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RESEARCH ARTICLE

Whole-Genome Random Sequencing
and Assembly of Haemophilus influenzae Rd
Circular representations of the Haemophilus influenzae Rd genome, which has now been sequenced and assembled. The illustration depicts the locations of genes color coded by their role in the organism, the underlying clone scaffolding used to confirm the structure, locations of ribosomal RNA operons, and other global characteristics of the DNA sequence. See page 538 and the related News story (page 468) and Research Article (page 496). [Image: A. R. Kerlavage]

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Indicates accompanying feature

OB protein weighs in

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