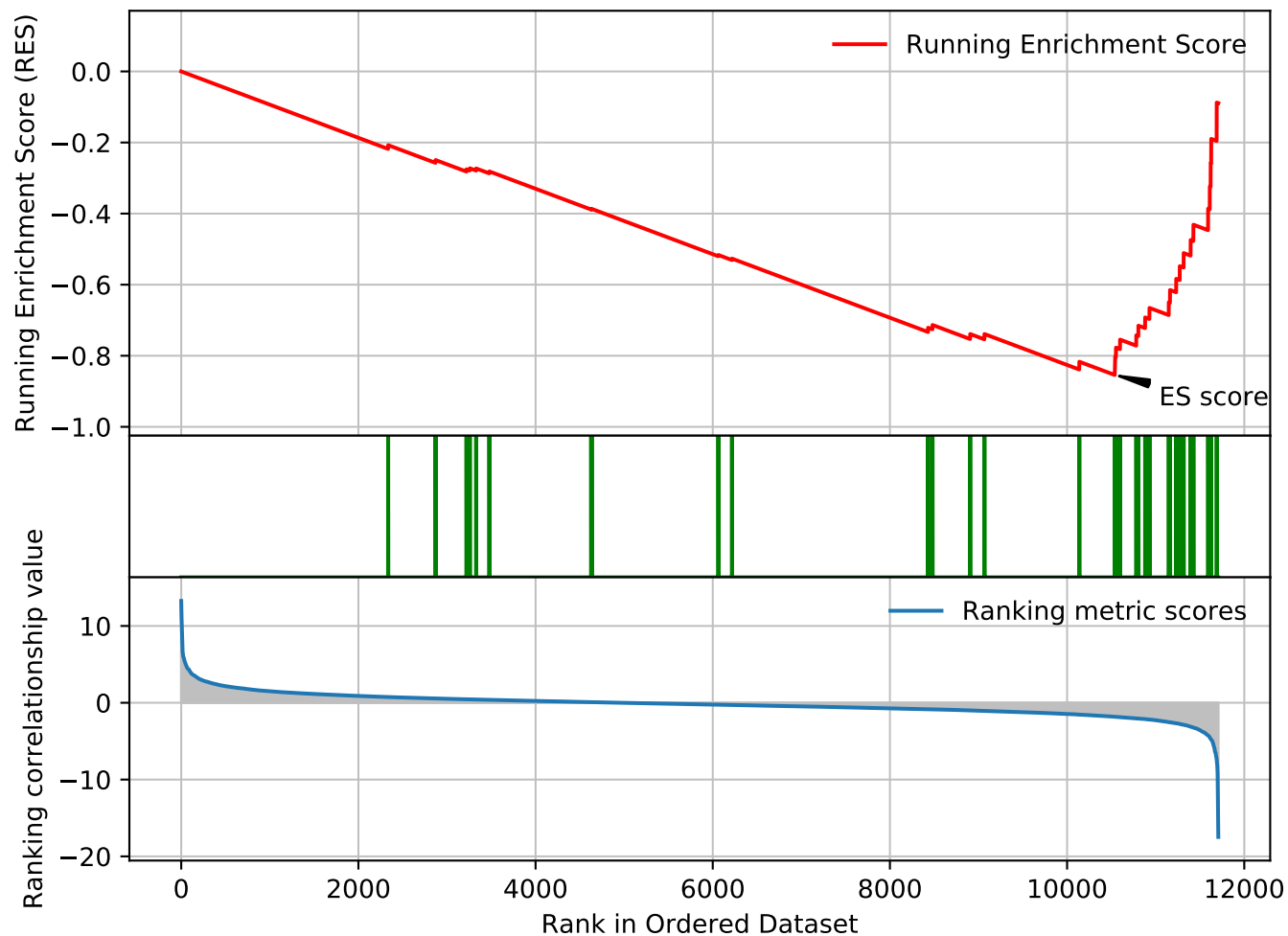
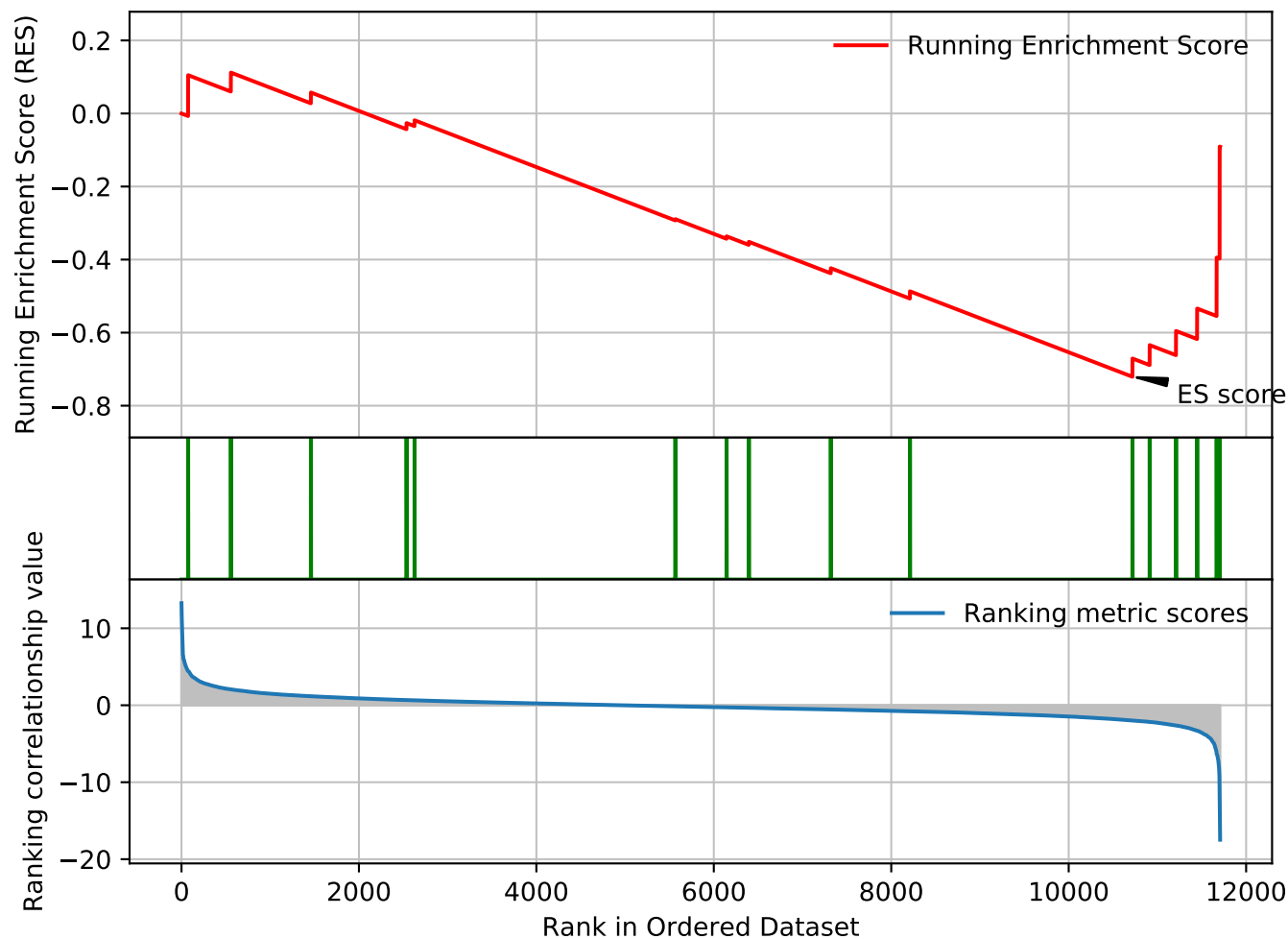
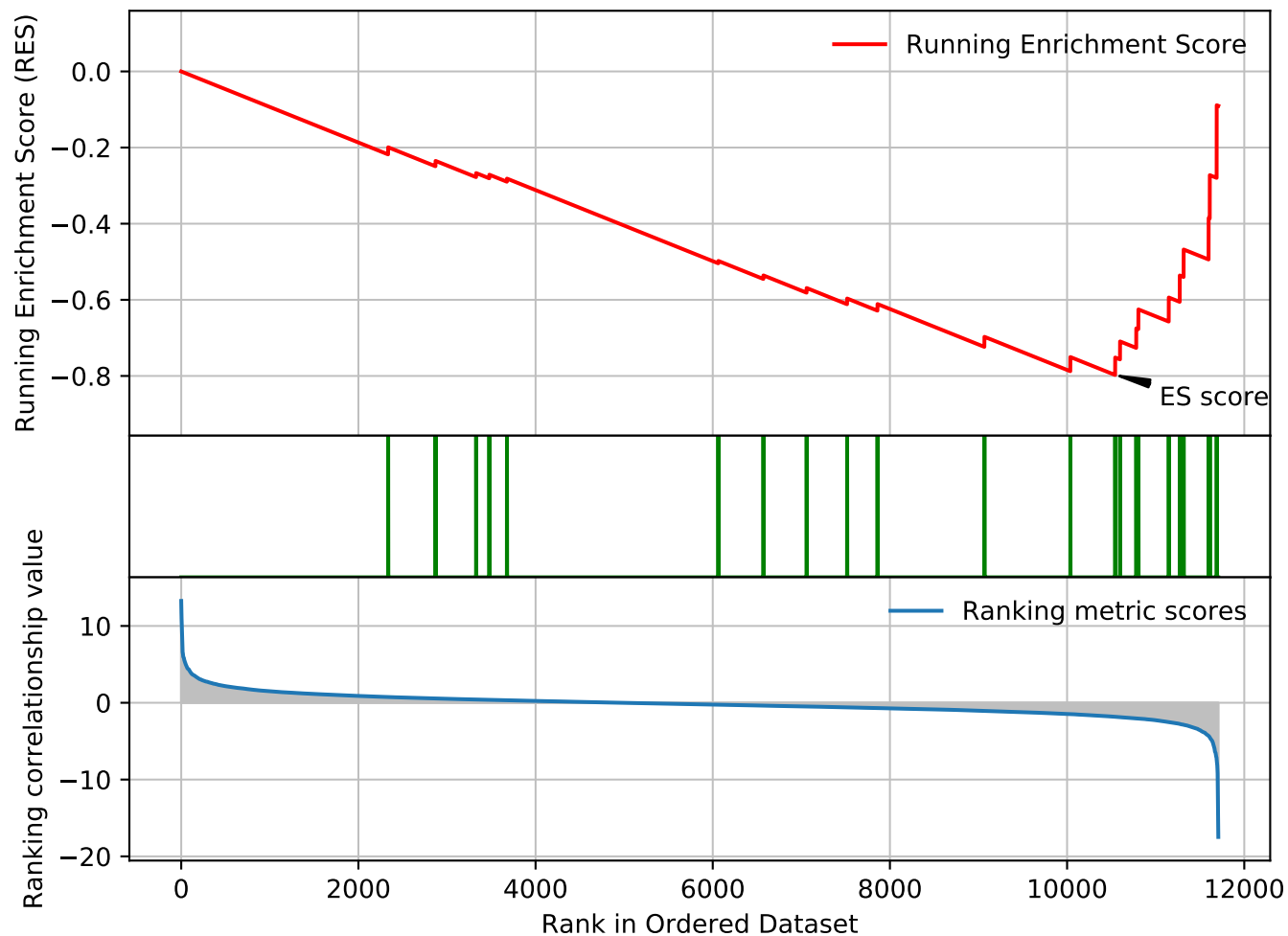


# WT mmu03030 (DNA replication) GSEA\_Result

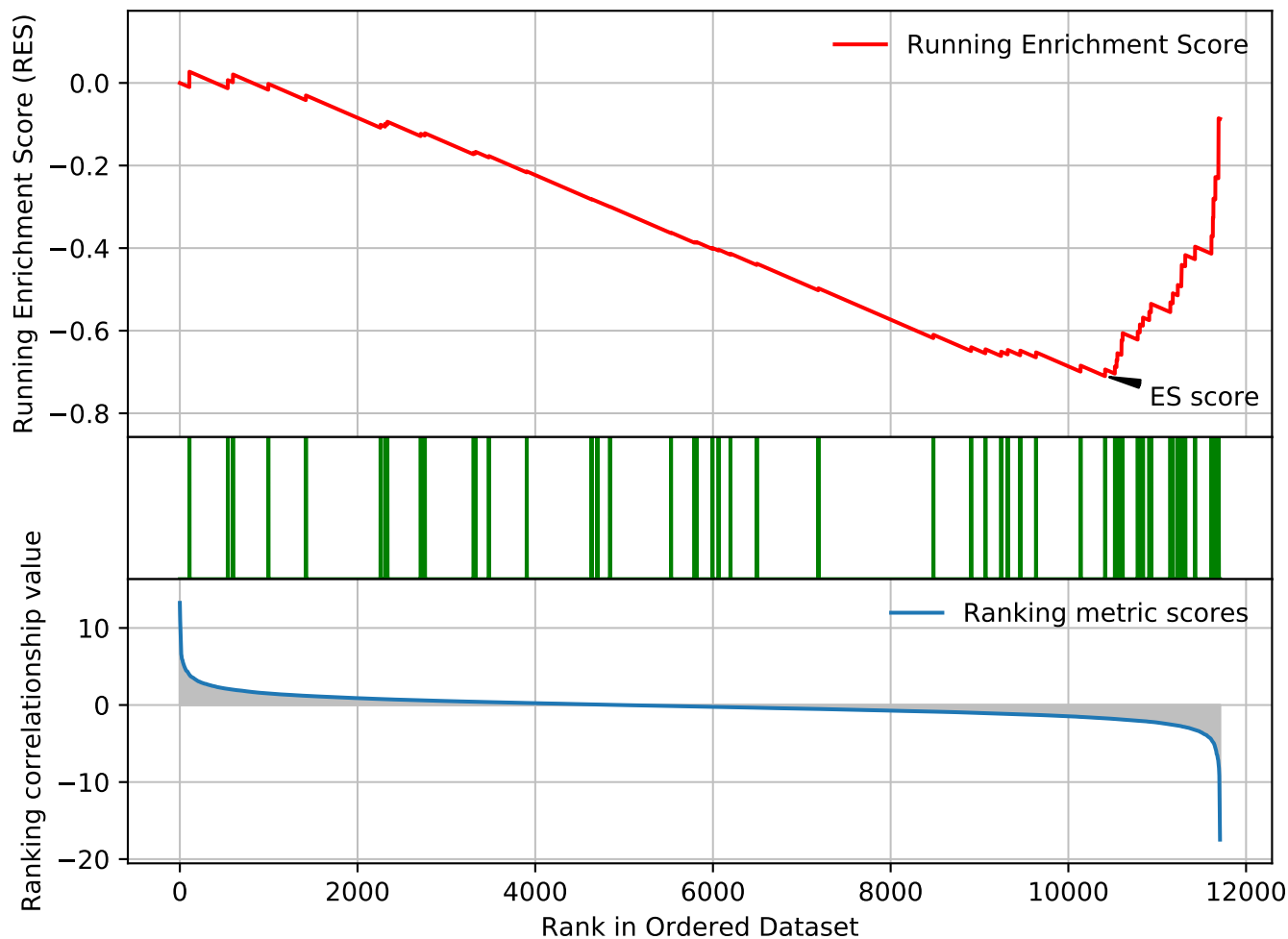




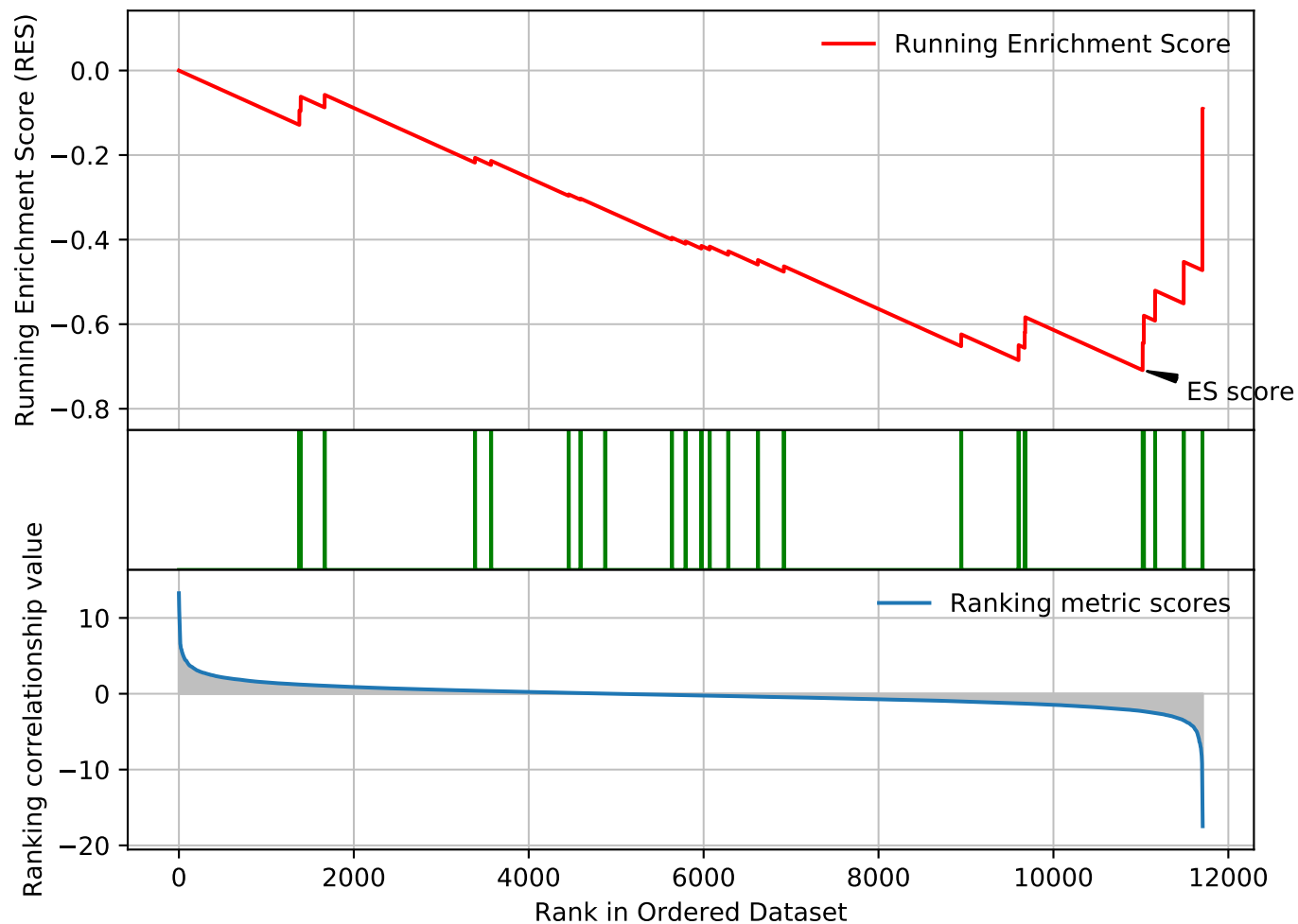
# WT mmu03430 (Mismatch repair) GSEA\_Result



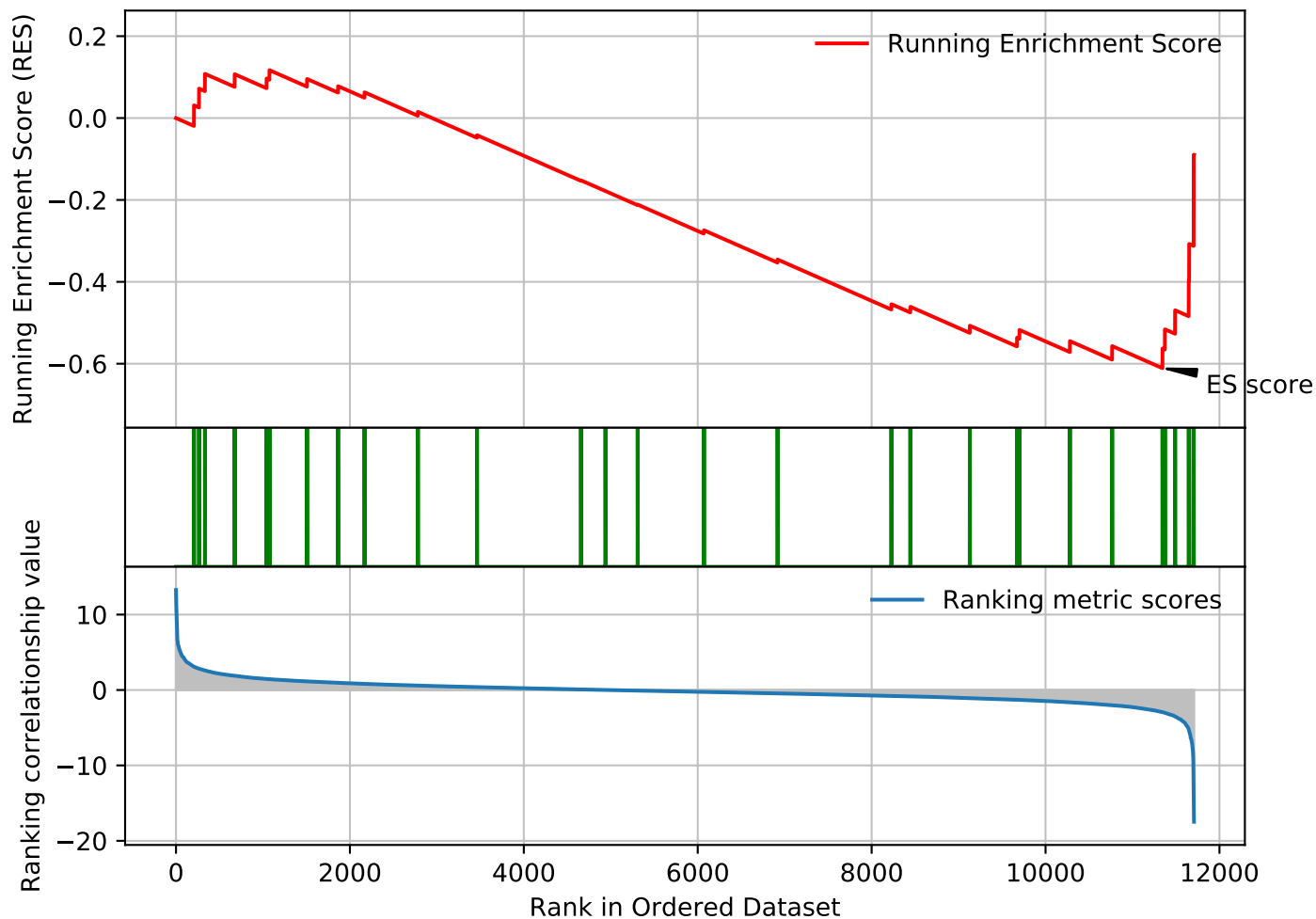
# WT R-MMU-73886 (Chromosome Maintenance) GSEA\_Result



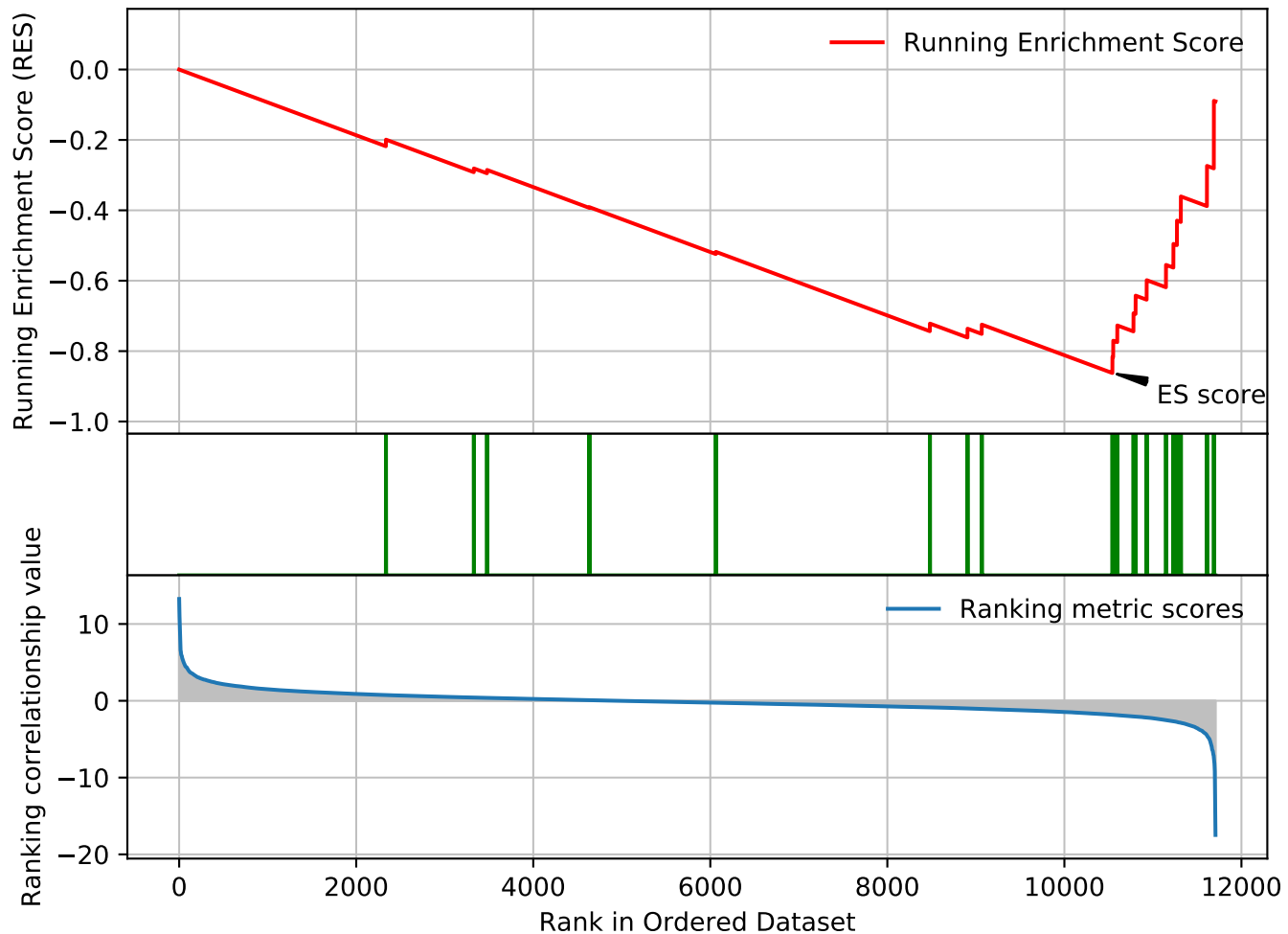
# WT P00020 (FAS signaling pathway) GSEA\_Results



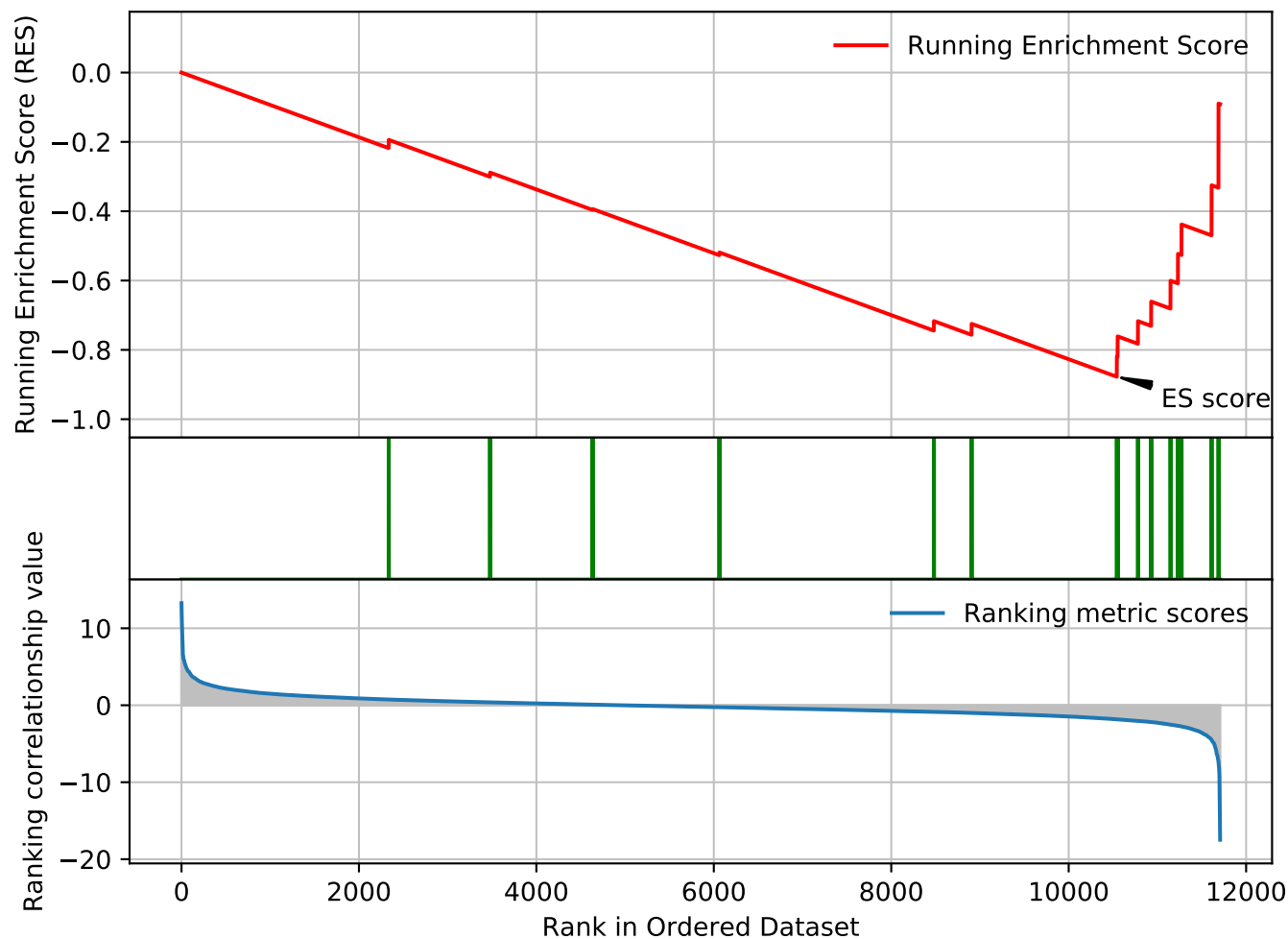
# WT R-MMU-109606 (Intrinsic Pathway for Apoptosis) GSEA\_Results



# WT R-MMU-69186 (Lagging Strand Synthesis) GSEA\_Result

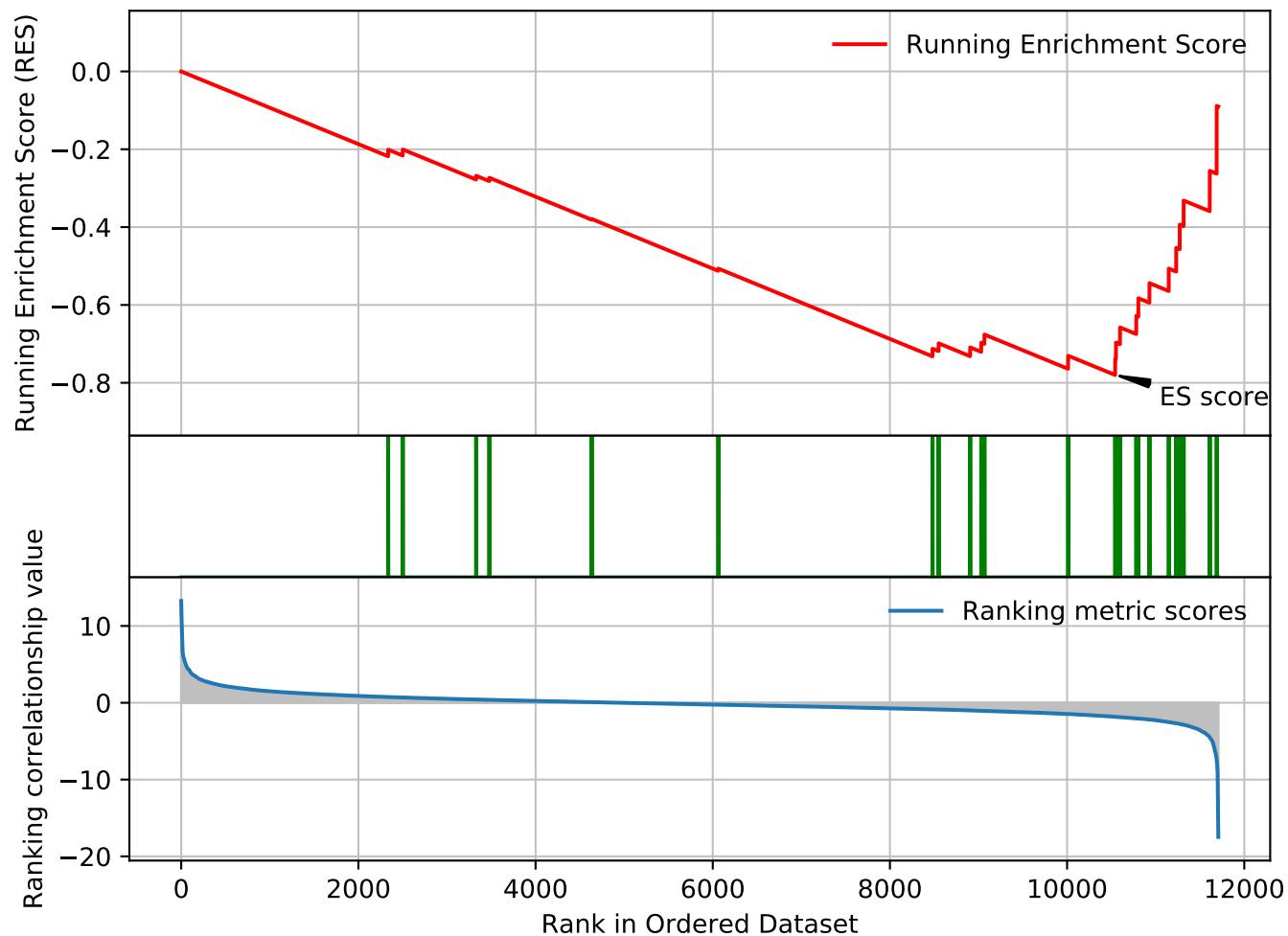


# WT R-MMU-69183 (Processive synthesis on the lagging strand) GSEA\_Result

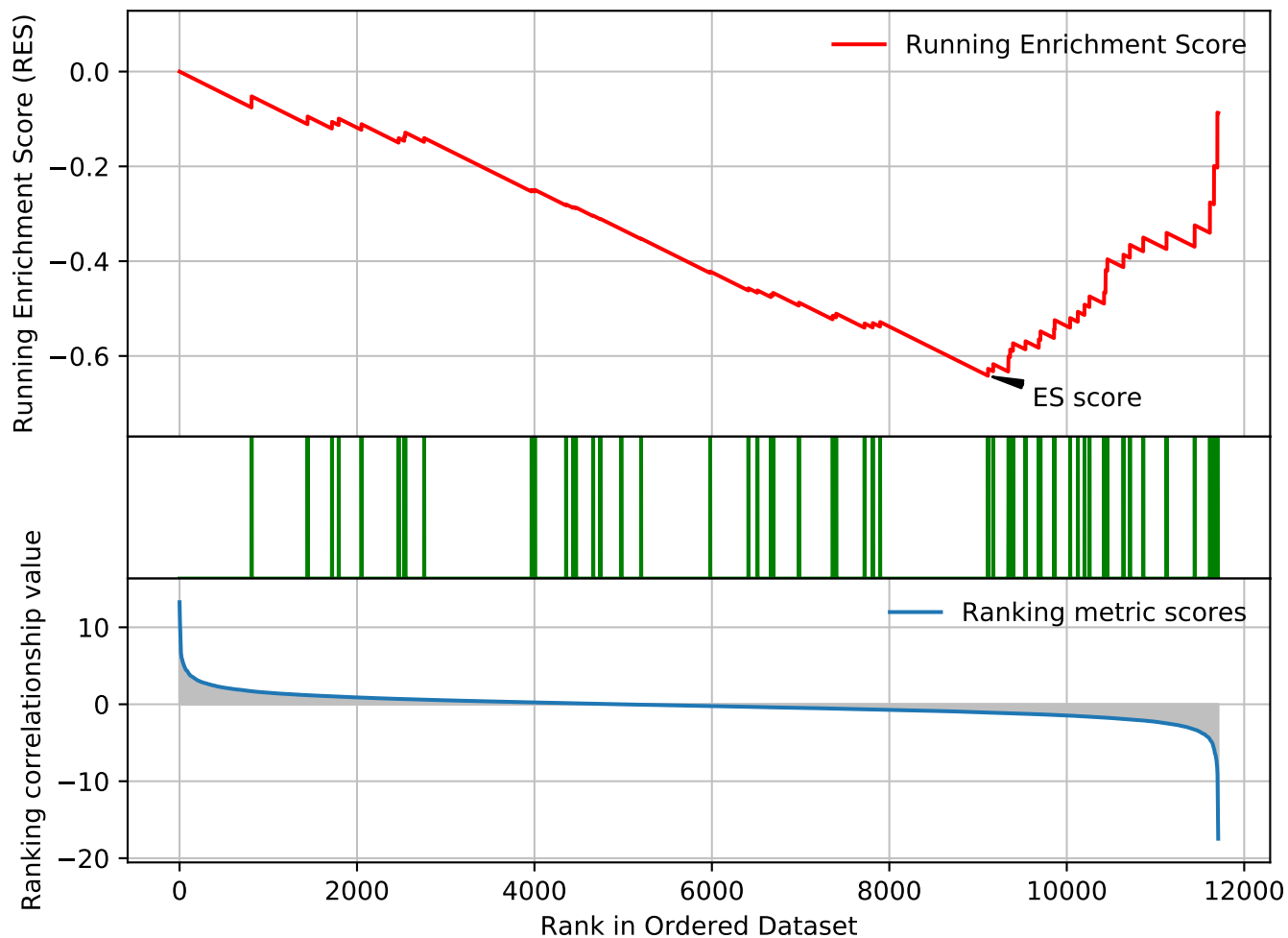




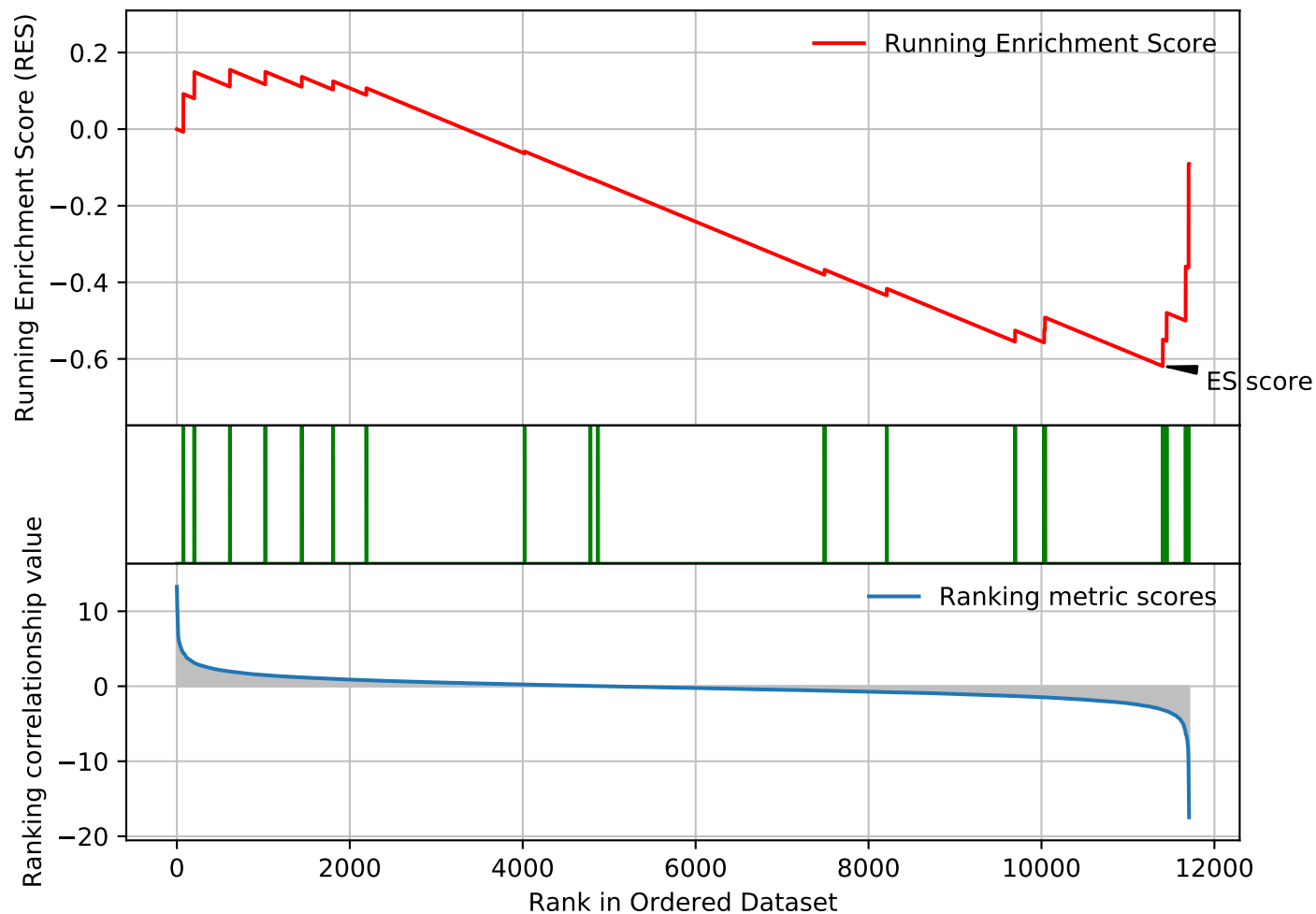
# WT R-MMU-69190 (DNA strand elongation) GSEA\_Result

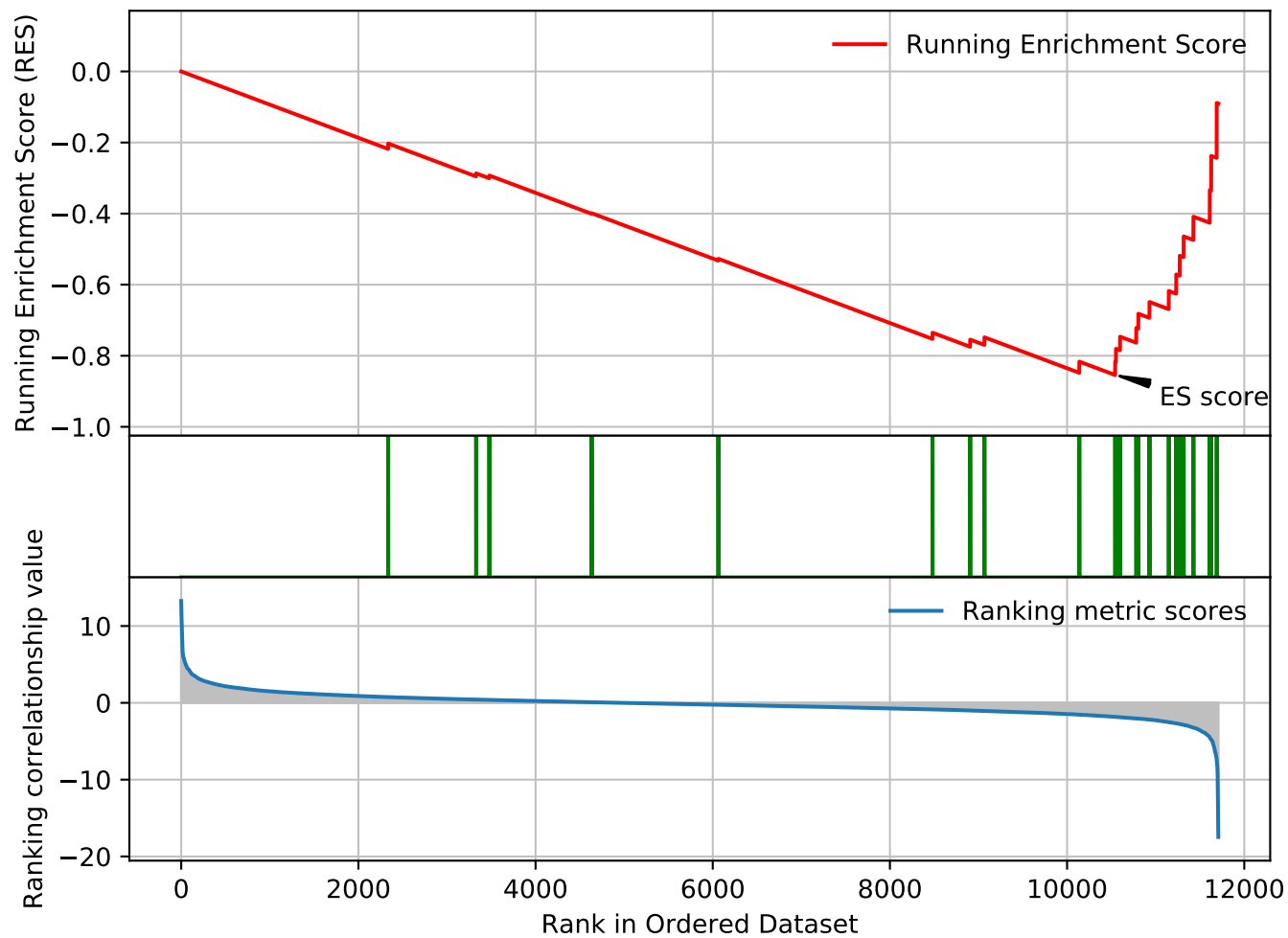


# WT mmu00562 (Inositol phosphate metabolism) GSEA\_Result

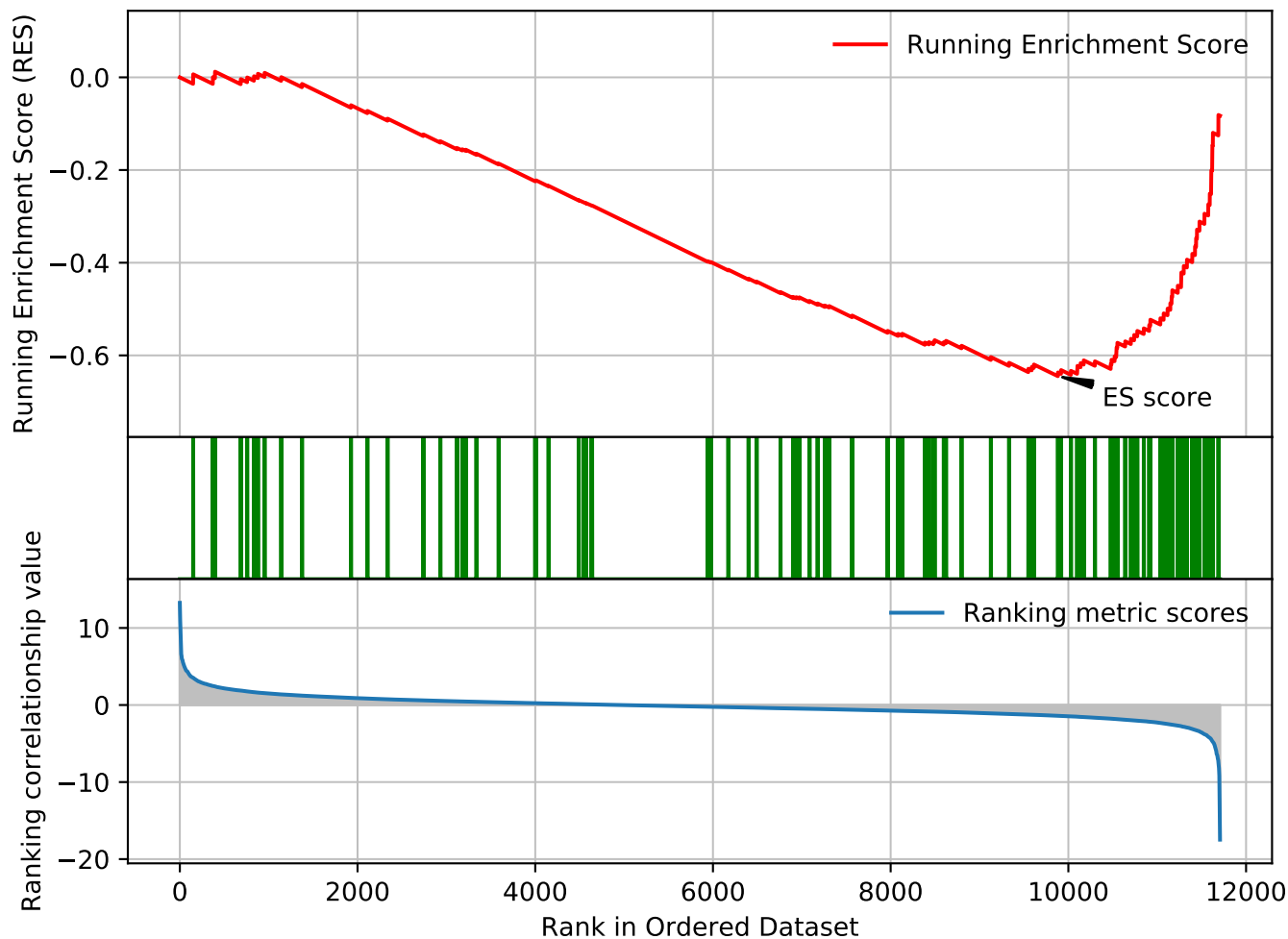


# WT P02738 (De novo purine biosynthesis) GSEA\_Result

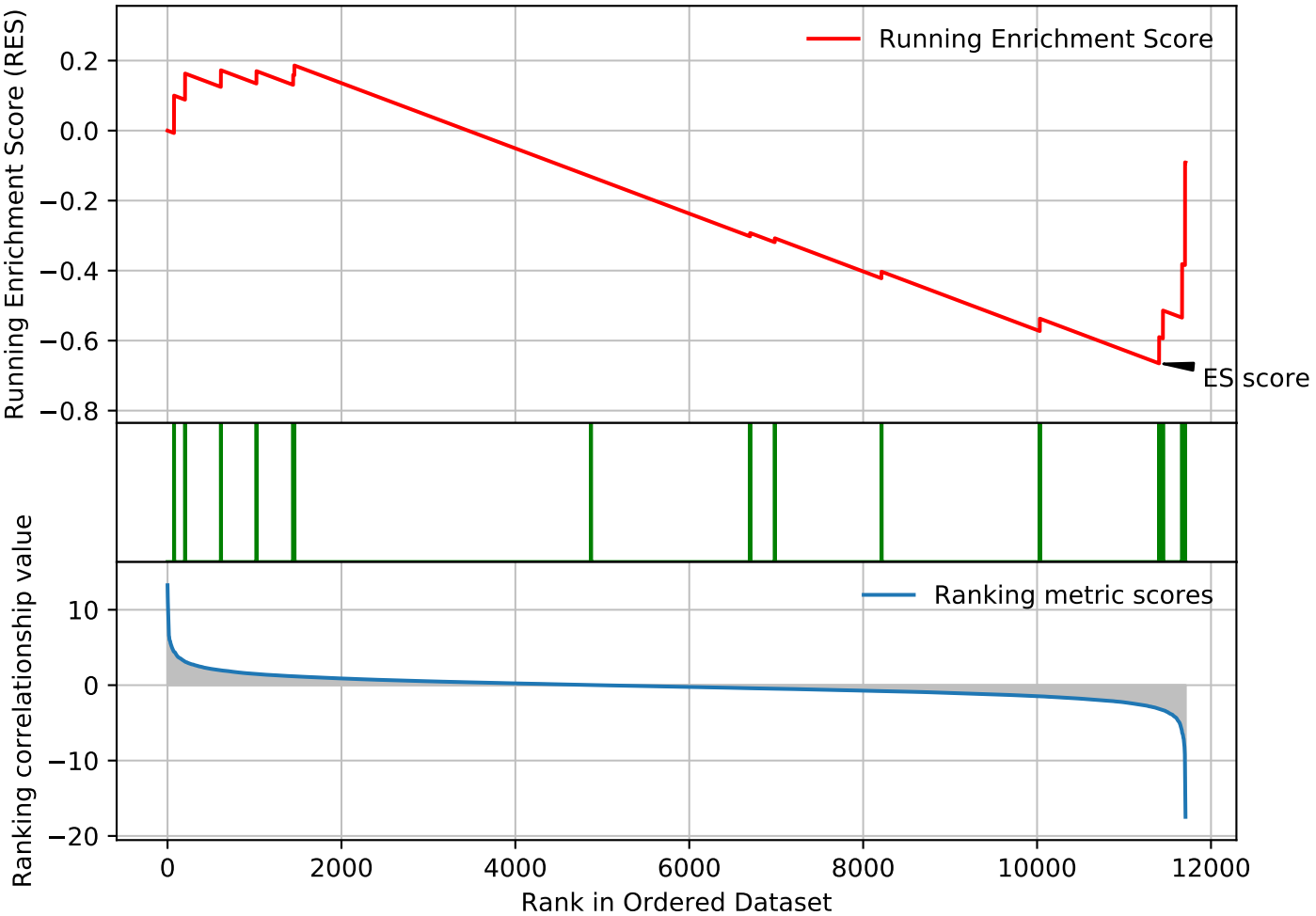




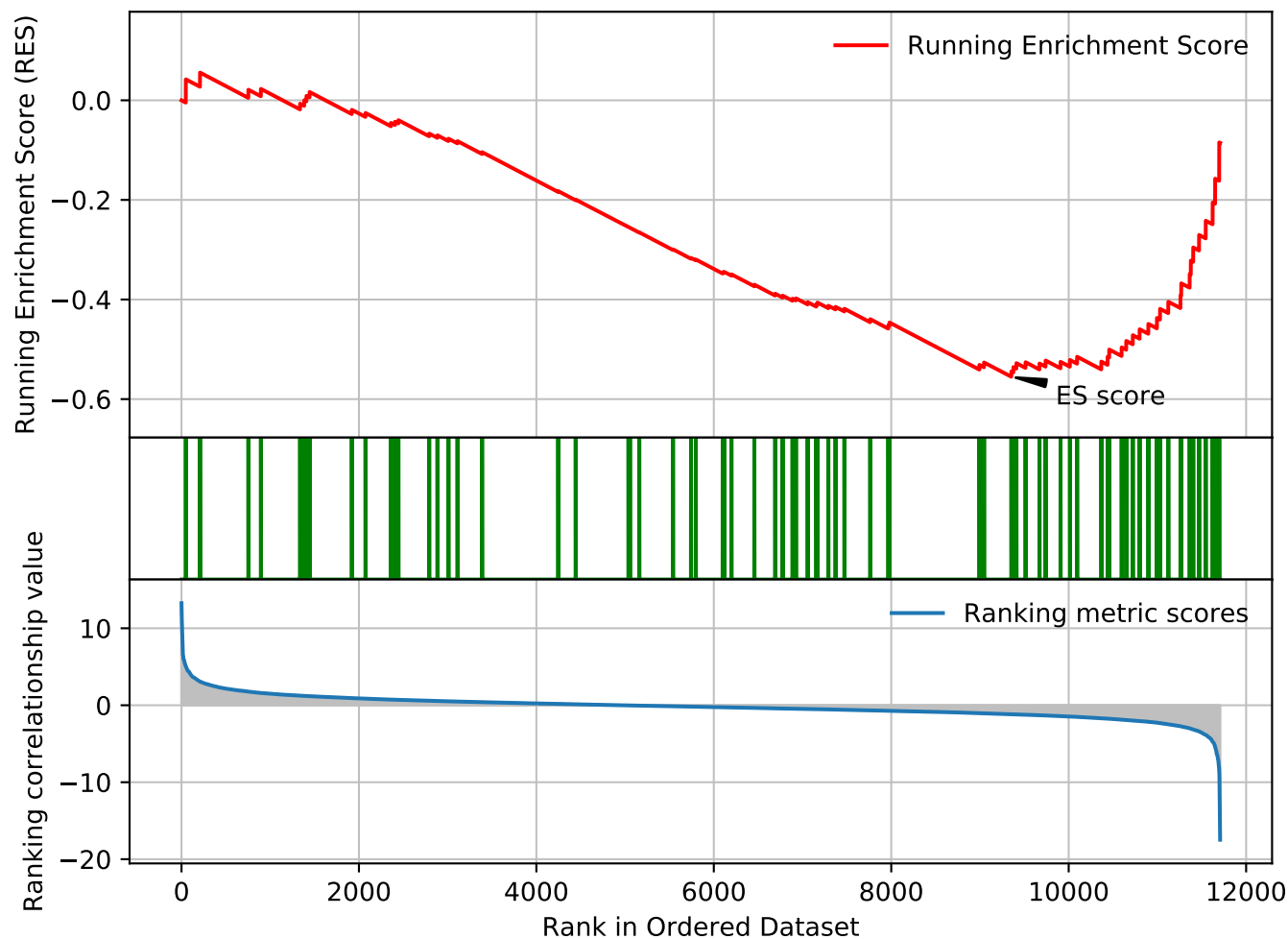
# WT R-MMU-69206 (G1/S Transition) GSEA\_Result

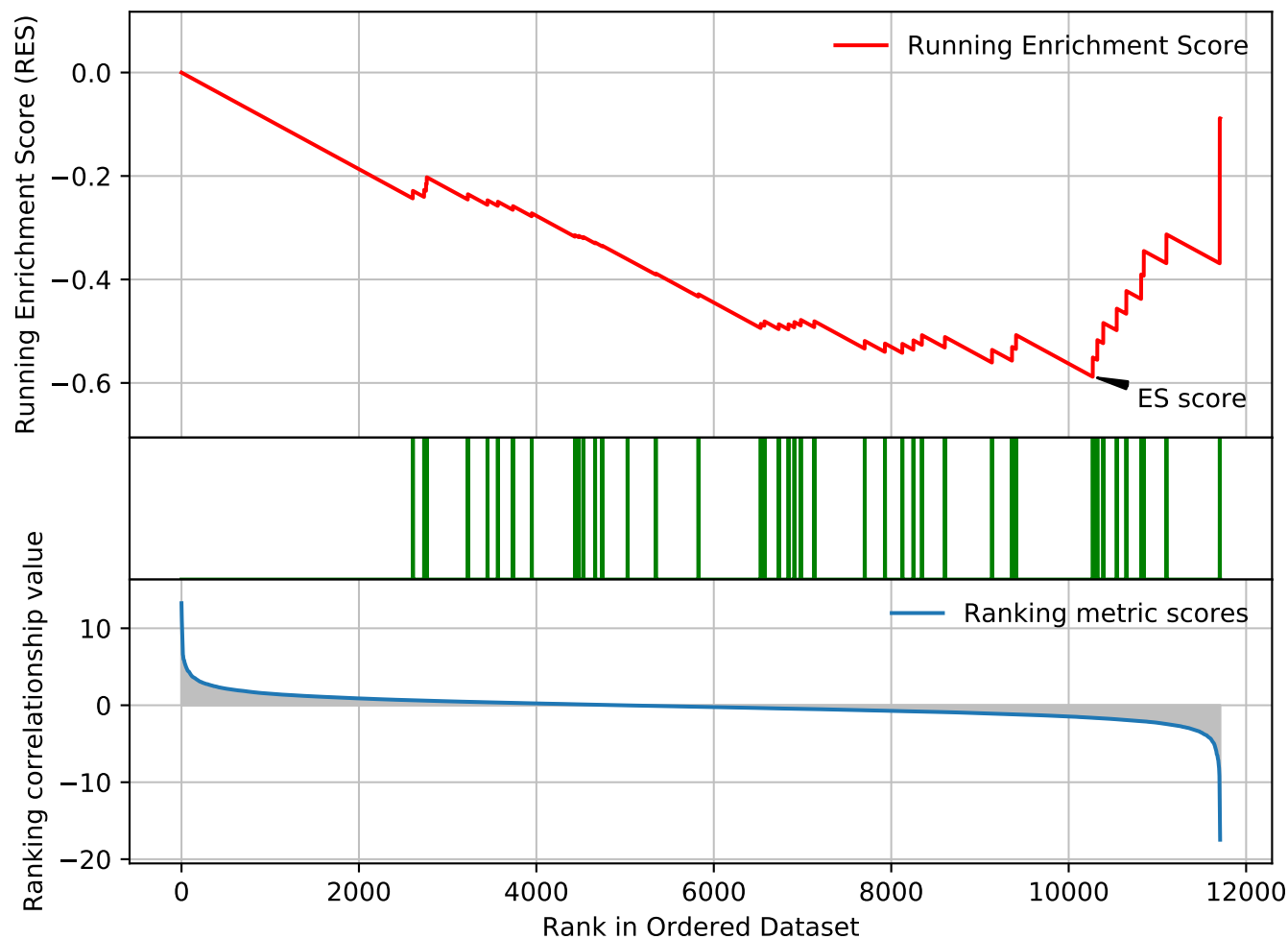


WT PWY-6126 (adenosine nucleotides de novo biosynthesis) GSEA\_Result



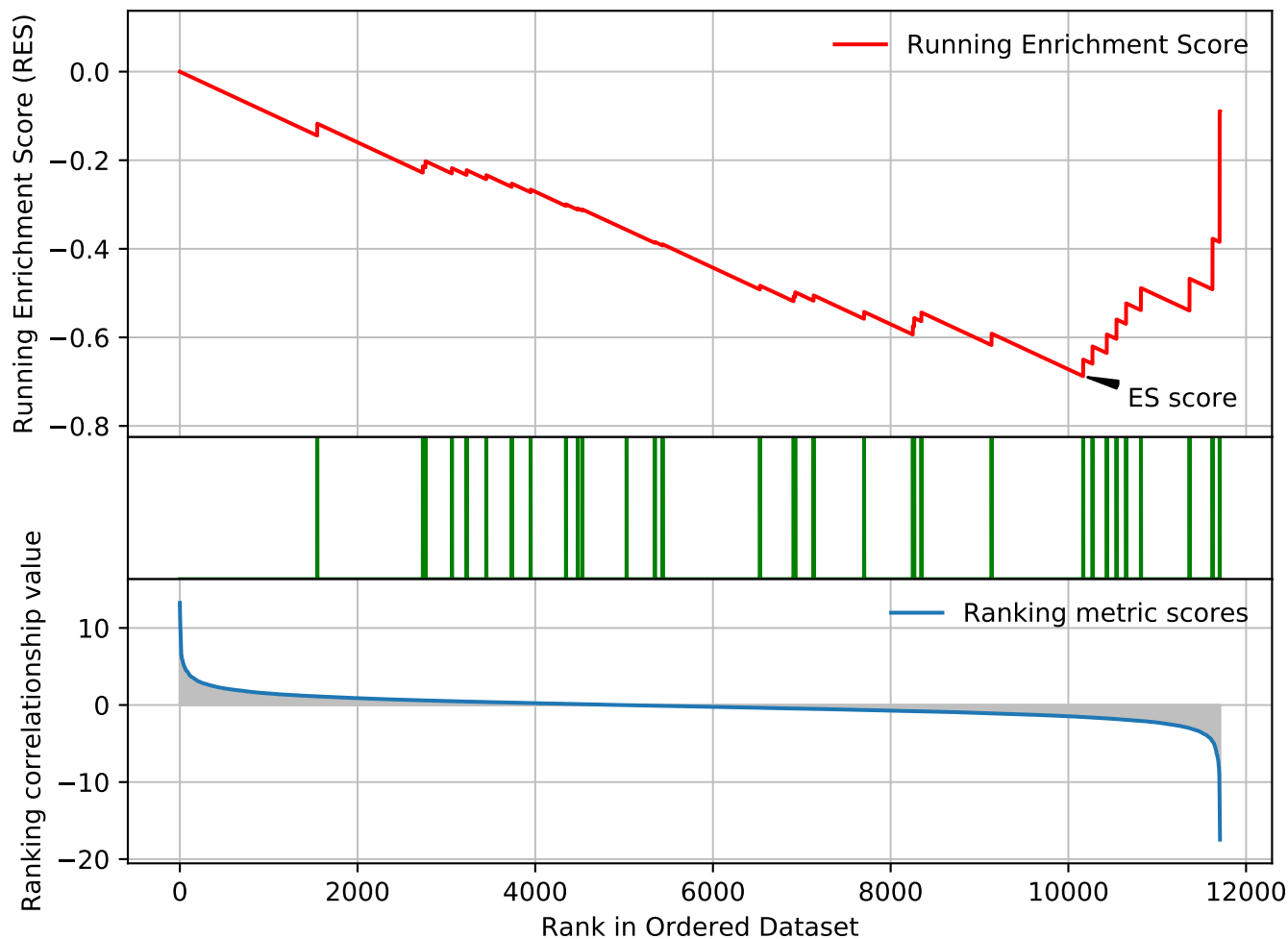
# 231 (Factors involved in megakaryocyte development and platelet production

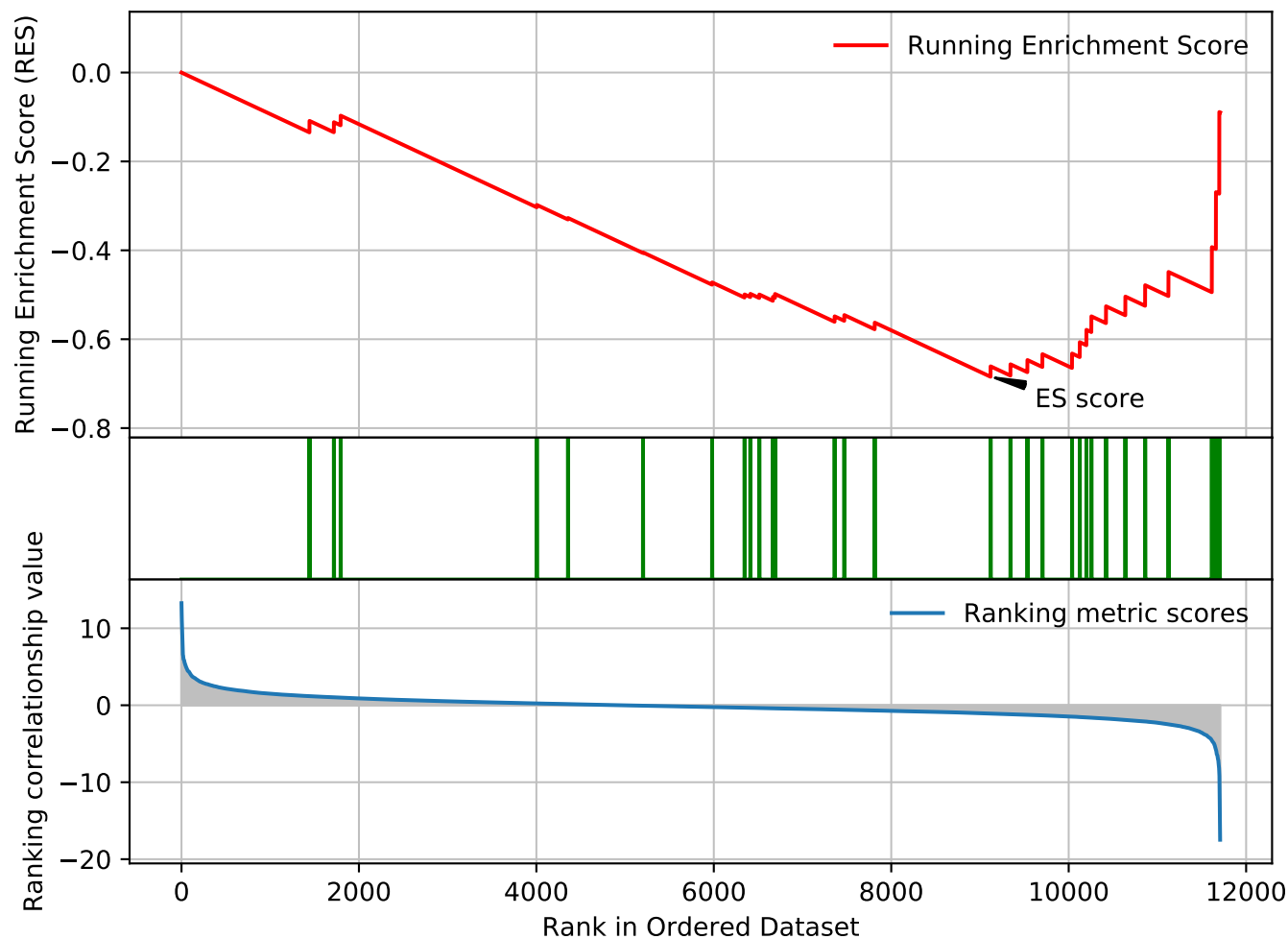




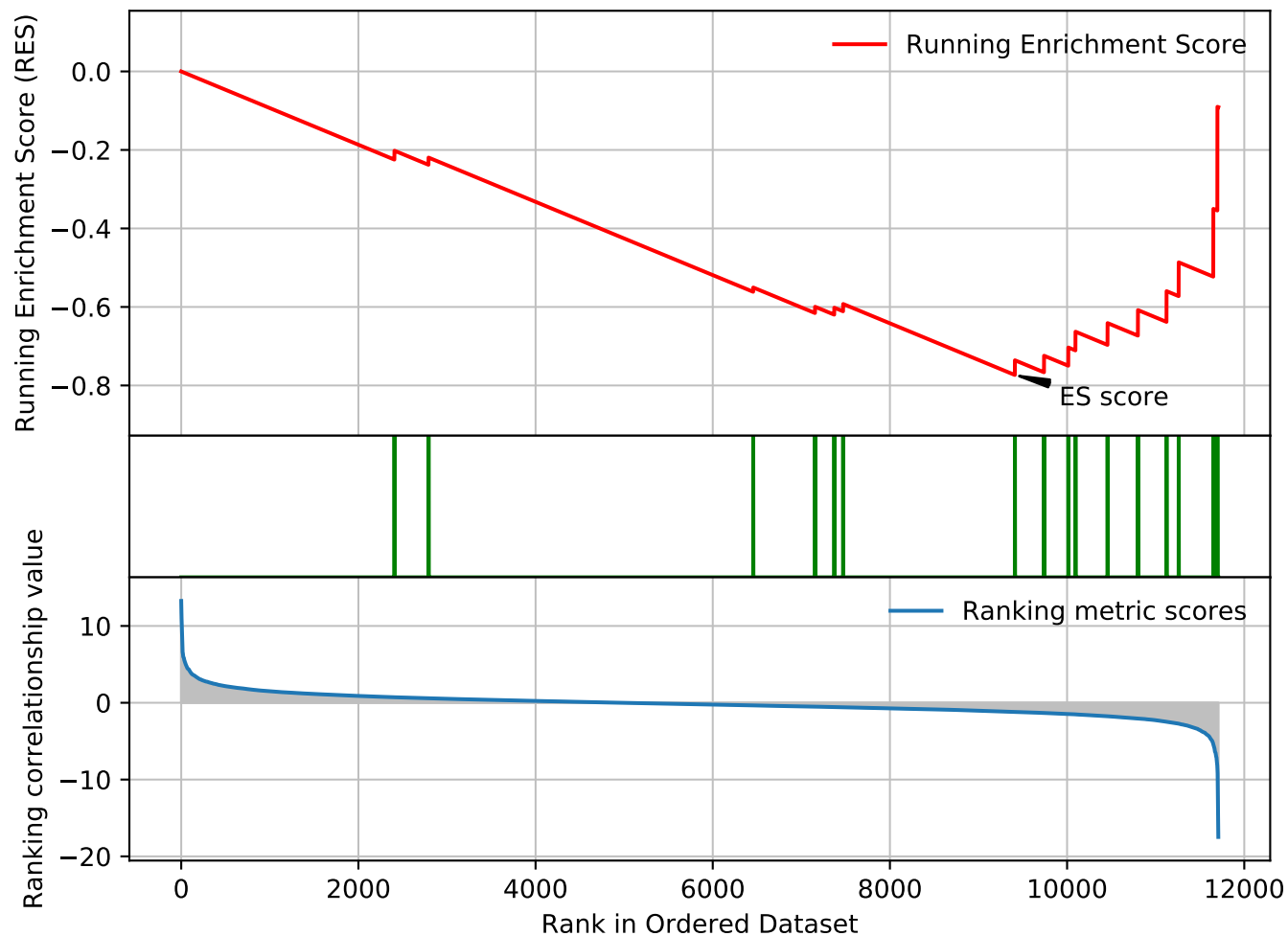


# WT P04378 (Beta2 adrenergic receptor signaling pathway) GSEA\_Result





# WT R-MMU-983189 (Kinesins) GSEA\_Result



# WT mmu03440 (Homologous recombination) GSEA\_Result

