

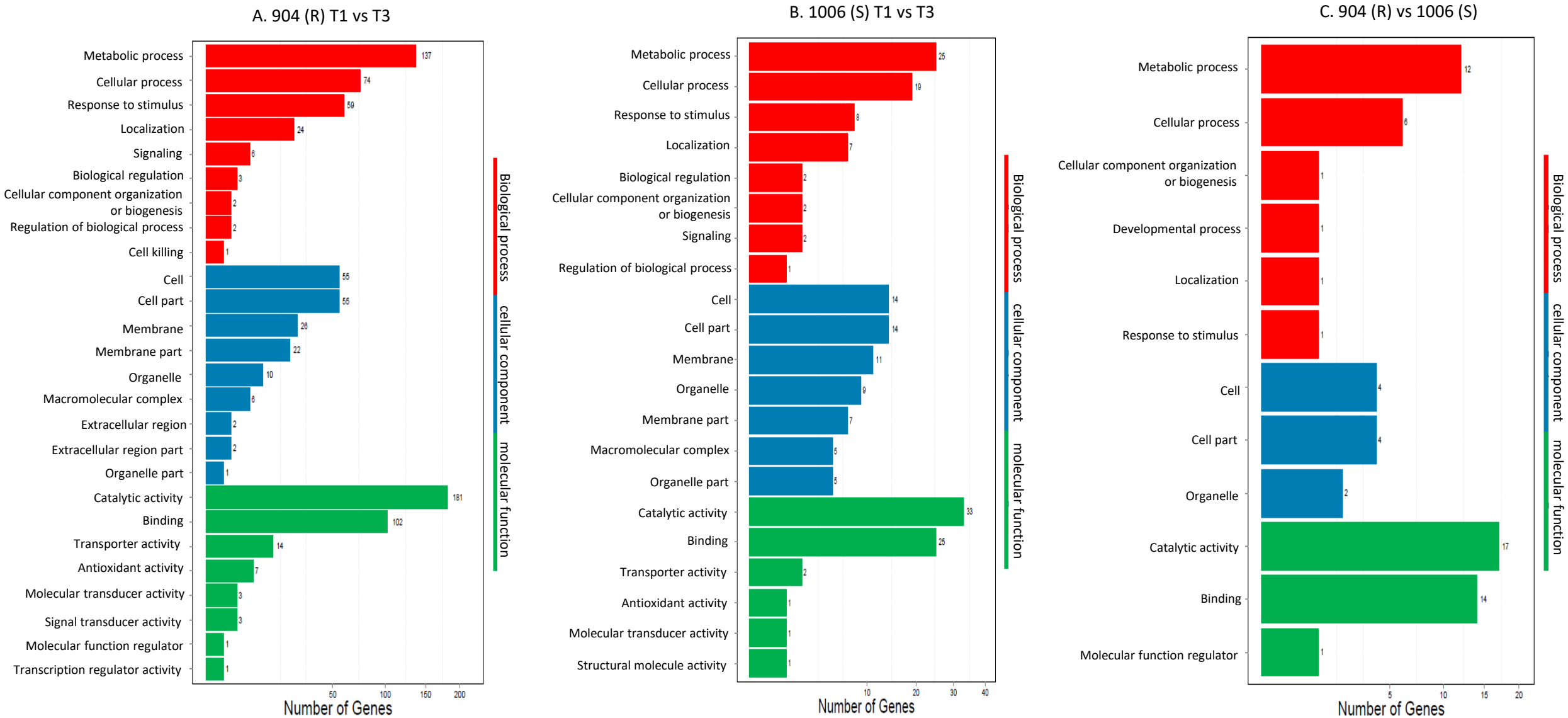


Fig. 4. Gene ontology (GO) enrichment analysis showed DEGs involved in three function categories. A. T1 (4 h) vs T3 (44 h) in R line; B. T1 vs T3 in S line; C. R vs S lines.