Visual Analysis of Probabilistic Infection Contagion in Hospitals Supplemental Material

M. Wunderlich, I. Block, T. von Landesberger, M. Petzold, M. Marschollek, and S. Scheithauer

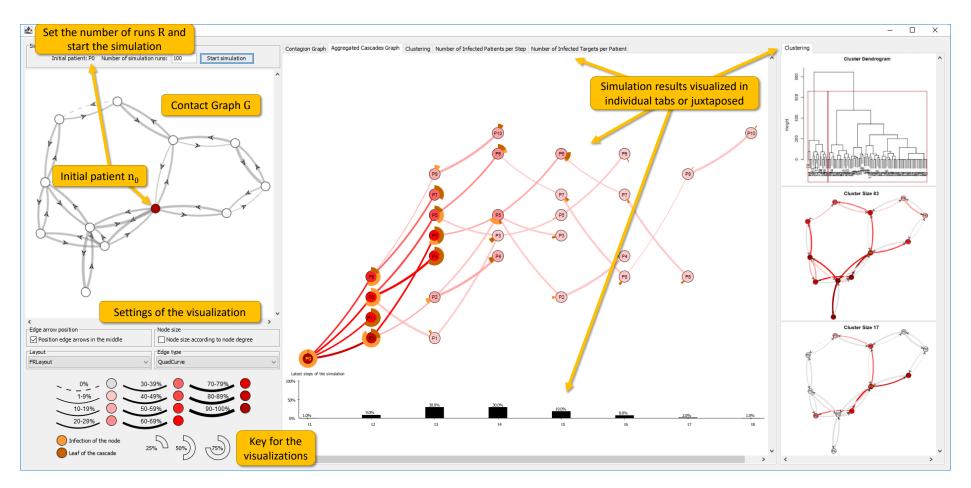


Figure 1: Main interface



Figure 2: Use Case: Load the contact graph and set the number of simulation runs R before starting the simulation

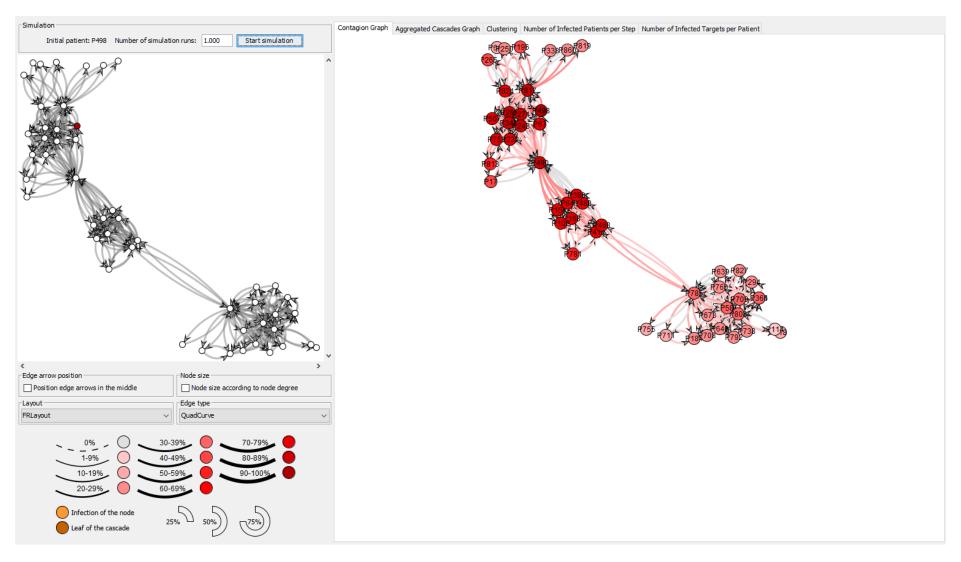


Figure 3: Use Case: The Contagion Graph View showing all simulation runs combined

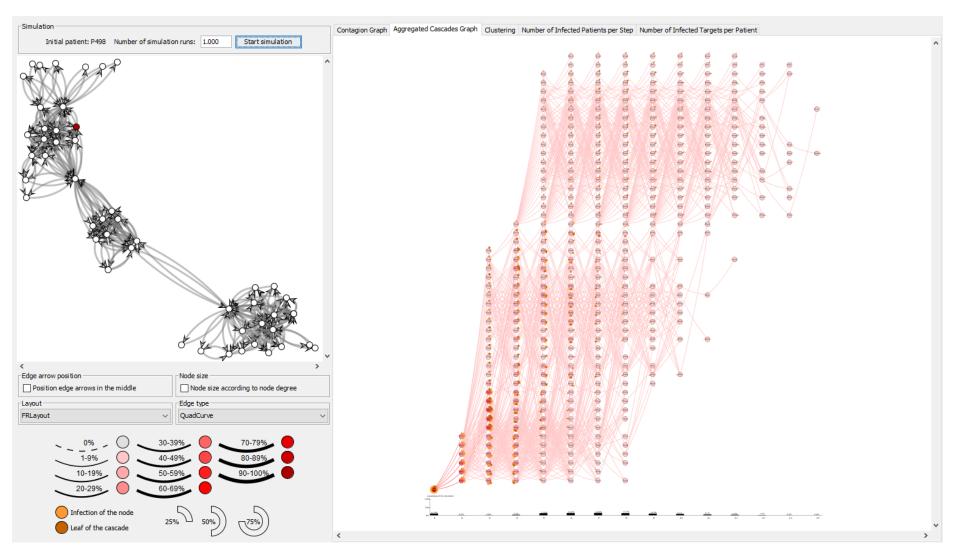


Figure 4: Use Case: The Aggregated Cascades Graph View showing all simulation runs combined (together with the supportive visualization of the Length of the Contagion)

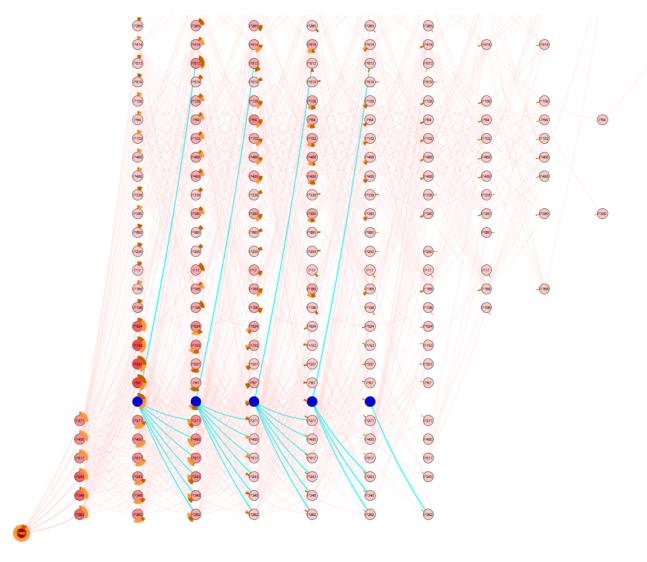


Figure 5: Use Case: The Aggregated Cascades Graph View showing all simulation runs combined (together with the supportive visualization of the Length of the Contagion) — Zoom in and highlighting of nodes with infected targets

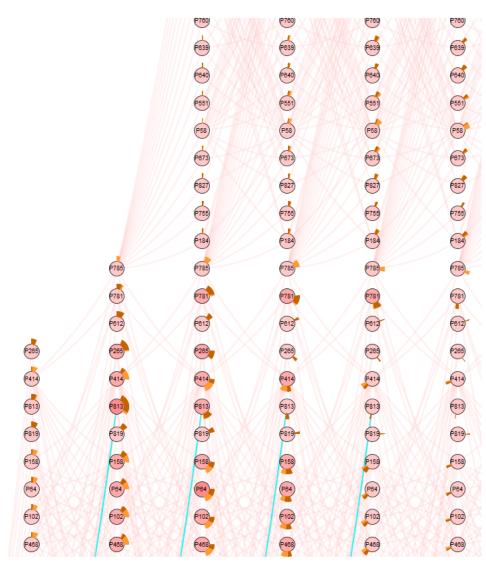


Figure 6: Use Case: The Aggregated Cascades Graph View showing all simulation runs combined (together with the supportive visualization of the Length of the Contagion) — Zoom in

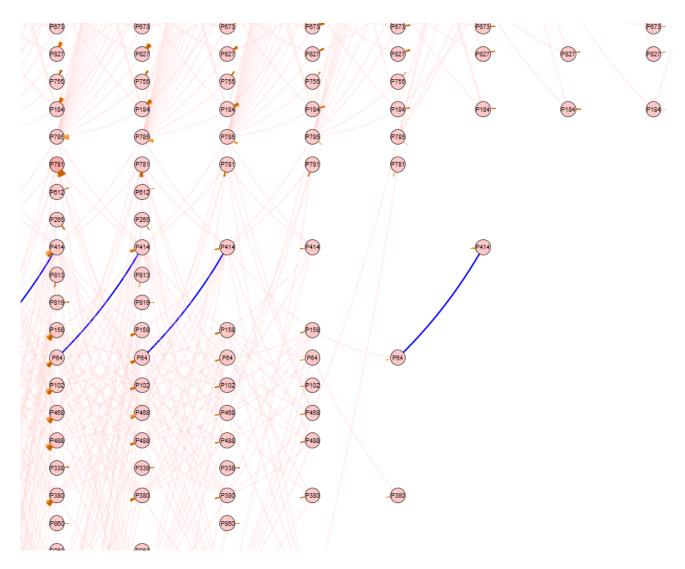


Figure 7: Use Case: The Aggregated Cascades Graph View showing all simulation runs combined (together with the supportive visualization of the Length of the Contagion) — Zoom in and highlighting of transmission pattern

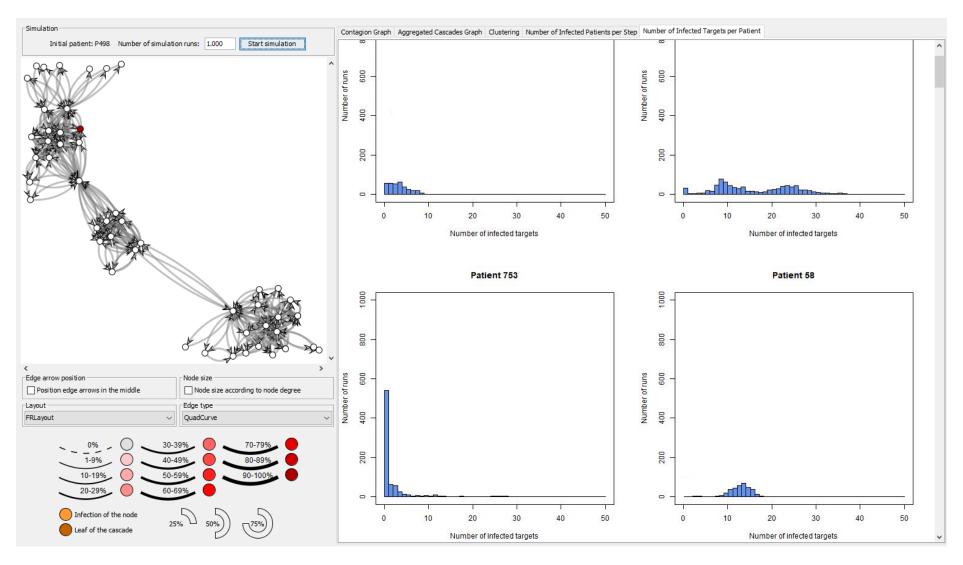


Figure 8: Use Case: The Number of Infected Targets per Patient

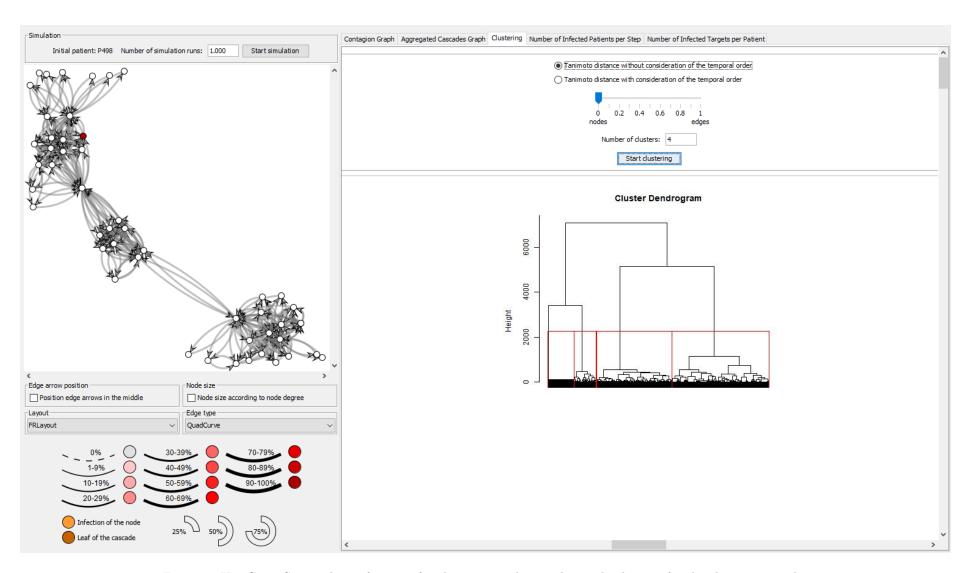


Figure 9: Use Case: Set similarity function for clustering and view cluster dendogram for the clustering results

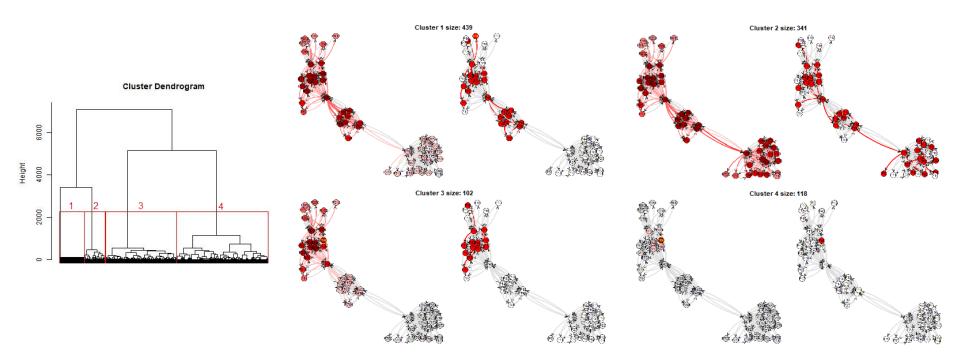


Figure 10: Use Case: Cluster dendogram, cluster summaries (left), and cluster representatives (right) visualized with the Contagion Graph View