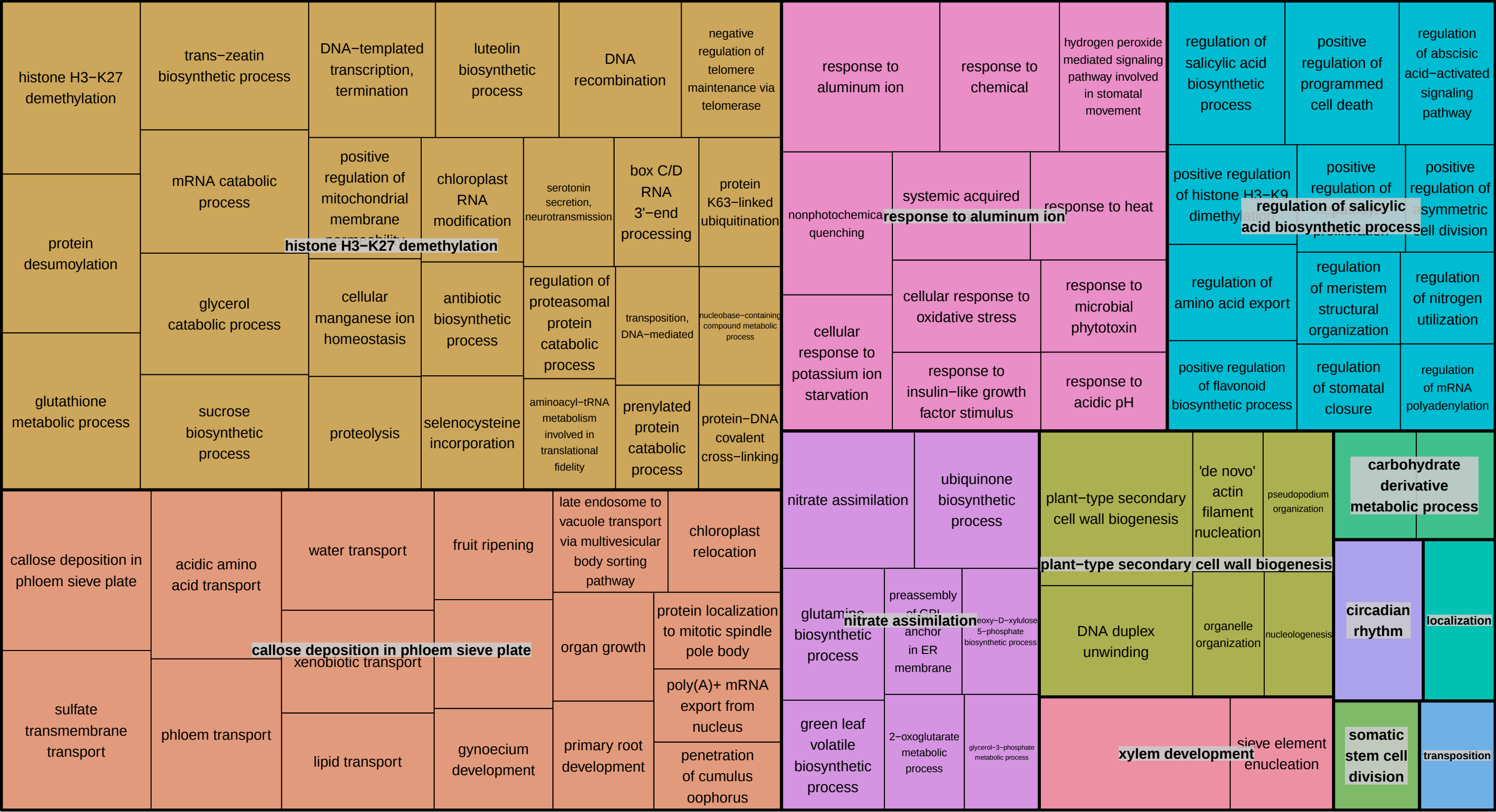


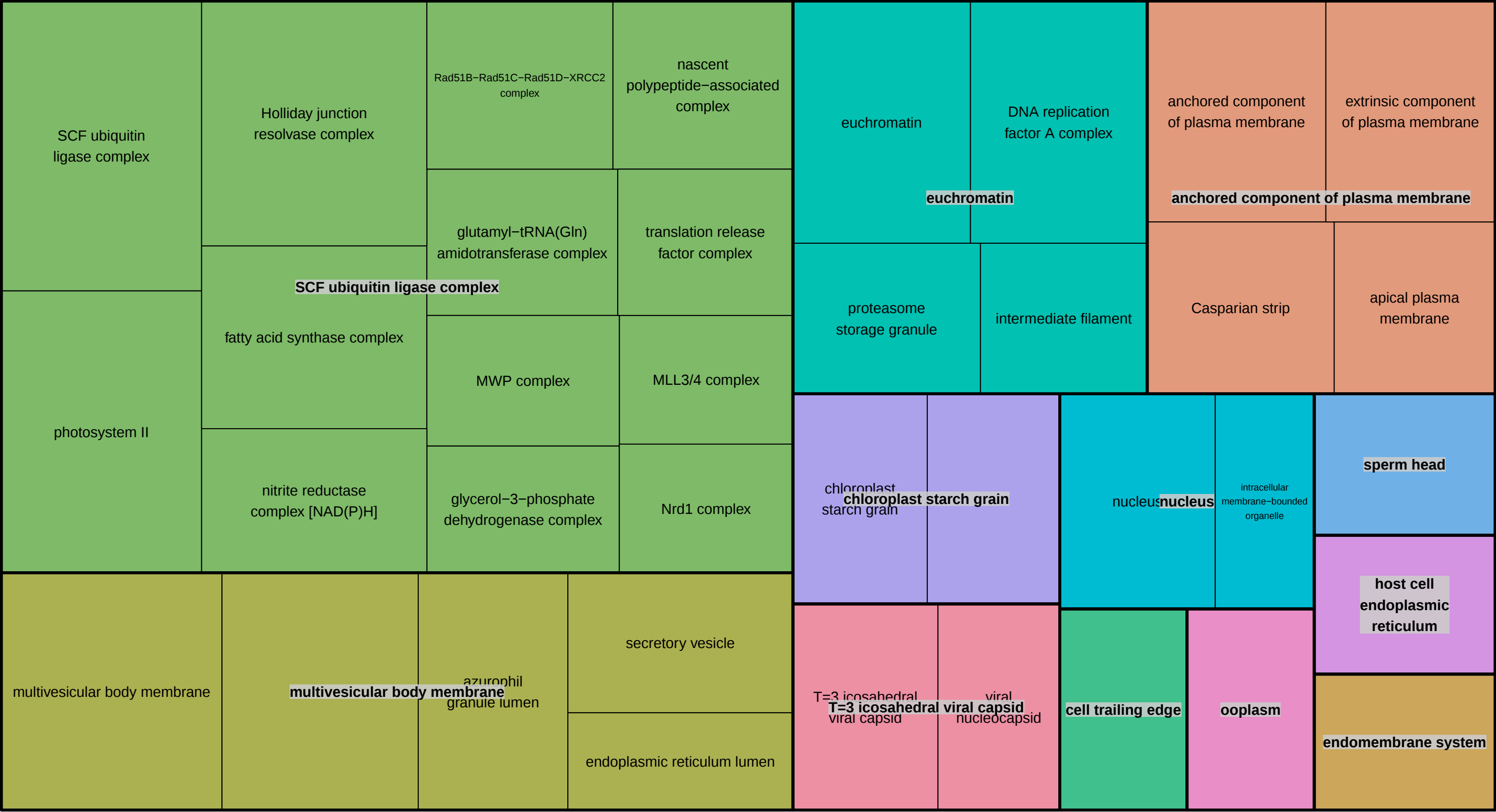
Additional file 13: “TreeMap” views of significant GO terms constructed with REVIGO (Supek et al., 2011). Each rectangle is a single cluster representative. The representatives are joined into ‘superclusters’ of loosely related terms, visualized with different colors. Size of the rectangles reflect the p -value.

Supek, Fran, et al. "REVIGO summarizes and visualizes long lists of gene ontology terms." PloS one 6.7 (2011): e21800.

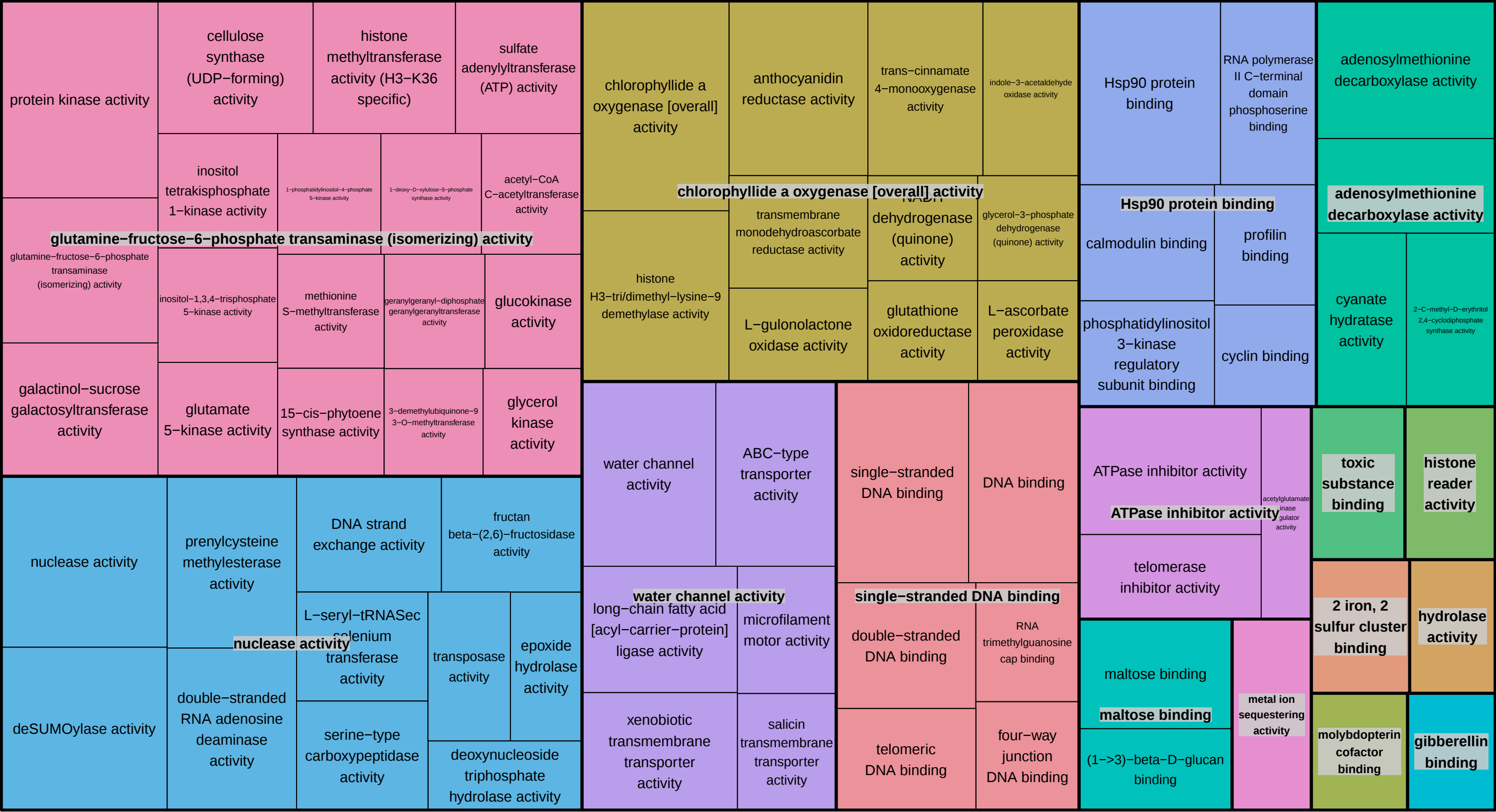
Revigo TreeMap – SAM TAP – Biological Process



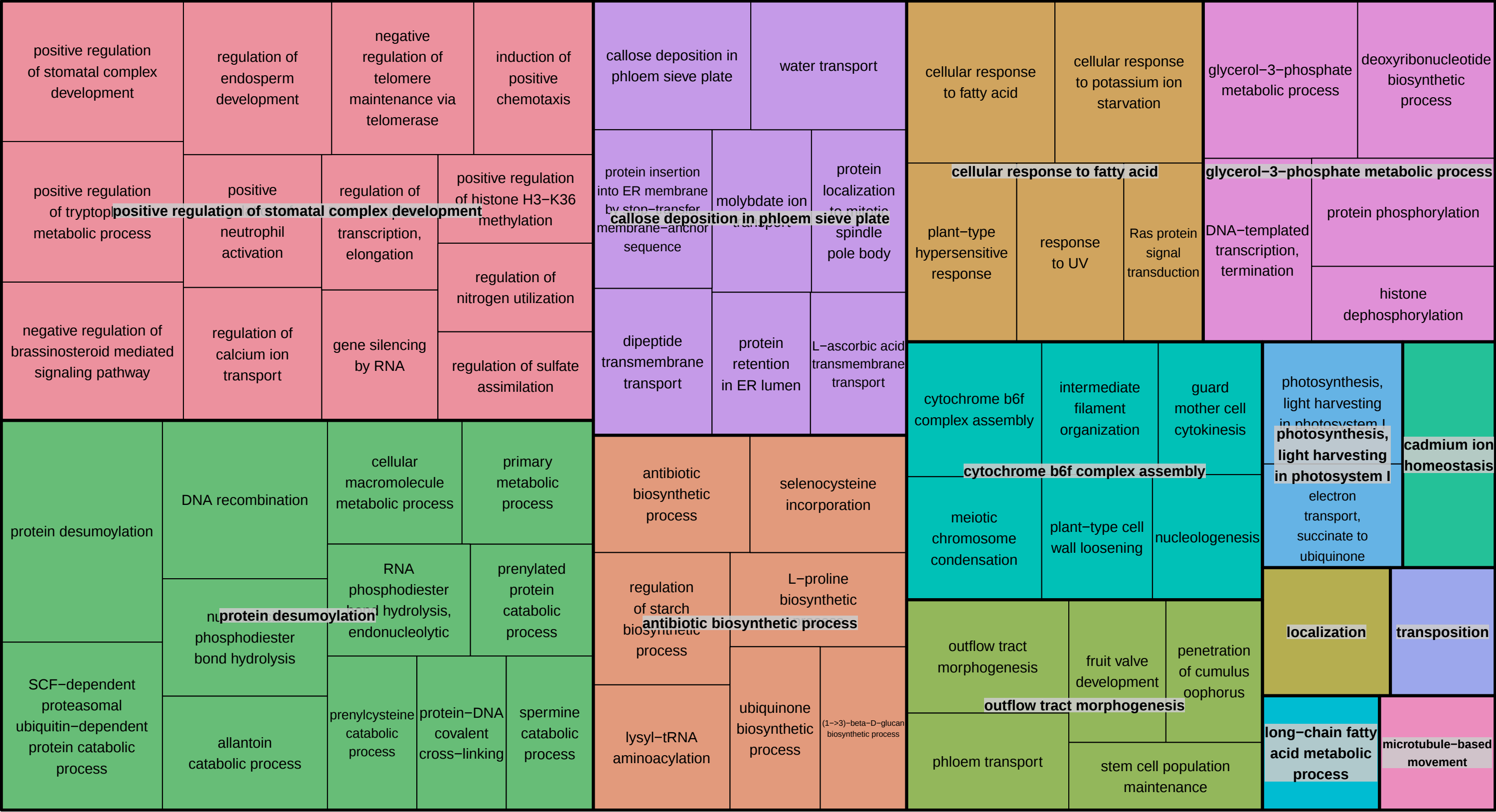
Revigo TreeMap – SAM TAP – Cellular Component



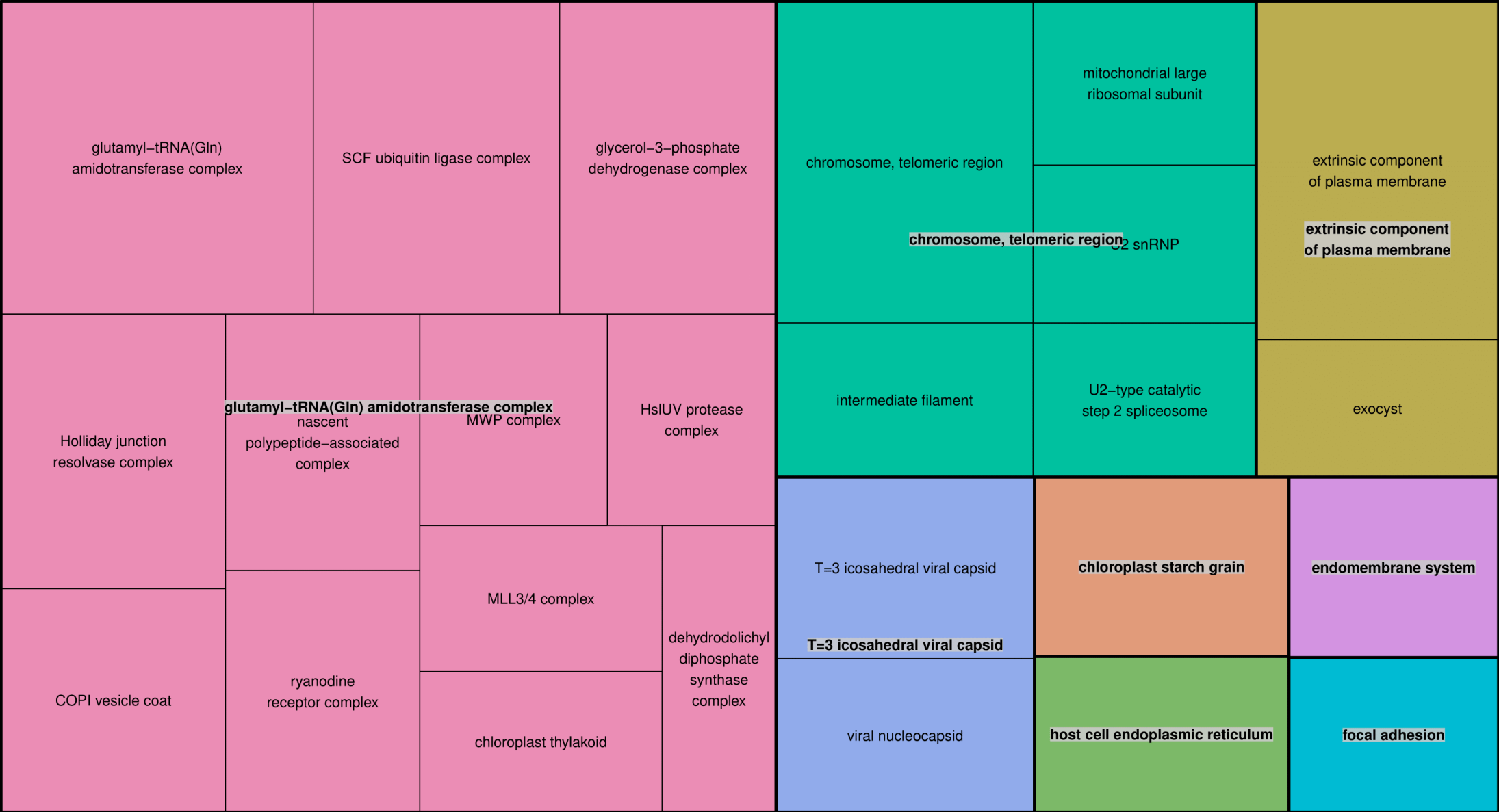
Revigo TreeMap – SAM TAP – Molecular Function



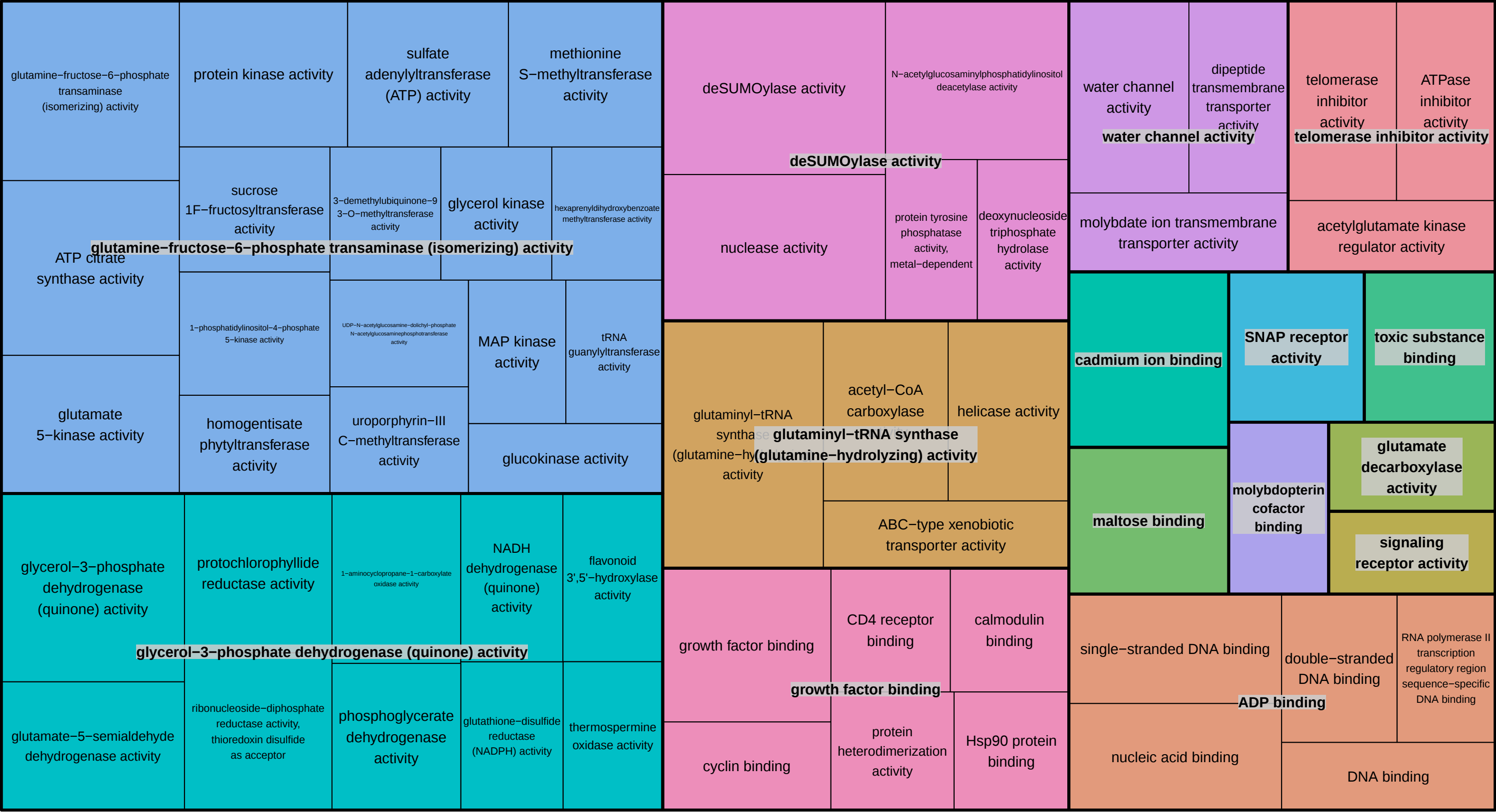
Revigo TreeMap – SAM DRS – Biological Process



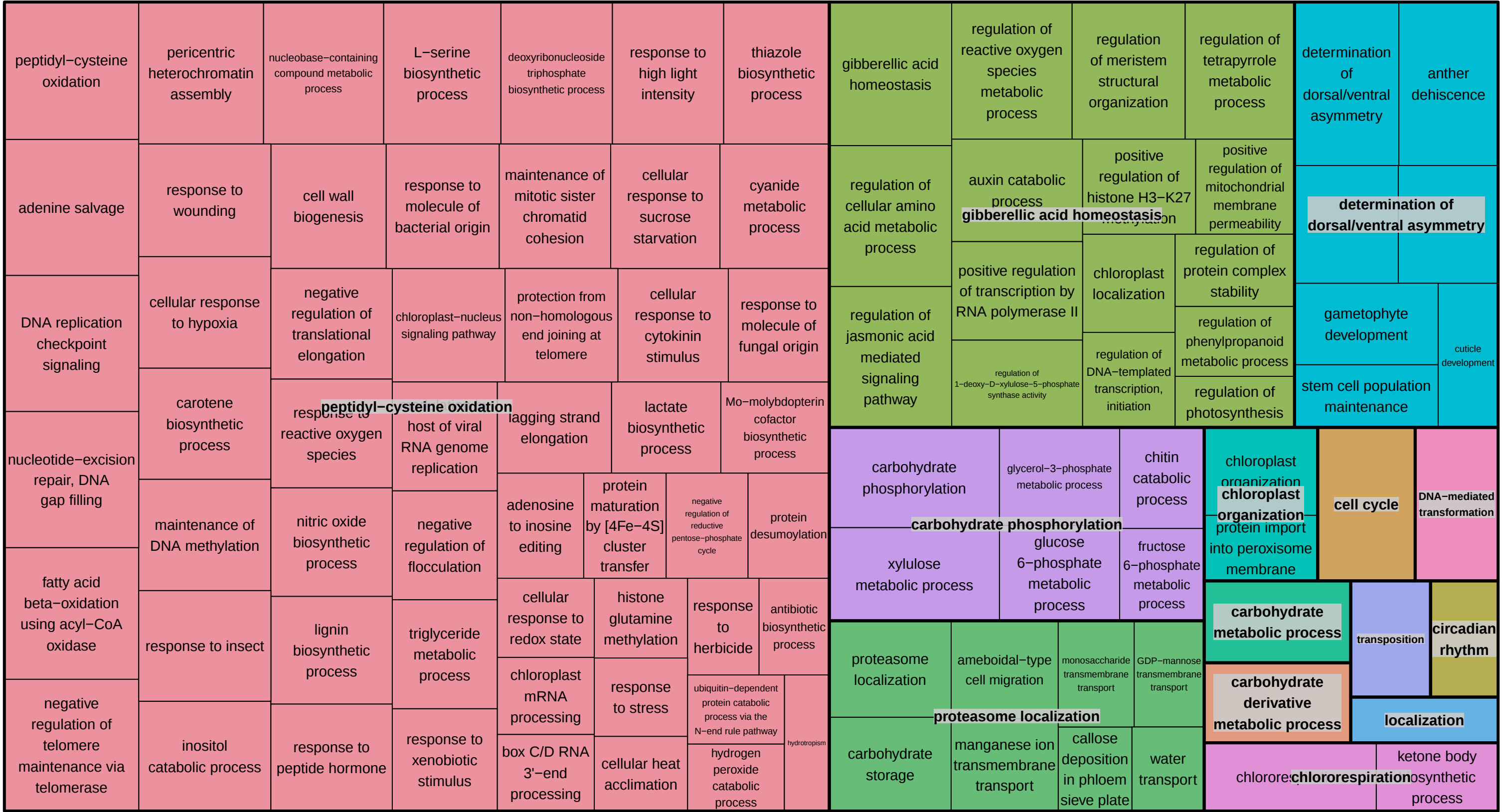
Revigo TreeMap – SAM DRS – Cellular Component



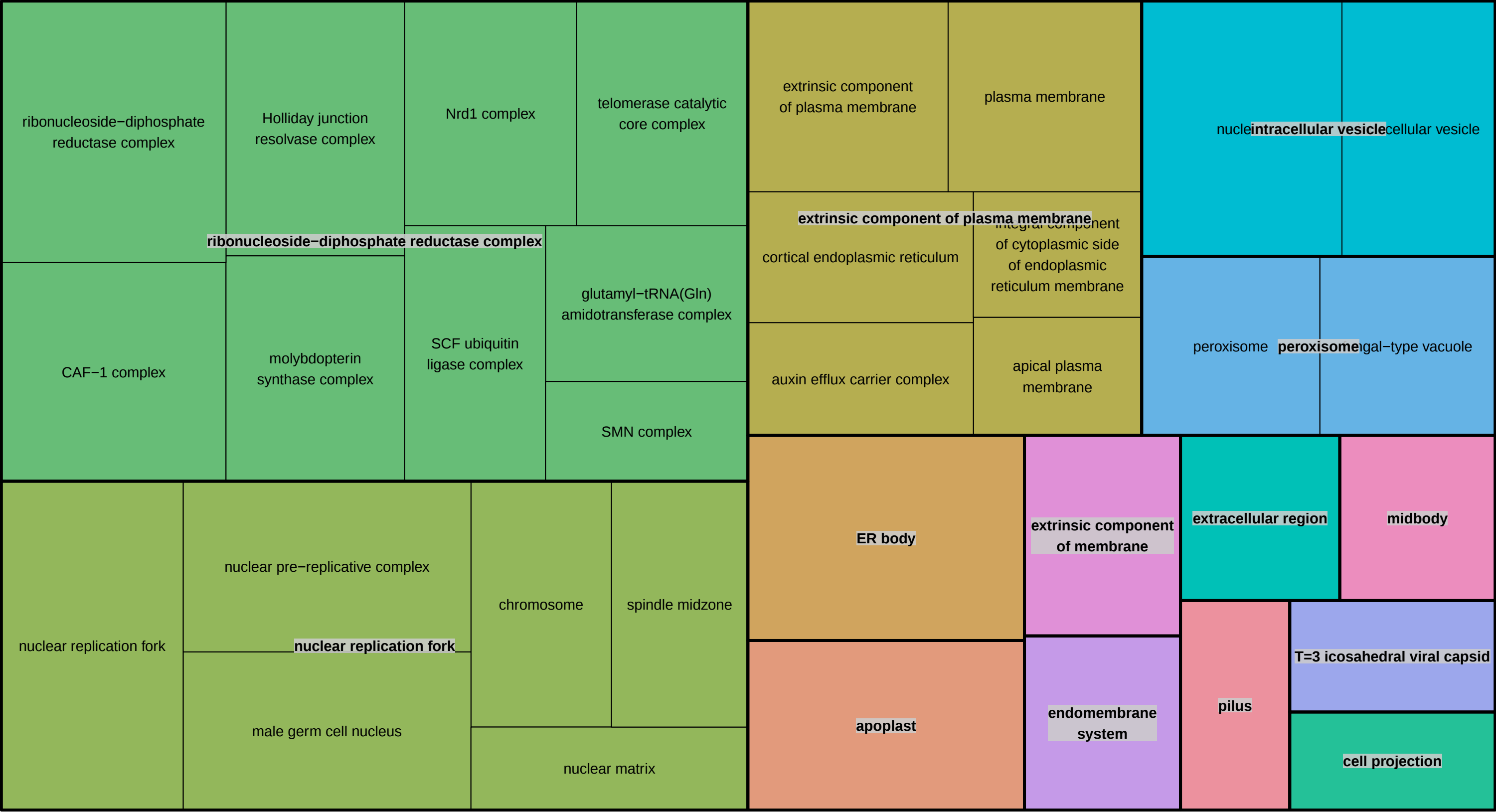
Revigo TreeMap – SAM DRS – Molecular Function



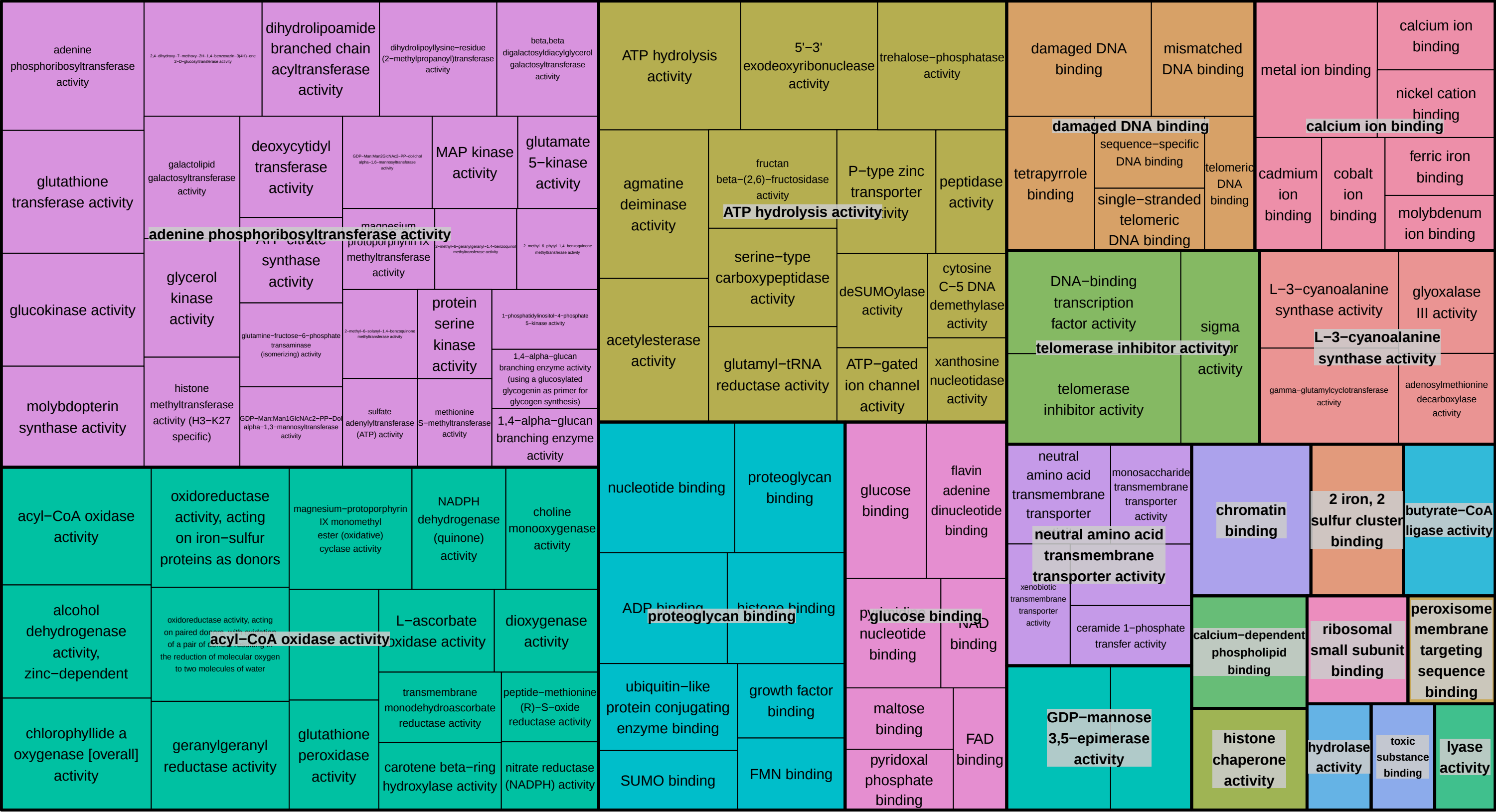
Revigo TreeMap – SAM LRP – Biological Process



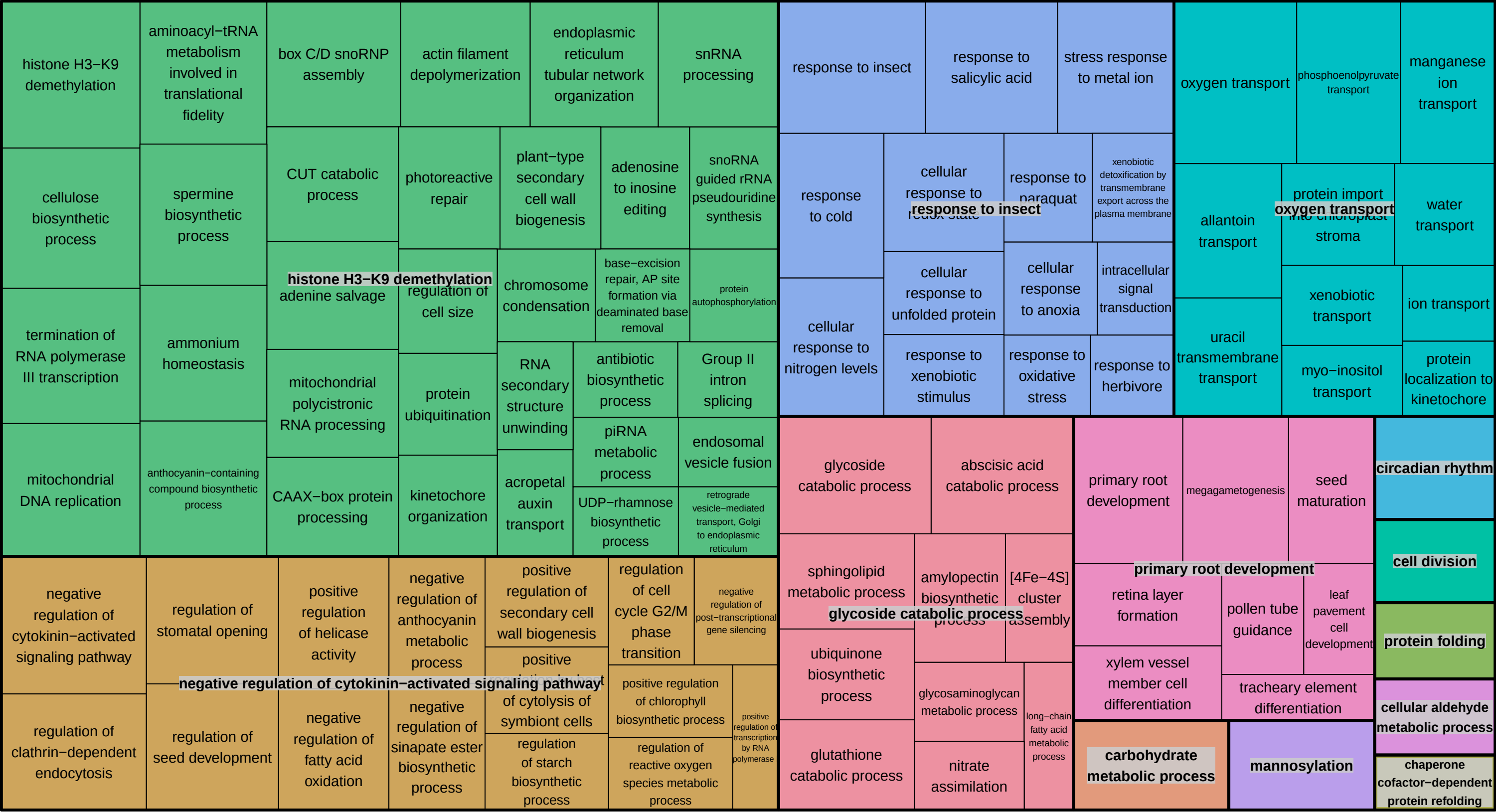
Revigo TreeMap – SAM LRP – Cellular Component



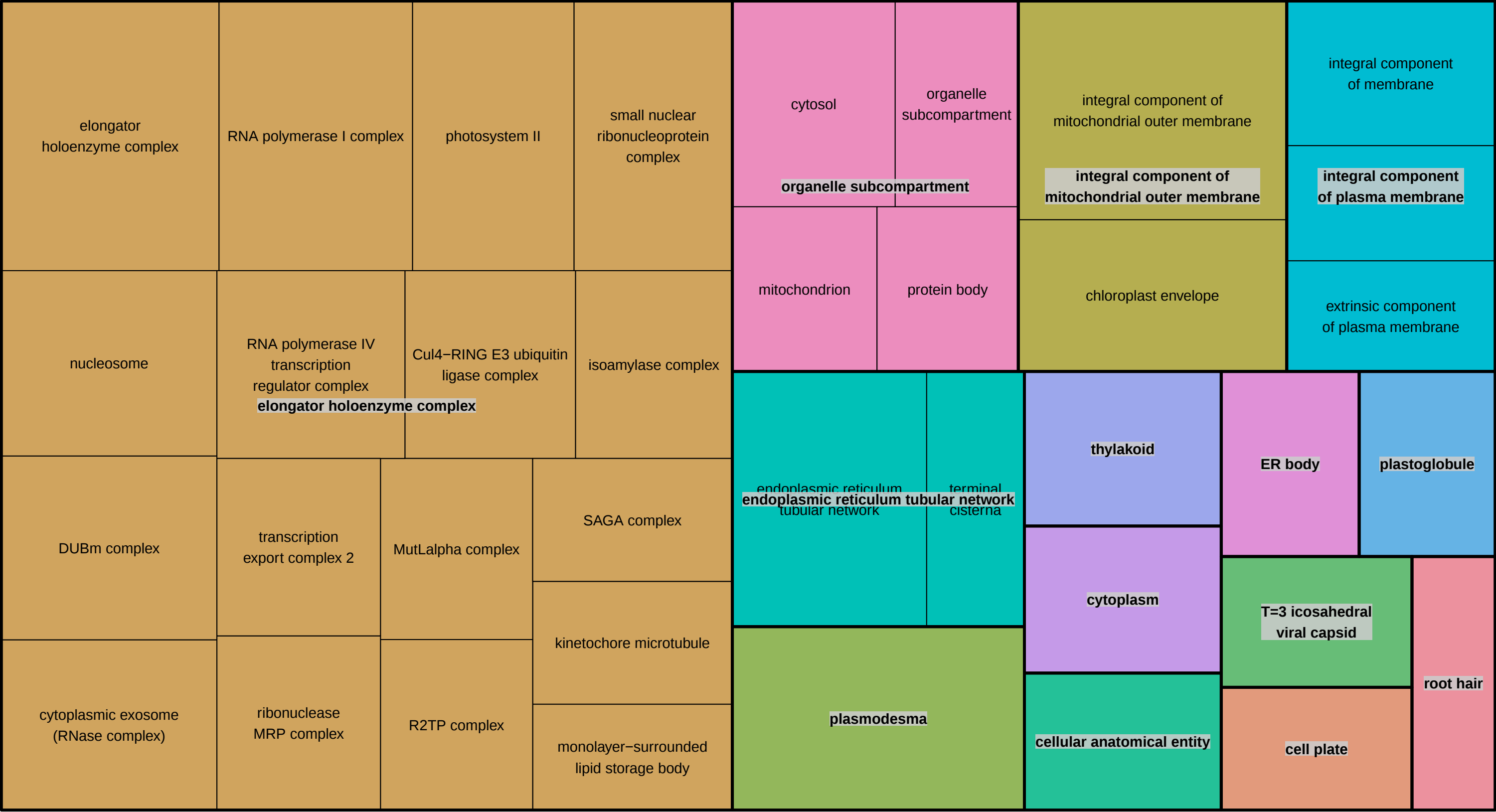
Revigo TreeMap – SAM LRP – Molecular Function



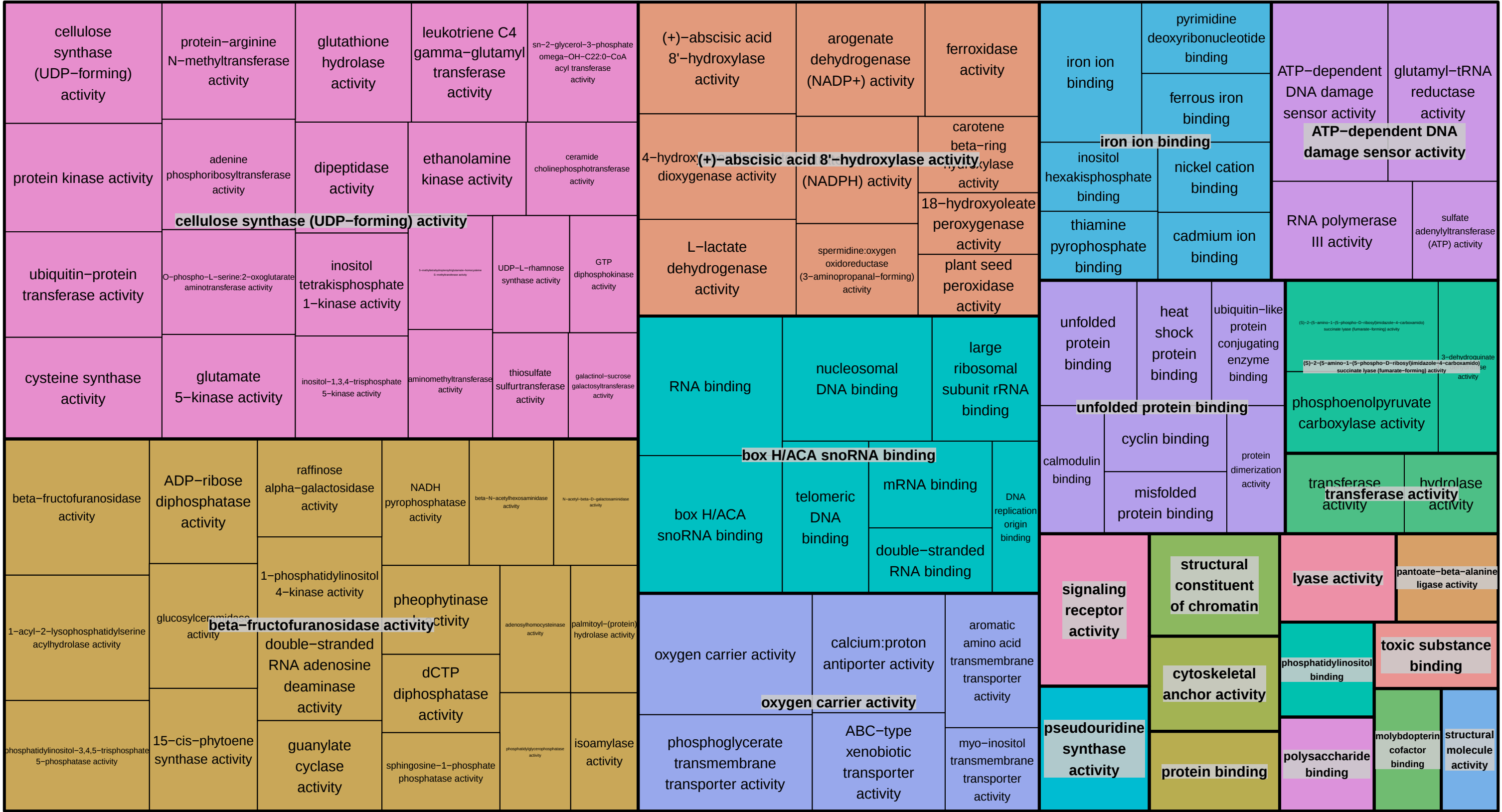
Revigo TreeMap – Leaves Tap – Biological Process



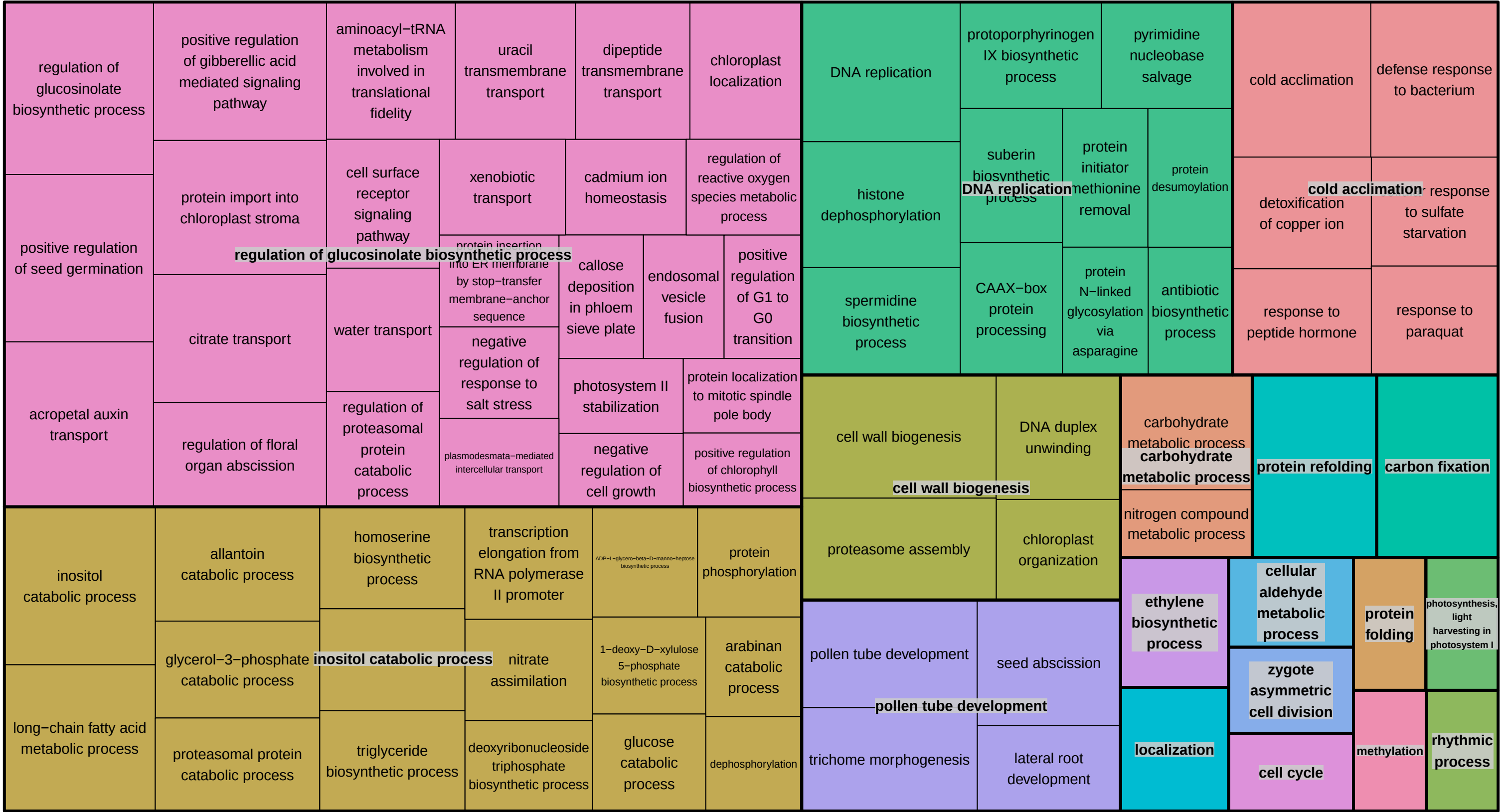
Revigo TreeMap – Leaves TAP – Cellular Component



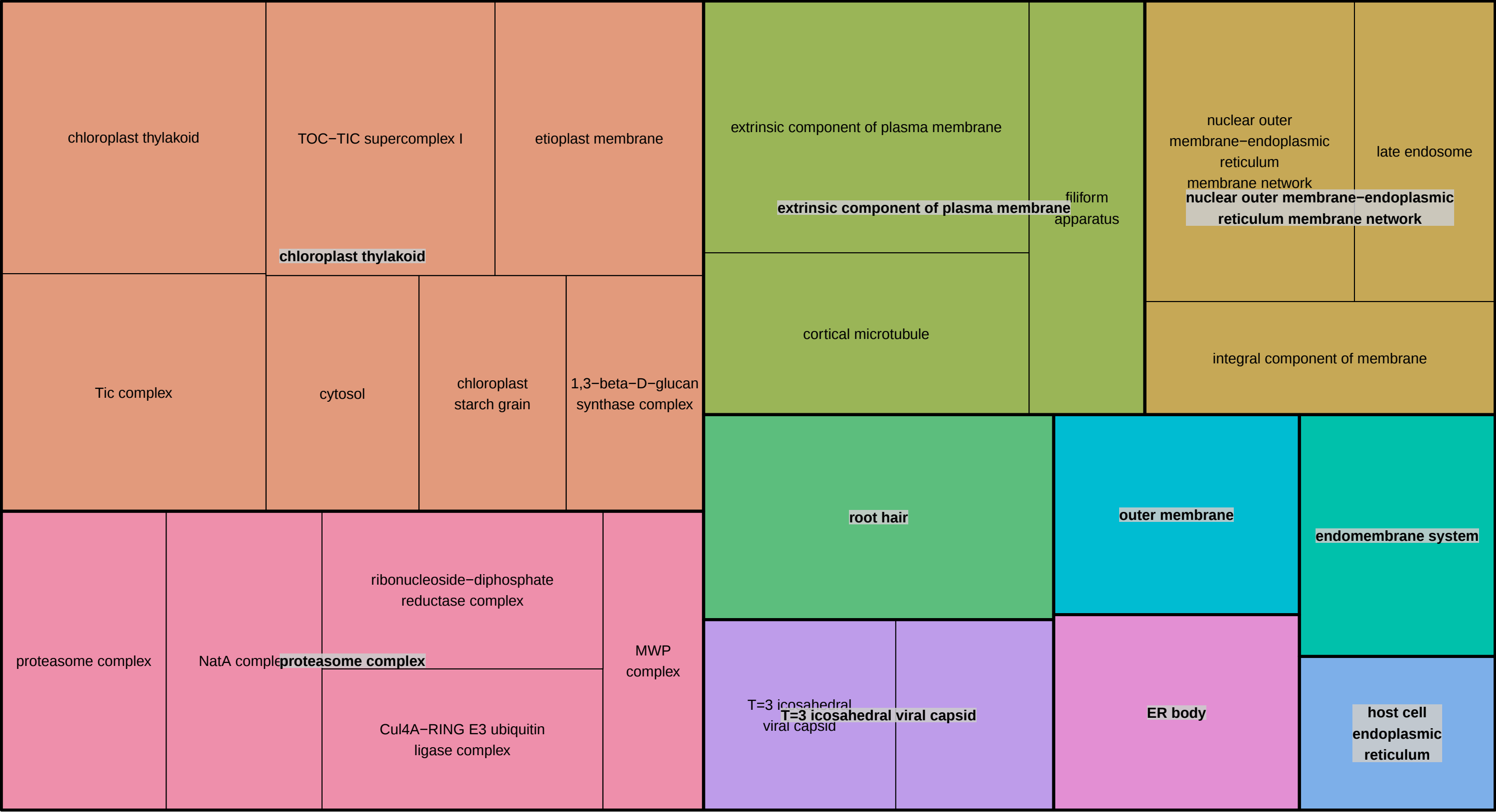
Revigo TreeMap – Leaves TAP – Molecular Function



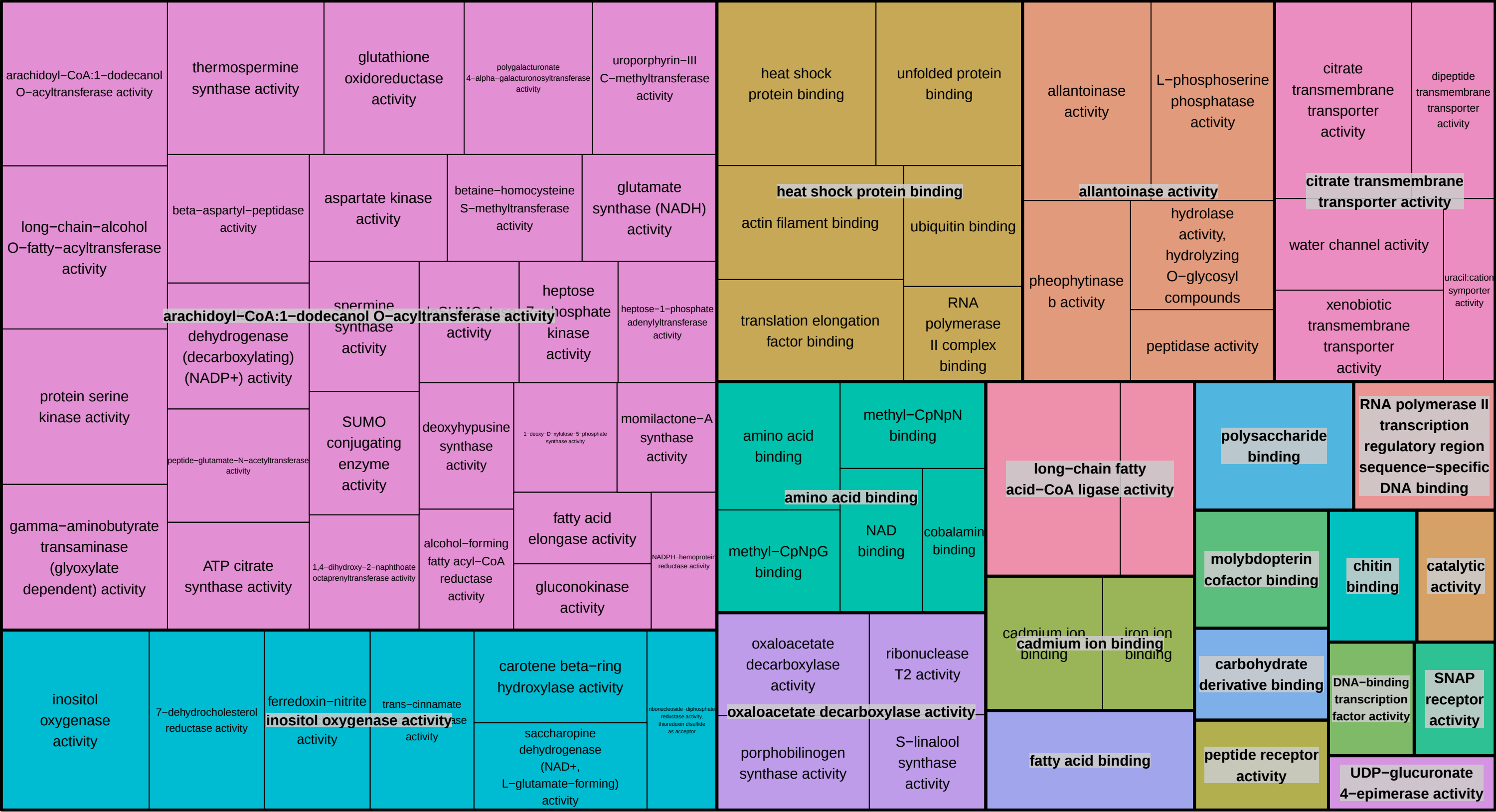
Revigo TreeMap – Leaves DRS – Biological Process



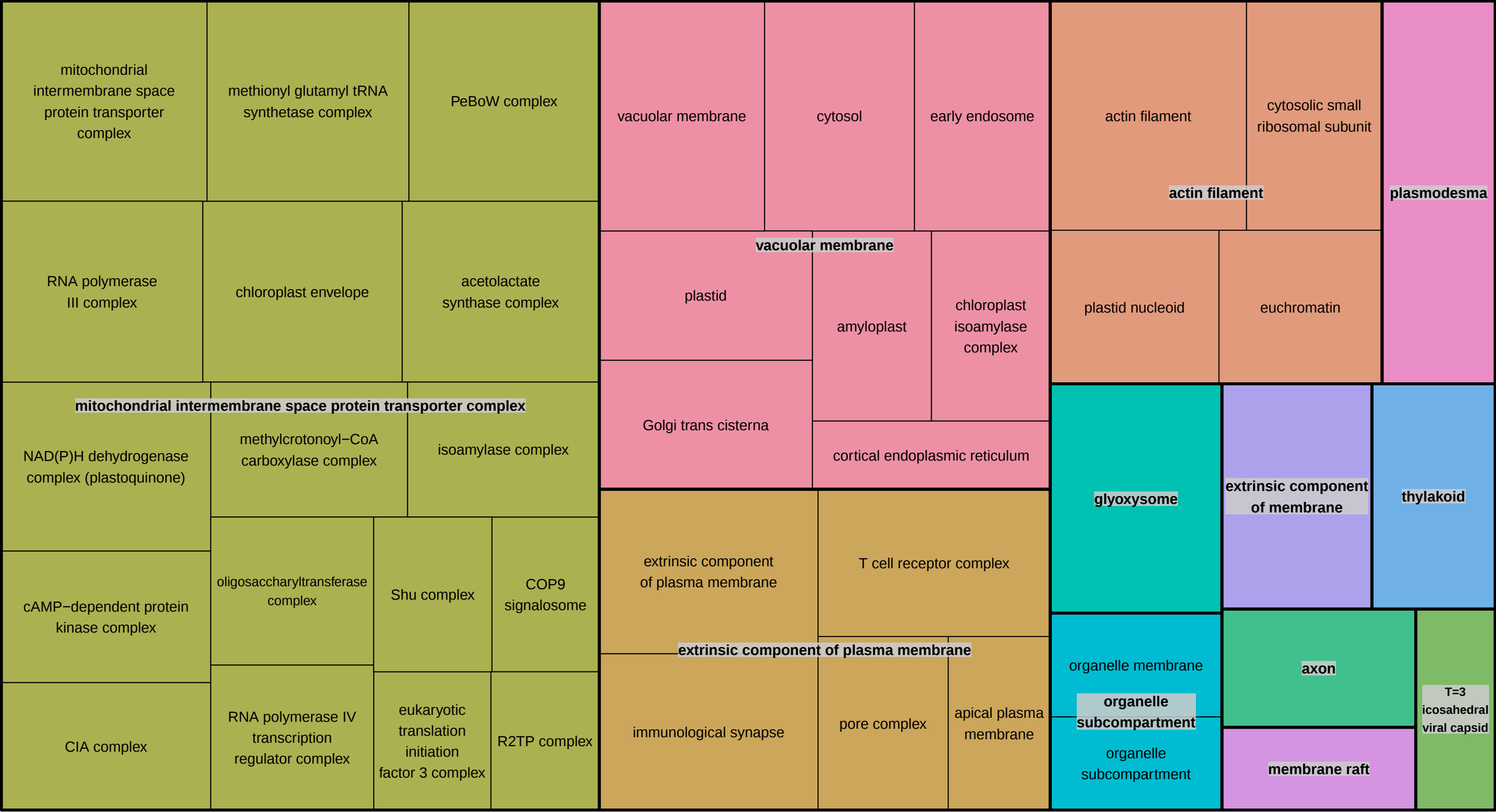
Revigo TreeMap – Leaves DRS – Cellular Component



Revigo TreeMap – Leaves DRS – Molecular Function



Revigo TreeMap – Leaves LRP – Cellular Component



Revigo TreeMap – Leaves LRP – Molecular Function

