

Supplementary Figure 5:

Sequence context of wrong base calls in Solexa reads from *Helicobacter acinonychis* and *Beta vulgaris*, considering two bases upstream and downstream of the wrong base calls. An “e” indicates the substituted base. The scatterplot shows the correlation of the relative frequencies (relating the frequency of 5-tuples at error positions to the frequency of all 5-tuples in the reads) for the two data sets.

