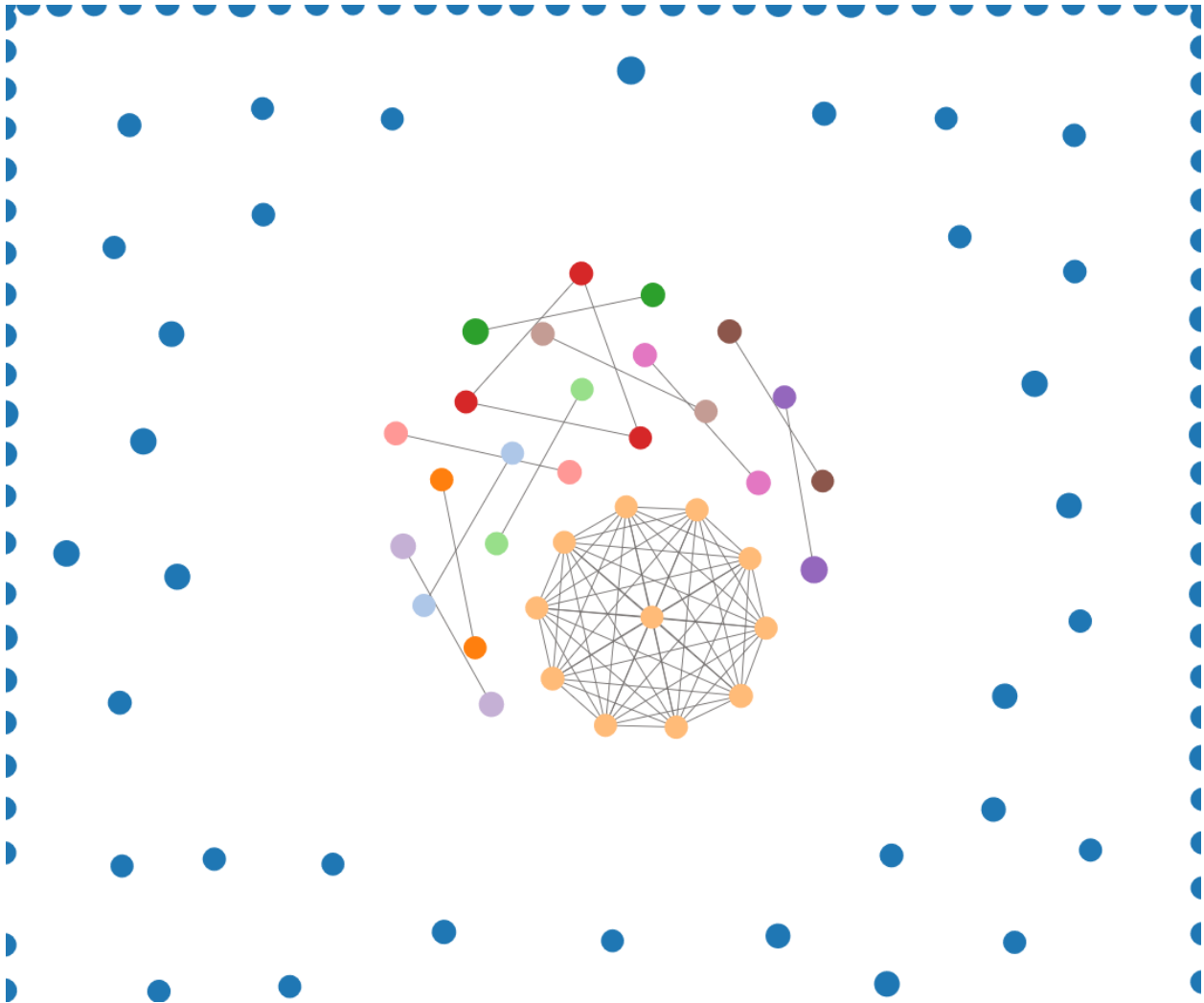


Michigan HCV GHOST Results (as of October 5, 2018)

NASTAD Technical Assistance Meeting October 10-12, 2018



Specimens Sequenced: 206

- Number of “Clusters” identified: 12 (one suspected outbreak)
- Number of Sequences with ≥ 1 match: 32 (16%)
- Number of Sequences without a match: 174 (84%)

Prioritization of Specimens:

- Part of a suspected outbreak
- Population-based surveillance:
 - Young – mean age 31.1 years
 - Rural/Suburban – 86 (41.7%)
 - White – 172 (83.5%)

Genotypes:

- 1a: 132 (64%)
- 1b: 6
- 2a: 2
- 2b: 16 (8%)
- 3a: 49 (24%)
- 6a: 1

Patient Risk Factors:

- IDU: 120 (58.3%)
- MSM: 12 (5.8%)
- Incarceration: 71 (34.5%)
- Unknown: 59 (28.6%)