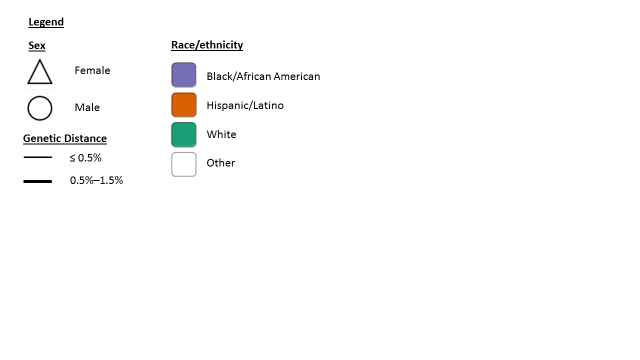
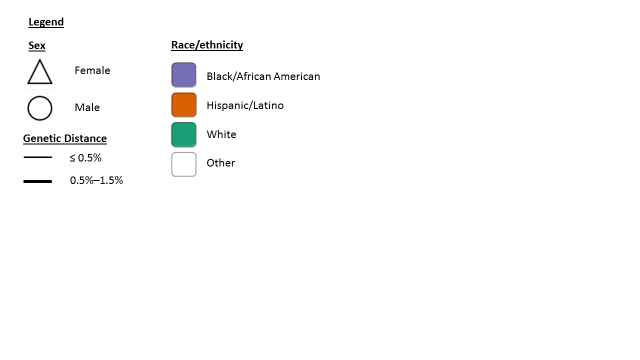
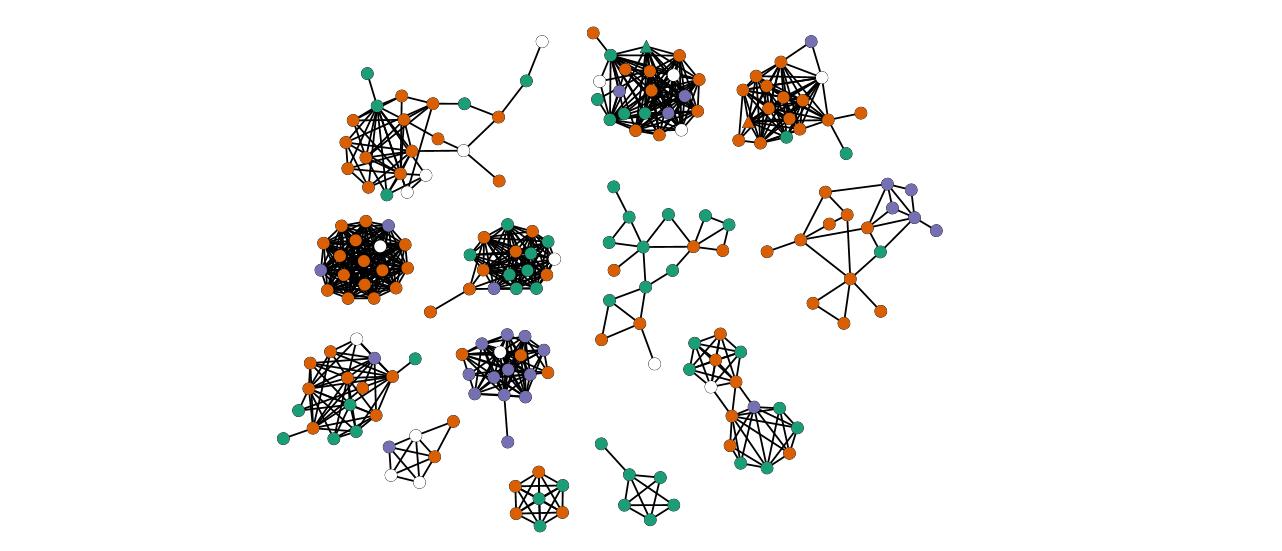
Supplemental Figure 1. Clusters inferred at 0.5% genetic distance threshold with 3 years of diagnoses and at least 5 diagnoses in the past 12 months.



Supplemental Table 1. Estimates of time to most recent common ancestor (tMRCA) for 13 clusters identified at 0.5% genetic distance threshold with 3 years of diagnoses and at least 5 diagnoses in the past 12 months, including median tMRCA and 95% highest posterior density (HPD).

|  |  |  |  |
| --- | --- | --- | --- |
| Cluster | Cluster Size | tMRCA median | tMRCA 95% HPD |
| 1 | 17 | 4.1 | 3.1–5.5 |
| 2 | 6 | 2.4 | 1.4–3.7 |
| 3 | 18 | 2.0 | 1.4–2.9 |
| 4 | 18 | 4.7 | 3.6–6.3 |
| 5 | 21 | 4.4 | 3.2–5.8 |
| 6 | 22 | 7.4 | 4.8–10.4 |
| 7 | 15 | 6.5 | 4.3–8.9 |
| 8 | 17 | 6.1 | 4.7–8.0 |
| 9 | 16 | 7.4 | 5.2–10.1 |
| 10 | 16 | 7.3 | 5.1–9.8 |
| 11 | 16 | 7.7 | 6.0–9.7 |
| 12 | 6 | 3.9 | 2.6–5.6 |
| 13 | 7 | 1.8 | 1.5–2.5 |