



FIGURE 19.25. Codon usage bias increases with level of gene expression in yeast, *Saccharomyces cerevisiae*. The proportion of sites occupied by the preferred (“major”) codon is plotted against the number of messenger RNA (mRNA) transcripts per cell.

19.25, redrawn from Akashi H., *Curr. Opin. Genet. Dev.* **11**: 660–666, © 2001 Elsevier