Table S3

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Network1 | Participants2 | Network measure | Phylogeny measure | Spearman correlation coefficient | p-value3 | Mean4 | SD5 |
| Baseline |  |  |  |  |  |  |  |
|  | Genotype 1a infection | geodesic distance | MCL distance | 0.132 | 0.080 | -0.001 | 0.045 |
|  | Genotype 3a infection | geodesic distance | MCL distance | -0.011 | 0.453 | -0.000 | 0.044 |
| Flattened |  |  |  |  |  |  |  |
|  | *Genotype 1a infection* | *geodesic distance* | *MCL distance* | *0.062* | *0.019* | *-0.000* | *0.063* |
|  | Genotype 3a infection | geodesic distance | MCL distance | -0.006 | 0.420 | 0.002 | 0.060 |