



Additional file 3. Distribution of distances (shown in nt) between the terminal monomer of the tandem array and the TE present at the TE/tandem array junction (X-axis).

(A) TE/pSc200 junctions, **(B)** TE/pSc250 junctions. The Y-axis plots the ration between the number of junctions harboring a given size of spacer DNA and the total number of junctions harboring the same TE.