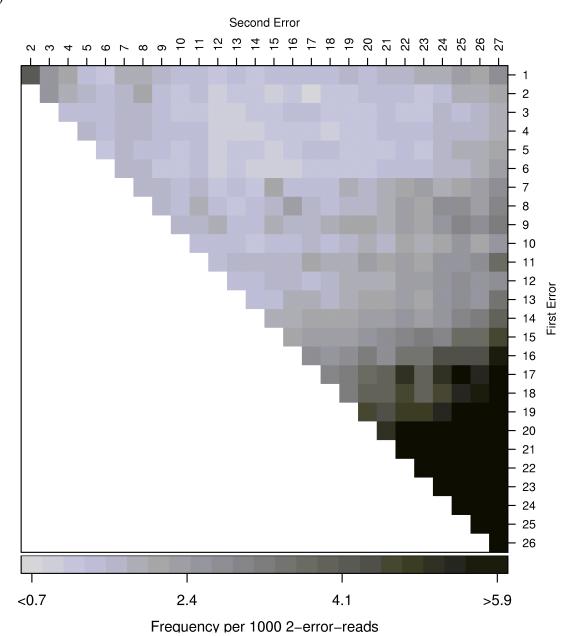
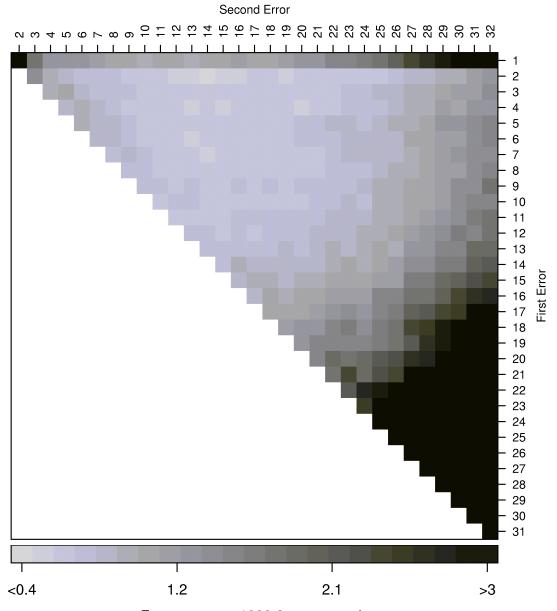
Supplementary Figure 3

Heatmaps illustrating error positions in Solexa reads containing two errors. Frequent error position combinations are shown in black, rare combinations in gray shades. The color bar at the bottom indicates gray shades and frequencies. For the *Beta* data set (a) and for the *Helicobacter* data set (b) we observe cumulations of errors towards the end and the beginning of the reads.

a)





Frequency per 1000 2-error-reads