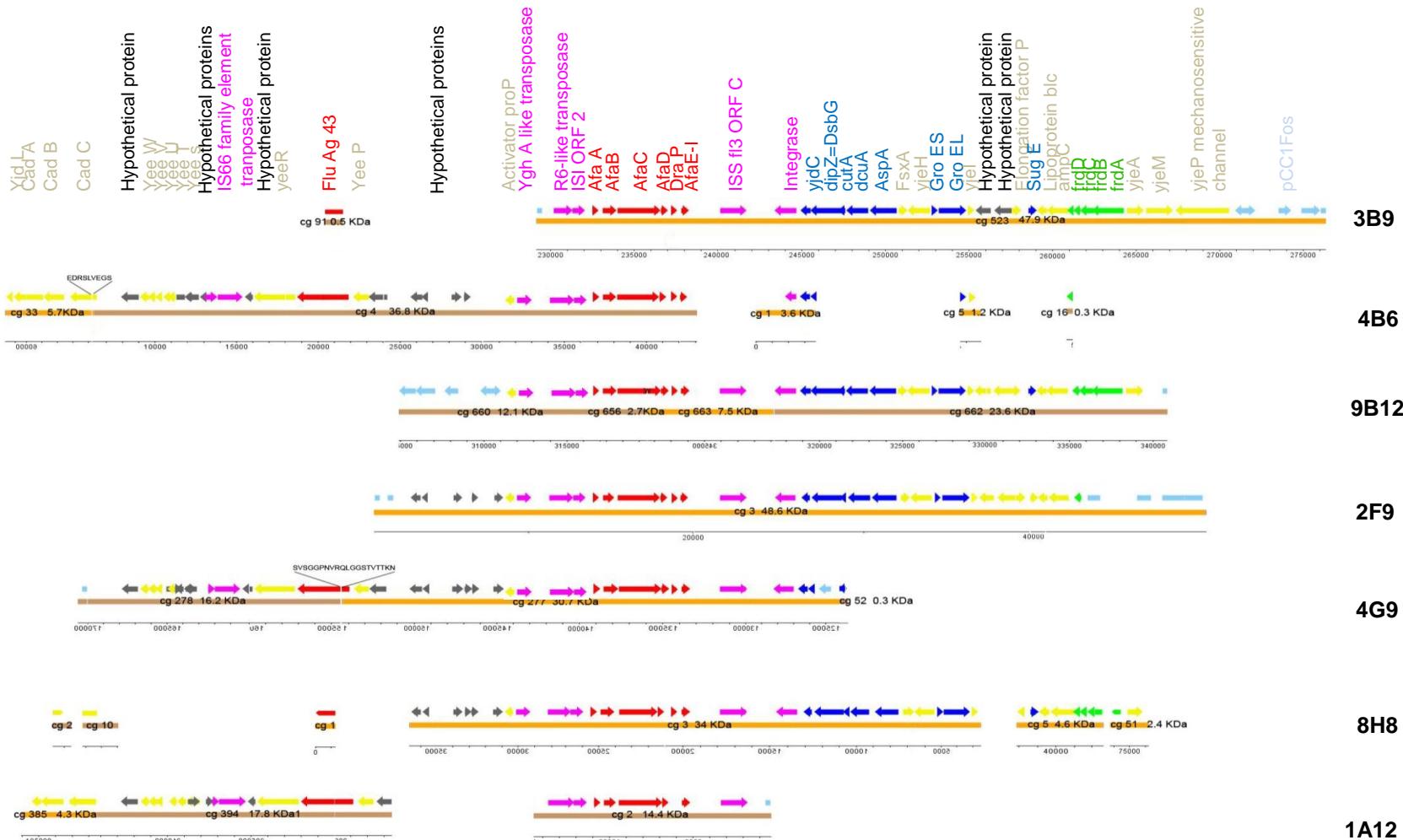


Supplementary file S3

Figure S3: 454 sequence alignment of fosmids extracted from haemagglutinin (HA) positive library clones. The *afa-1* operon coding for afimbral adhesin was identified as an element common to 7 of 8 HA-positive fosmids*. Sequence information for the eighth fosmid (2F4) was not interpretable due to excessive noise but presence of the *afa* operon was subsequently confirmed by PCR.



*The nucleotide sequence of the *afa-1* cluster has been submitted to GenBank; Accession number JN688153