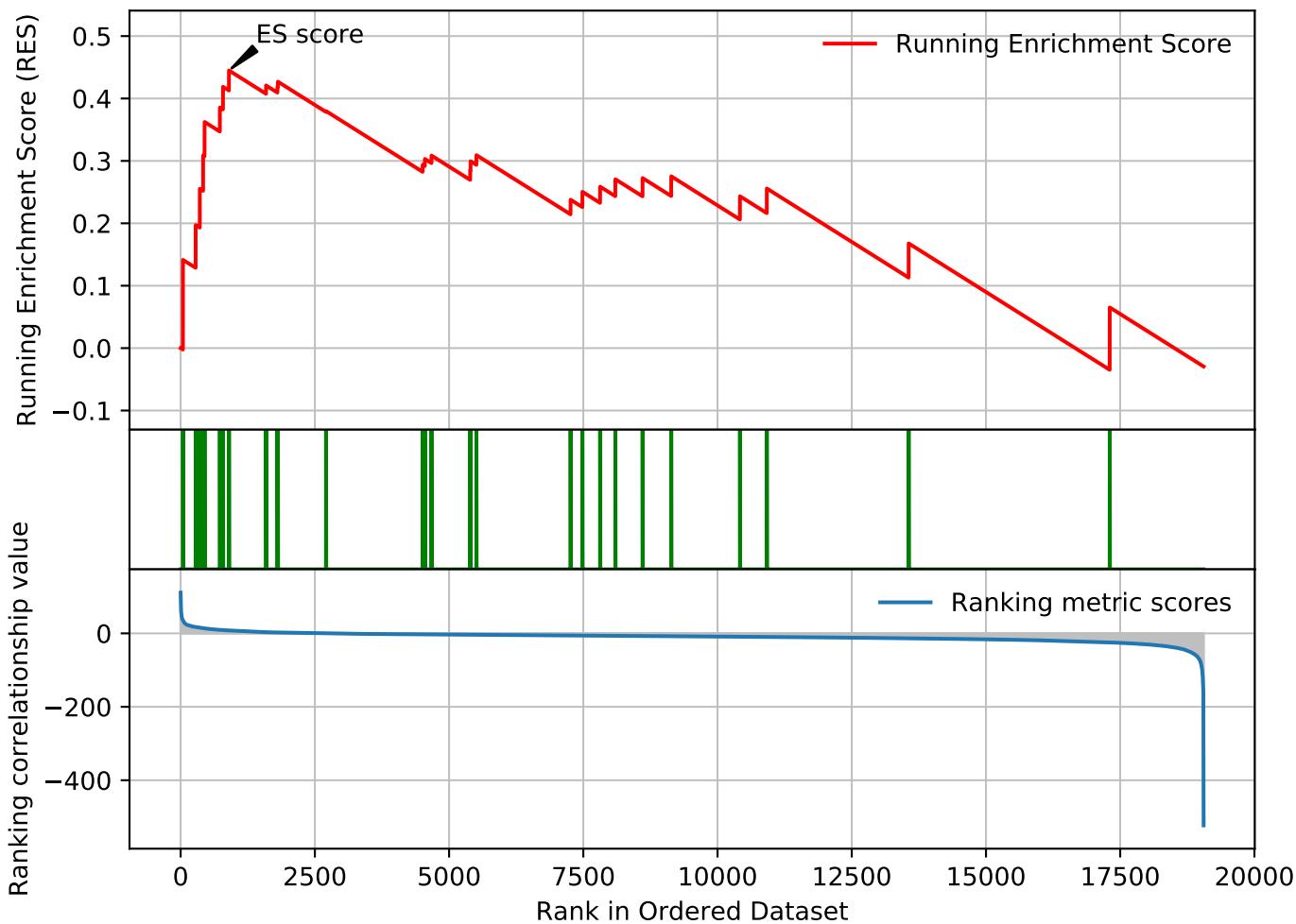
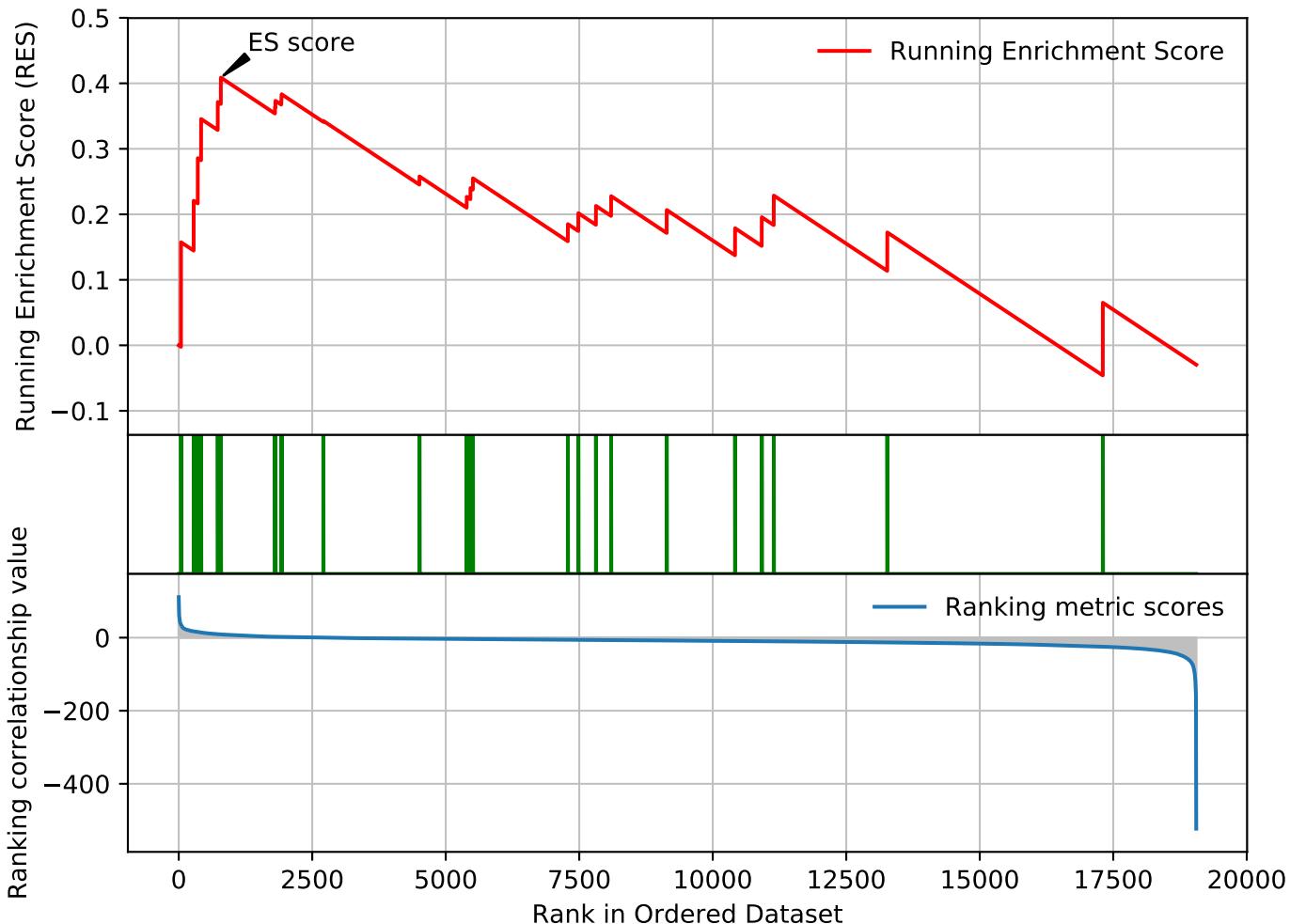


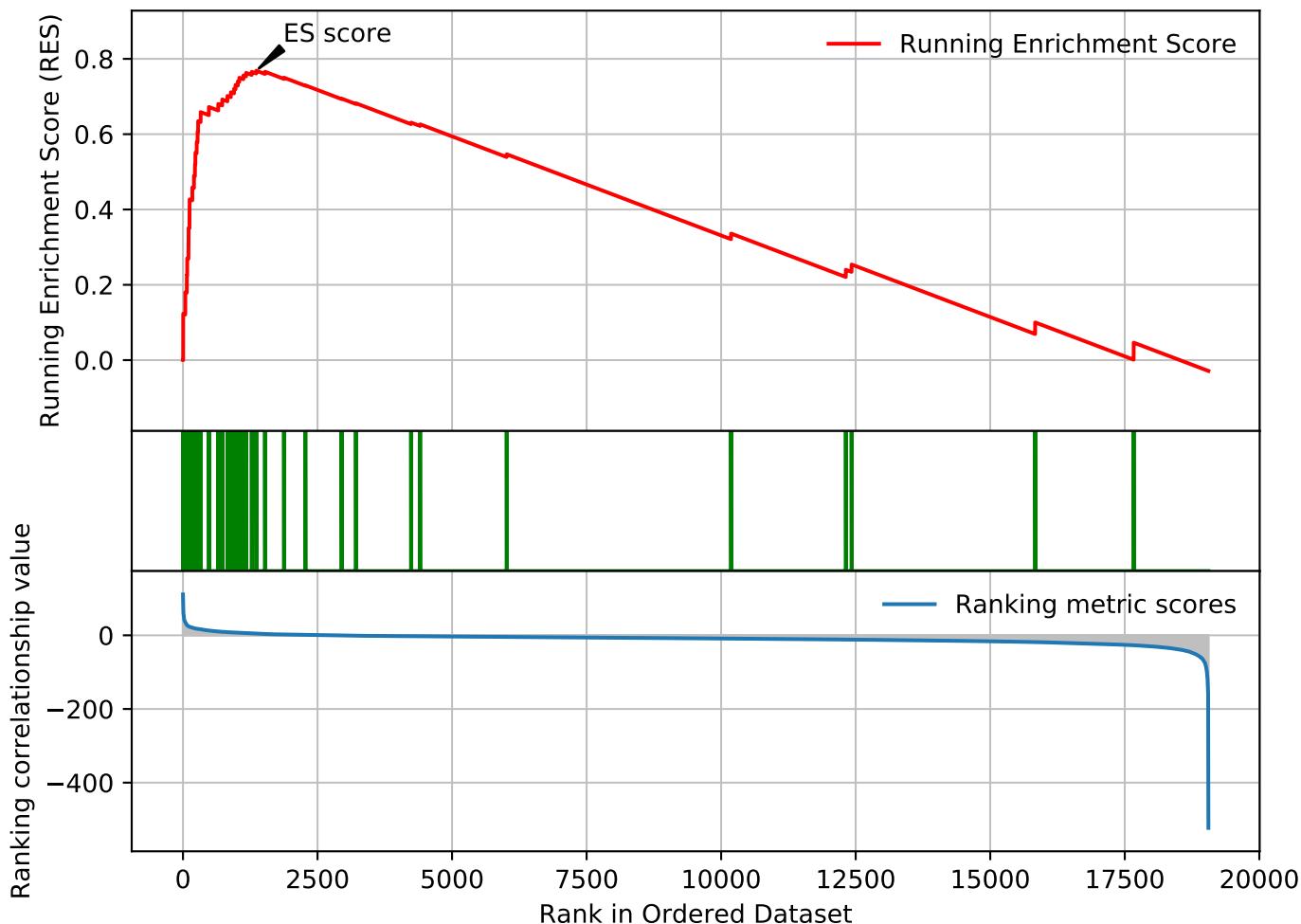
shoot ath00062 (Fatty acid elongation) GSEA_Result



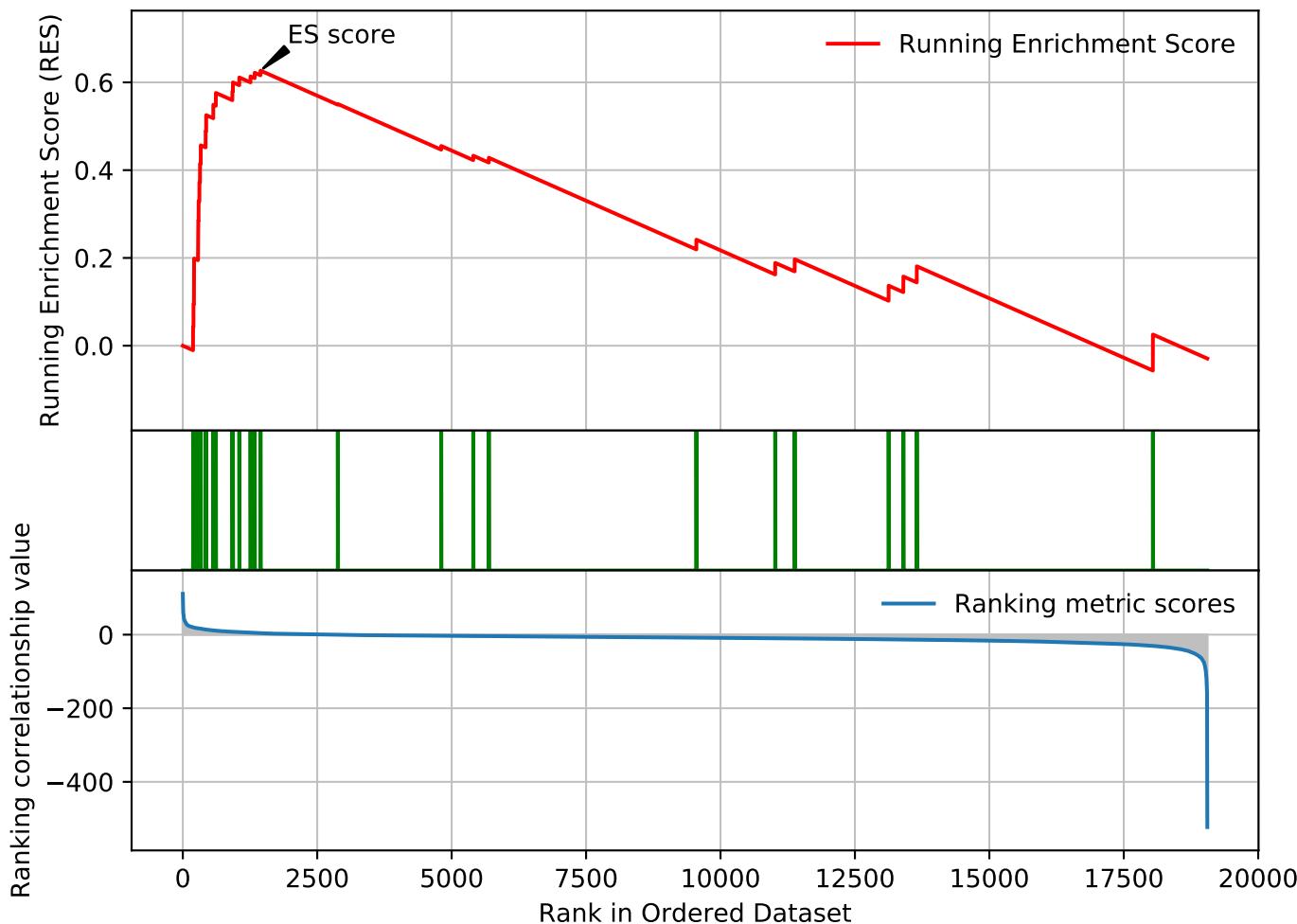
shoot PWY-5080 (very long chain fatty acid biosynthesis I) GSEA_Result



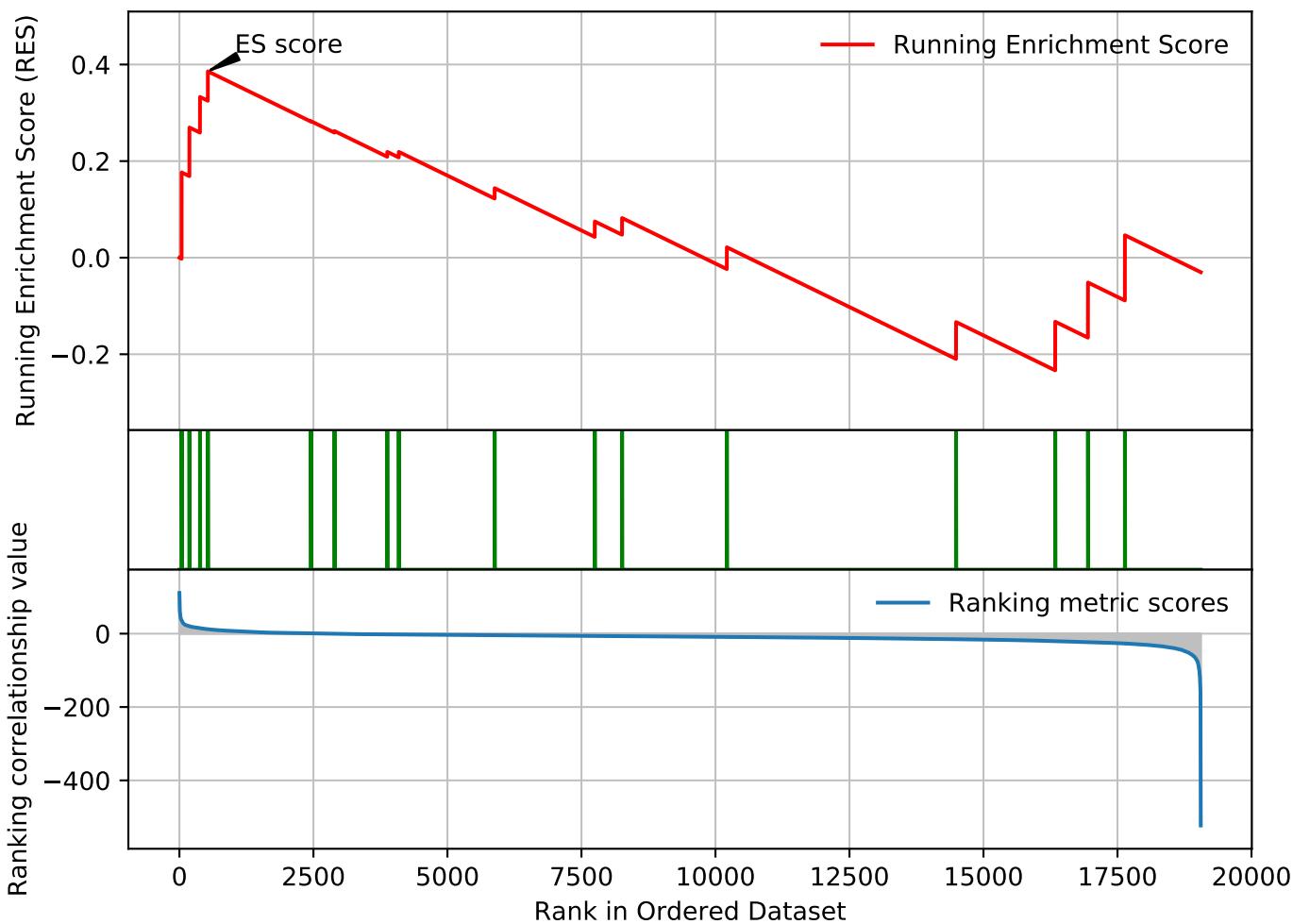
shoot ath00860 (Porphyrin and chlorophyll metabolism) GSEA_Result



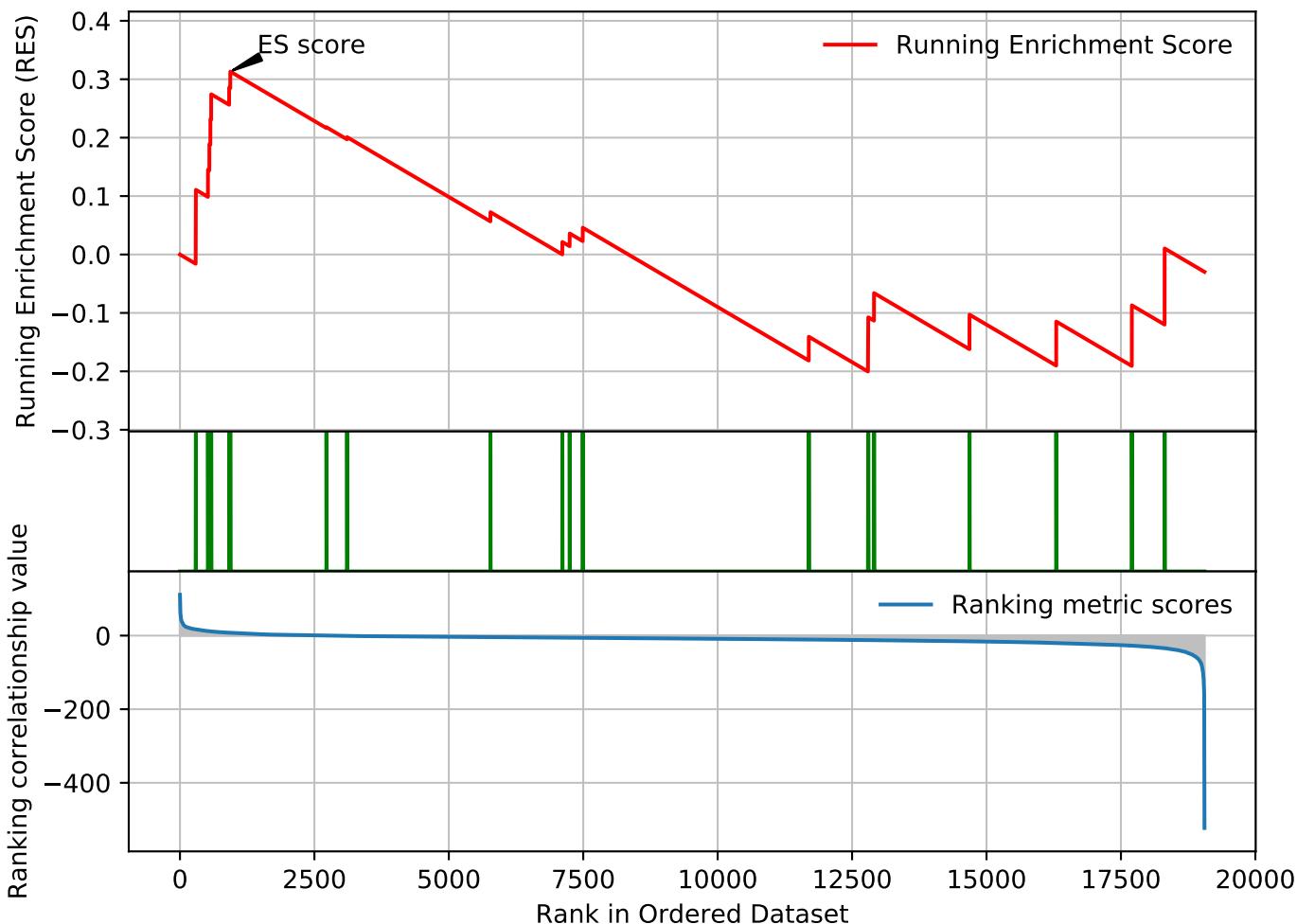
shoot PHOTOALL-PWY (oxygenic photosynthesis) GSEA_Result



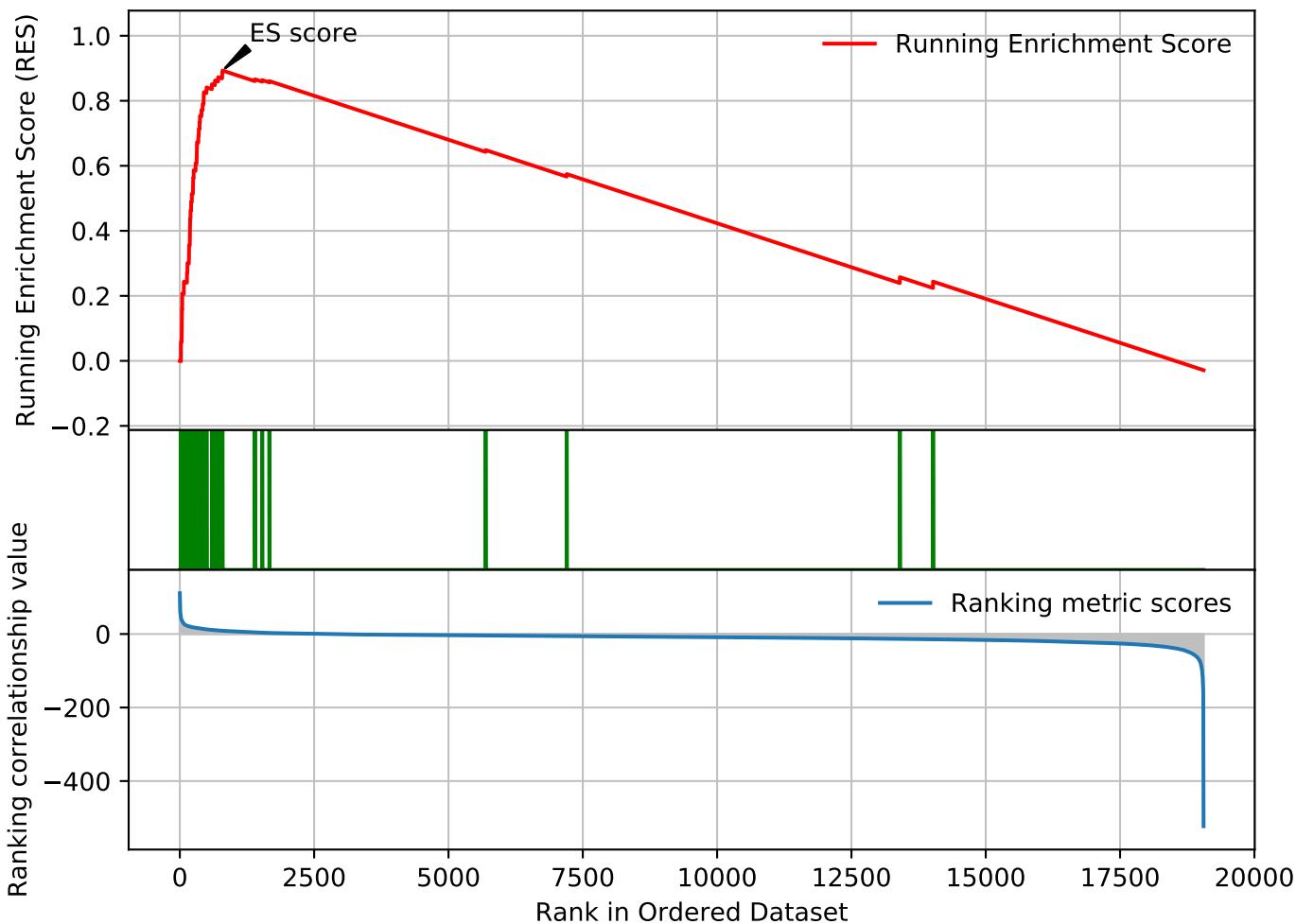
shoot PWY-622 (starch biosynthesis) GSEA_Result



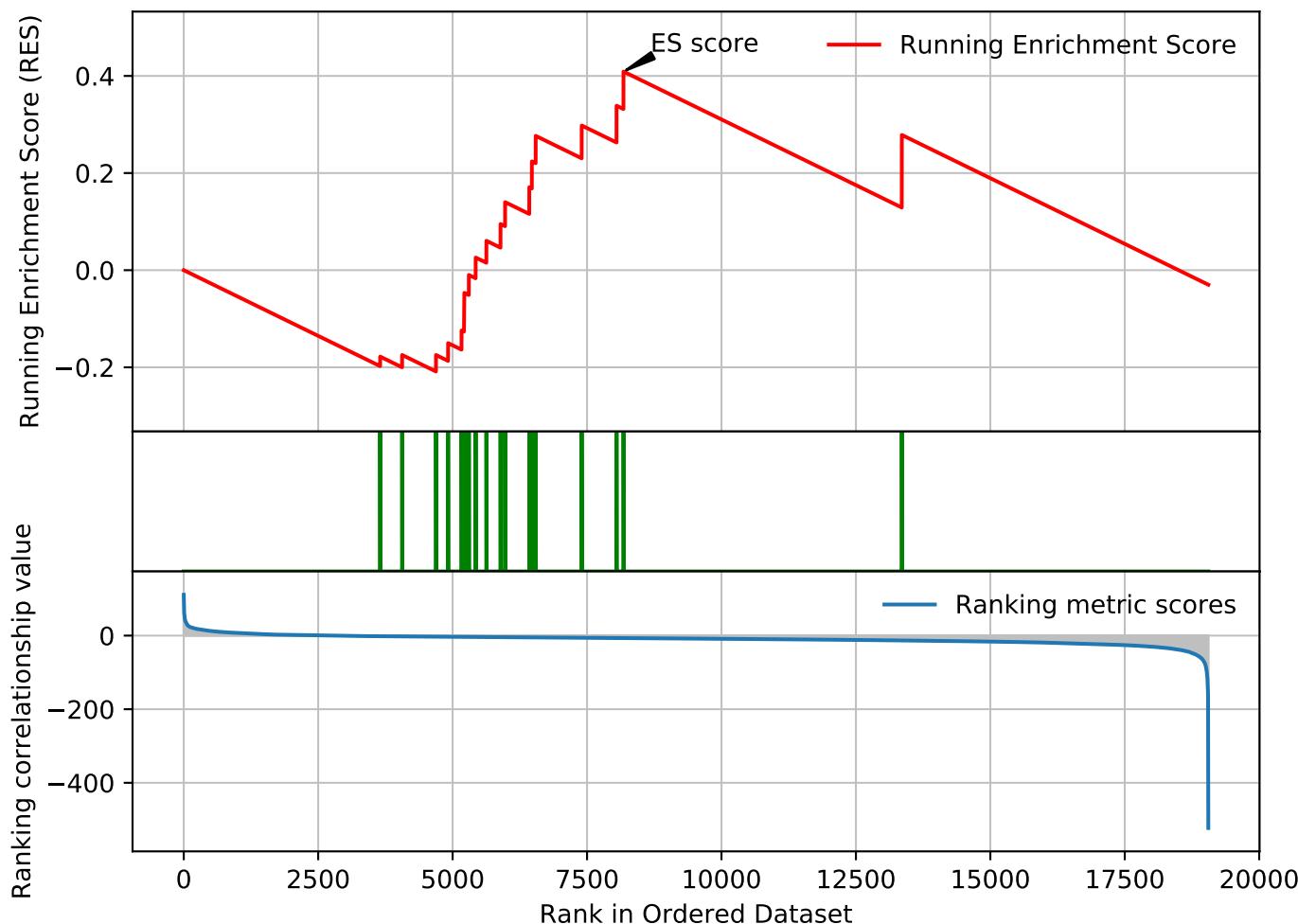
mot R-ATH-389661 (Glyoxylate metabolism and glycine degradation) GSEA_Results



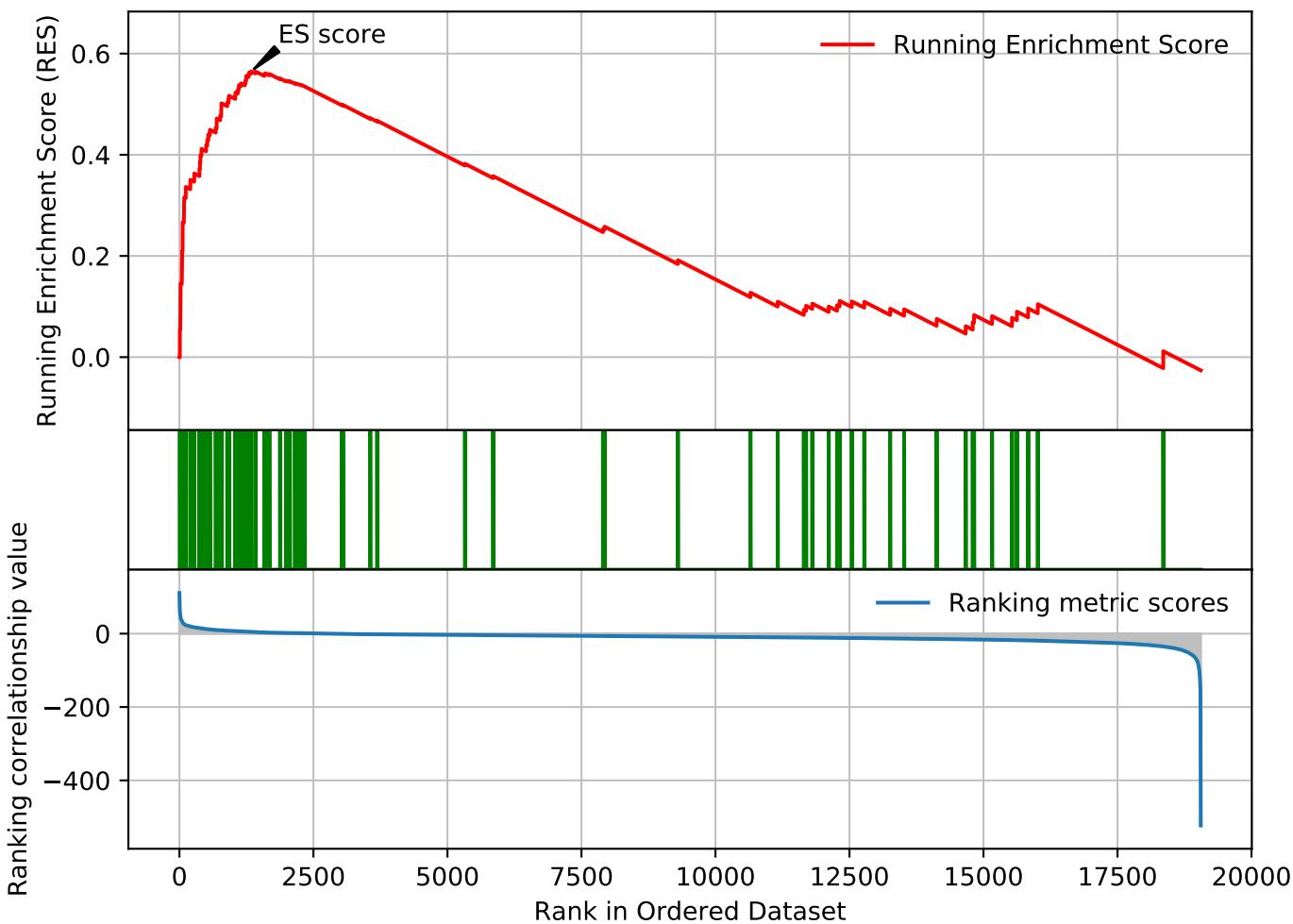
shoot ath00195 (Photosynthesis) GSEA_Result



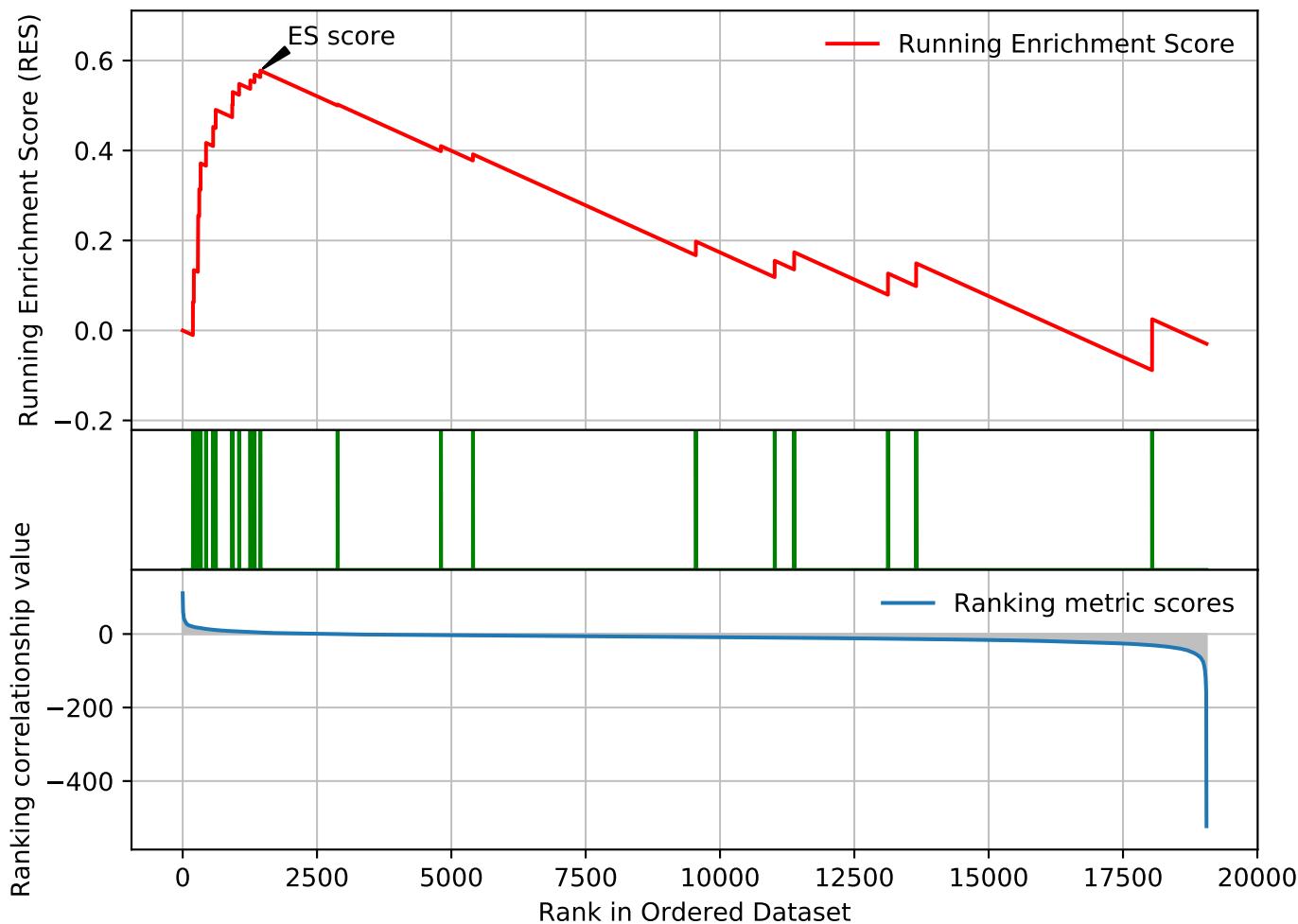
shoot R-ATH-68867 (Assembly of the pre-replicative complex) GSEA_Result



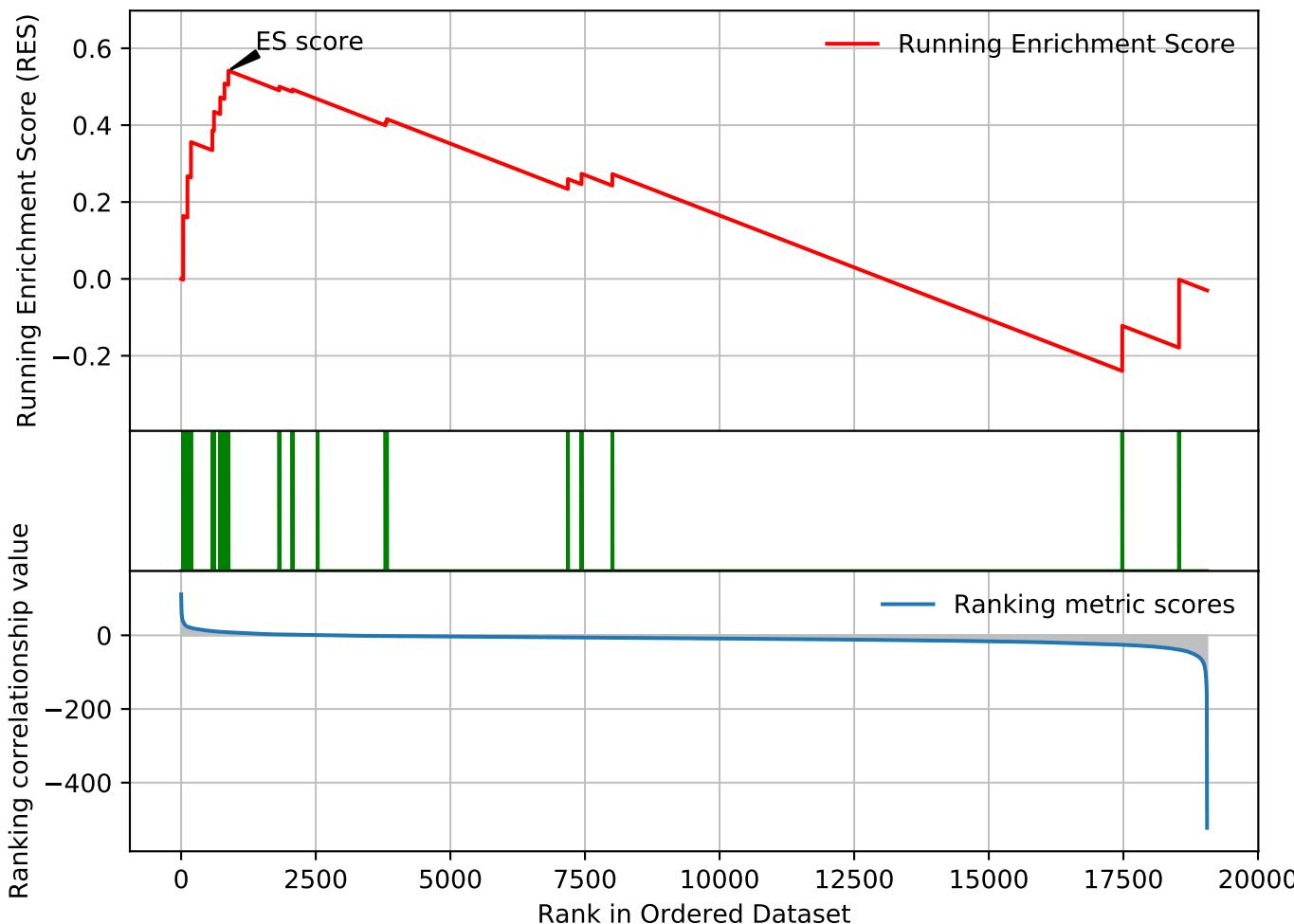
shoot ath00970 (Aminoacyl-tRNA biosynthesis) GSEA_Result



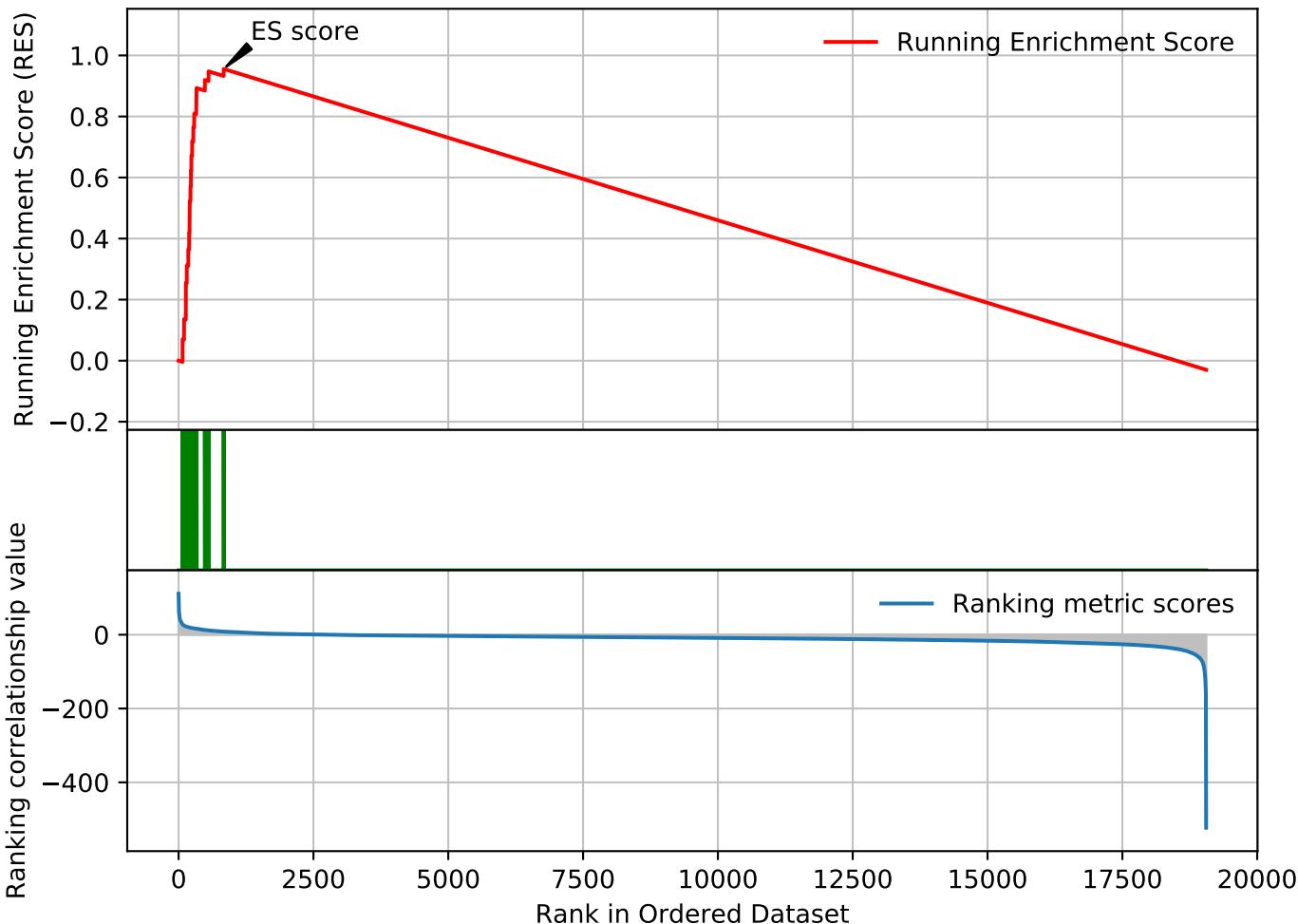
shoot CALVIN-PWY (Calvin-Benson-Bassham cycle) GSEA_Result

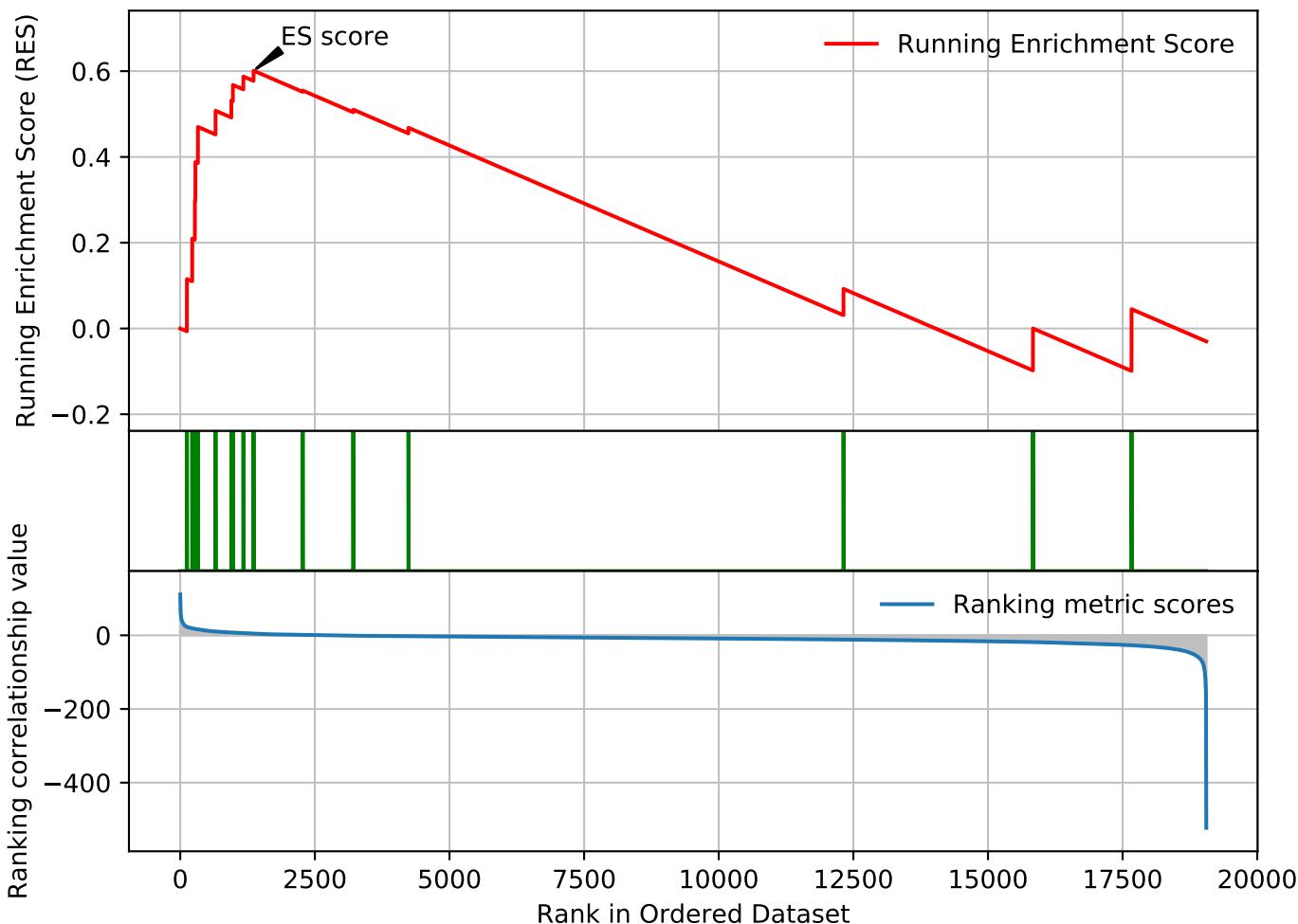


shoot PWY-7450 (anthocyanidin modification (Arabidopsis)) GSEA_Result

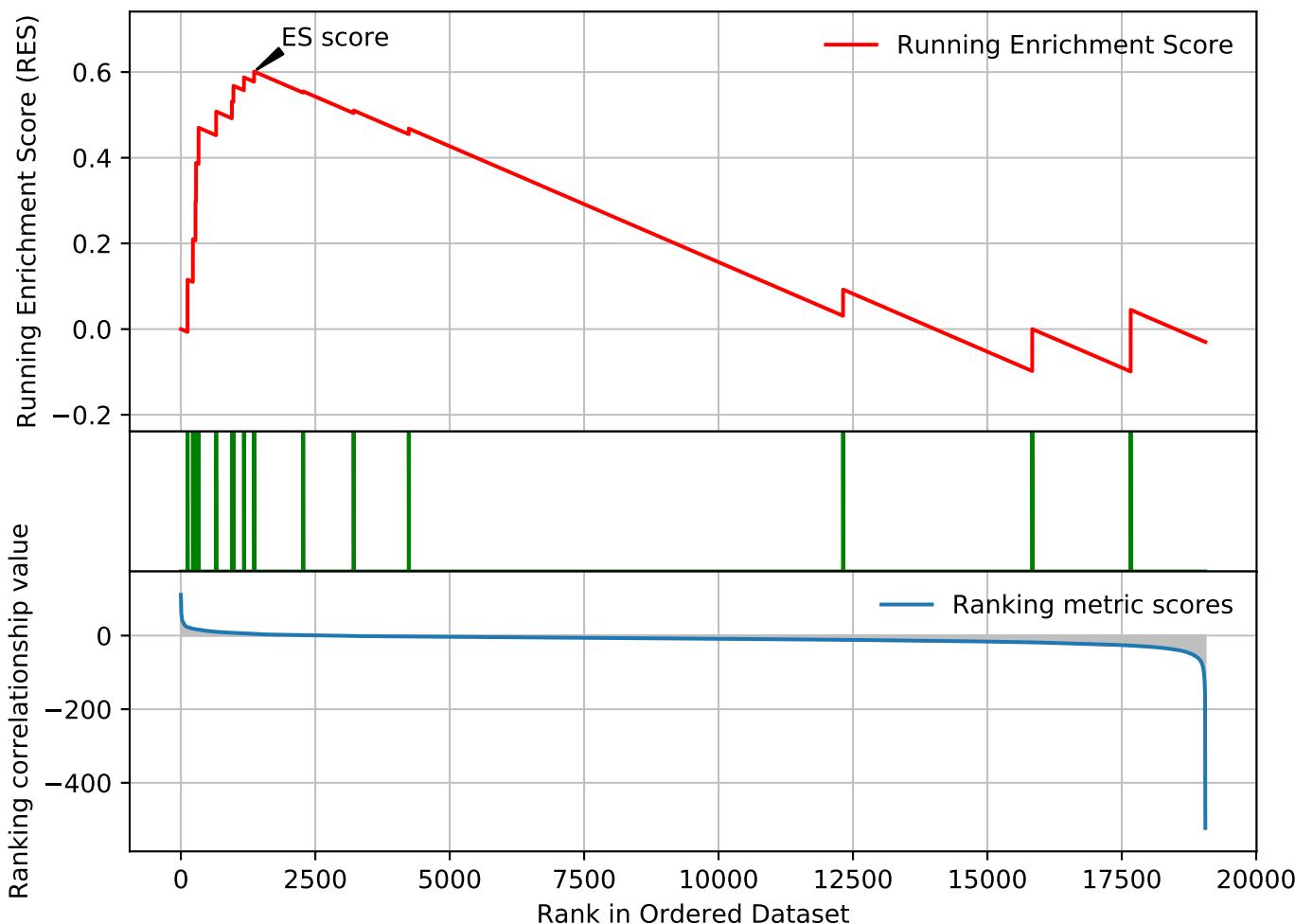


shoot ath00196 (Photosynthesis - antenna proteins) GSEA_Result

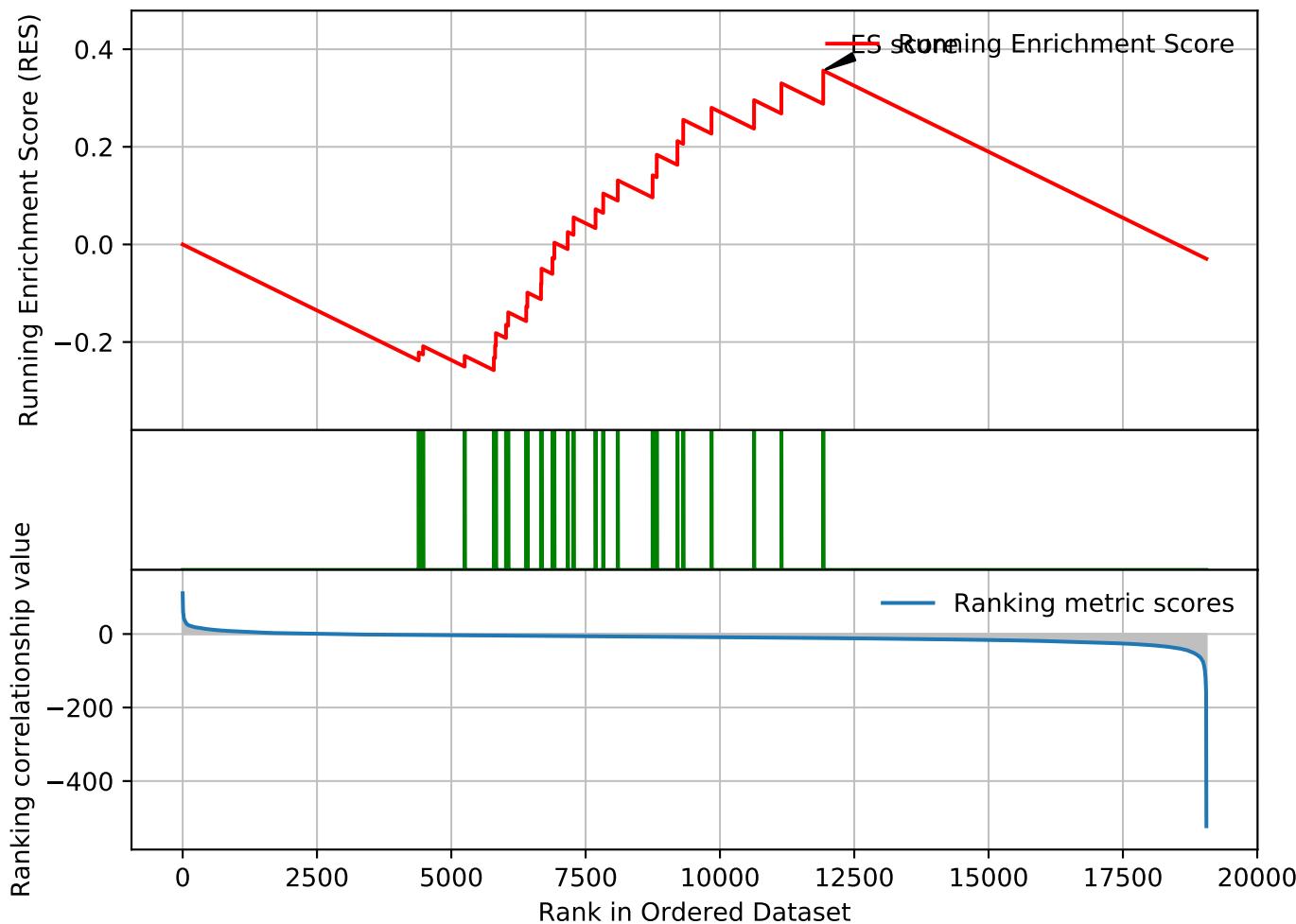




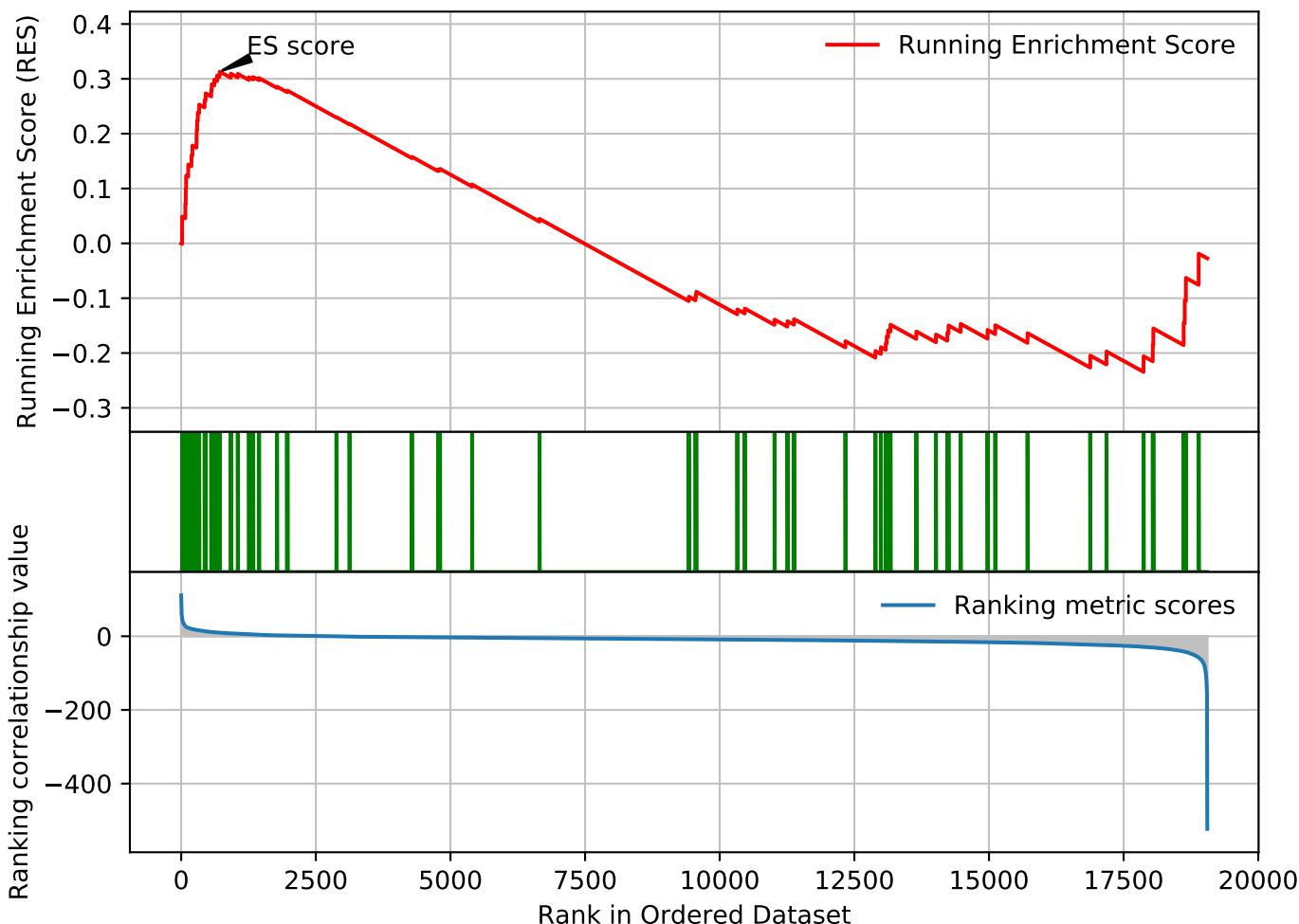
root PWY-5918 (superpathay of heme biosynthesis from glutamate) GSEA_Results



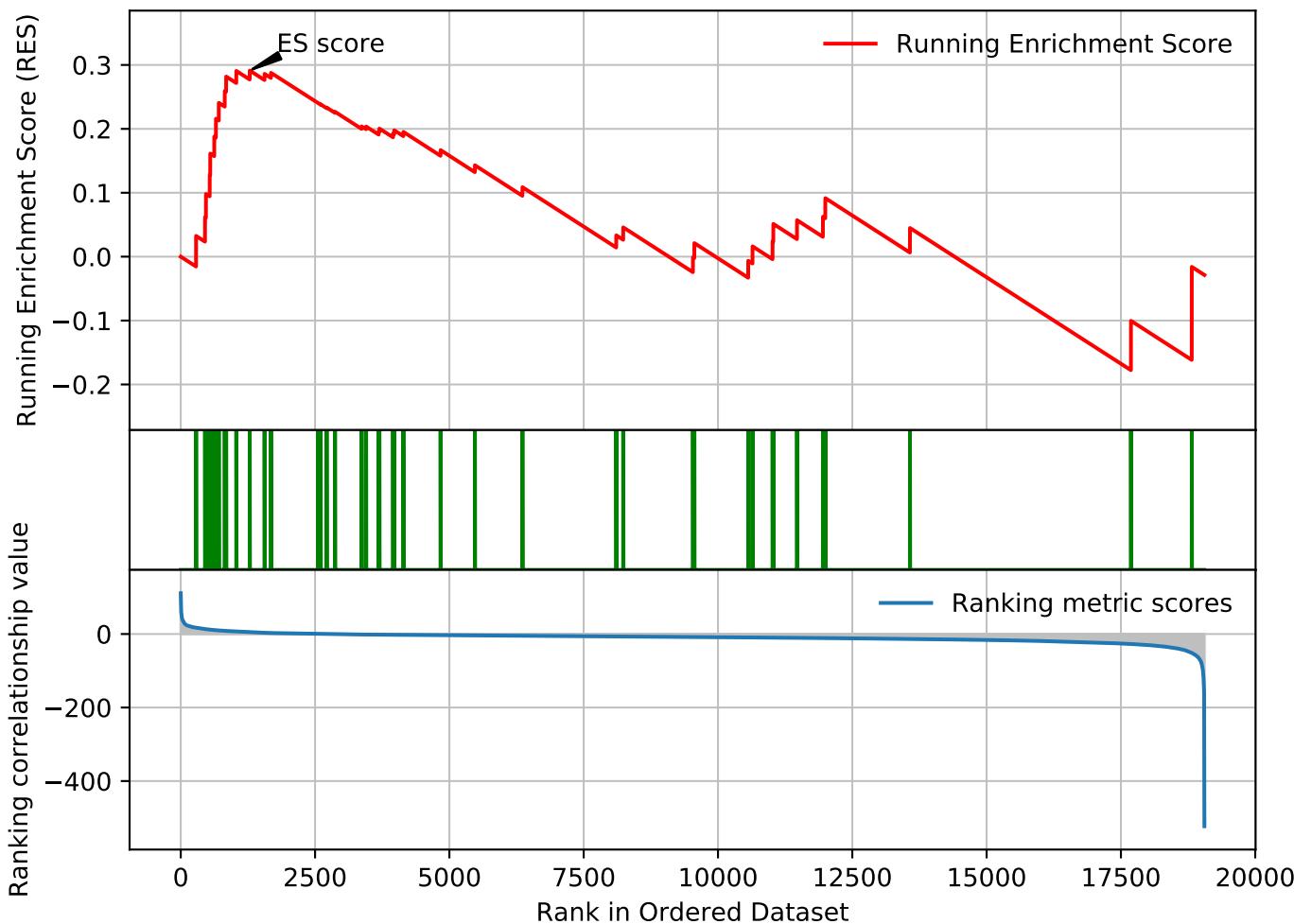
shoot R-ATH-69190 (DNA strand elongation) GSEA_Result



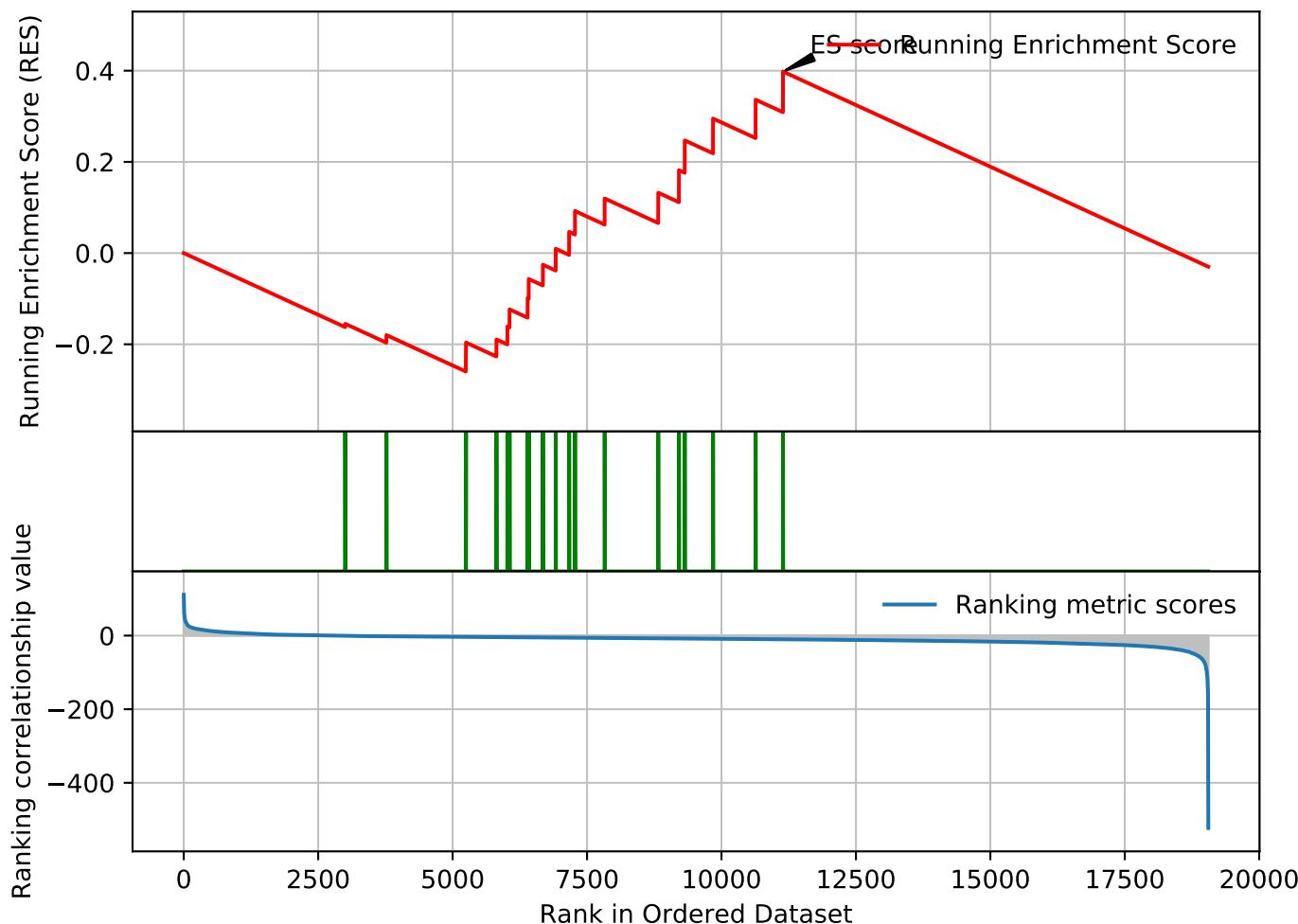
shoot ath00710 (Carbon fixation in photosynthetic organisms) GSEA_Result



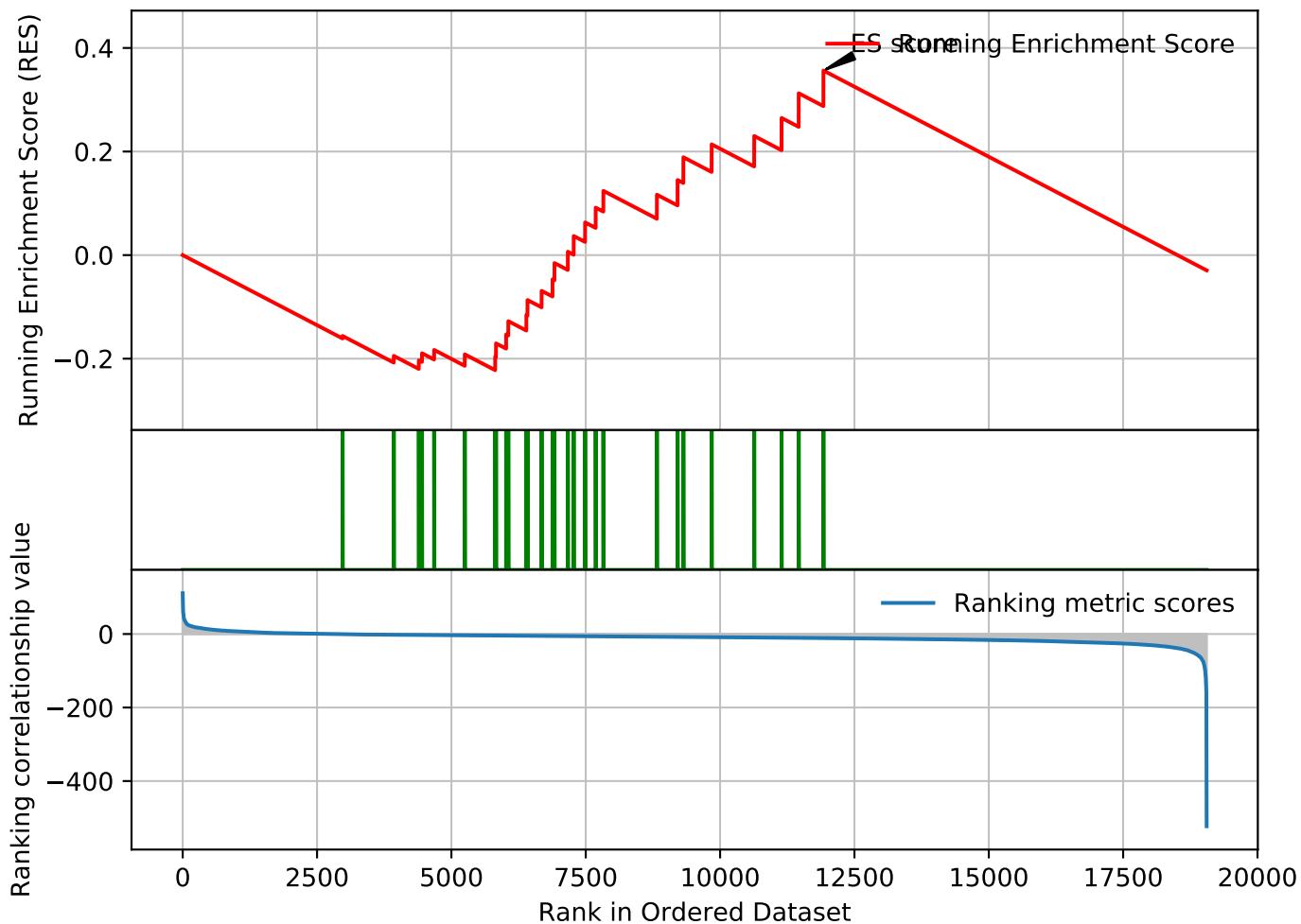
shoot LIPAS-PWY (triacylglycerol degradation) GSEA_Result



R-ATH-5651801 (PCNA-Dependent Long Patch Base Excision Repair) GSEA_F



shoot R-ATH-180786 (Extension of Telomeres) GSEA_Result



shoot R-ATH-73886 (Chromosome Maintenance) GSEA_Result

