

Lyme, Informatics, & Evolution

[Qiu Lab @Belfer](#)

Department of Biological Sciences
Hunter College, City University of New York

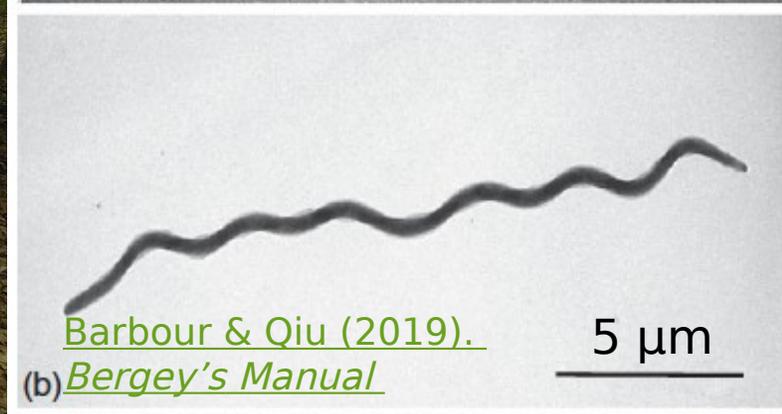
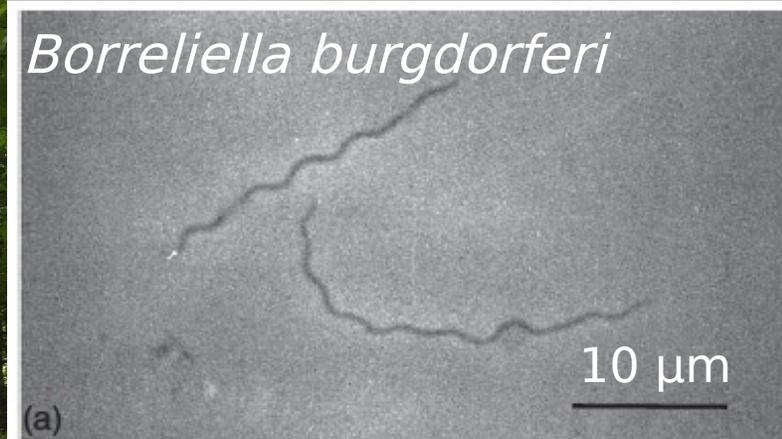
Ixodes scapularis



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Photo-credit: Pedro Peg



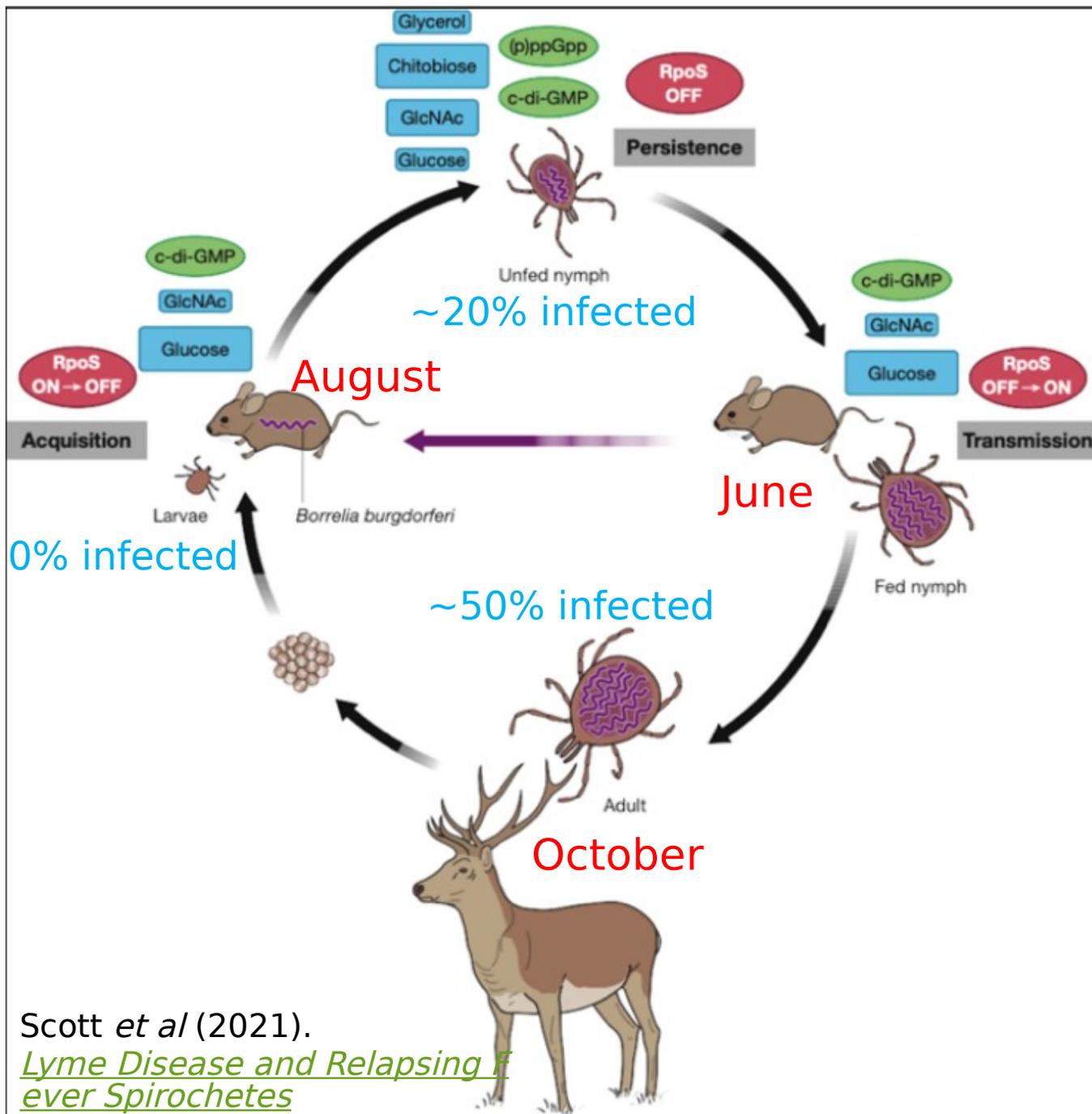
Borrelia burgdorferi



[Barbour & Qiu \(2019\).](#)

(b) [Bergey's Manual](#)

Lyme Disease

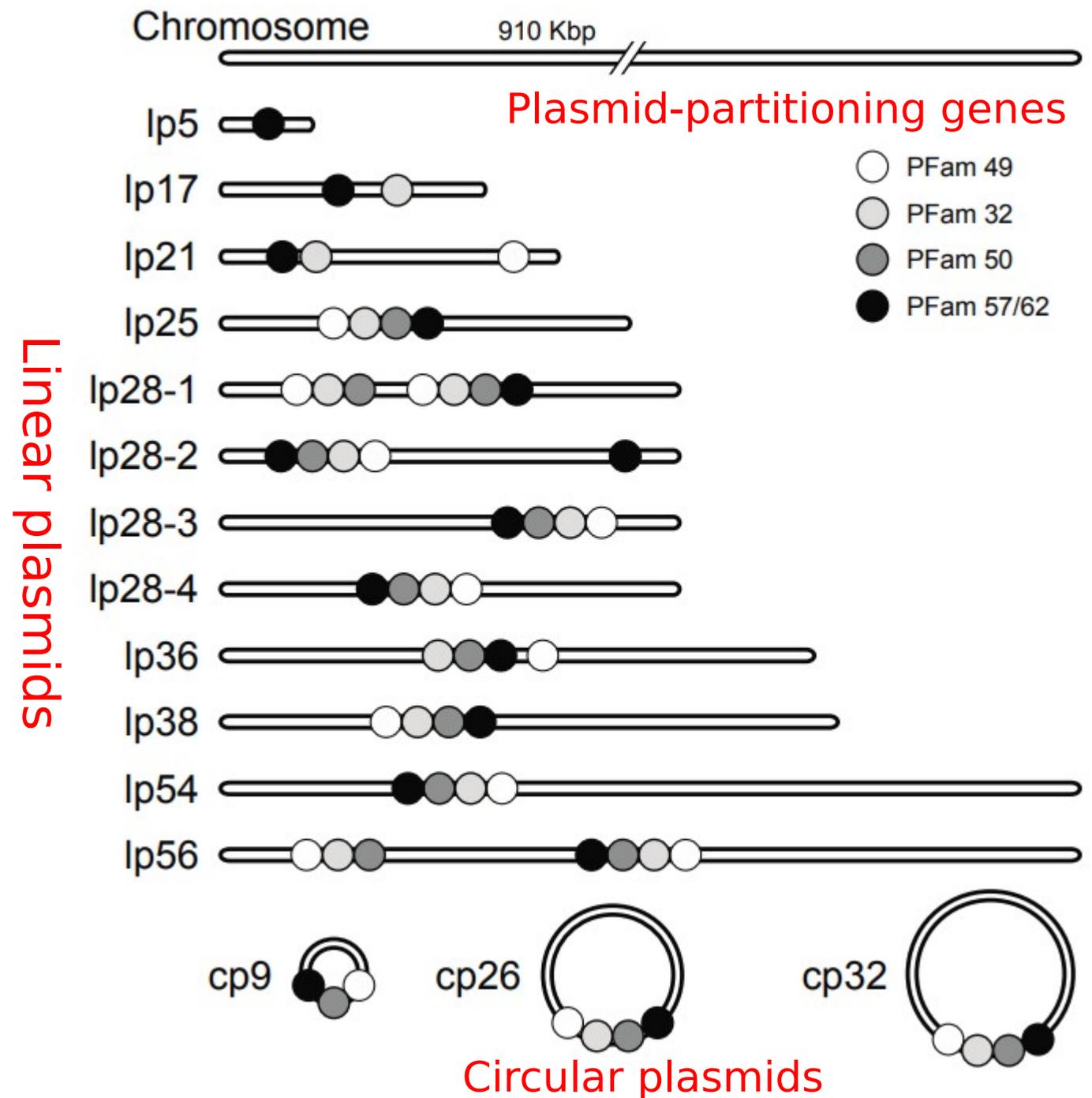


- Symptoms
 - Stage I. Skin infections
 - Stage II. Arthritis
 - Stage III. Chronic infections
- Most common vector-borne disease in US and Europe; 300,000 annual cases in the US (Northeast & Midwest)
- Poor/Controversial diagnostics
- No human-use vaccine
- No targeted antibiotics

Scott *et al* (2021).
Lyme Disease and Relapsing Fever Spirochetes

Borrelia Genomics

- **Multipartite Genomes**
- **Bi-phasal** (tick & mammalian) switching



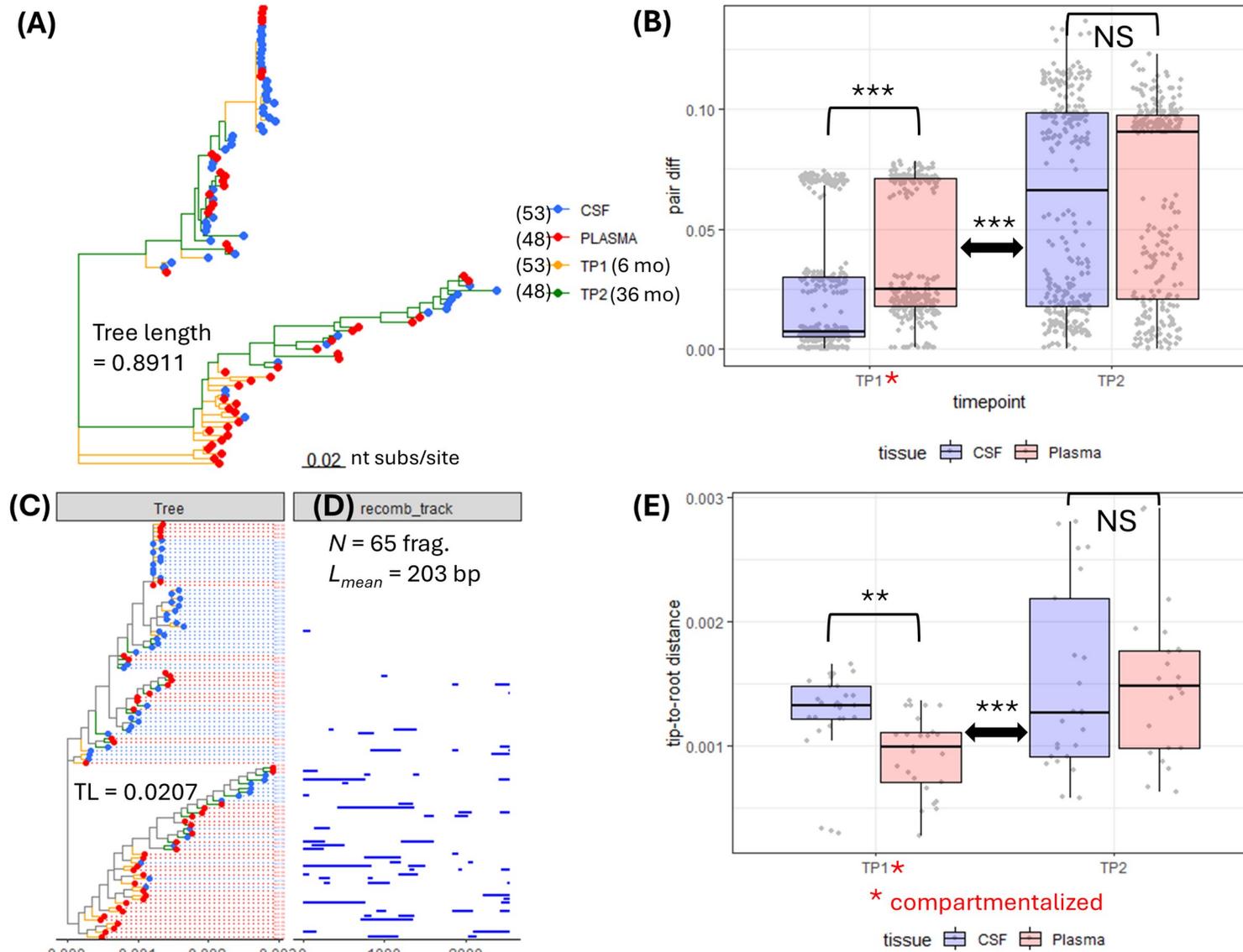
Schwartz *et al* (2021).

Lyme Disease and Relapsing Fever Spirochetes

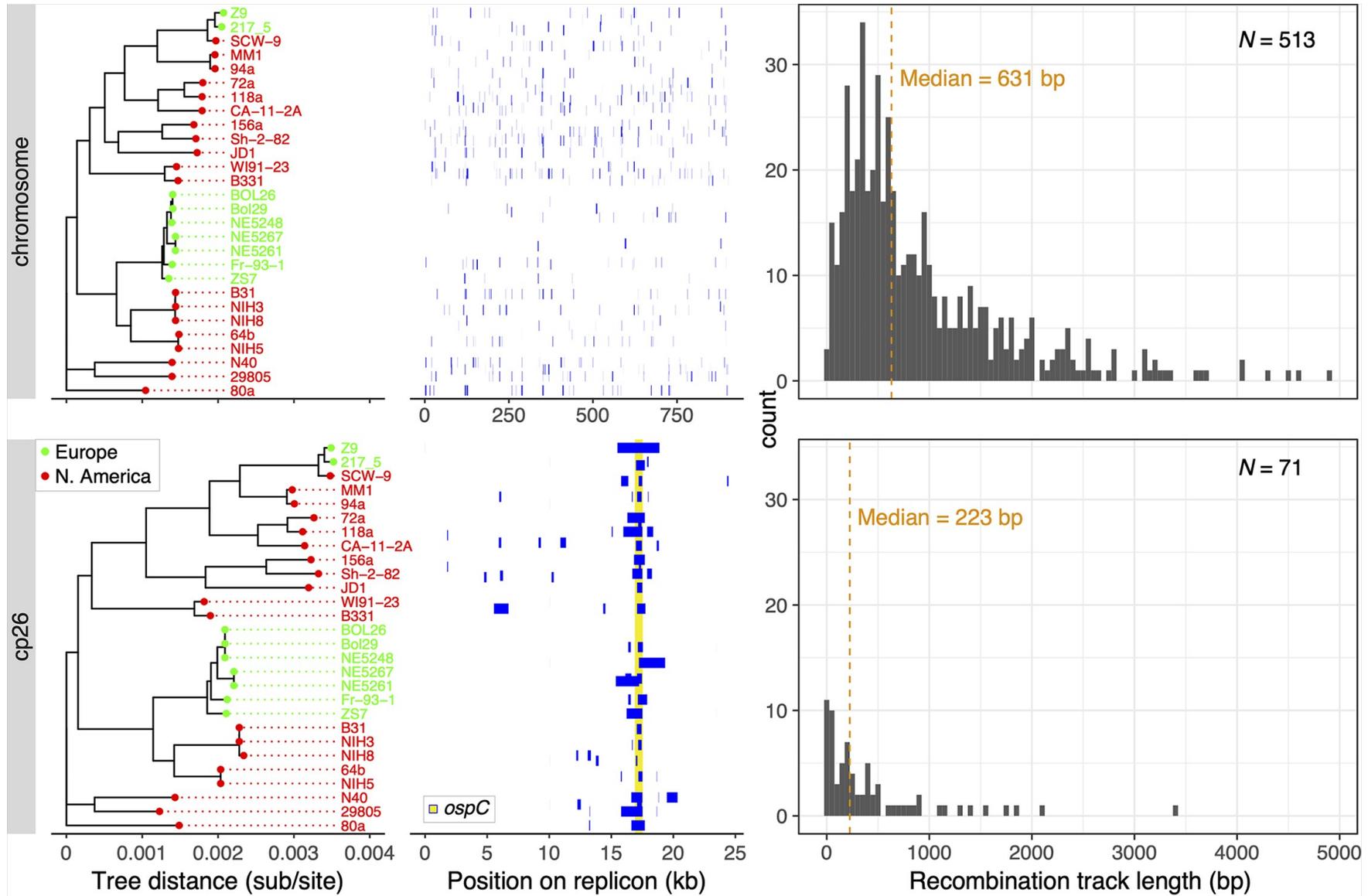
Microbes Recombine (not clonal)

Pathogenicity & Virulence

HIV evolution within a single host



Borrelia diversification



Evolutionary History & Mechanisms



8 | Editor's Pick | Genomics and Proteomics | AAM Contribution-Research Article

Natural selection and recombination at host-interacting lipoprotein loci drive genome diversification of Lyme disease and related bacteria

Saymon Akther,¹ Emmanuel F. Mongodin,² Richard G. Morgan,³ Lia Di,¹ Xiaohua Yang,⁴ Maryna Golovchenko,⁵ Natalie Rudenko,⁵ Gabriele Margos,⁶ Sabrina Hepner,⁶ Volker Fingerle,⁶ Hiroki Kawabata,⁷ Ana Cláudia Norte,⁸ Isabel Lopes de Carvalho,⁹ Maria Sofia Nuncio,⁹ Adriana Marques,¹⁰ Steven E. Schutzer,¹¹ Claire M. Fraser,² Benjamin J. Luft,⁴ Sherwood R. Casjens,¹² Weigang Qiu^{1,13}

- Phylogenetic coverage: 47 new genomes, all 23 species, 15 major US pathogen strains
- Leading institutions: U. Maryland, U. Utah, Stony Brook U, NJ Medical School, Hunter/CUNY
- Participants: Japan, Germany, Portugal, Czech Republic

BpWrapper: a Command-Line Utilities

