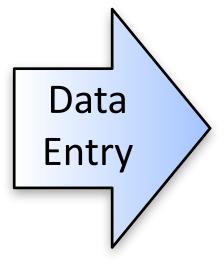


1. Swab (3 min)



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

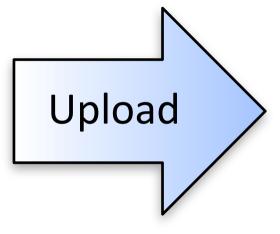


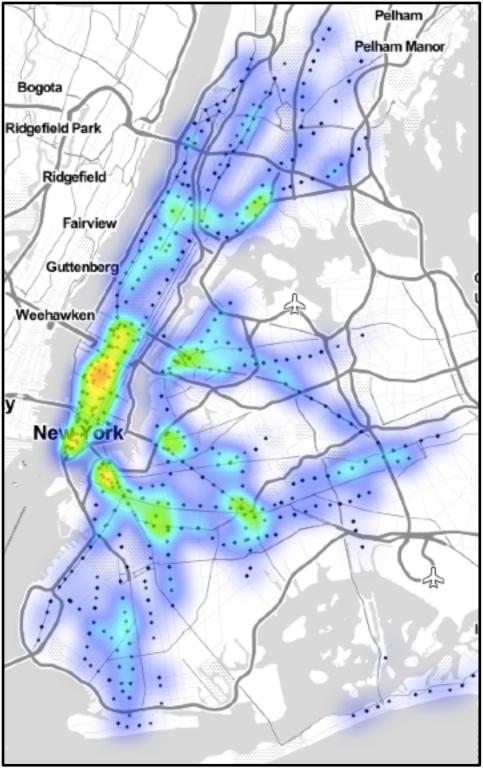






Confirm with MetaPhIAN





Pseudomonas Density

Half of the world under our fingertips is unknown

