Supplementary Figure 6. Detailed schematic representation of *C. hominis* chromosome VI. *C. hominis* chromosome VI sequence aligned to its *C. parvum* homolog (shown as scales above panels). Tracks indicate: a) *C. hominis* contigs (blue), sequence gaps (white) and physical gaps (red); b) HAPPY markers (blue, see manuscript for reference) and positions of the octamers TGGCGCCA (red) and TGCATGCA (green); c) Gene Ontology classification for the predicted genes (see key); d) tRNAs (blue) and rRNAs (magenta); e) percent identity between *C. hominis* and *C. parvum* in 5 kb windows; and f) BAC clone coverage (overlapping clones collapsed to a single line).