Gene Set 1694: DAWSON_METHYLATED_IN_LYMPHOMA_TCL1 **Gene Set Null Distribution Heat Map for Genes in Gene Set** Malignant Benign Class
MACROD2
EPHAP
TO THE MACROD2
EPHAP
TO THE MACROD2
EPHAP
TO THE MACROD2
LPHIND
CDKN2A
FAM 18AA
HMGN1
HOXD13
FAM 18AA
HMGN1
HOXD13
FAM 18AA
FAM 18AA
FAM 18AA
FAM 18AA
HMGN1
FAM 18AB
FAM 18 Gene Set Null Density Observed Gene Set ES value 0.0 Running Enrichment Score (RES) $^{\circ}$ P(ES) Zero crossing at 8459 -0.5 Peak at 19568 0 Neg. ES "Malignant" Pos. ES: "Benign" "Benign" "Malignant" GSM721625 GSM721593 GSM721594 GSM721595 GSM721596 GSM721598 GSM721619 GSM721620 GSM721623 GSM721624 GSM721597 GSM721621 GSM721622 5000 10000 15000 20000 25000 -0.4-0.2 0.0 0.2 0.4 Gene List Index Number of genes: 25691 (in list), 56 (in gene set) ES = -0.437 NES = -1.66 Nom. p-val= 0 FWER= 0.87 FDR= 1