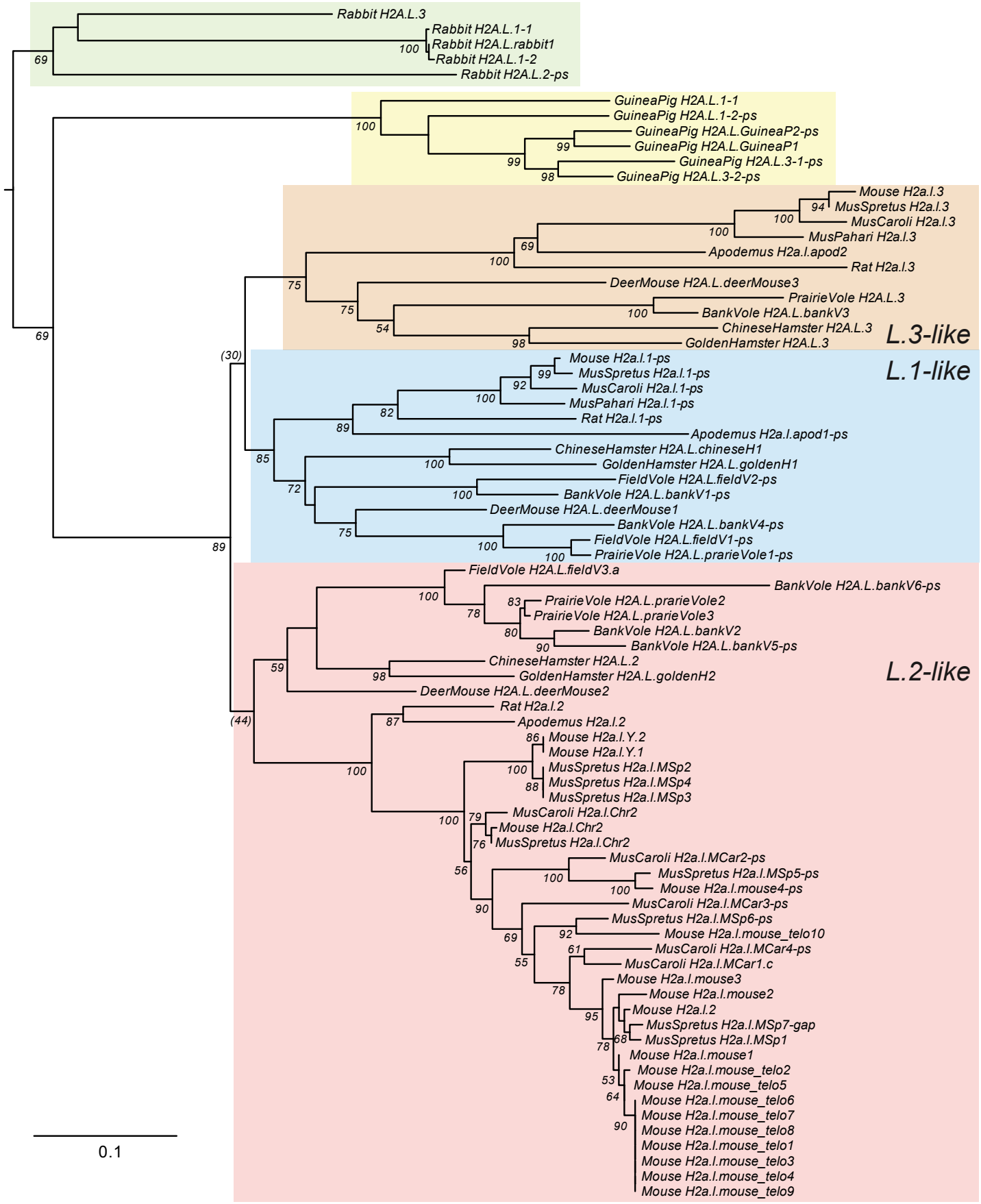

Supplemental Fig S8: Nucleotide phylogeny of rodent *H2A.L* genes.
Evolutionary analysis shows that the three ancestral *H2A.L* loci evolve on independent trajectories in rodents. We show all bootstrap values >50%, with selected other lower values shown in parentheses. Many genes in the *H2A.L.1* locus are pseudogenes in rodents. The gene at the *H2A.L.2* locus has experienced additional duplications to dispersed genomic locations in the Mus lineage.

Supplemental Fig S8
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