



Figure S6. T cell activation results in partitioning of genome topology. Pie chart representation of the distribution of (A) TADs in resting CD4⁺ T cells overlapping 0, 1, 2 or ≥ 3 TADs in resting CD8⁺ T cells, (B) TADs in resting CD8⁺ T cells overlapping 0, 1, 2 or ≥ 3 TADs in resting CD4⁺ T cells (C) TADs in resting CD4⁺ T cells overlapping 0, 1, 2 or ≥ 3 TADs in activated CD4⁺ T cells, and (D) TADs in resting CD8⁺ T cells overlapping 0, 1, 2 or ≥ 3 TADs in activated CD8⁺ T cells.