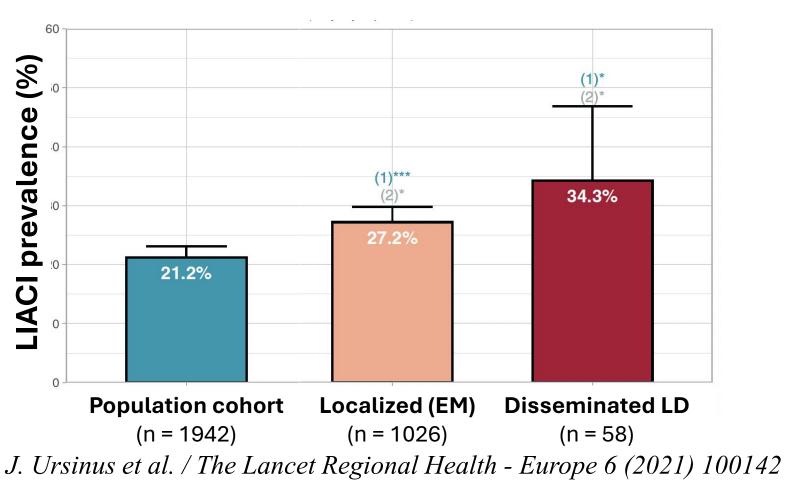
Assessing contribution of *B. burgdorferi* genomic diversity to disseminated infection and persistent symptoms of LD

Research for Lyme Infection-Associated Chronic Illnesses Treatment Broadening the Lens – Committee Meeting 2 July 11, 2024 John Leong, MD, PhD





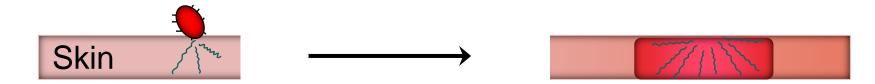
A. Persistent symptoms

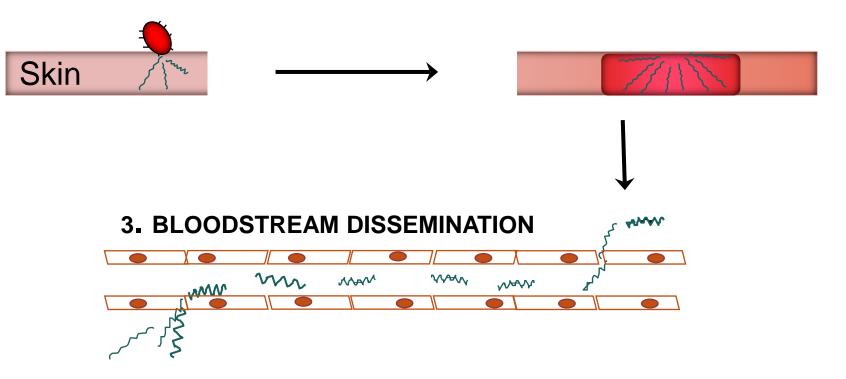


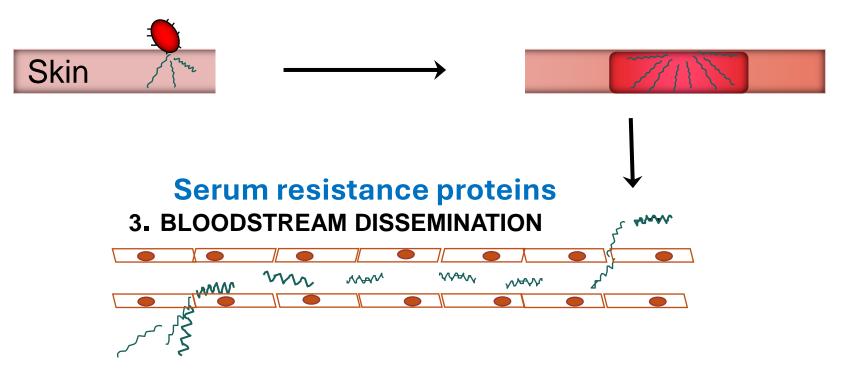
A. Persistent symptoms B. Specific *Bb* genotypes ospC 60 % isolates (%) genotype o 20 40 60 80 100 $(1)^{3}$ Α prevalence Η 34.3% Ν 27.2% **Disseminated** B 21.2% Localized IACI Κ Ε Localized (EM) Population cohort **Disseminated LD** (n = 1942)(n = 1026) (n = 58) G J. Ursinus et al. / The Lancet Regional Health - Europe 6 (2021) 100142 F

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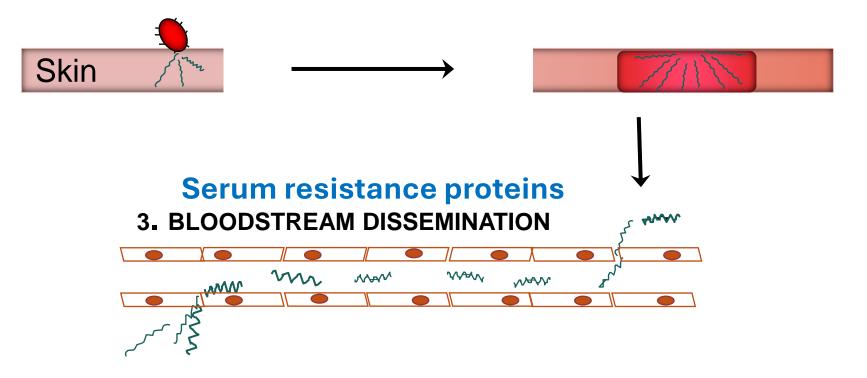
Does microbial genotype contribute to Lyme IACI and if so, how?







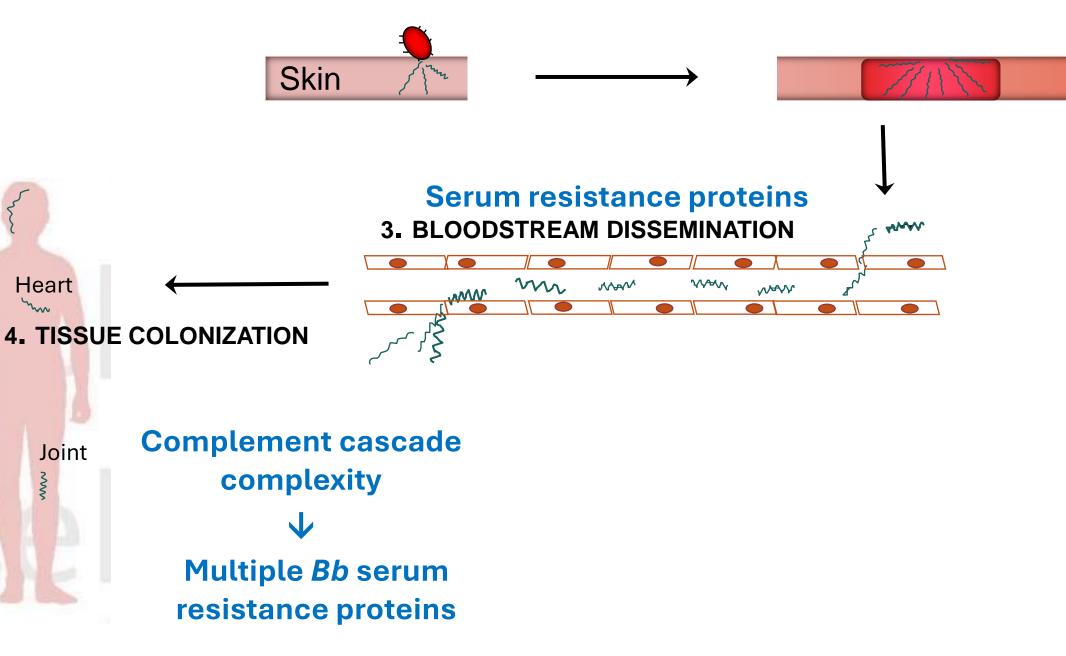
2. SKIN INFECTION



Complement cascade complexity ↓ Multiple *Bb* serum resistance proteins

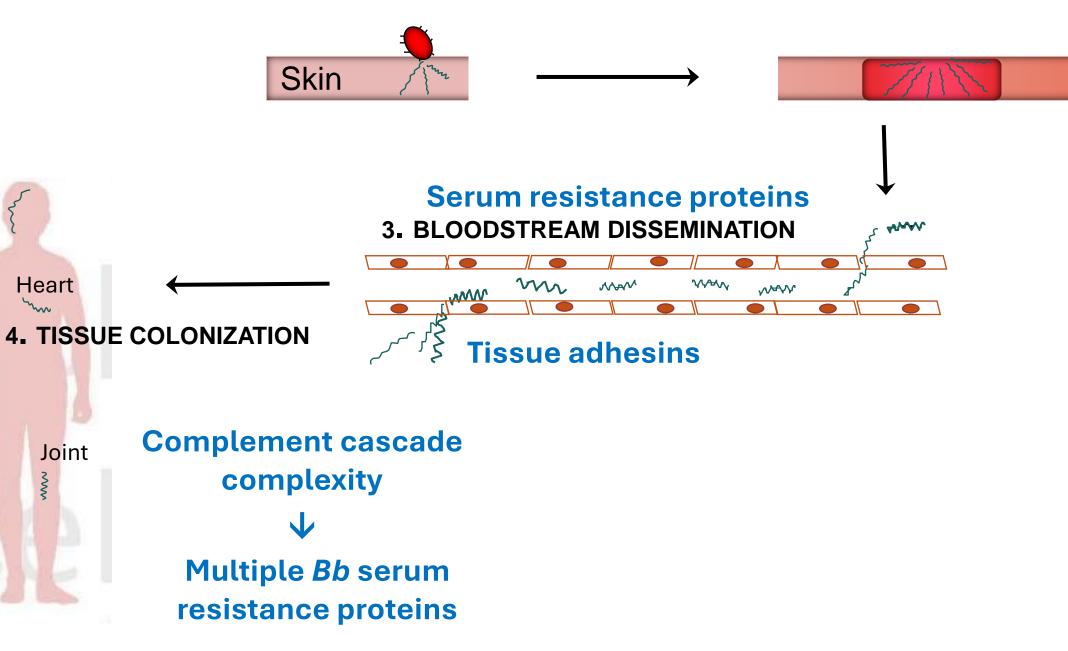
CNS

Skins



CNS

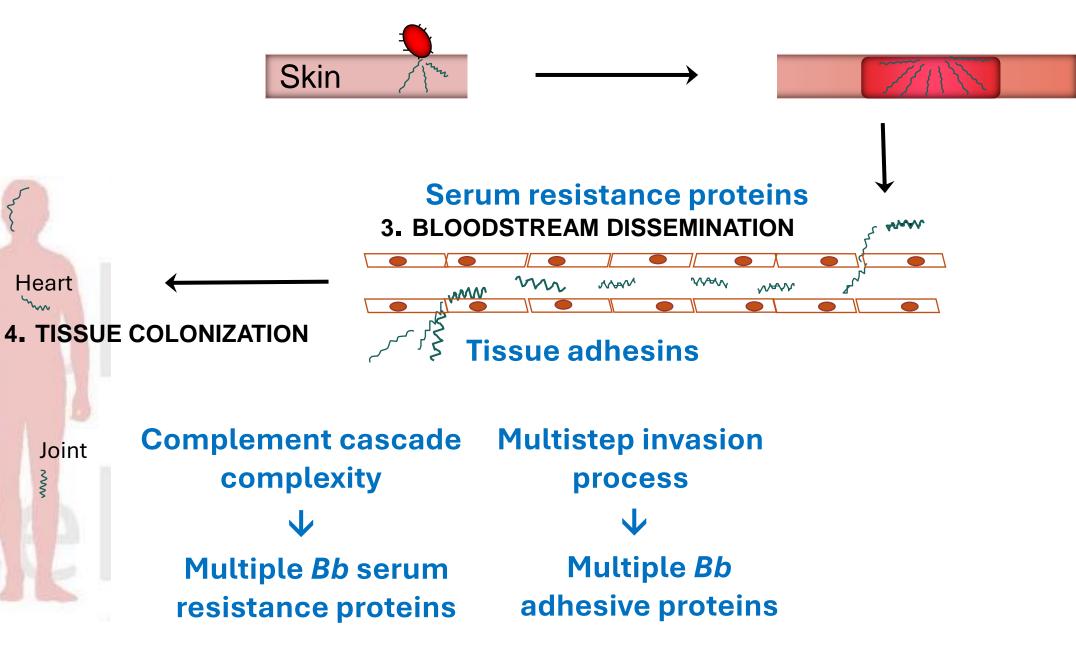
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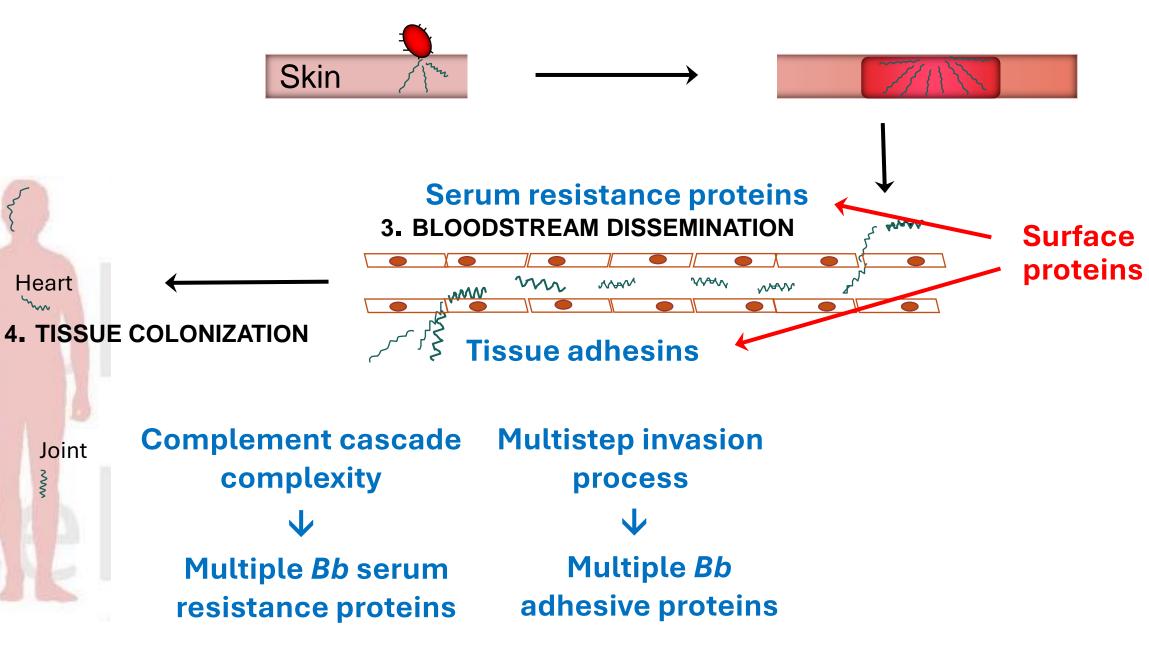
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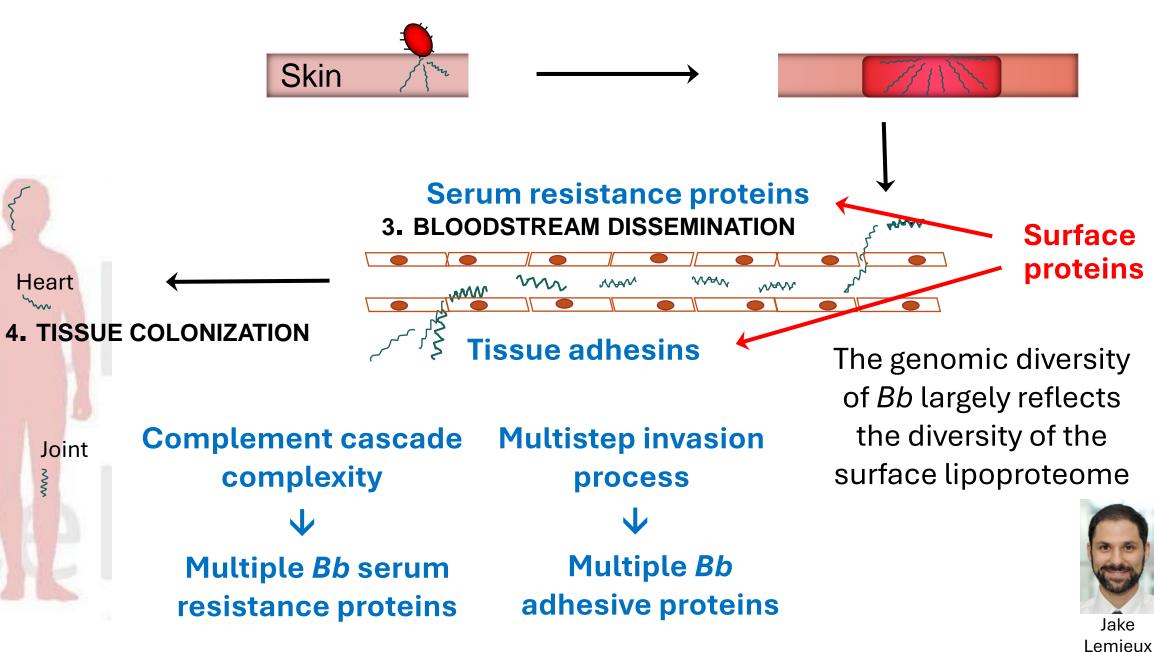


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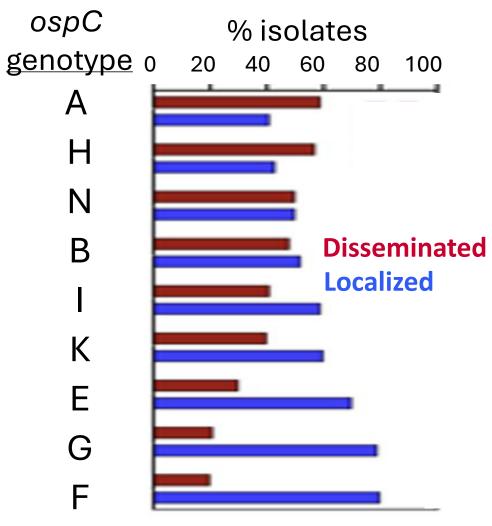




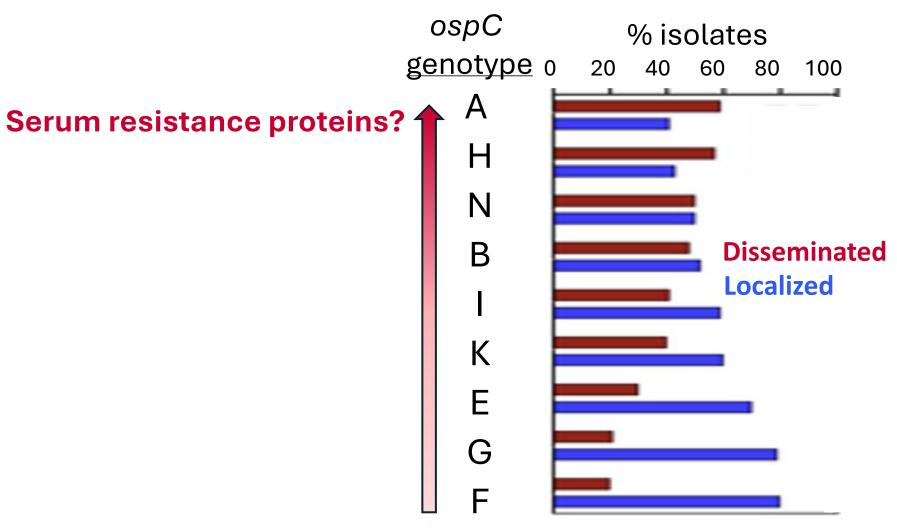
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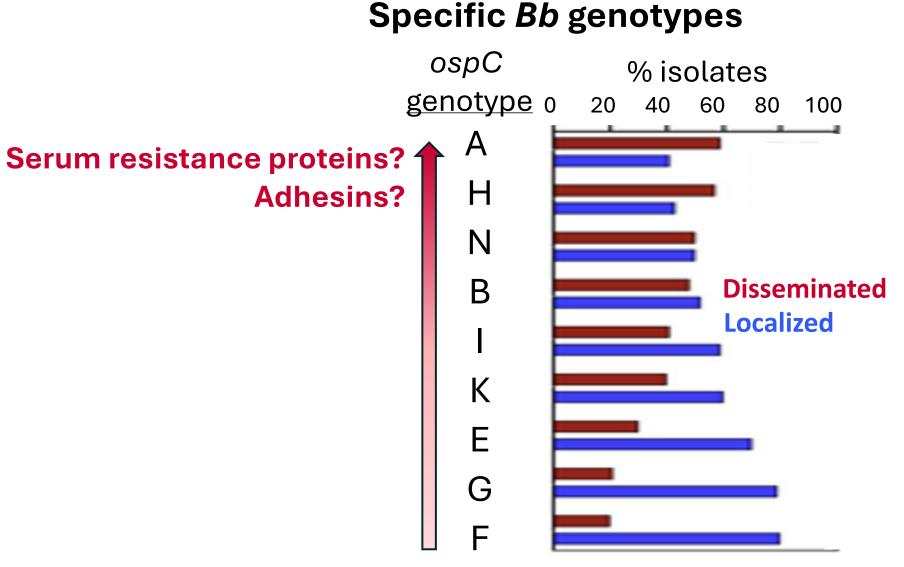


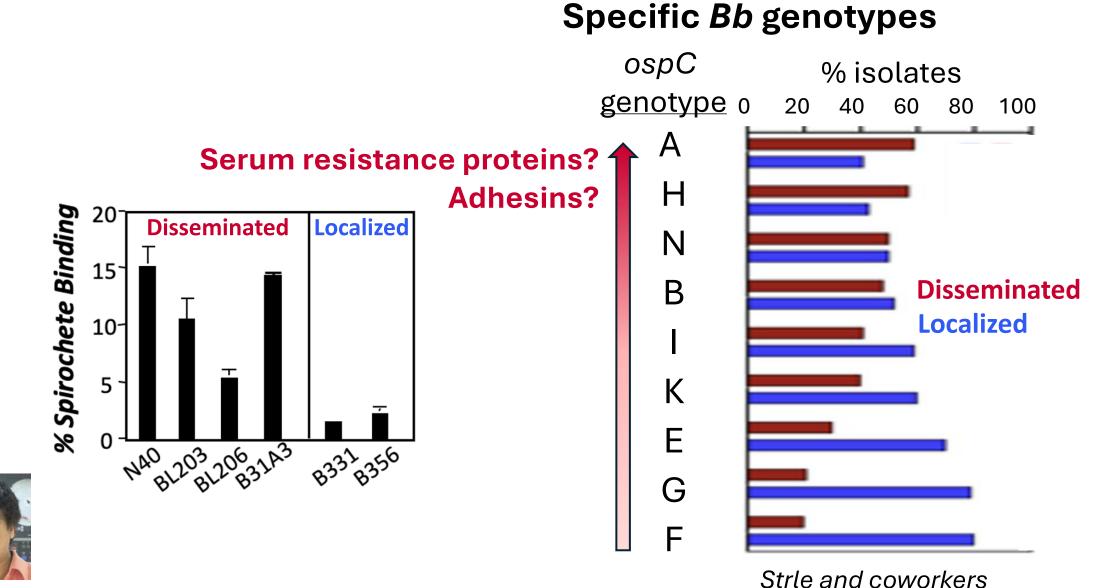
#### Specific *Bb* genotypes





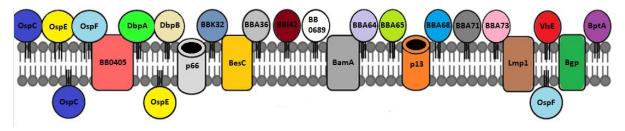




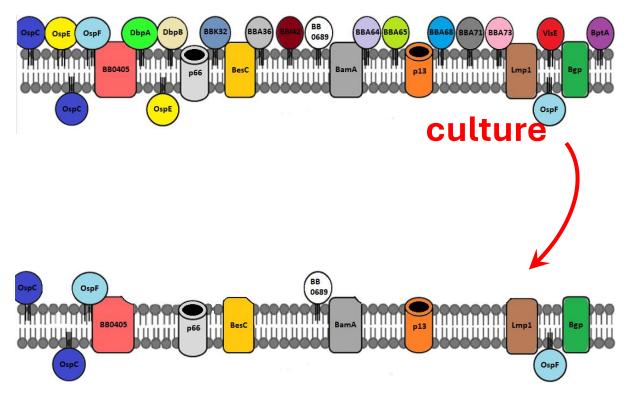


Ira Nikhat Schwartz Parveen

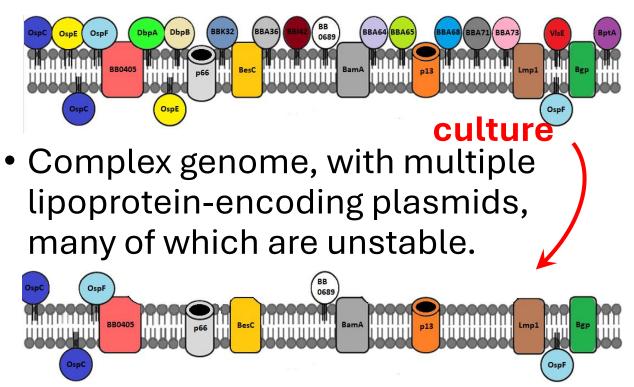
#### **Experimental Challenges**



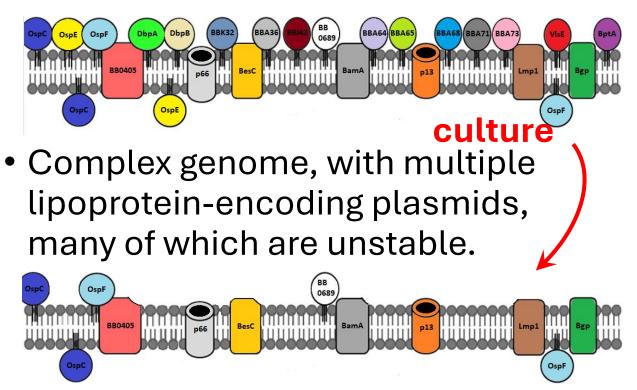
#### **Experimental Challenges**



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#### **Experimental Challenges** Approaches



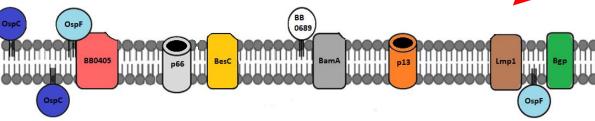
#### **Experimental Challenges**

 >80 surface lipoproteins, many with multiple functions, leading to functional redundancy

 Complex genome, with multiple lipoprotein-encoding plasmids, many of which are unstable.

BesC

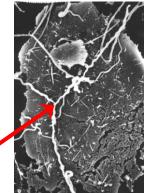
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#### Approaches

• In vitro cell binding assays

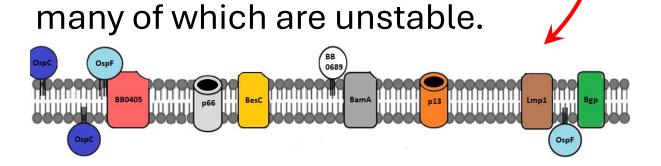
Bound Bb



#### **Experimental Challenges**

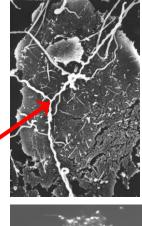
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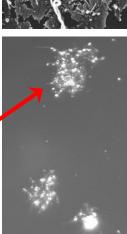


#### Approaches

- In vitro cell binding assays
   Bound Bb
- In vitro serum killing assays



Dead Bb <



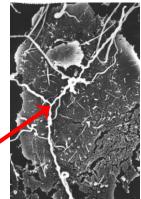
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BesC BamA

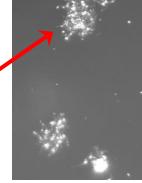
#### Approaches

In vitro cell binding assays
 Bound Bb



• In vitro serum killing assays

"Gain-of-function" Bb strains

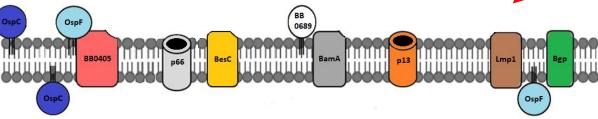


**Dead Bb** 

#### **Ectopic production**

• Complex genome, with multiple lipoprotein-encoding plasmids, many of which are unstable.

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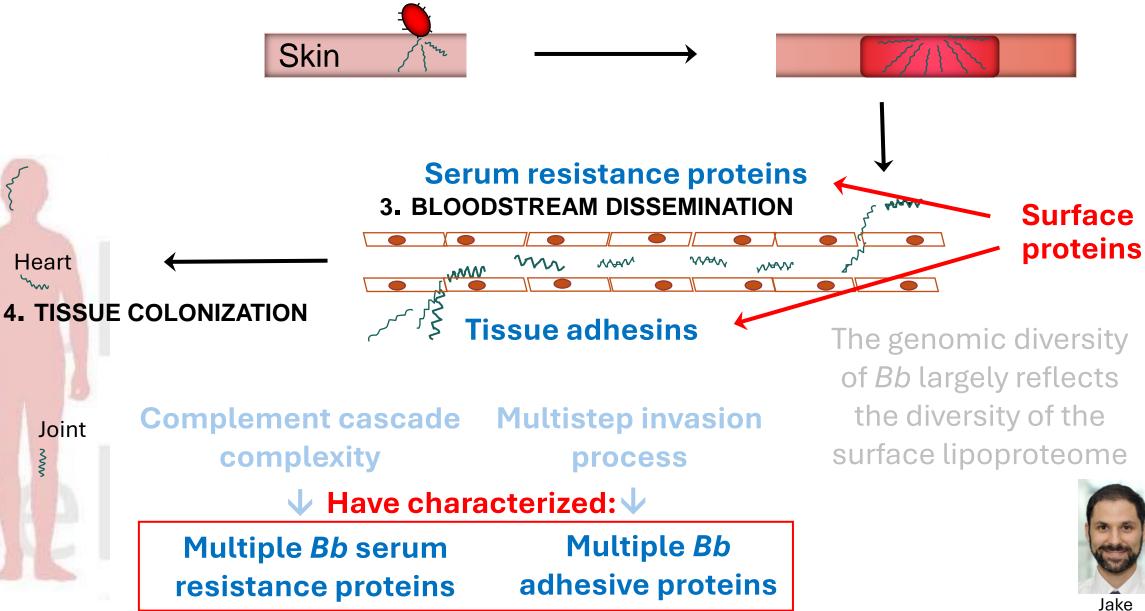
culture



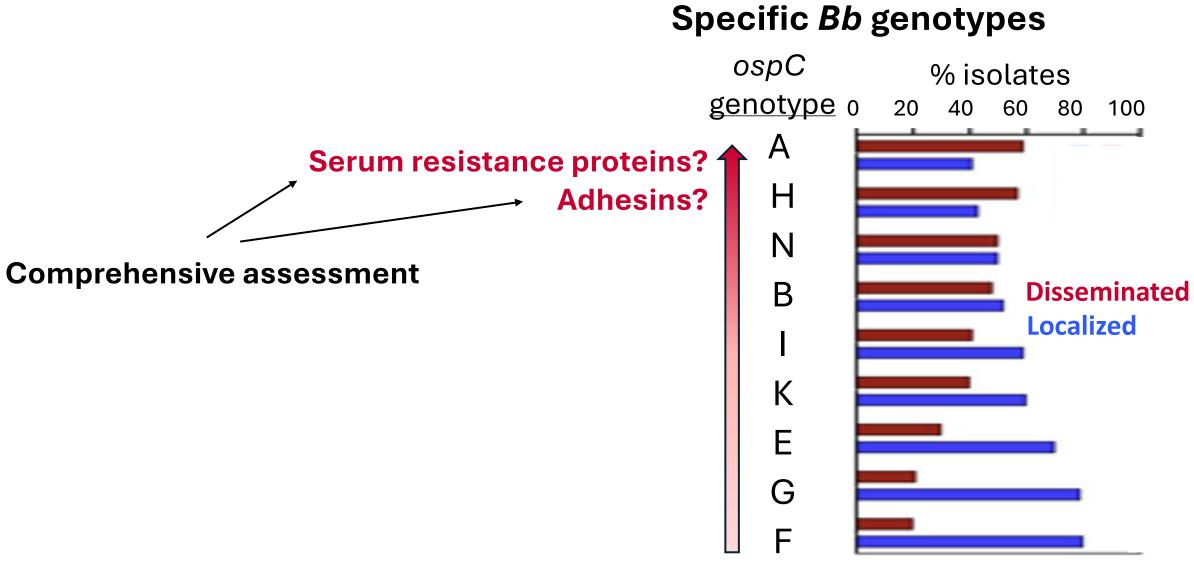
Skins

3

2. SKIN INFECTION



Lemieux



**Experimental Challenges** 

#### **Experimental Challenges**

- Enumeration of *Bb* is labor intensive.
  - Plating efficiency variable.
  - Staining/OD-based methods unreliable.

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• Automated, stain-free, flow cytometry-based counting



Loranne Magoun

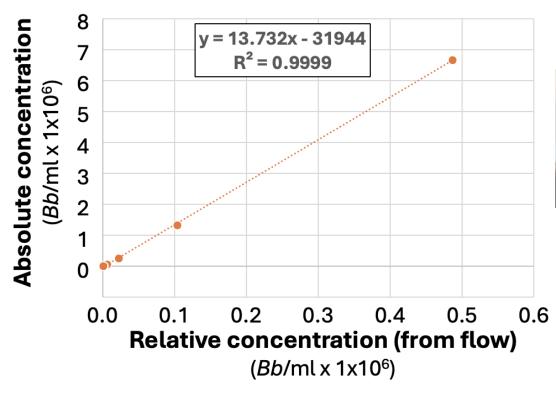


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Loranne Magoun



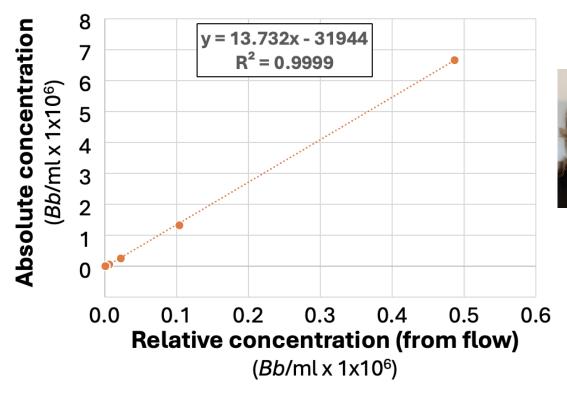
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- Enumeration of *Bb* is labor intensive.
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  - Staining/OD-based methods unreliable.

Permits rapid, large-scale, microtiter well-based screening of surface proteins for serum resistance or adhesive activity

#### Approaches

• Automated, stain-free, flow cytometry-based counting

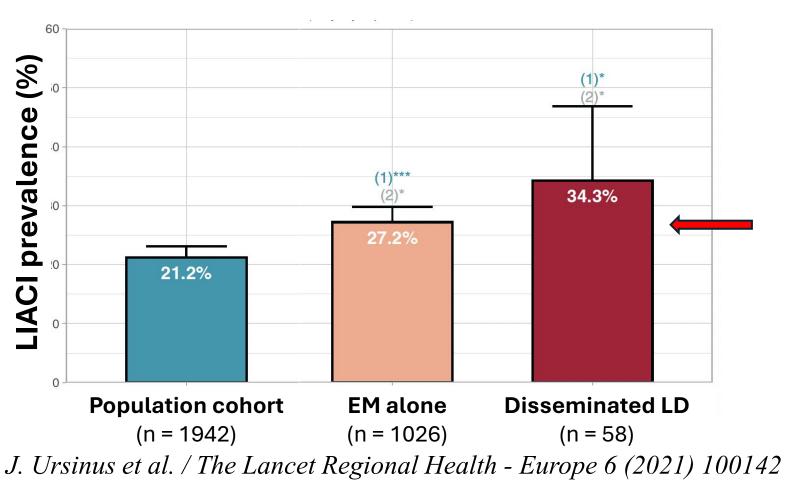






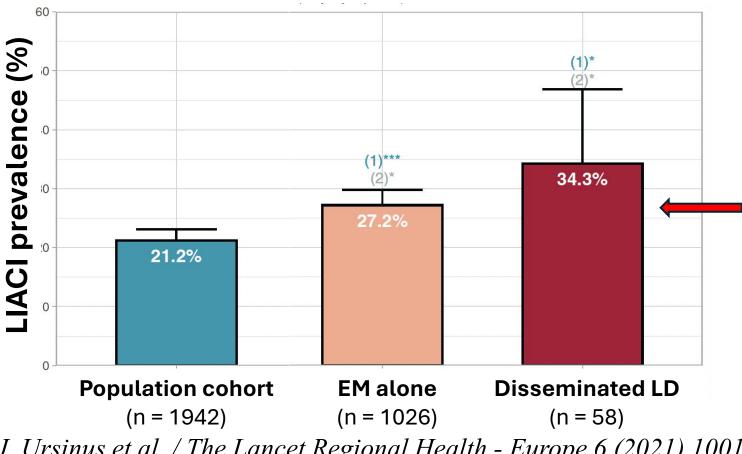
Linda Xu

#### **A. Persistent symptoms**



# Does microbial genotype contribute to Lyme IACI and if so, how?

#### **A. Persistent symptoms**



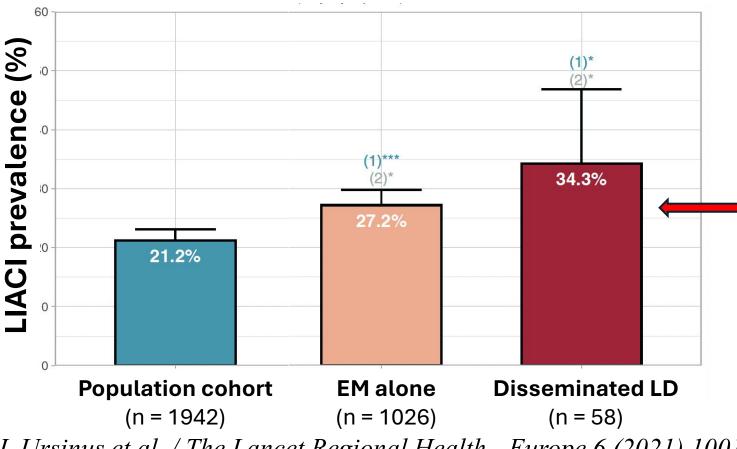
#### J. Ursinus et al. / The Lancet Regional Health - Europe 6 (2021) 100142

#### **Does microbial genotype** contribute to Lyme IACI and if so, how?

#### **Future:**

Collaborate with clinicians to analyze Bb isolates from LD patients with localized or disseminated infection and/or infection-associated chronic illness.

#### A. Persistent symptoms



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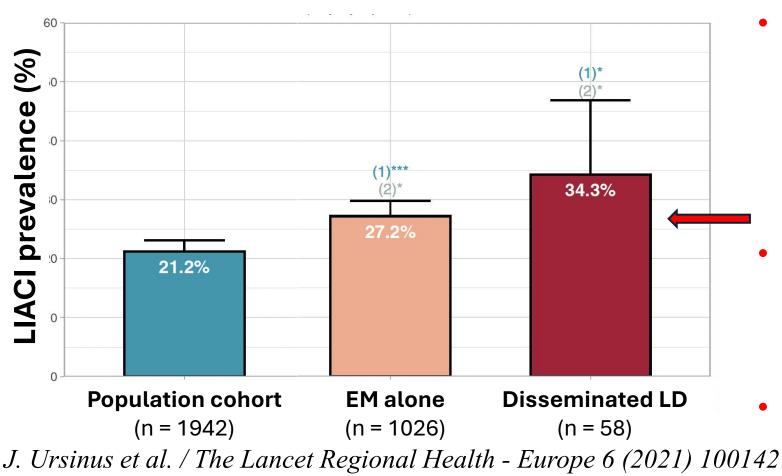
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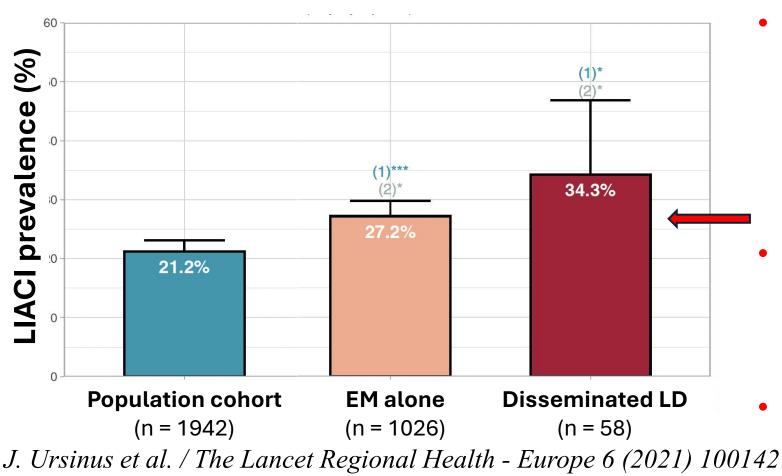


# Does microbial genotype contribute to Lyme IACI and if so, how?

#### **Future:**

- Collaborate with clinicians to analyze Bb isolates from LD patients with localized or disseminated infection and/or infection-associated chronic illness.
- Apply new screening methods to identify their serum resistance and tissue adhesive proteins.
  - Assess associations of these factors with disseminated infection and infectionassociated chronic illness.

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Loranne Magoun

Linda Xu Mass General Hospital Harvard University



Jake Lemieux



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Carol Lyn Piazza and Graham Willsey in the Mantis lab, Wadsworth Institute