

Supplementary File for “Constrained Multi-objective Optimization-Based Temporal Network Observability for Critical Biomarker Identification of Individual Patients”

S-I. Supplementary contents

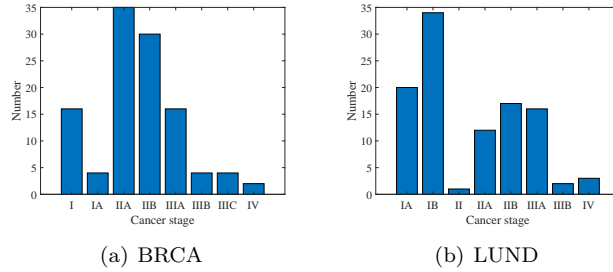


Fig. S-1. Number of patients in each stages for BRCA and LUND.

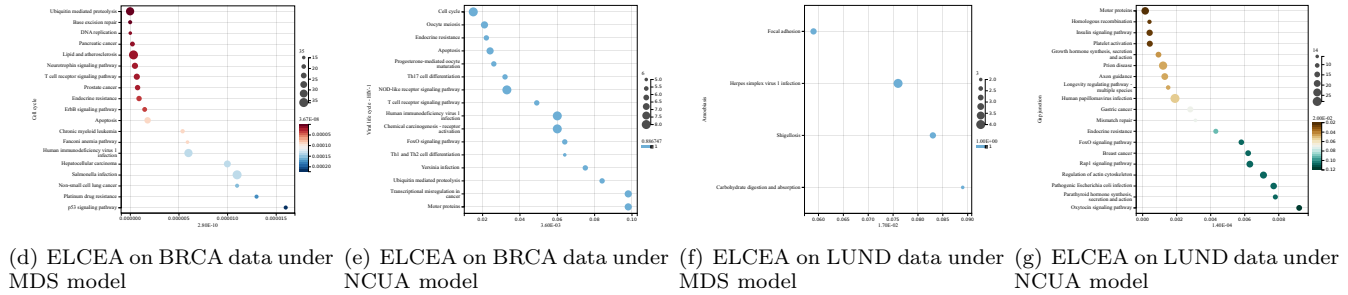
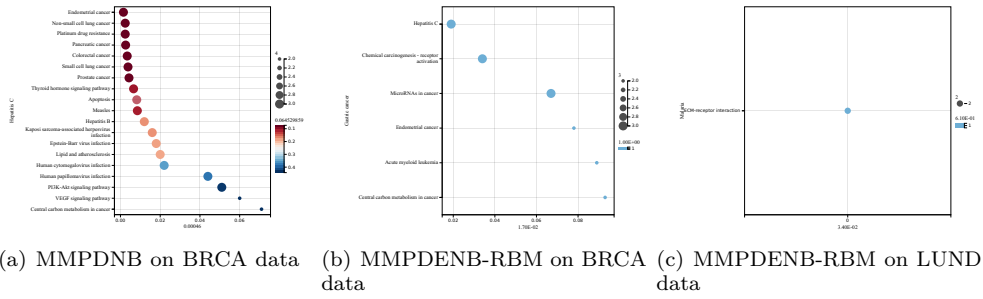


Fig. S-2. Pathway enrichment analysis. Since the result of MMPDNB on LUND data is not shown because it only finds one pathway.

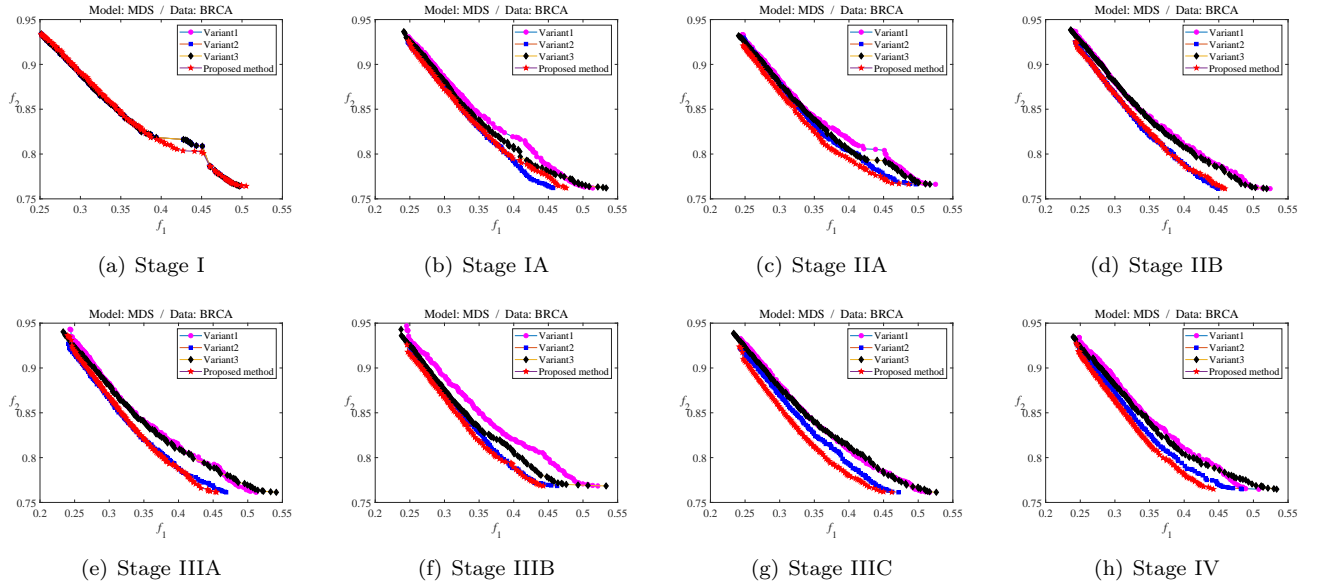


Fig. S-3. Populations of algorithms on the first patient data for each stage of BRCA data under MDS framework.