FIGURE 13.22. Regions under higher functional constraint show lower diversity in a survey of 213 human genes. Genes are ranked in order of nucleotide diversity, with most diverse on the left and least diverse on the right. The 5′ flanking region and introns are much more diverse than coding sequences and exons.

13.22, modified from Livingston R.J. et al., Genome Res. 14: 1821–1831, © 2004 CSHLP

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