Supplementary Figures

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Figure 1: **Top:** Distribution of node *degree* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *degree* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 2: **Top:** Distribution of node *degree* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *degree* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 3: **Top:** Distribution of node *degree* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *degree* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 4: **Top:** Distribution of node *degree* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *degree* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 5: **Top:** Distribution of node *degree* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *degree* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 6: **Top:** Distribution of node *betweenness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *betweenness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 7: **Top:** Distribution of node *betweenness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *betweenness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 8: **Top:** Distribution of node *betweenness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *betweenness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 9: **Top:** Distribution of node *betweenness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *betweenness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 10: **Top:** Distribution of node *betweenness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *betweenness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 11: **Top:** Distribution of node *closeness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *closeness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 12: **Top:** Distribution of node *closeness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *closeness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 13: **Top:** Distribution of node *closeness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *closeness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 14: **Top:** Distribution of node *closeness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *closeness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 15: **Top:** Distribution of node *closeness* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *closeness* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 16: **Top:** Distribution of node *eigenvector* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *eigenvector* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 17: **Top:** Distribution of node *eigenvector* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *eigenvector* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 18: **Top:** Distribution of node *eigenvector* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *eigenvector* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 19: **Top:** Distribution of node *eigenvector* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *eigenvector* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.



Figure 20: **Top:** Distribution of node *eigenvector* centrality in the complete ConsensusPathDB (minimum edge confidence 95%) for nodes that are associated with at least one disease and **Bottom:** Distribution of node *eigenvector* centrality in the subgraph of the ConsensusPathDB that consists only of nodes that are associated with at least one disease.

References

[1] D. Ellinghaus, L. Jostins, S. L. Spain, A. Cortes, J. Bethune, B. Han, Y. R. Park, S. Raychaudhuri, J. G. Pouget, M. Hübenthal, T. Folseraas, Y. Wang, T. Esko, A. Metspalu, H.-J. Westra, L. Franke, T. H. Pers, R. K. Weersma, V. Collij, M. DAmato, J. Halfvarson, A. B. Jensen, W. Lieb, F. Degenhardt, A. J. Forstner, A. Hofmann, The International IBD Genetics Consortium (ibdgc), International Genetics of Ankylosing Spondylitis Consortium (igas), International PSC Study Group (ipscsg), Genetic Analysis of Psoriasis Consortium (gapc), Psoriasis Asso- ciation Genetics Extension (page), S. Schreiber, U. Mrowietz, B. D. Juran, K. N. Lazaridis, S. Brunak, A. M. Dale, R. C. Trembath, S. Weidinger, M. Weichen- thal, E. Ellinghaus, J. T. Elder, J. N. W. N. Barker, O. A. Andreassen, D. P. McGovern, T. H. Karlsen, J. C. Barrett, M. Parkes, M. A. Brown, and A. Franke, "Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci", *Nature Genetics*, vol. advance online publication, Mar. 2016.