Supplementary Materials

The Robustness of Pathway Analysis in Identifying Potential Drug Targets in Non-Small Cell Lung Carcinoma

Andrew Dalby and Ian Bailey

Figure S1. The pathway analysis between normal and adenocarcinoma in the gcrma normalized G-GEOD-6044 dataset.
Figure S2. The pathway analysis between normal and squamous cell carcinoma in the rma normalized G-GEOD-6044 dataset.
Figure S3. The pathway analysis between normal and squamous cell carcinoma in the gcrma normalized G-GEOD-6044 dataset.
Figure S4. The pathway analysis between normal and squamous cell carcinoma in the farms normalized G-GEOD-6044 dataset.