Figure 2 – High consistency of log-ratio measurements across a 20-fold range of hybridized material. Heatmap of mean-centered log₂(Cy5/Cy3) ratios for 6 genes representing a 32-fold intensity range (see Fig. 1A) across a 20-fold range of hybridized material. For each gene the maximum fold-difference ($D_{\text{max}}$) between any two of the triplicate averages is also shown in percent.