Figure. Number of predicted tryptic peptides per *E. coli* protein as a function of mRNA signal intensity: a) bar graph; b) scatter plot. To get the x-axis, the 4291 protein-coding genes were first sorted from lowest to highest mRNA signal intensity. For chart (a), they were then divided into 10 equal classes. Class 1 was the 10% of genes with the lowest intensities, and so on. The ranges of mRNA signal intensities for classes 1 through 10 were, respectively, 1) 5-61, 2) 61-112, 3) 113-176, 4) 176-262, 5) 262-394, 6) 394-611, 7) 612-994, 8) 996-1745, 9) 1747-3689, 10) 3698-58,952.