

What: **MicrobeTrace** is a web-based tool to visually render networks from surveillance, epi investigations, and genetic analyses. Using simple point-and-click interfaces, data can be readily imported, integrated and visualized...no coding required!

Why: Networks are rich with sensitive data that are hard to visualize without specialized programming skills or tools that required sending data over the web. **MicrobeTrace** provides ready-access to visuals for fact sheets, publications and communiques.

How: **MicrobeTrace** is delivered by a modern* web browser, but runs locally on your computer or laptop. This means data and analyses are not sent over the web.

Where: **MicrobeTrace:** <https://microbetrace.herokuapp.com>
Documentation: <https://github.com/CDCgov/MicrobeTRACE/wiki>
Example Files: <https://github.com/CDCgov/MicrobeTrace/tree/master/demo>
Support: microbetrace@cdc.gov

Who: Developed by the Molecular Epidemiology and Bioinformatics Team in the Laboratory Branch of the Division of HIV/AIDS Prevention at CDC.

Other interactive and interoperable views

- 2D Network
- 3D Network
- Histogram
- Table
- Aggregation
- Bubbles
- Flow Diagram
- Scatterplot
- Waterfall
- Map
- Globe
- Gantt
- Timeline
- Heatmap
- Phylogenetic Tree
- Sequences

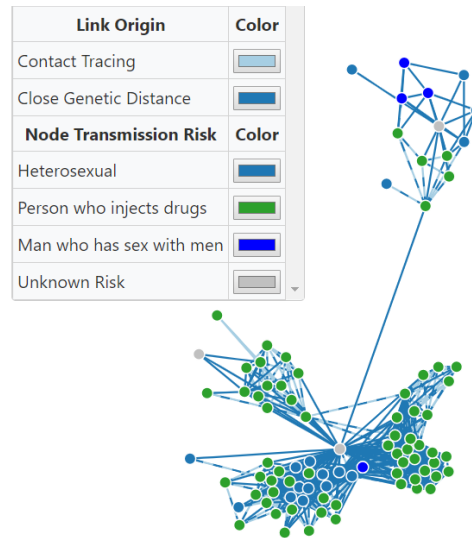
*Developed for Google Chrome, does not work in Internet Explorer

Input Files

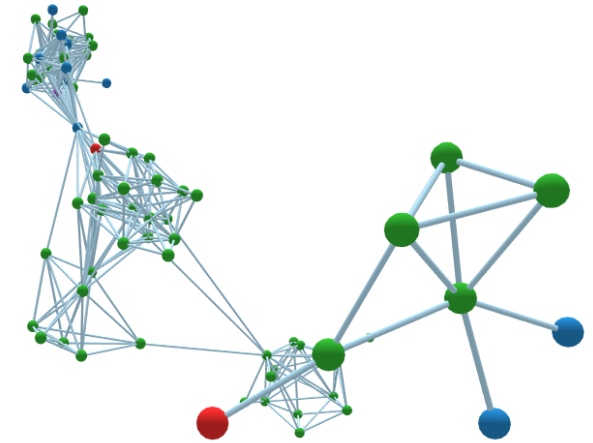
(any combination):

- Node Lists(s)
line-list of people or places
- Edge List(s)
connections between people or places
- Sequence file(s)
microbial genomic sequences
- Distance matrix
pairwise genetic distances
- Tree file
independent phylogenetic analysis

2D Network



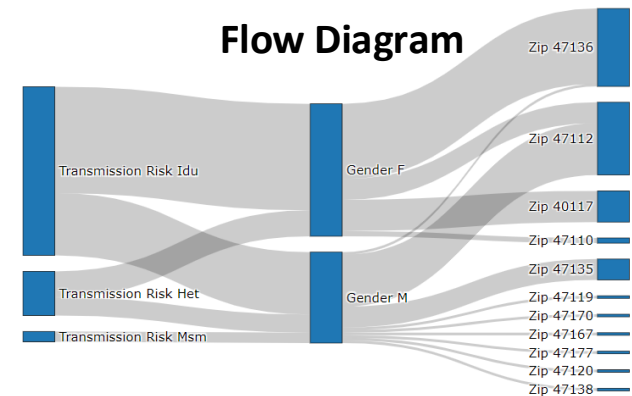
3D Network



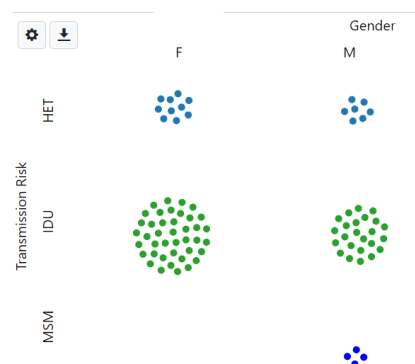
Global Network



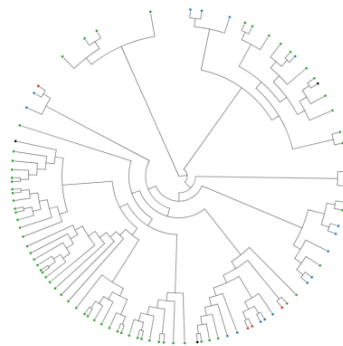
Flow Diagram



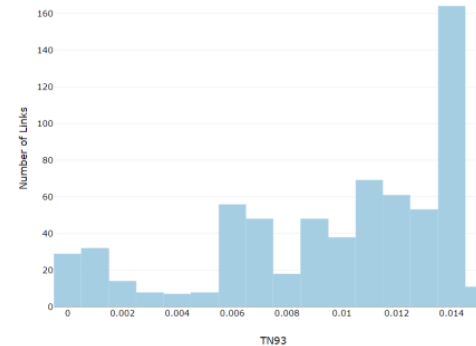
Bubbles Matrix



Phylogenetic Tree



Custom Histogram



Custom Aggregate Report

