

Supplementary Table S2. Genomic sequence corresponding to 3' untranslated sequence of H3 gene transcripts. Each sequence below begins immediately following the termination codon. The position of a strongly conserved segment at the 3' end of the canonical (H3.1) H3s important for RNA processing and regulation (Marzluff *et al.* 2008) is underlined.

H3F3A	GAATCCACTATGATGGGAAACATTCTCAAAAAAAAAAAAAAATTCTCTTCTTC...
H3F3B	GTGAAGGCAGTTTATGGCGTTTAGTAAATTCTGTAAAACACTTGGTTAATTGT...
H3F3C	GTGAAGGCAGTTTATGGCATTGTAGTAAATTCTGTAAAACACTTGGTTAATTGGT...
HIST1H3A	TTACTGTGGTCTCTGACGGTCCA <u>AAGCAAAGGCT</u> CTTTCAGAGC...
HIST1H3B	ATGTAAAGTCACCTTTCATCAGTCTAA <u>ACCCAAAGGCT</u> CTTTCAGAGC...
HIST1H3C	GTCTGCCGTTCTCCTCATT <u>GAAAAGGCT</u> CTTTCAGAGC...
HIST1H3D	ATTGTCTGTGAATGTGTGCTAAC <u>CCAAAGGCT</u> CTTTCAGAGC...
HIST1H3E	ATTGTTTGAGTACAAACCTTAA <u>ATCCAAAGGCT</u> CTTTCAGAGC...
HIST1H3F	TTTGTACTGAGGGTGTGCGCCA <u>ACTTAAACCCAAAGGCT</u> CTTTCAGAGC...
HIST1H3G	AGGTTTCCGGGAGTAATTCTAT <u>CCAACCTTAAAGGCT</u> CTTTAAGAGC...
HIST1H3H	GTCTCAAGGACTCA <u>TGATTACATA</u> CCCAAAGGCTCTTTCAGAGC...
HIST1H3I	ATATCCCTAT <u>CAGCCAAAGAAGT</u> CTGTCCAACCCAAAGGCTCTTTCAGAGC...
HIST1H3J	TCCCCTGCTATCTTGGTTCTTAATTGCTTCAAG <u>GCTTCCAAGGCT</u> CTTTCAAGAGC...
HIST2H3A/C	GAAGTGGCGGTT <u>CGGCCGGAGGTTCCATCGTATCCAAAGGCT</u> CTTTCAGAGC...
HIST2H3D	GGCATATTTTAAGTGGTCGAT <u>CTAAAGGCT</u> CTTTCAGAGC...
HIST2H3PS2	GTATATTTTAAGTGGTCGAT <u>CTAAAGGCT</u> CTTTCAGAGC...
HIST3H3	GAGGGCTAT <u>CTGCCACCTGAGAGGTT</u> CGCAACGTT <u>CCCCAAAGGCT</u> CTTTAAGAGC...