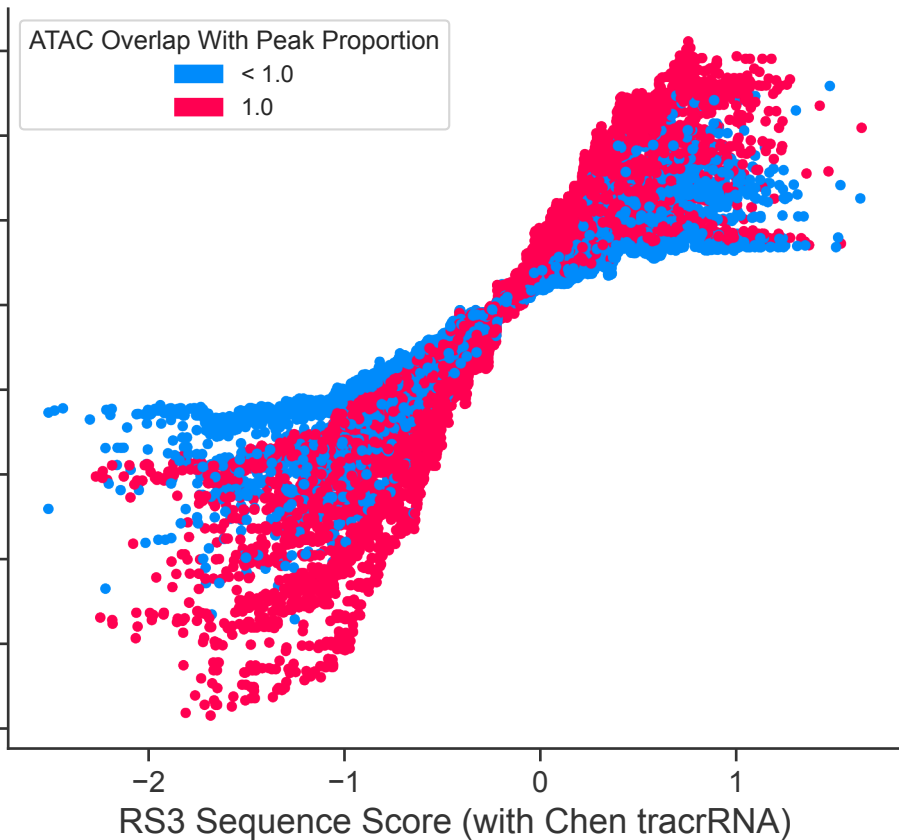


SHAP value for RS3 Sequence Score

ATAC Overlap With Peak Proportion



0.6
0.4
0.2
0.0
-0.2
-0.4
-0.6
-0.8
-1.0



RS3 Sequence Score (with Chen tracrRNA)

ATAC overlap with peak proportion 1

1.0
0.5
0.0