**S2 Table.** DAVID Term Enrichment for DNA Copy Number Deconvolution. The table provides significantly enriched \((p \leq 0.05)\) terms, identified by source repository and term, Benjamini-corrected p-values, and associated vertices of the inferred simplicial complex.

<table>
<thead>
<tr>
<th>Source</th>
<th>Term</th>
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