Declaration

I, Manaswita Saikia, hereby declare that the thesis entitled "**Biomarker Identification for Critical Diseases using Machine Learning Techniques**" submitted to the Department of Computer Science and Engineering under the School of Engineering, Tezpur University, in partial fulfillment for the award of the degree of Doctor of Philosophy in Computer Science and Engineering is a bona-fide work carried out by me. The results presented in this thesis have not been submitted in part or in full, to any other University or Institute for the award of any degree or diploma.

Date:

Place: Tezpur University, Napaam, Tezpur

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Certificate

This is to certify that the thesis entitled "**Biomarker Identification for Critical Diseases using Machine Learning Techniques**" submitted to the Department of Computer Science and Engineering under the School of Engineering, Tezpur University, in partial fulfillment for the award of the degree of Doctor of Philosophy in Computer Science and Engineering is a record of research work carried out by Ms. Manaswita Saikia under our supervision and guidance.

All helps received by her from various sources have been duly acknowledged. No part of this thesis has been submitted elsewhere for award of any other degree.

Signature of Supervisor

Signature of Co-supervisor

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LIST OF TABLES

| 2.1 | Summary of the microarray datasets, GSE20347 and GSE23400, and the | |
|------|---|-----|
| | bulk RNA-Seq dataset, GSE130078 for ESCC | 35 |
| 2.2 | Summary of the scRNA-Seq dataset, GSE160269 for ESCC | 37 |
| 3.1 | Eight chosen biclustering methods: A Comparison | 57 |
| 3.2 | Summary of the biclusters detected by BicGenesis in all three datasets | 70 |
| 3.3 | Subset of Normal and Disease Biclusters | 74 |
| 3.4 | Preservation analysis of modules in the microarray datasets, GSE20347 | |
| | and GS23400, and the bulk RNA-Seq dataset, GSE130078 | 77 |
| 3.5 | Top 20 hub-genes for each extracted MoI in the two microarray and one | |
| | bulk RNA-Seq datasets | 79 |
| 3.6 | Percentages of genes in each MoT that are annotated to the Gene Ontol- | |
| | ogy (GO) databases and KEGG pathways | 82 |
| 3.7 | Summary of BCGs detected by BicGenesis in the microarray dataset, | |
| | GS20347, that are annotated to top 3 GO terms in the three GO databases. | 86 |
| 3.8 | Summary of BCGs detected by BicGenesis in the microarray dataset, | |
| | GSE23400, that are annotated to top 3 GO terms in the three GO databases. | 88 |
| 3.9 | Summary of BCGs detected by BicGenesis in the bulk RNA-Seq dataset, | |
| | GS130078, that are annotated to top 3 GO terms in the three GO databases | 89 |
| 3.10 | Summary of BCGs detected by BicGenesis in the two microarray and | |
| | one bulk RNA-Seq datasets that have been annotated to the top 5 KEGG | |
| | enriched pathways | 90 |
| 3.11 | Summary of potential biomarkers identified by BicGenesis | 96 |
| 3.12 | Summary of potential ESCC biomarkers identified by BicGenesis using | |
| | the biomarker criteria | 100 |
| 4.1 | DE methods for Microarray and bulk RNA-Seq data | 112 |
| 4.2 | Summary of detected DEGs by the three RNA-Seq methods and the | |
| | three microarray methods for three datasets 1 | 120 |
| 4.3 | Preservation analysis of modules detected by our Integrative DEA method | |
| | in the two microarray and one RNA-Seq datasets 1 | 126 |
| 4.4 | Top 20 hub-genes for each extracted MoI in the two microarray and one | |
| | RNA-Seq datasets | 127 |

| 4.5 | Percentages of genes in each MoI that are annotated to the Gene Ontol- |
|--|---|
| | ogy (GO) databases and KEGG pathways |
| 4.6 | DEGs that are annotated to most enriched GO term in all three GO |
| | databases (BP, CC and MF) as well as the most enriched pathway 130 |
| 4.7 | Summary of BCGs detected by Integrative DEA in the microarray dataset, |
| | GS20347, that are annotated to top 3 GO terms in the three GO databases. 133 |
| 4.8 | Summary of BCGs detected by Integrative DEA in the microarray dataset, |
| | GSE23400, that are annotated to top 3 GO terms in the three GO databases.134 |
| 4.9 | Summary of BCGs detected by Integrative DEA in the bulk RNA-Seq |
| | dataset, GS130078, that are annotated to top 3 GO terms in the three GO |
| | databases |
| 4.10 | Summary of BCGs detected by our method, Integrative DEA, in the |
| | three datasets |
| 4.11 | Summary of potential biomarkers identified by our framework, Integra- |
| | tive DEA |
| 4.12 | Summary of potential ESCC biomarkers identified by Integrative DEA |
| | using the biomarker criteria |
| 4.13 | Comparison of our method, Integrative DEA with two recent works that |
| | employ DEA on ESCC datasets |
| | |
| 5.1 | Centrality Measures for hub-gene finding employed in CBDCEM 157 |
| 5.1 5.2 | Centrality Measures for hub-gene finding employed in CBDCEM 157 Comparison of the seven centrality measures employed by CBDCEM . 159 |
| | |
| 5.2 | Comparison of the seven centrality measures employed by CBDCEM . 159 |
| 5.2 5.3 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.25.35.45.5 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.25.35.45.5 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.25.35.45.5 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.25.35.45.55.6 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.25.35.45.55.6 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 5.5 5.6 5.7 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 5.5 5.6 5.7 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 5.5 5.6 5.7 5.8 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 5.5 5.6 5.7 5.8 5.9 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 5.5 5.6 5.7 5.8 5.9 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |
| 5.2 5.3 5.4 5.5 5.6 5.7 5.8 5.9 5.10 | Comparison of the seven centrality measures employed by CBDCEM . 159 Symbols used in proposed Hub-gene finding algorithm |

| 5.12 | Summary of potential ESCC biomarkers identified by CBDCEM using | |
|------|---|-----|
| | the biomarker criteria | 190 |
| 5.13 | Summary of potential biomarkers detected by CBDCEM and four other | |
| | hub-gene finding methods | 192 |
| 5.14 | Summary of performance of CBDCEM vs. four other methods in terms | |
| | of proportion of modules | 198 |
| 6.1 | Centrality Measures for hub-gene finding employed in scDiffCoAM | 205 |
| 6.2 | Comparison of the four of the seven measures employed by scDiffCoAM. | 207 |
| 6.3 | Preservation Analysis (Z _{summary}) of CD45+ modules in CD45- dataset | |
| | and vice versa | 222 |
| 6.4 | Top 20 hub genes for each extracted MoIs in CD45+ and CD45- datasets | |
| | using our hub-gene finding algorithm | 227 |
| 6.5 | Percentage of genes in each MoI that are annotated in the GO databases | |
| | and KEGG pathways | 230 |
| 6.6 | Summary of hub-genes detected by scDiffCoAM that have been anno- | |
| | tated to the Top 20 KEGG enriched pathways in the CD45+ cell types . | 231 |
| 6.7 | Summary of hub-genes detected by scDiffCoAM that have been anno- | |
| | tated to the Top 20 KEGG enriched pathways in CD45- cell types | 232 |
| 6.8 | Summary of hub-genes detected by scDiffCoAM that have been anno- | |
| | tated to the top enriched GO terms in the three GO databases for CD45+ | |
| | cell types | 233 |
| 6.9 | Summary of hub-genes detected by scDiffCoAM that have been anno- | |
| | tated to the top enriched GO terms in the three GO databases for CD45- | |
| | cell types | 235 |
| 6.10 | Summary of potential biomarkers candidates identified by scDiffCoAM. | |
| | Here, All 3 under GO databases imply all three databases, BP, CC, and | |
| | MF | 247 |
| 6.11 | Summary of potential ESCC biomarkers identified by scDiffCoAM us- | |
| | ing the biomarker criteria | 251 |
| 6.12 | Summary of potential biomarkers detected by scDiffCoAM and three | |
| | other hub-gene finding methods | 254 |
| 6.13 | Ranking all potential biomarkers for ESCC identified by all four pro- | |
| | posed frameworks | 261 |

LIST OF FIGURES

| 1.1 | Steps involved in the Central Dogma of molecular biology | 2 |
|------|---|-----|
| 2.1 | Pre-processing pipeline employed by our proposed frameworks for the three types of gene expression data | 37 |
| 2 1 | | 42 |
| 3.1 | Biclustering Approaches: A Taxonomy | 43 |
| 3.2 | Proposed Biclustering Analysis Framework | 61 |
| 3.3 | Pipeline for DCA | 64 |
| 3.4 | Hierarchical trees for normal and disease in the microarray datasets GSE203 | |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | 71 |
| 3.5 | Soft thresholds for normal and disease in the microarray datasets GSE20347 | |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | 72 |
| 3.6 | Dendrograms for normal and disease in the microarray datasets GSE20347 | |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | 75 |
| 3.7 | Zsummary plots for normal and disease in the microarray datasets GSE2034 | 7 |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | 76 |
| 3.8 | GRN for normal modules a) <i>skyblue</i> and b) <i>white</i> in GSE20347, disease | |
| | modules c) yellowgreen, d) white e) salmon4, and f) purple in GSE20347 | 84 |
| 3.9 | GRN for normal modules a) brown2 in GSE23400 and b)lightcyan in | |
| | GSE130078 | 85 |
| 3.10 | GRN for disease module <i>orange</i> in GSE130078 | 85 |
| 4.1 | Proposed Integrative Differential Expression Analysis Framework | 115 |
| 4.2 | Pipeline for DCA | 17 |
| 4.3 | Dendrograms for normal and disease in the microarray datasets, GSE20347 | |
| | and GSE23400, and bulk RNA-Seq dataset, GSE130078 for ESCC \square | 123 |
| 4.4 | Zsummary plots for normal and disease in the microarray datasets GSE2034 | 7 |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | 125 |
| 4.5 | GRN for normal module a) <i>pink</i> and disease modules b) <i>greenyellow</i> in | |
| | GSE20347, disease modules c) darkgreen, d) lightsteelbluel e) black in | |
| | GSE20347. GRN for disease module f) magenta in GSE23400 | 131 |
| 4.6 | GRN for normal modules a) <i>purple</i> and b) <i>greenyellow</i> in GSE20347, | |
| | and disease modules c) <i>blue</i> in GSE23400. GRN for disease modules d) | |
| | lightyellow e) violet, and f) steelblue in GSE130078 | 132 |

| 5.1 | Proposed Centrality Based DCA Framework, CBDCEM | 161 |
|------|---|-----|
| 5.3 | Dendrograms for normal and disease in the microarray datasets GSE20347 | |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | |
| 5.4 | Heiarchical trees for module detection for normal and disease in the mi- | |
| | croarray datasets GSE20347 and GSE23400, and bulk RNA-Seq dataset | |
| | GSE130078 for ESCC | 172 |
| 5.5 | Zsummary plots for normal and disease in the microarray datasets GSE2034 | 47 |
| | and GSE23400, and bulk RNA-Seq dataset GSE130078 for ESCC | 173 |
| 5.6 | GRN for normal module a) <i>paleturquoise</i> , disease modules b) <i>dark</i> - | |
| | turquoise and c) orange in GSE20347. GRN for disease module d) | |
| | <i>grey60</i> in GSE23400 | 179 |
| 5.7 | GRN for disease module a) <i>lightcyan</i> , d) <i>tan</i> e) <i>green</i> in GSE23400, and | |
| | disease module f) <i>salmon</i> in GSE130078 | 180 |
| | | |
| 6.1 | Steps involved in WGCNA analysis for high dimensional data using hd- | |
| | WGCNA | |
| 6.2 | Proposed framework for DCA on scRNA-Seq Dataset, scDiffCoAM | |
| 6.3 | Violin Plots for CD45+ and CD45 | |
| 6.4 | Elbow Plots for CD45+ and CD45 | |
| 6.5 | Soft Thresholds for three CD45+ and three CD45- cell types | 219 |
| 6.6 | Dendrograms for three CD45+ cell types and one CD45- cell type | 220 |
| 6.7 | Dendrograms for two CD45- cell types | 221 |
| 6.8 | $Z_{summary}$ plot for Tcell (CD45+) in Epithelial (CD45-) and Endothelial | |
| | (CD45-) and vice versa | 223 |
| 6.9 | Z _{summary} plot for Tcell (CD45+) in Fibroblast (CD45-) and Bcell (CD45+) | |
| | in Epithelial (CD45-) and Endothelial (CD45-), and vice versa | 224 |
| 6.10 | Z _{summary} plot for Bcell (CD45+) in Fibroblast (CD45-) and Myeloid | |
| | (CD45+) in Epithelial (CD45-) and Endothelial (CD45-), and vice versa. | 225 |
| 6.11 | $Z_{summary}$ plot for Myeloid (CD45+) in Fibroblast (CD45-) and vice versa. | 226 |
| 6.12 | GRN for two modules in CD45+ cell type, Tcell | 237 |
| 6.13 | GRN for one, two, and two modules in CD45+ cell types, Tcell, Bcell, | |
| | and Myeloid and one module CD45- cell types, Epithelial | 238 |
| 6.14 | GRN for three and one modules in CD45- cell types, Endothelial and | |
| | Fibroblast | 239 |
| 6.15 | GRN for module <i>turquoise</i> in CD45- cell type, Fibroblast | 239 |
| 6.16 | Summary of performances of scDiffCoAM vs. four other methods | 258 |

Glossary

| BC | Bcell |
|-------|---|
| BCG | Biomarker Candidate Gene |
| CEN | Co-expression Network |
| СРМ | Counts Per Million |
| DAVID | Database for Annotation, Visualisation, and Integrated Dis- |
| | covery |
| DCA | Differential Co-expression analysis |
| DCE | Differential Co-Expression |
| DCG | Differentially Co-expressed Gene |
| DEA | Differential Expression Analysis |
| DEG | Differentially Expressed Gene |
| DNA | Deoxyribonucleic Acid |
| EBAM | Empirical Bayes analysis of Microarrays |
| EN | Endothelial |
| EP | Epithelial |
| ESCC | Esophageal Squamous Cell Carcinoma |
| FDR | False Discovery Rate |
| FI | Fibroblast |
| GO | Gene Ontology |
| GO_BP | Gene Ontology Biological Processes |
| GO_CC | Gene Ontology Cellular Componenets |
| GO_MF | Gene Ontology Molecular Funxtions |
| GRN | Gene Regulatory Network |
| GSEA | Gene Set Enrichment Analysis |
| HNSCC | Head and Neck Squamous Cell Carcinoma |
| KEGG | Kyoto Encyclopedia of Genes and Genomes |
| LaSCC | Laryngeal Squamous Cell Carcinoma |
| IFDR | Local False Discovery Rate |
| lgEGo | List of Enriched GO Terms |
| lgEP | List of Enriched Pathways |
| LSCC | Lung Squamous Cell Carcinoma |

| MoI | Module of Interest |
|-----------|--|
| MY | Myeloid |
| MSR | Mean Squared Residue |
| OSCC | Oral Squamous Cell Carcinoma |
| PCA | Princpal Component Analysis |
| PCC | Pearson Correlation Coefficient |
| PEGoT | Percentage of Enriched GO Terms |
| PEP | Percentage of Enriched Pathways |
| RNA | Ribonucleic Acid |
| RNA-Seq | RNA Sequencing |
| SAM | Significance Analysis of Microarrays |
| SCC | Squamous Cell Carcinoma |
| scRNA-Seq | Single cell RNA-Sequencing |
| TC | Tcell |
| TED | Top Enriched DEG |
| TF | Transcription Factor |
| TG | Target Gene |
| TOM | Topological Overlap Matrix |
| TSCC | Tongue Squamous Cell Carcinoma |
| WGCNA | Weighted Gene Co-expression Network Analysis |