You asked:
Are 'MEFB1 (BLOC segments)' overlapping 'SINE (Repeating elements)', more than expected by chance?

Simplistic answer:
No support from data for this conclusion in any bin

Precise answer:
0 significant bins out of 19, at 10% FDR*

A collection of FDR-corrected p-values per bin was computed. Not able to compute a global p-value for this analysis.

* False Discovery Rate: The expected proportion of false positive results among the significant bins is no more than 10%.

In each bin, the test of
H0: The segments of track 1 are located independently of the segments of track 2 with respect to overlap
vs
H1: The segments of track 1 tend to overlap the segments of track 2 was performed.

P-values were computed under the null model defined by the following preservation and randomization rules:

Preserve segments of T2, segment and inter-segment lengths of T1, randomize positions (MC)

The test statistic used is:
The number of base pairs that are inside segments of both tracks