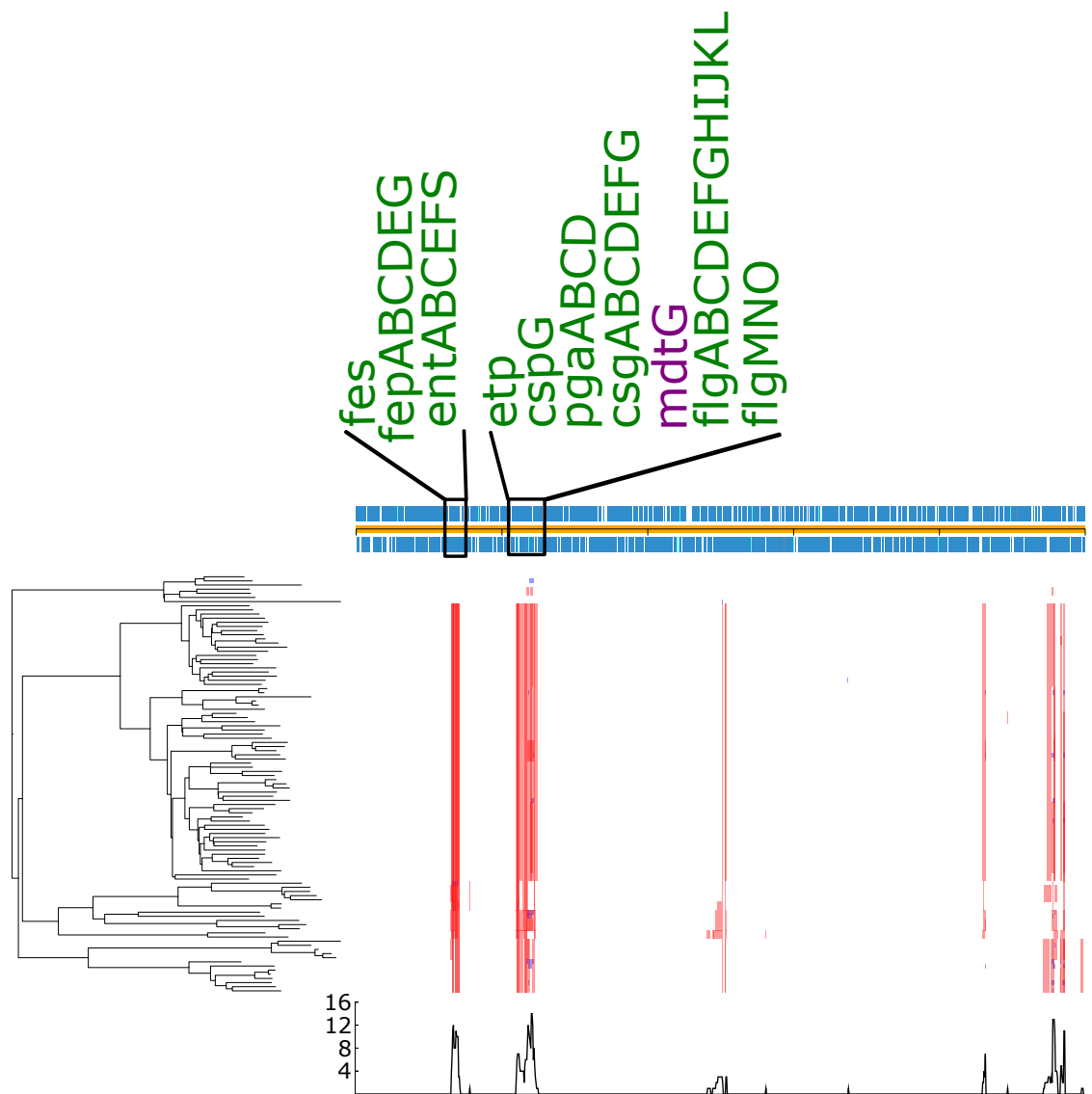


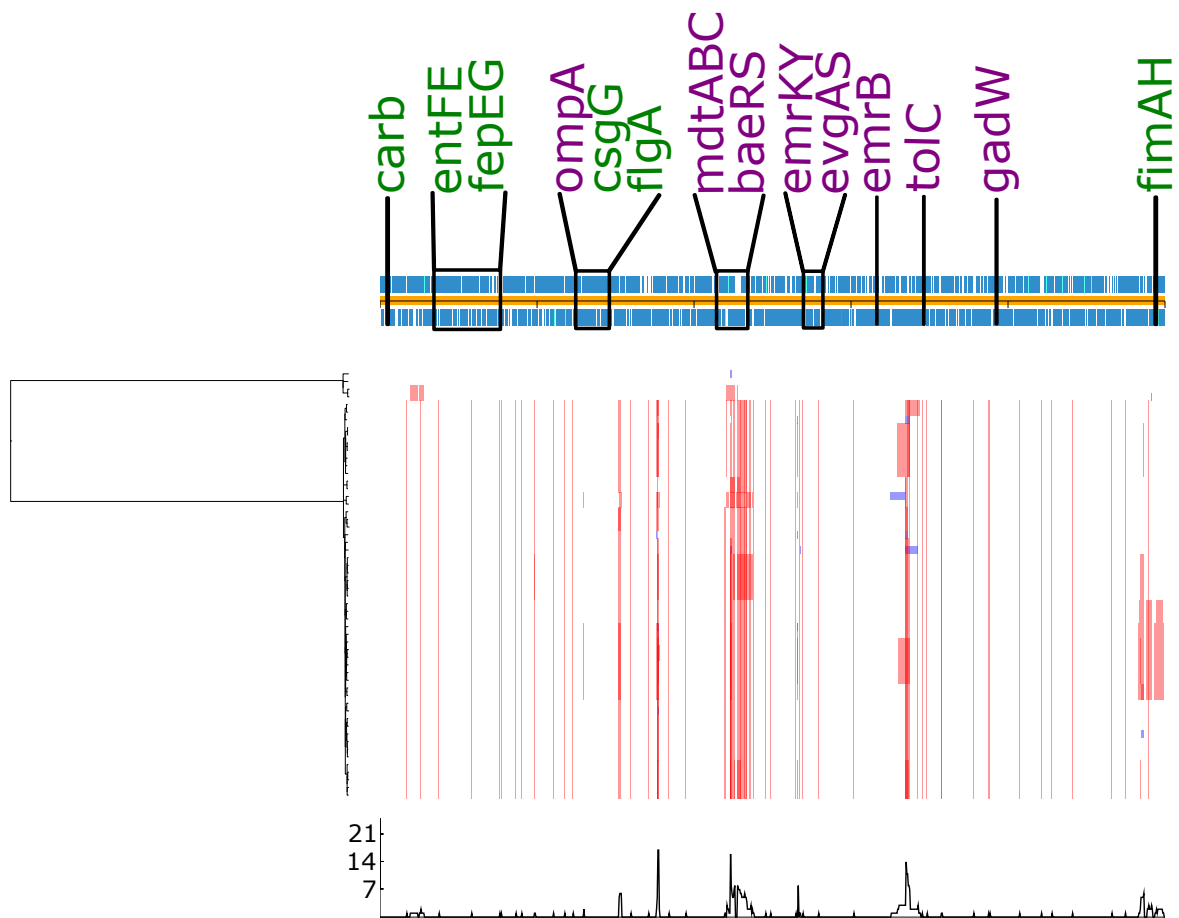
Lineage-specific variation in frequency and hotspots of recombination in invasive *Escherichia coli*

Kathryn R. Piper, Stephanie S.R. Souza, Odion O. Ikhimiukor, Adrienne Workman, Isabella W. Martin, Cheryl P. Andam

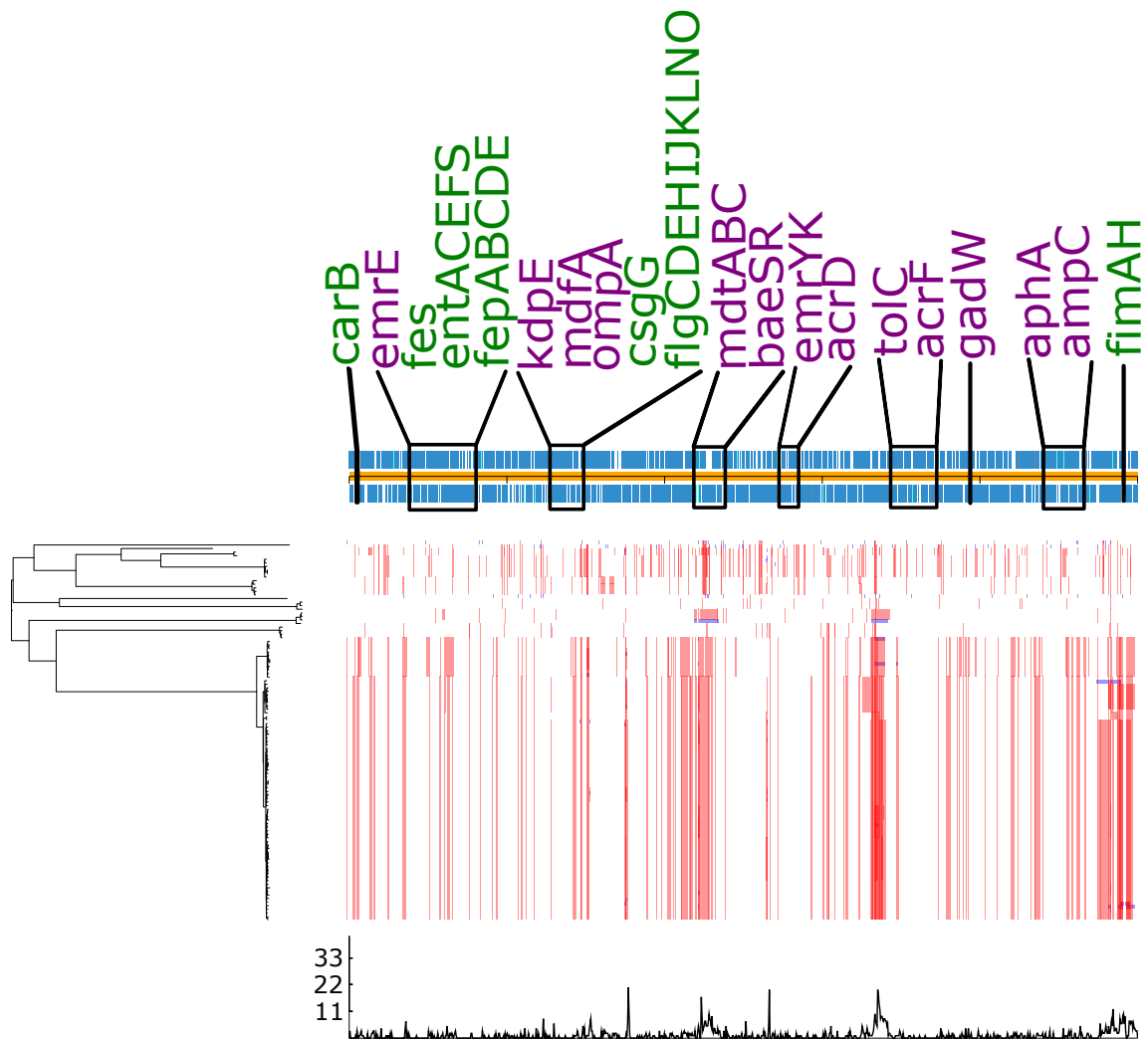
Supplementary Material



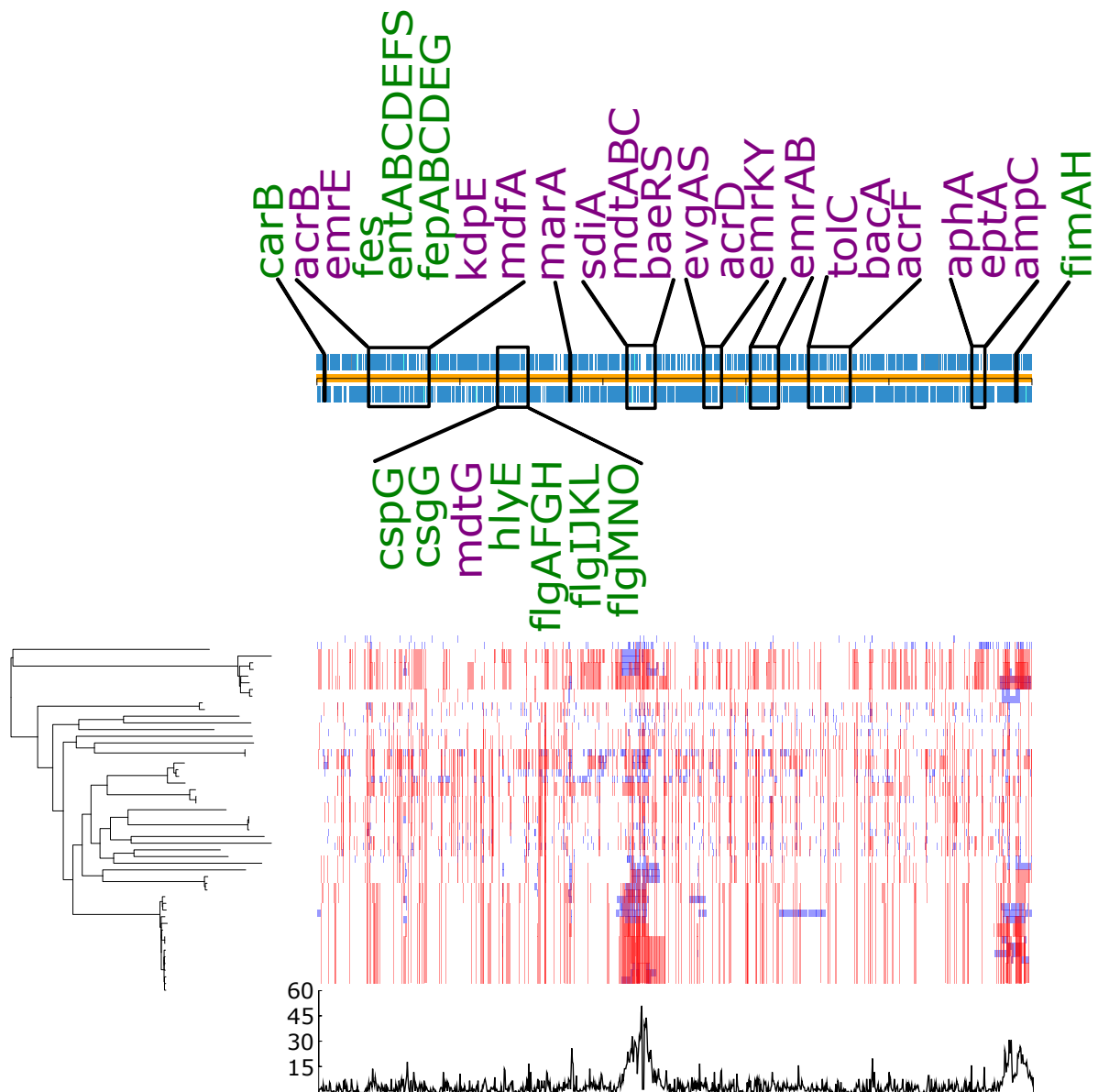
Supplementary Figure S1. Recombination hotspots in sequence cluster BAPS1 inferred by Gubbins.



Supplementary Figure S2. Recombination hotspots in sequence cluster BAPS4 inferred by Gubbins.



Supplementary Figure S3. Recombination hotspots in sequence cluster BAPS10 inferred by Gubbins.



Supplementary Figure S4. Recombination hotspots in sequence cluster BAPS14 inferred by Gubbins.