11.1_STRINGweighted_CTS_cardiac_network.R

Inputs:

- BioTIP.res.RData (data)
- sce E.825 uncorrected.RData (data)
- 10090.protein.aliases.v12.0.txt.gz (data/PPIN)
- 10090.protein.info.v12.0.txt.gz (data/PPIN)
- 10090.protein.links.v12.0.txt.gz (data/PPIN)

Outputs:

- DEG_perState_min.prop0.25_lfc0.6_FDFR0.05.rds (data)
- markers.up all ttest.rds (data)
- markers.up ttest min.prop0.25.rds (data)
- GSE87038_STRING_graph_perState_notsimplified.rds (results)

11.1.1_check_vertex_duplication.R

Inputs:

- GSE87038_STRING_graph_perState_notsimplified.rds (results)
- DEG_perState_min.prop0.25_lfc0.6_FDFR0.05.rds (data)
- 10090.protein.aliases.v12.0.txt.gz (data/PPIN)
- 10090.protein.info.v12.0.txt.gz (data/PPIN)
- 10090.protein.links.v12.0.txt.gz (data/PPIN)

Outputs:

correct n edges HiG STRING2.14.0.rds (results)

11.2.1 CTS cardiac network degreeDistribution.R

Inputs:

- GSE87038 STRING graph perState notsimplified.rds (results)
- correct_n_edges_HiG_STRING2.14.0.rds (results)

- degree_distribution_w_vsize.pdf (results)
- normalized degree distribution.pdf (results)
- degree_GSE87038.pdf (results)
- boxplot_normalized_degree_GSE87038.pdf (results)
- V deg dis.rds (results)

11.2.0_update_network_weights_clean_max.R

Inputs:

- sce E.825 uncorrected.RData (data)
- GSE87038 STRING graph perState notsimplified.rds (results)
- correct n edges HiG STRING2.14.0.rds (results)

Dependencies:

- celltype_specific_weight_v9.R (code)
- BioTIP update 06162025.R (code)

Outputs:

- network_specificity_list.rds (results/PPI_<max>weight)
- GSE87038_STRING_graph_perState_simplified_combinedweighted.rds (results/PPI <max>weight)
- GSE87038_STRING_graph_perState_simplified_ratioweighted.rds (results/PPI <max>weight)
- GSE87038_STRING_graph_perState_simplified_zscoreweighted.rds (results/PPI_<max>weight)
- GSE87038_STRING_graph_perState_simplified_diffweighted.rds (results/PPI <max>weight)
- compare specificity method vs PPIscores.pdf (results/PPI <max>weight)
- compare_specificity_method_HiGCTS_8_vs_PPIscores.pdf (results/PPI_<max>weight)

11.2.1 CTS cardiac network strengthDistribution.R

Inputs:

- sce E.825 uncorrected.RData (data)
- GSE87038 STRING graph perState notsimplified.rds (results)
- correct_n_edges_HiG_STRING2.14.0.rds (results)

Dependencies:

celltype_specific_weight_v9.R (code)

- strength distribution w vsize.pdf (results)
- normalized_strength_distribution.pdf (results)
- strength GSE87038.pdf (results)
- boxplot normalized strength GSE87038.pdf (results)

11.2.1_weighted_network_robustness.simulation.R (requires High Performance Computing)

Inputs:

 GSE87038_STRING_graph_perState_simplified_combinedweighted.rds (results/PPI <max>weight)

Dependencies:

celltype_specific_weight_v9.R (code)

Outputs:

- failure.edge 100 simplified combinedweighted.rds (results/PPI <max>weight)
- failure.vertex 100 simplified combinedweighted.rds (results/PPI <max>weight)
- AUC_compt.pct_attac_V_random_1000runs_combinedweighted.rds (results/PPI_<max>weight)
- AUC_compt.pct_attac_E_random_1000runs_combinedweighted.rds (results/PPI_<max>weight)

11.2_CTS_cardiac_network_robustness.R (requires High Performance Computing)

Inputs:

- GSE87038_STRING_graph_perState_simplified_combinedweighted.rds (results/PPI_<max>weight)
- failure.vertex 100 simplified combinedweighted.rds (results/PPI <max>weight)
- failure.edge_100_simplified_combinedweighted.rds (results/PPI_<max>weight)
- AUC_compt.pct_attac_V_random_1000runs_combinedweighted.rds (results/PPI_<max>weight)
- AUC_compt.pct_attac_E_random_1000runs_combinedweighted.rds (results/PPI_<max>weight)

Dependencies:

celltype_specific_weight_v9.R (code)

- vertex_attack_GSE87038.pdf (results/PPI_<max>weight)
- attack.vertex.btwn.rds (results/PPI_<max>weight)
- attack.edge.btwn.rds (results/PPI_<max>weight)
- attack GSE87038.pdf (results/PPI <max>weight)
- box wilcox-test attack GSE87038.pdf (results/PPI <max>weight)
- line.adjp wilcox attack GSE87038.pdf (results/PPI <max>weight)
- box wilcox-test attack AUC GSE87038.pdf (results/PPI <max>weight)
- Vertex_All_AUC_density_plots.pdf (results/PPI_<max>weight)

Edge_All_AUC_density_plots.pdf (results/PPI_<max>weight)

11.3_CTS_cardiac_network_ANND_pagerank.R

Inputs:

- GSE87038_STRING_graph_perState_simplified_combinedweighted.rds (results/PPI <max>weight)
- CHD Cilia Genelist.rds (data)

Outputs:

- GSE87038_PageRank_Pvalue_by_rewiring.rds (results/PPI_<max>weight)
- GSE87038_annd_Pvalue_by_rewiring.rds (results/PPI_<max>weight)
- df PAGERANK strength ANND.rewring.P.rds (results/PPI <max>weight)
- df PAGERANK strength ANND.rewring.P.tsv (results/PPI <max>weight)
- df_betweeness.tsv (results/PPI_<max>weight)
- table top5 Betweenness perPPI.tsv (results/PPI <max>weight)
- BetweennessCentrality_GSE87038_v2.pdf (results/PPI_<max>weight)
- table_top5_PageRank_perPPI.tsv (results/PPI_<max>weight)
- PageRank_GSE87038_v2.pdf (results/PPI_<max>weight)
- annd_GSE87038_v2.pdf (results/PPI_<max>weight)
- normalized.node.strength_GSE87038_v2.pdf (results/PPI_<max>weight)
- gene ranked by importance dotBoxPlot 6panel.pdf (results/PPI <max>weight)

11.5 visualization.R

Inputs:

- GSE87038_STRING_graph_perState_simplified_combinedweighted.rds (results/PPI_<max>weight)
- GSE87038_STRING_graph_perState_notsimplified.rds (results)
- CHD_Cilia_Genelist.rds (data)

- network view PPI GSE87038.pdf (results/PPI <max>weight)
- network_HiG_view_weight_shrink_GSE87038.pdf (results/PPI_<max>weight)