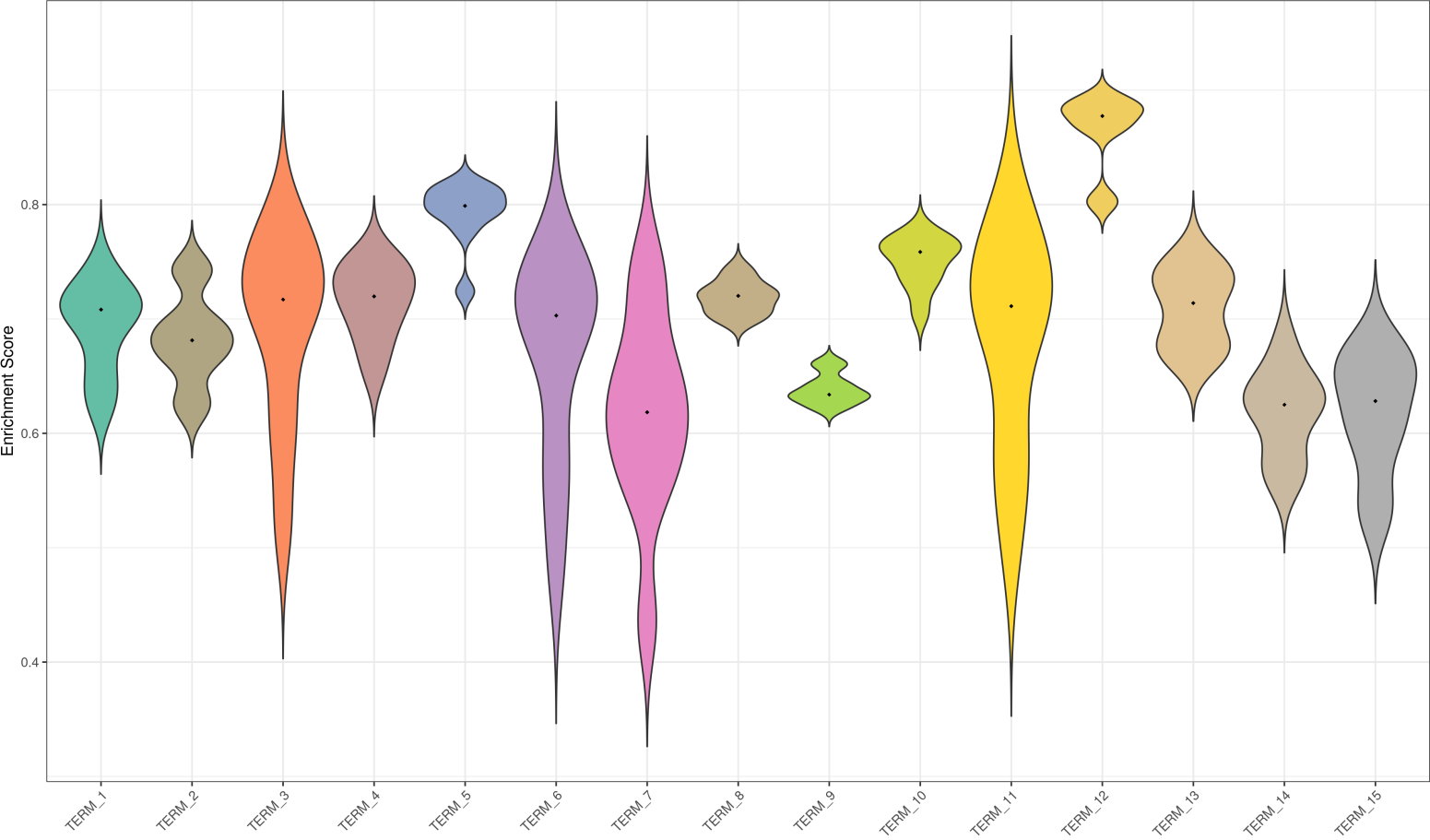


Single-sample Gene Set Enrichment Analysis

H: Hallmark Gene Sets

Term	Term_ID
HALLMARK_MITOTIC_SPINDLE	TERM_1
HALLMARK_TGF_BETA_SIGNALING	TERM_2
HALLMARK_G2M_CHECKPOINT	TERM_3
HALLMARK_ANDROGEN_RESPONSE	TERM_4
HALLMARK_PROTEIN_SECRETION	TERM_5
HALLMARK_INTERFERON_ALPHA_RESPONSE	TERM_6
HALLMARK_INTERFERON_GAMMA_RESPONSE	TERM_7
HALLMARK_UNFOLDED_PROTEIN_RESPONSE	TERM_8
HALLMARK_PI3K_AKT_MTOR_SIGNALING	TERM_9
HALLMARK_MTORC1_SIGNALING	TERM_10
HALLMARK_E2F_TARGETS	TERM_11
HALLMARK_MYC_TARGETS_V1	TERM_12
HALLMARK_OXIDATIVE_PHOSPHORYLATION	TERM_13
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY	TERM_14
HALLMARK_UV_RESPONSE_DN	TERM_15

Distribution of significant genesets for H



The diagram illustrates a hierarchical structure, likely representing a tree or a flowchart. It consists of several levels of rectangles. The top level has a single wide rectangle. This rectangle is divided into two main branches. The left branch further divides into two sub-branches, and the right branch divides into two sub-branches. The bottom-most rectangles are colored in a gradient from dark purple to light blue, while the upper levels are white with black outlines.

