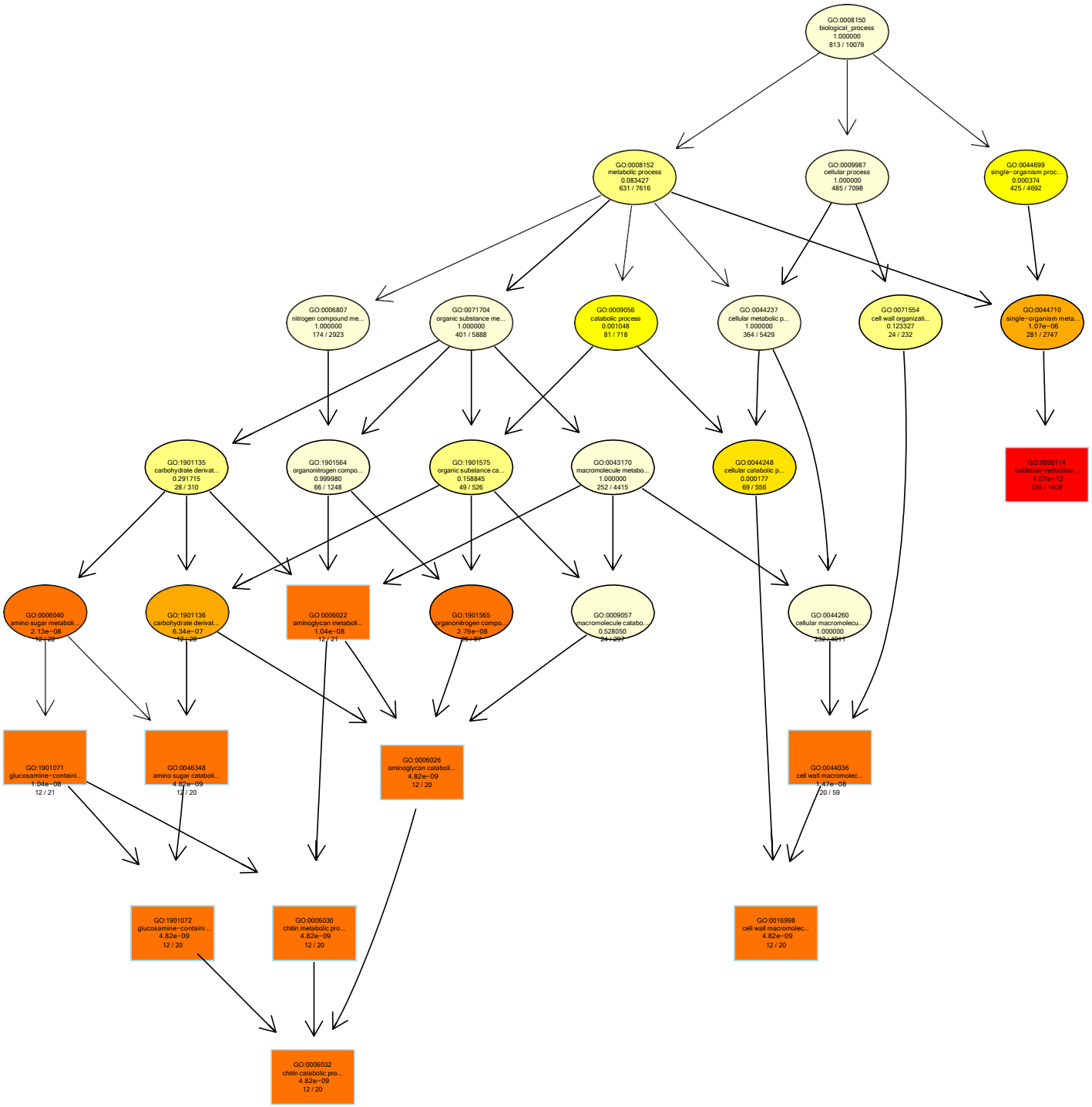

Figure S1: Biological processes that DEGs between etiolated and green wheat seedlings on vernalization 10 d involved in, revealed by topGO enrichment analysis.

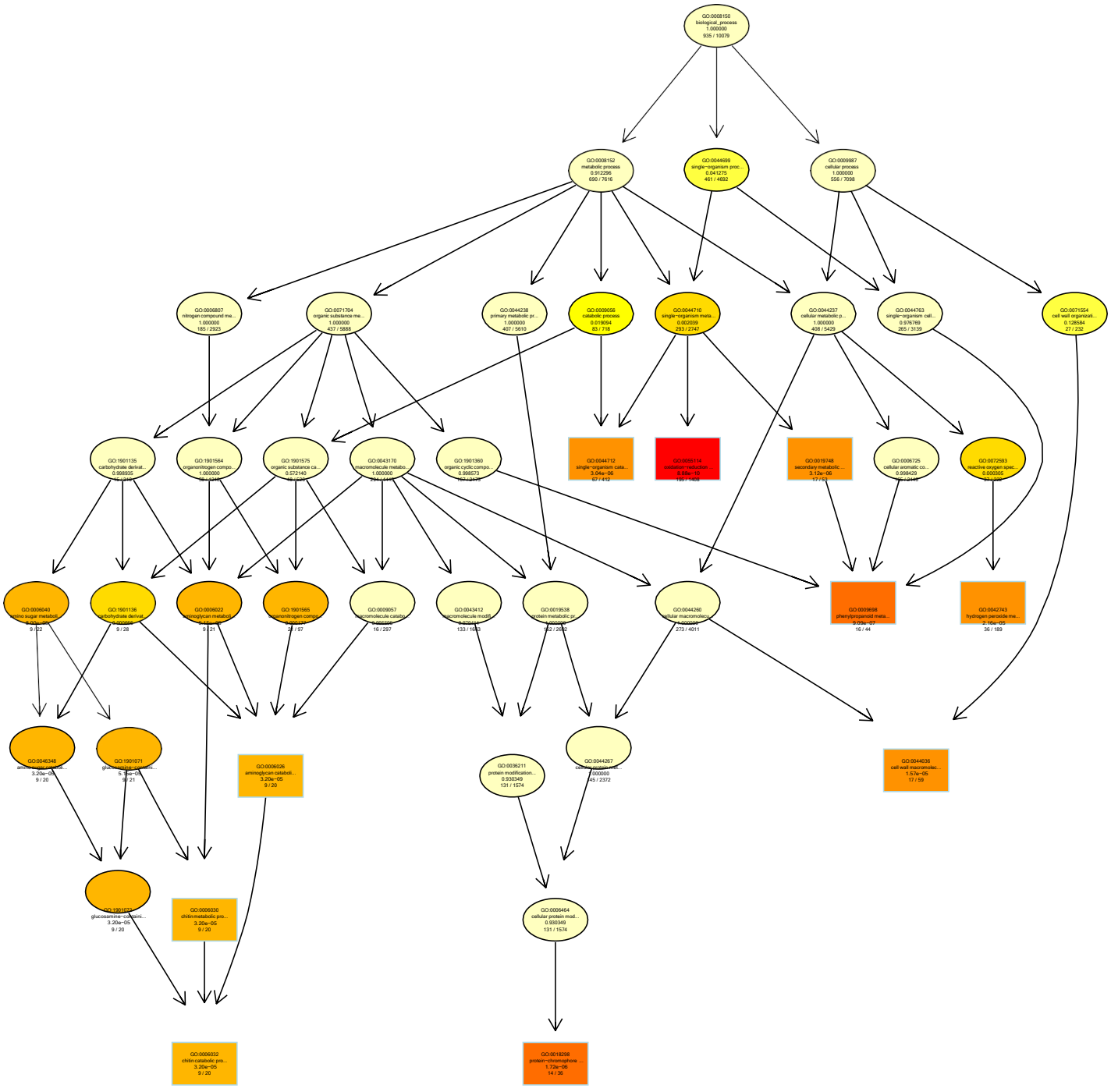

Figure S2: Biological processes that DEGs between etiolated and green wheat seedlings on vernalization 20 d involved in, revealed by topGO enrichment analysis.

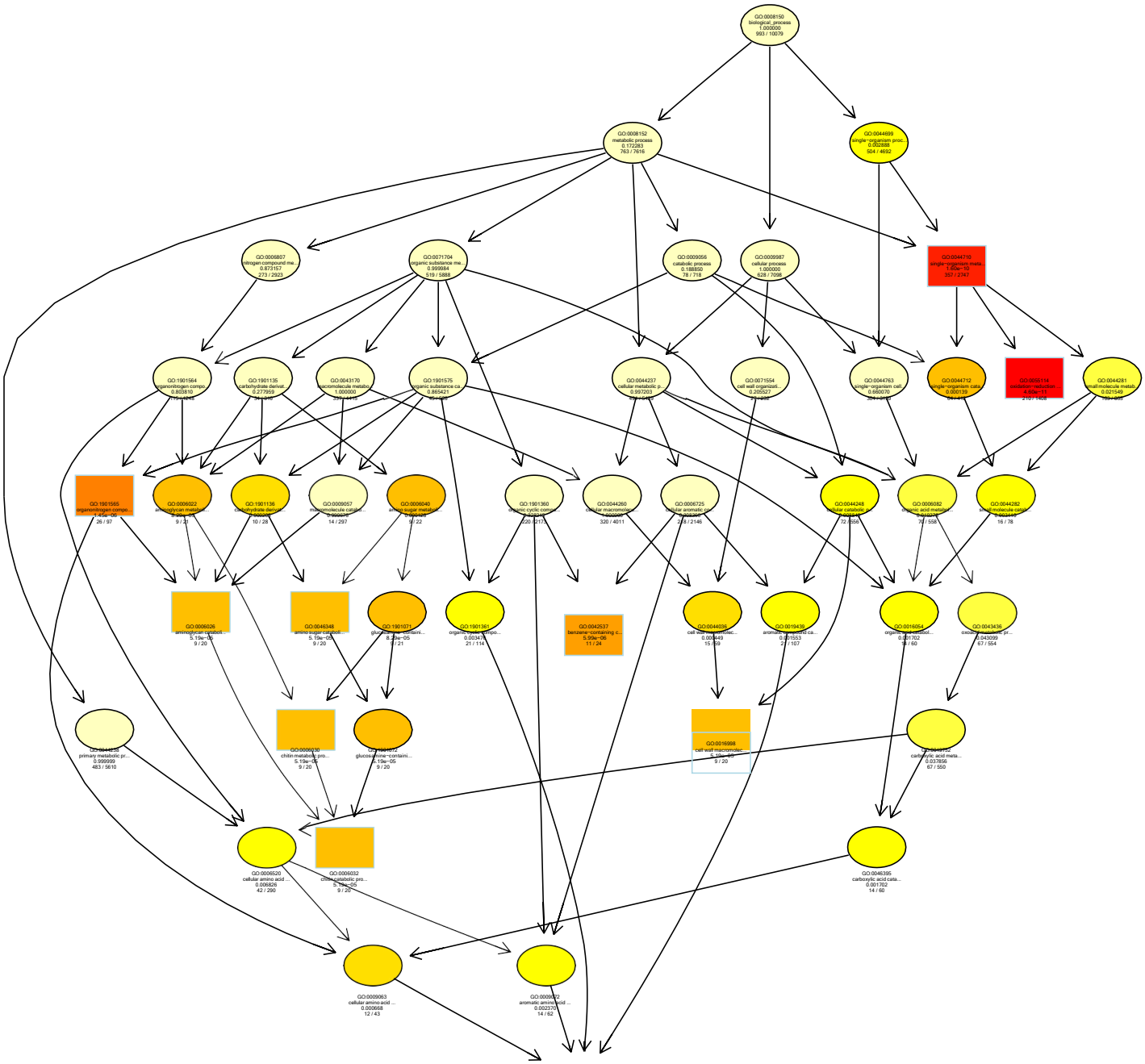

Figure S3: Biological processes that DEGs between etiolated and green wheat seedlings on vernalization 30 d involved in, revealed by topGO enrichment analysis.

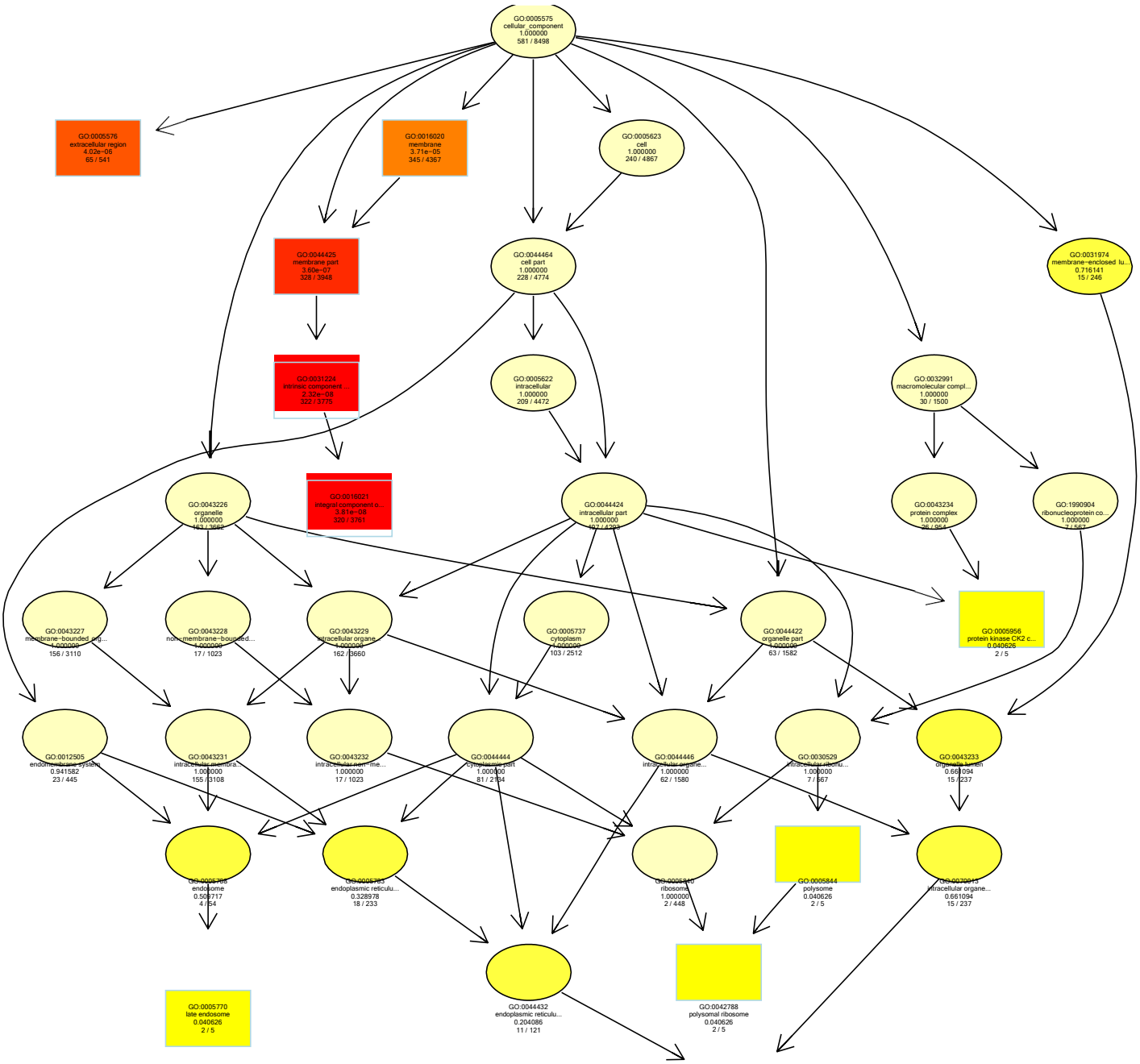

Figure S4: Cellular component where DEGs were found, revealed by topGO enrichment analysis. The DEGs were between etiolated and green wheat seedlings on vernalization 10 d.

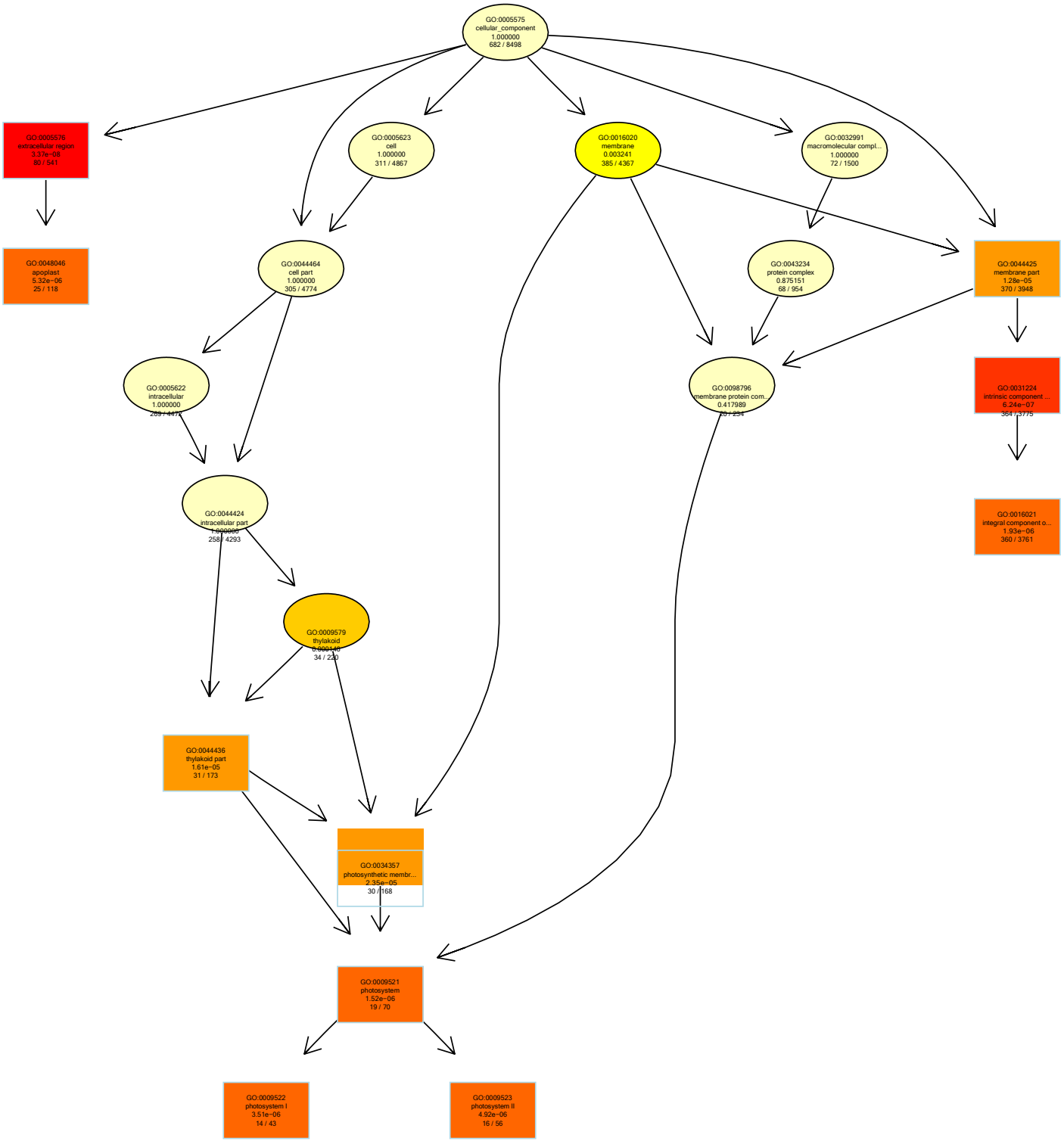

Figure S5: Cellular component where DEGs were found, revealed by topGO enrichment analysis. The DEGs were between etiolated and green wheat seedlings on vernalization 20 d.

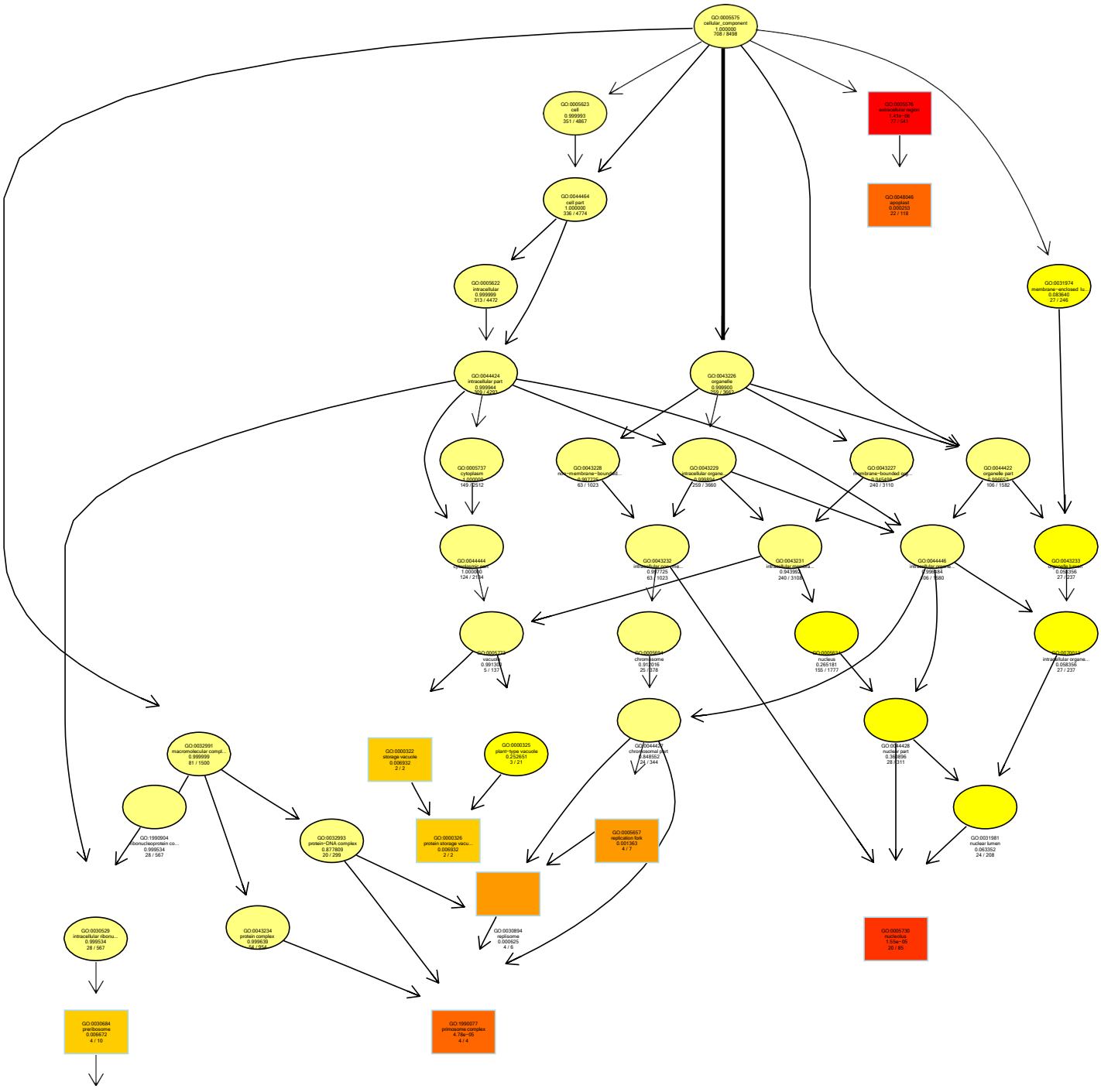

Figure S6: Cellular component where DEGs were found, revealed by topGO enrichment analysis. The DEGs were between etiolated and green wheat seedlings on vernalization 30 d.

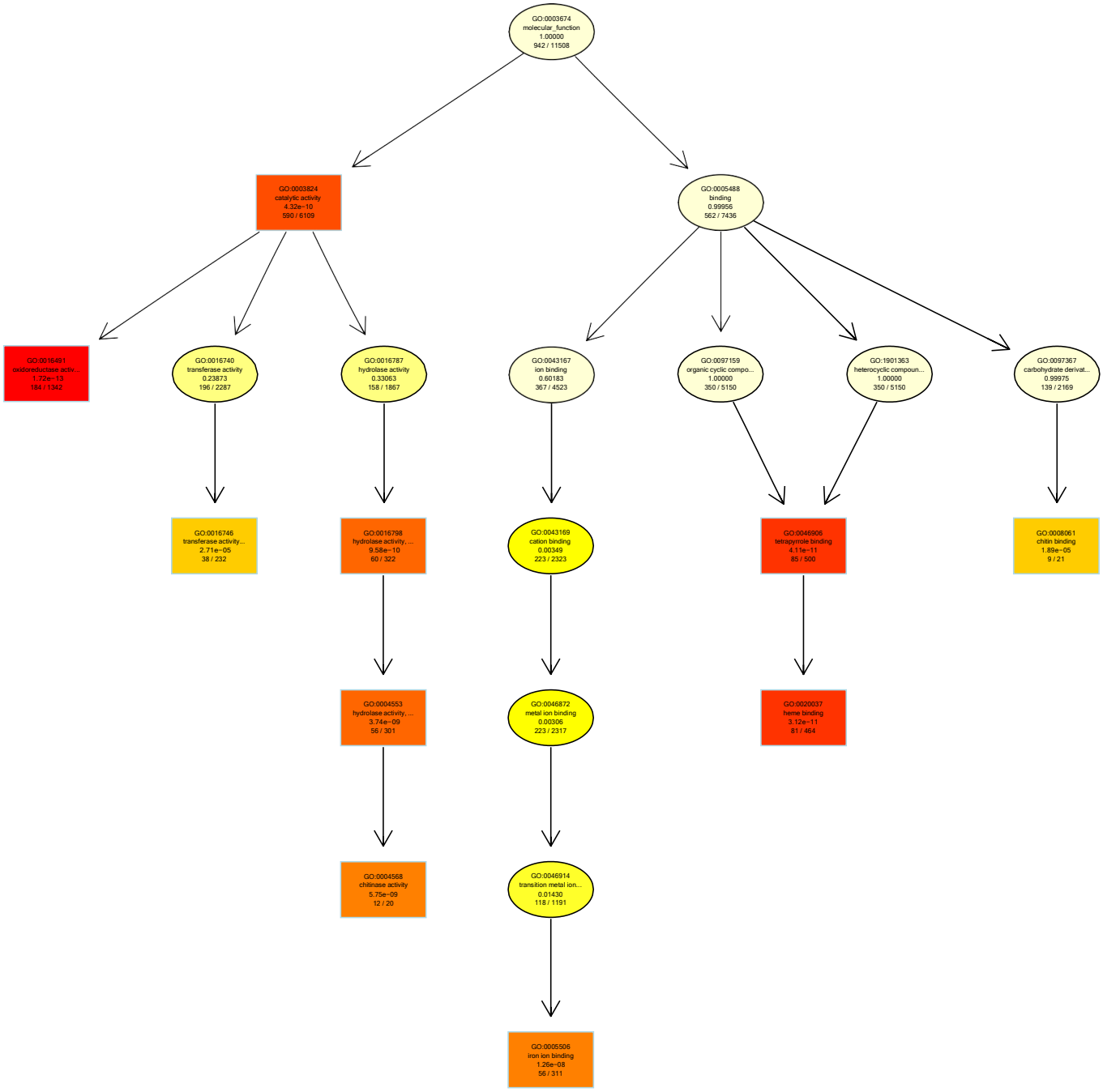

Figure S7: Molecular functions of DEGs between etiolated and green wheat seedlings on vernalization 10 d, revealed by topGO enrichment analysis.

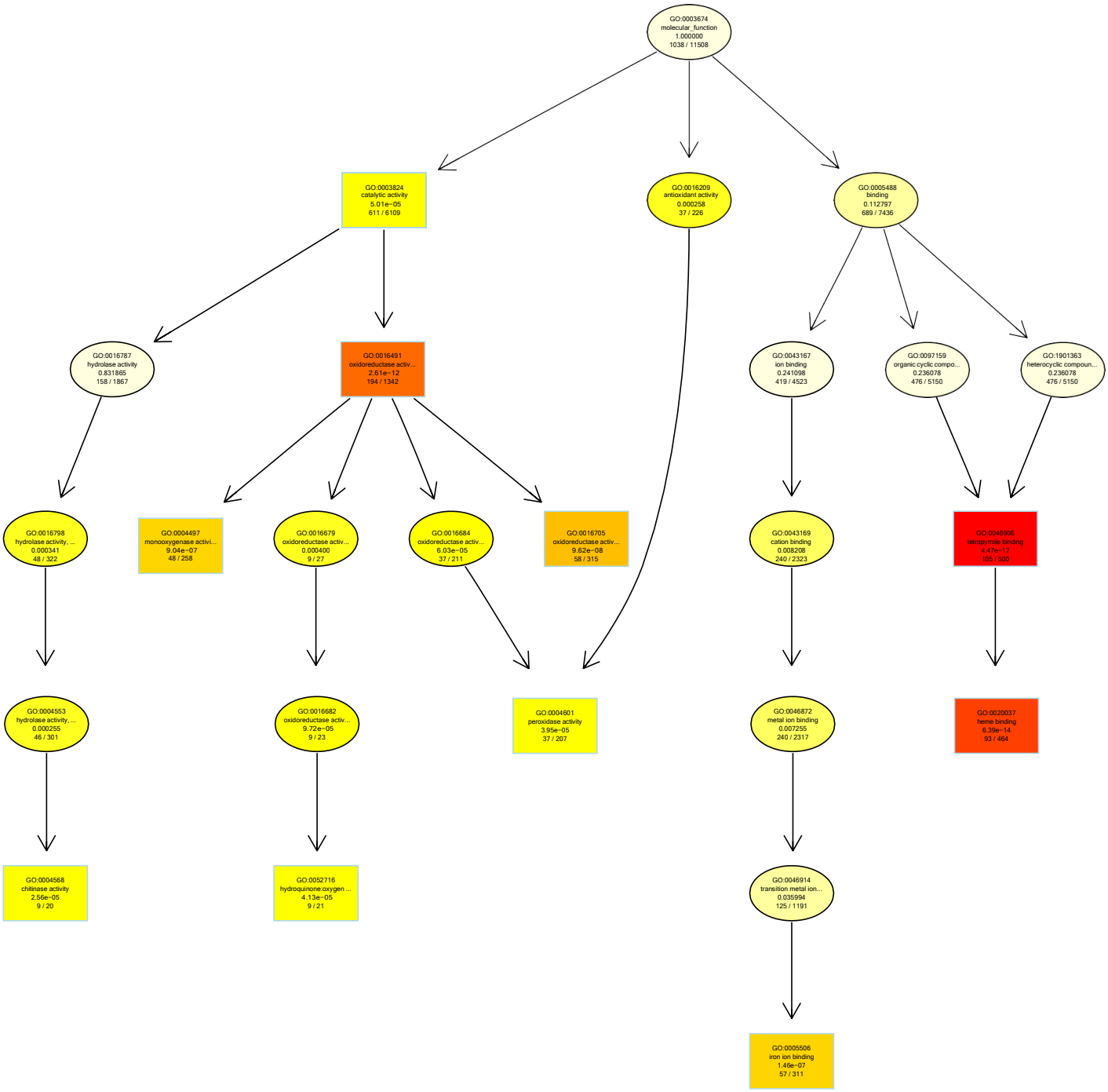

Figure S8: Molecular functions of DEGs between etiolated and green wheat seedlings on vernalization 20 d, revealed by topGO enrichment analysis.

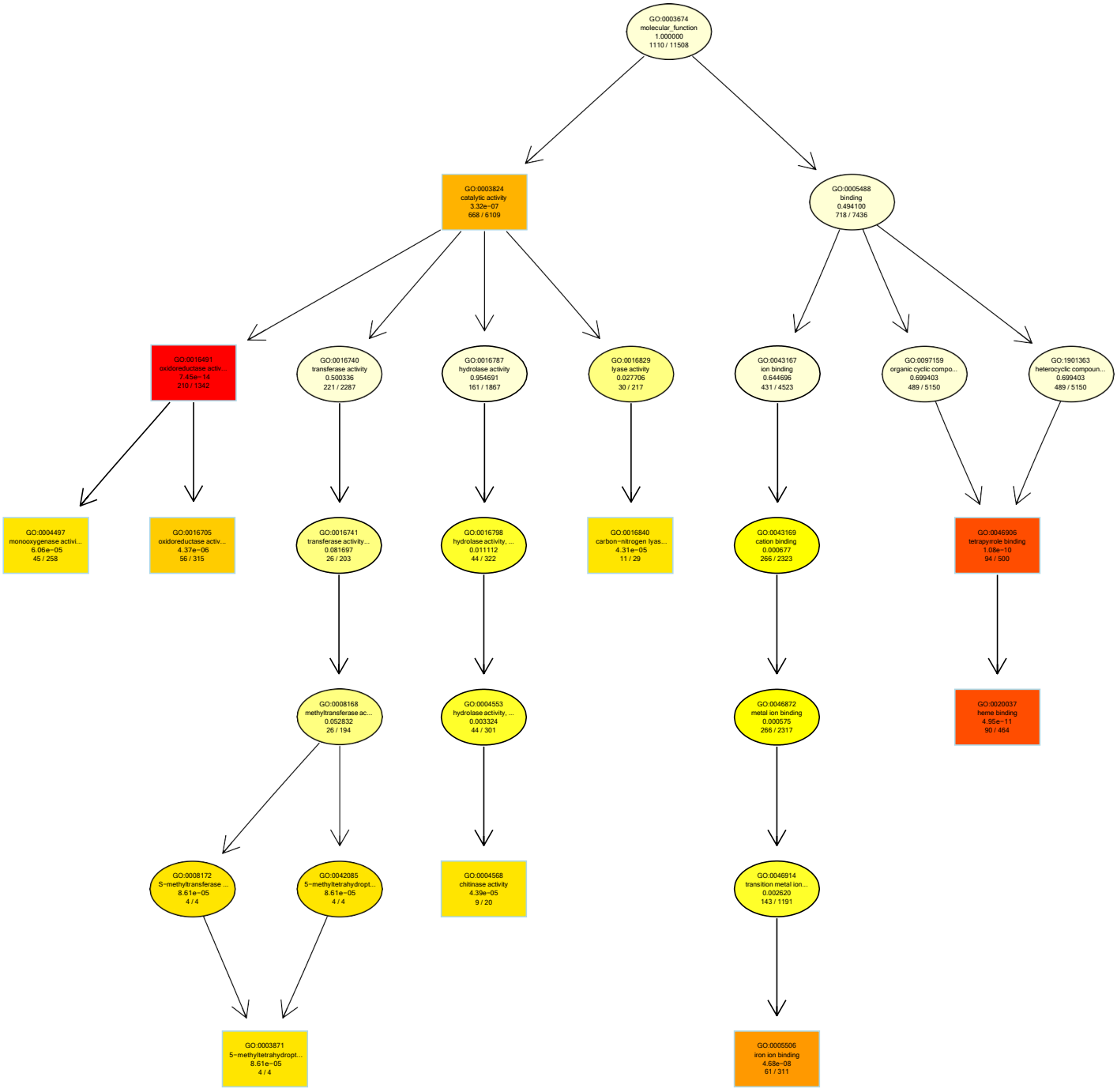

Figure S9: Molecular functions of DEGs between etiolated and green wheat seedlings on vernalization 30 d, revealed by topGO enrichment analysis.

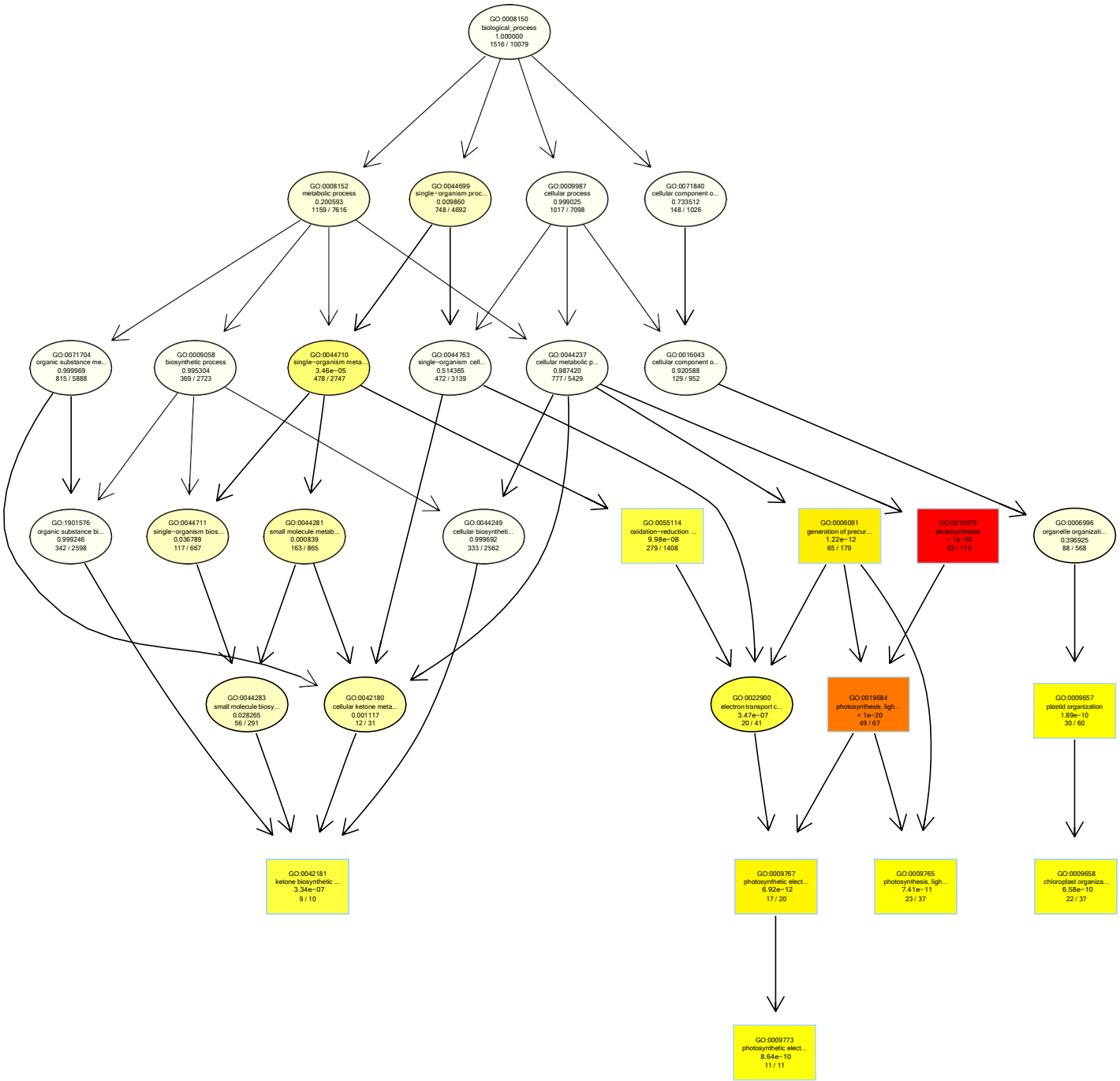

Figure S10: Biological processes that DEGs between etiolated and green wheat seedlings on vernalization 40 d involved in, revealed by topGO enrichment analysis.

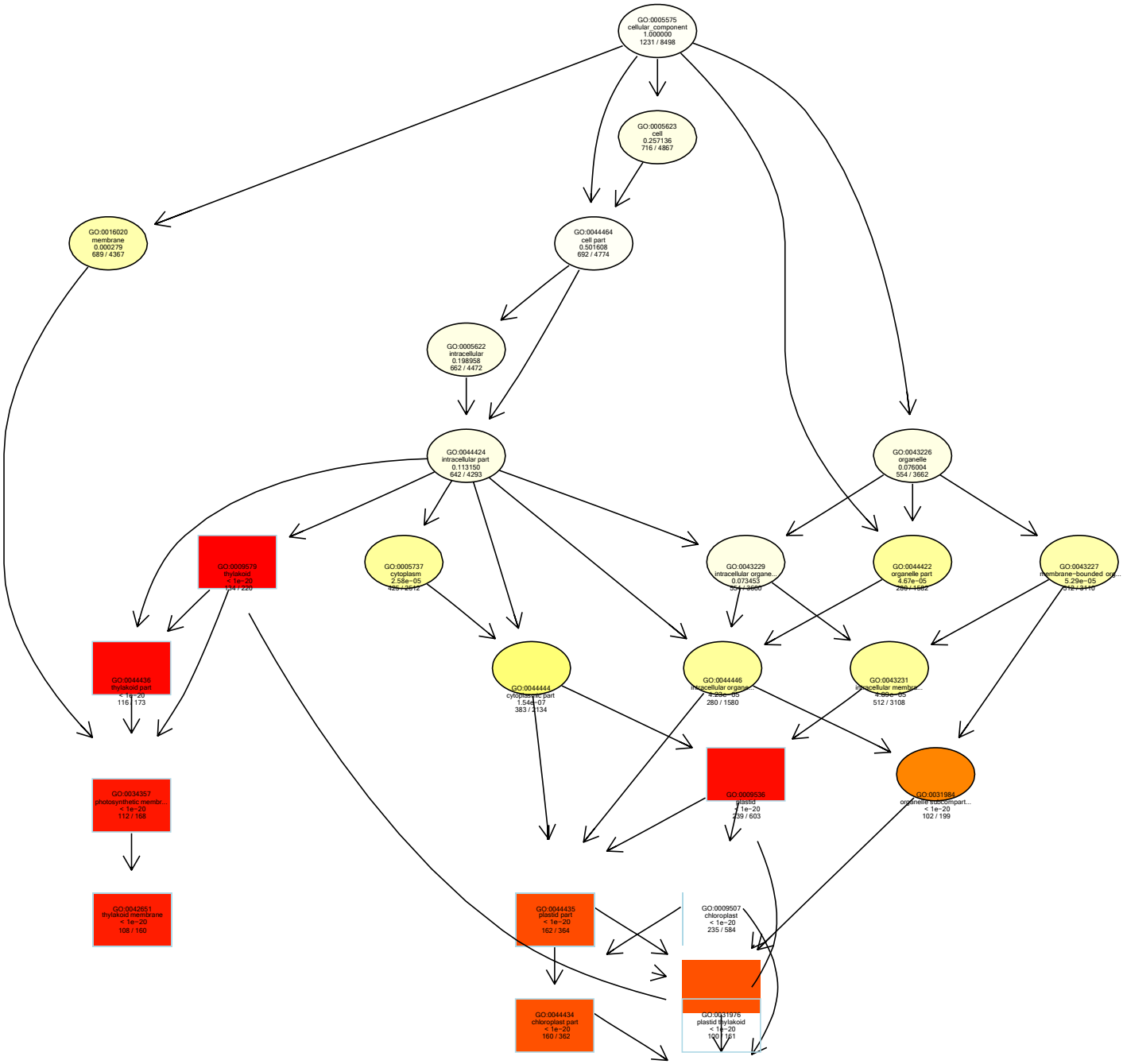

Figure S11: Cellular component where DEGs were found, revealed by topGO enrichment analysis. The DEGs were between etiolated and green wheat seedlings on vernalization 40 d.

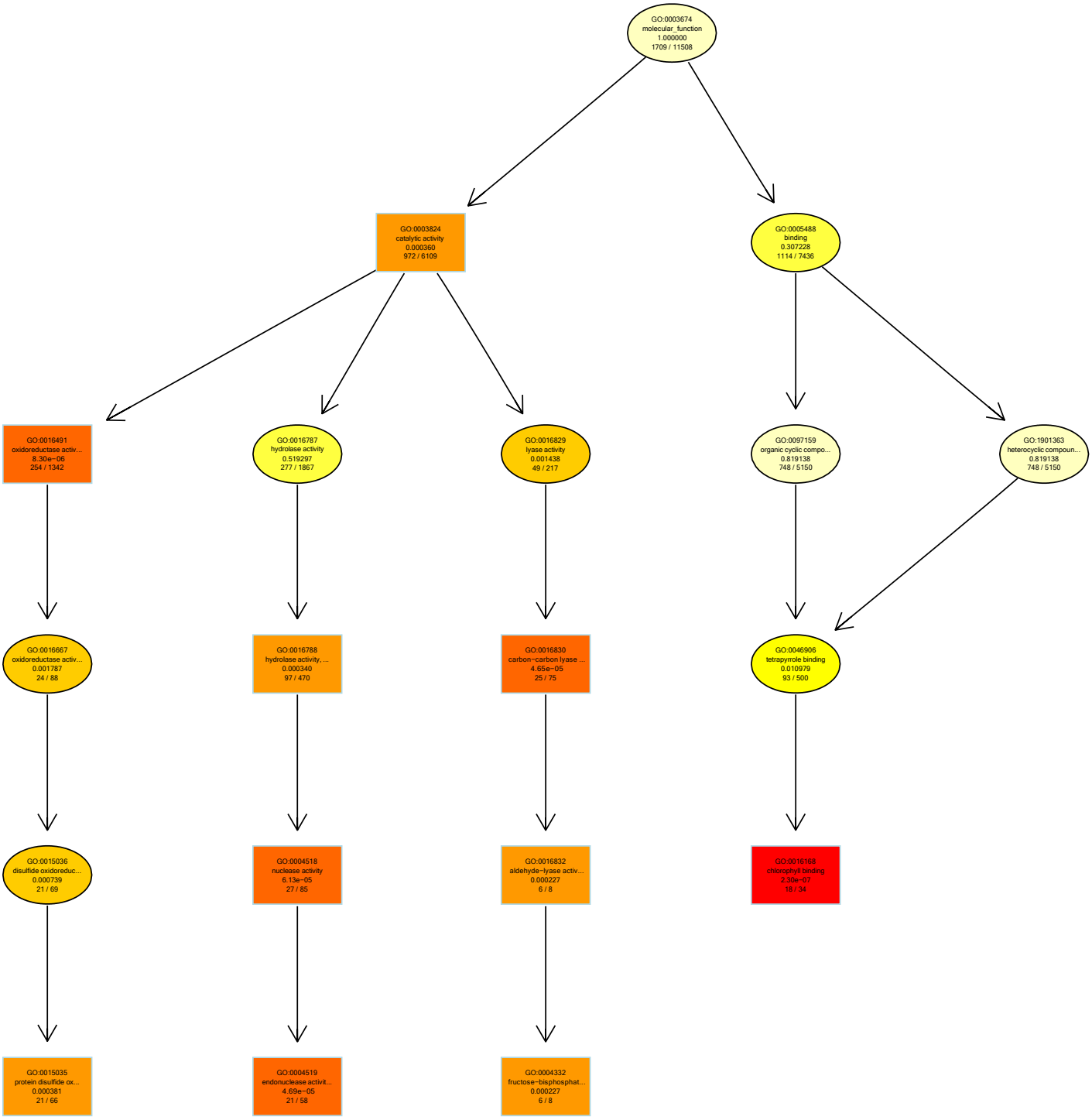

Figure S12: Molecular functions of DEGs between etiolated and green wheat seedlings on vernalization 40 d, revealed by topGO enrichment analysis.