40 *Populus deltoides* Microbiome Genomes

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- To aid in the investigation of the *Populus deltoides* microbiome, we generated draft genome sequences for twenty-one *Pseudomonas* and nineteen other diverse bacteria isolated from *Populus deltoides* roots
- Draft genome sequences for isolates similar to *Acidovorax, Bradyrhizobium, Brevibacillus, Caulobacter, Chryseobacterium, Flavobacterium, Herbaspirillum, Novosphingobium, Pantoea, Phyllobacterium, Polaromonas*, *Rhizobium, Sphingobium* and *Variovorax* were generated and comparative genomic studies are underway.

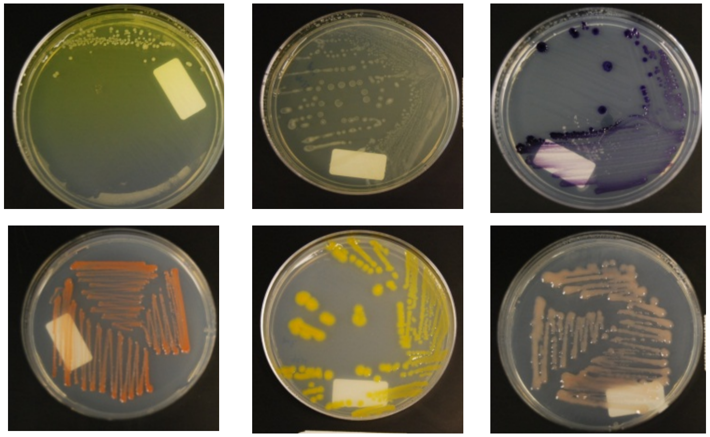
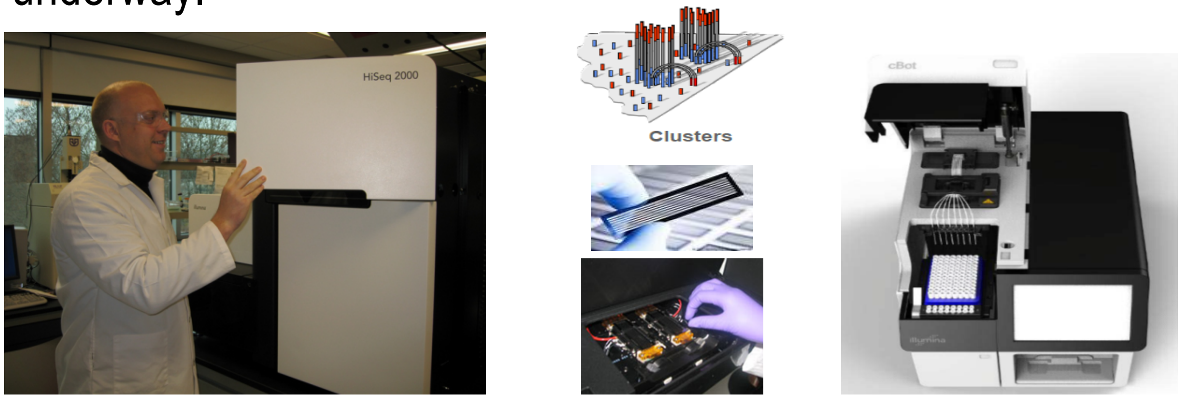


Figure 1. Bacterial isolates from the *Populus* Rhizosphere and Endosphere



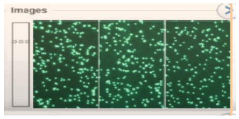


Figure 3. Cluster generation and sequencing.

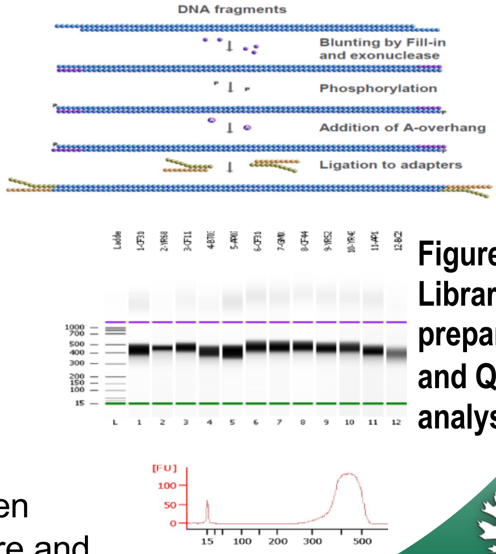


Figure 2. Library preparation and QC analysis

Brown et al., Twenty One *Pseudomonas* Genomes and Nineteen Genomes from Diverse Bacteria Isolated from the Rhizosphere and Endosphere of *Populus deltoides.* **Journal of Bacteriology, in press**