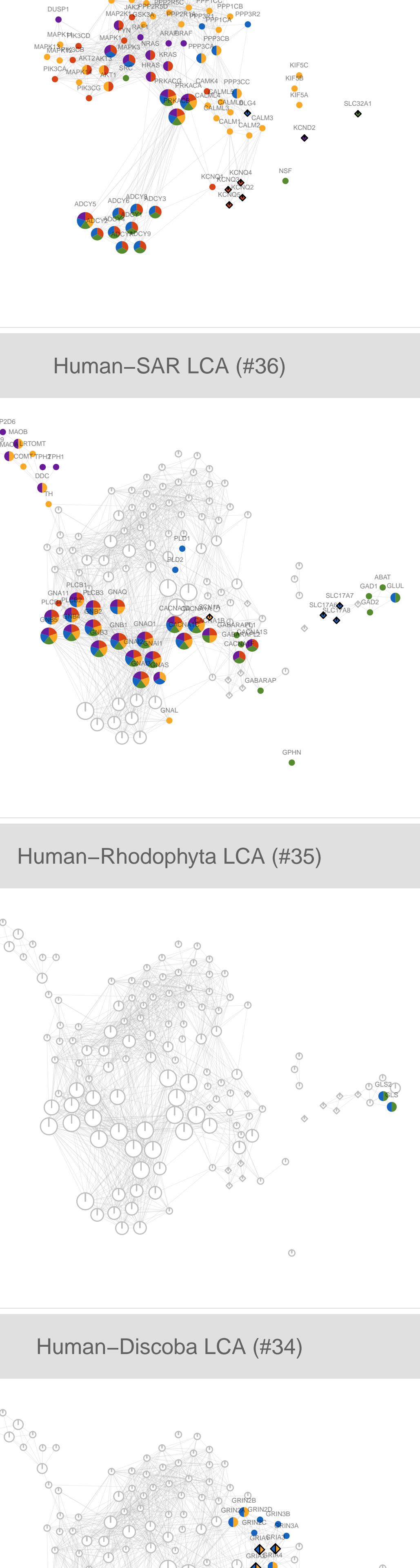
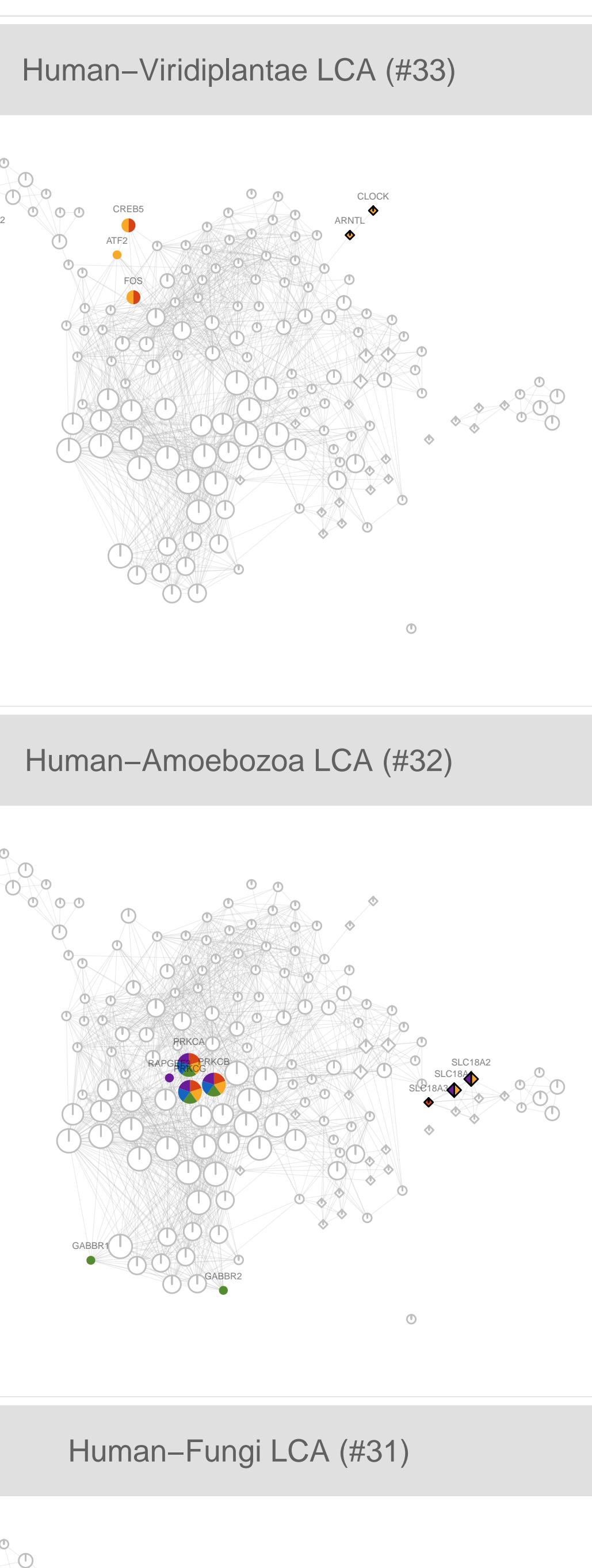


The figure displays a network graph with nodes representing proteins. Nodes are colored by species: Human (blue), Metamonada (orange), and Unassigned (grey). Node sizes vary, indicating interaction frequency or strength. The network shows numerous interactions between the Human and Metamonada protein sets.



A network graph visualization showing connections between nodes. The nodes are represented by circles and diamonds. A large cluster of nodes is located in the upper left, connected by many lines. Another smaller cluster of diamond-shaped nodes is in the upper right. A single node is located at the bottom center. A few isolated nodes are scattered in the lower right area.



A network graph illustrating interactions between genes. The central node is labeled *SLC6A3*. It is connected to numerous other nodes, which are represented by circles. Most of these nodes also contain the label *SLC6A3*, indicating a high density of self-interactions or strong internal connections for this gene. A few nodes are represented by diamond shapes, suggesting different types of entities or specific interaction partners.

