

Supplementary Figure 1:

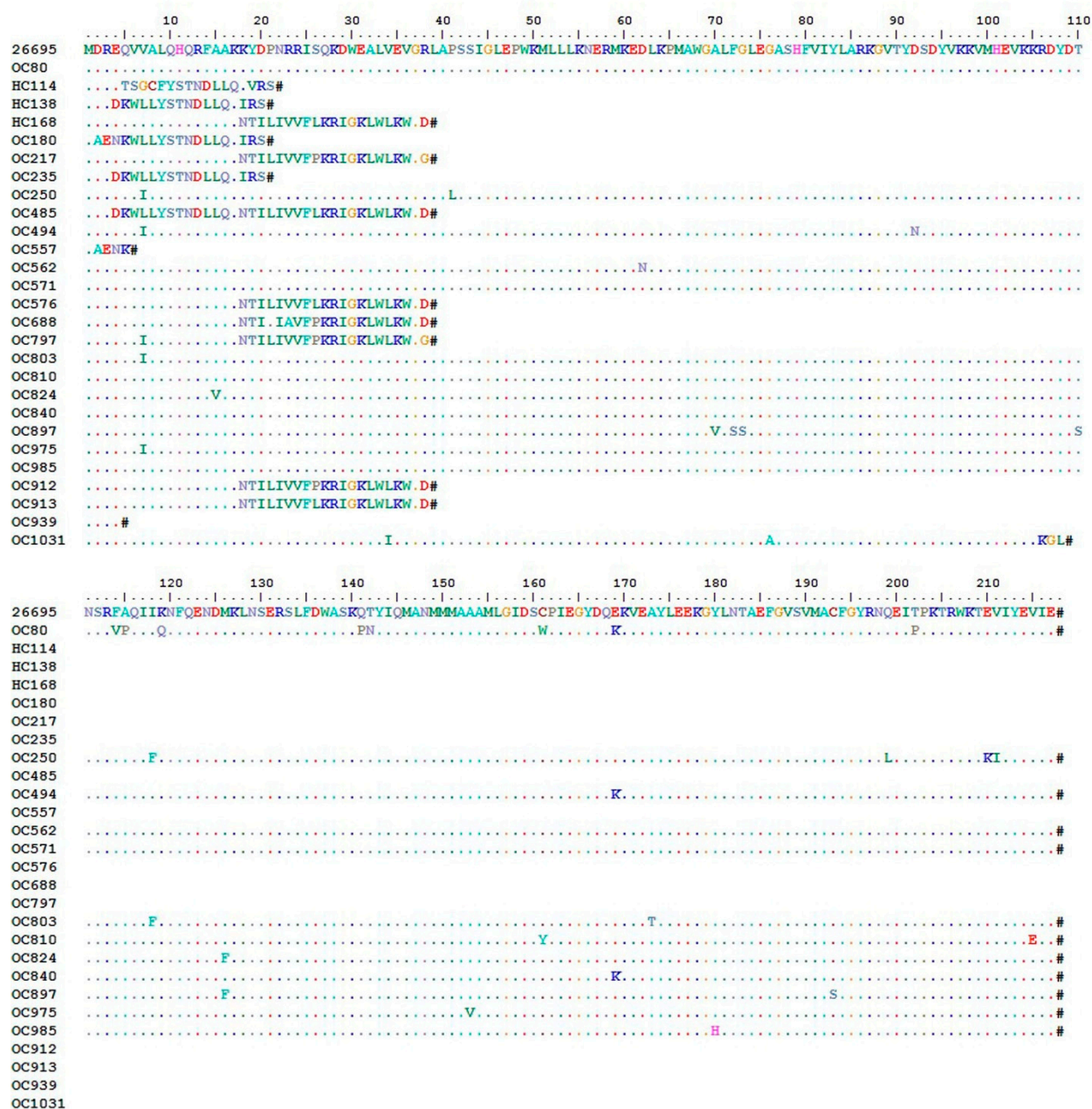


Figure S1. Amino acid sequence alignment of FrxA in metronidazole-susceptible and -resistant *H. pylori* isolates (n = 27) as compared with *H. pylori* reference strain 26695. The stop codon and premature truncation in peptide translation is indicated with (#).

Supplementary Figure 2:



Figure S2. Amino acid sequence alignment of RdxA in metronidazole-susceptible and -resistant *H. pylori* isolates (n = 28) as compared with *H. pylori* reference strain 26695. The stop codon and premature truncation in peptide translation is indicated with (#).

Supplementary Figure 3:



Figure S3. Amino acid sequence alignment of GyrA in fluoroquinolone-susceptible and -resistant *H. pylori* isolates (n = 31) as compared with *H. pylori* reference strain 26695.

Supplementary Figure 4:



Figure S4. Amino acid sequence alignment of GyrB in fluoroquinolone-susceptible and -resistant *H. pylori* isolates (n = 27) as compared with *H. pylori* reference strain 26695.

Supplementary Figure 5:



Figure S5. Nucleotide sequence alignment of 23S rRNA in clarithromycin-susceptible and -resistant *H. pylori* isolates (n = 24) as compared with *H. pylori* reference strain 26695.