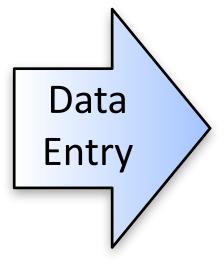


1. Swab (3 min)



Annotaate

3. GPS-tag/timestamp

Extract DNA (n=1,457 samples)



96-plex TruSeq/Qiagen Libraries

10.2 billion 125x125 DNA Seqs.

Quality Trim (Q20)

MegaBLAST-LCA alignment

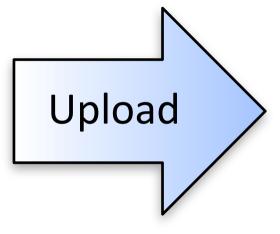


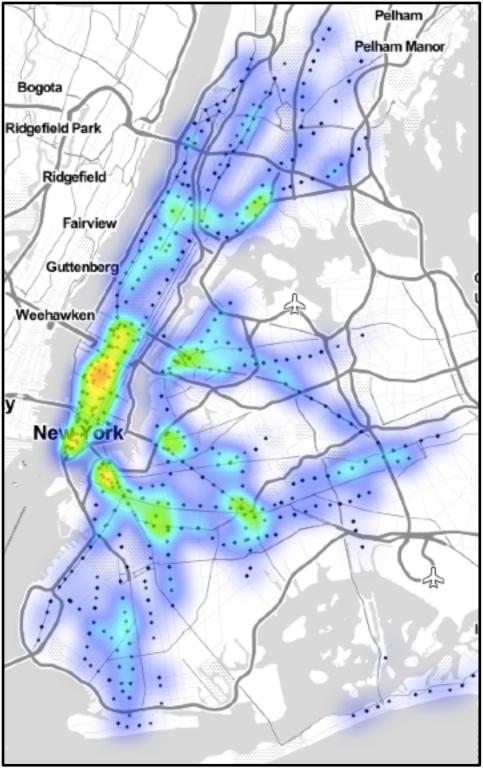






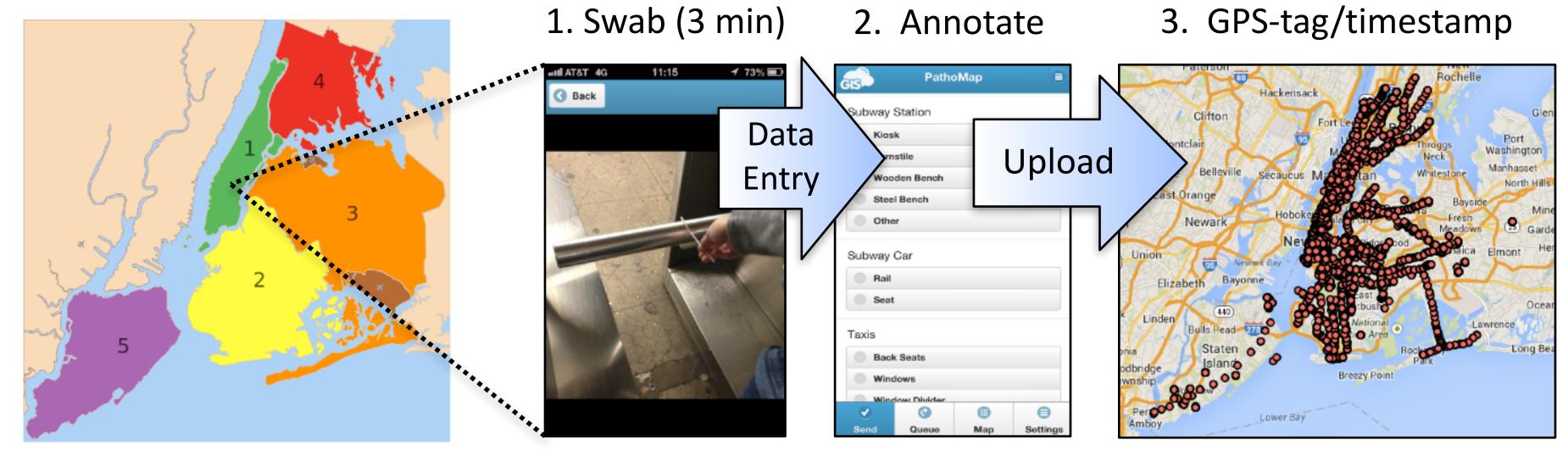
Confirm with MetaPhIAN

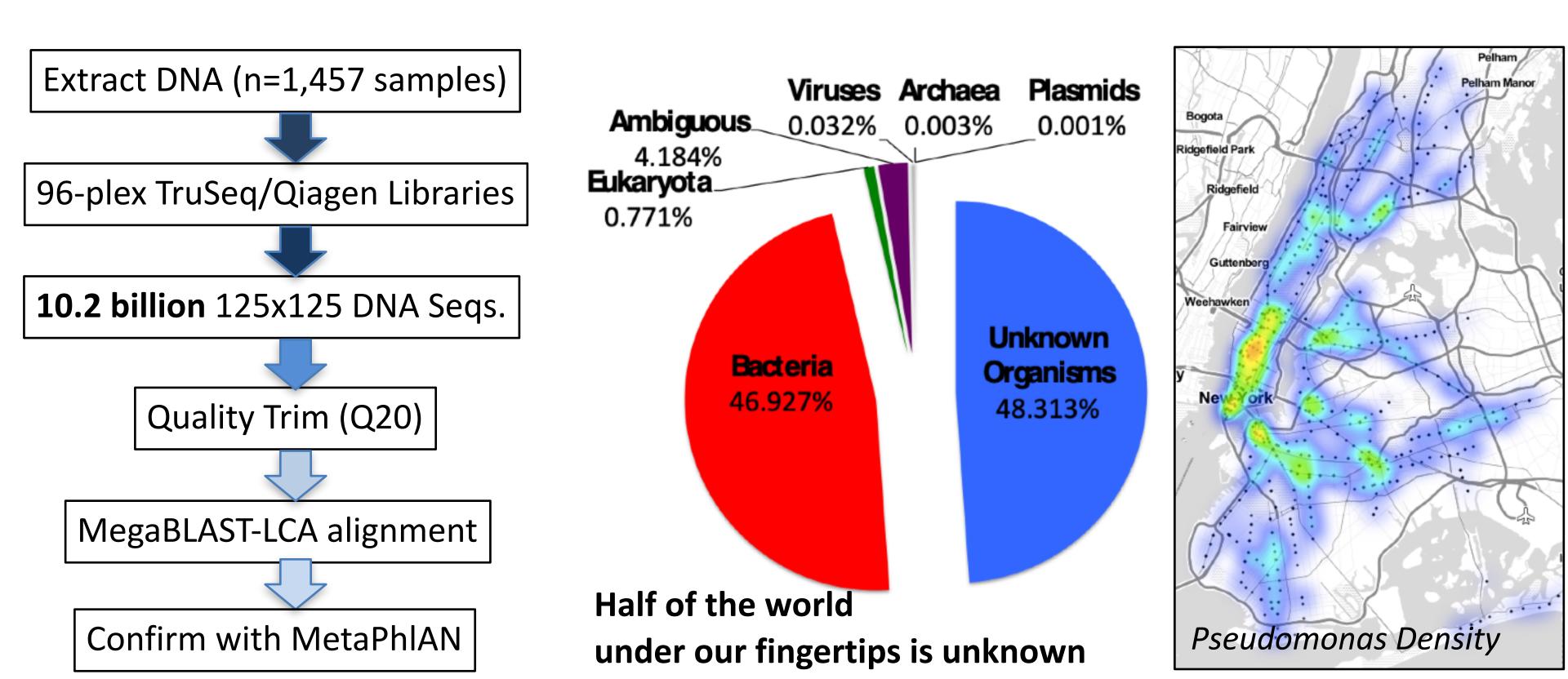


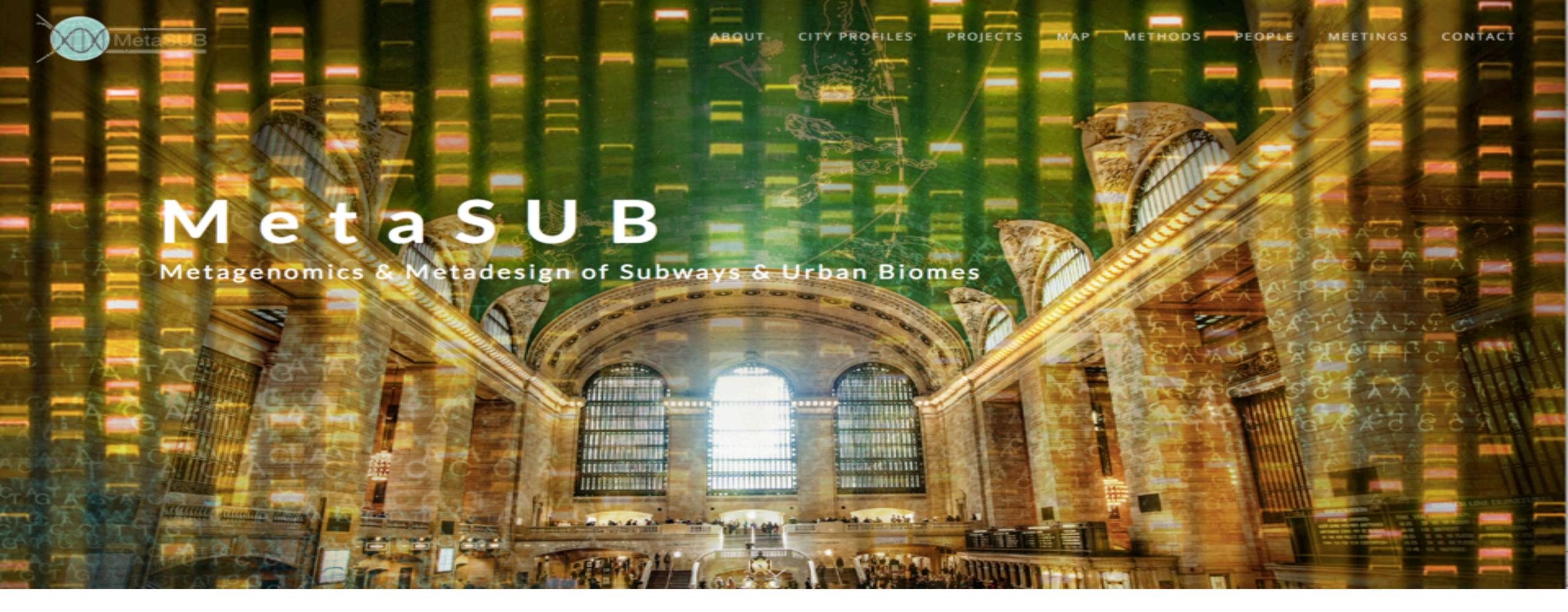


Pseudomonas Density

Half of the world under our fingertips is unknown







3 Goals:

- 1. Geospatial Metagenomic and Forensic Maps
- 2. Anti-microbial resistance (AMR) marker tracking by genetics and epigenetics
- 3. New Biosynthetic Gene Clusters (BGCs); new drugs

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