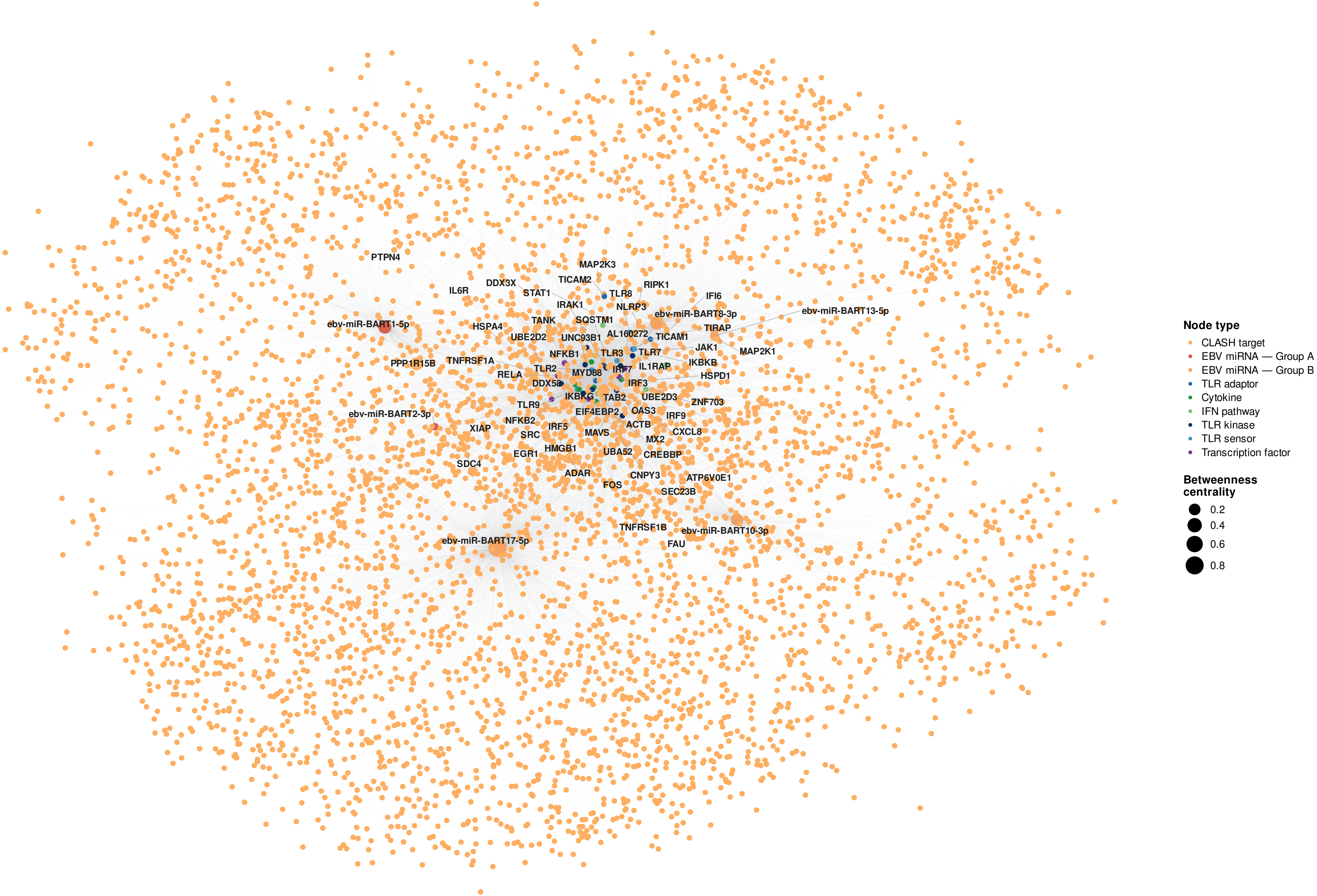


EBV exosomal miRNA–TLR signalling interaction network

5292 nodes · 16027 edges | Group A: FDR < 0.05 | Group B: logFC > 0.8



Node type

- CLASH target
- EBV miRNA — Group A
- EBV miRNA — Group B
- TLR adaptor
- Cytokine
- IFN pathway
- TLR kinase
- TLR sensor
- Transcription factor

Betweenness centrality

- 0.2
- 0.4
- 0.6
- 0.8