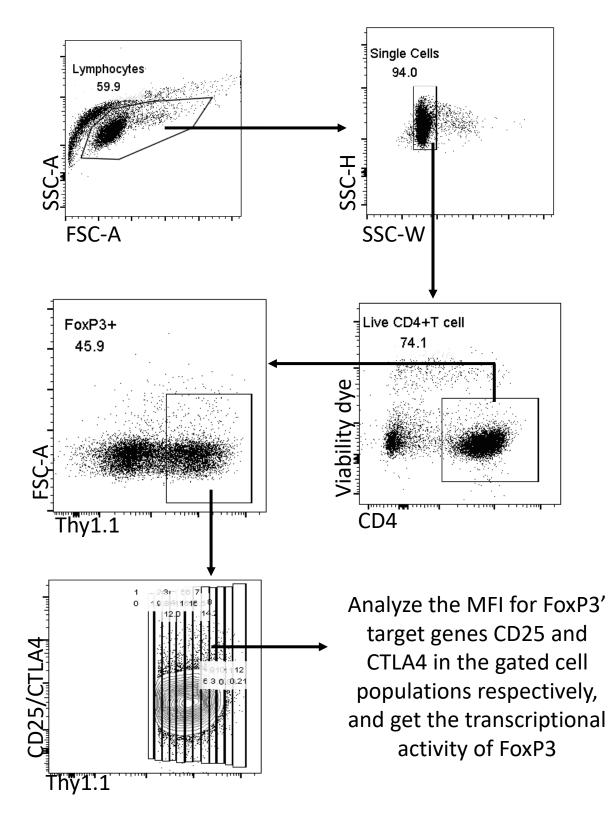
# **Supplementary information**

# FOXP3 recognizes microsatellites and bridges DNA through multimerization

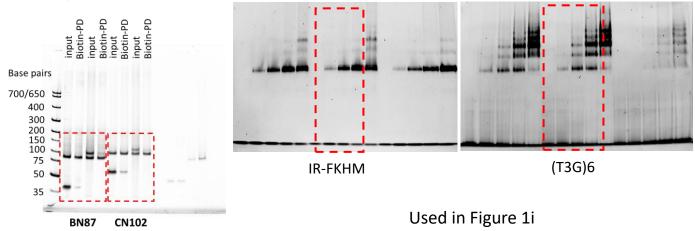
In the format provided by the authors and unedited

## **Supplementary Figure 1**

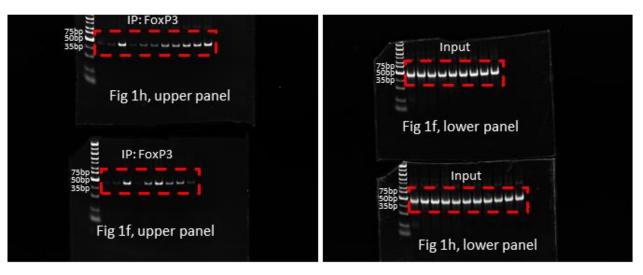


# **Supplementary Figure 2**

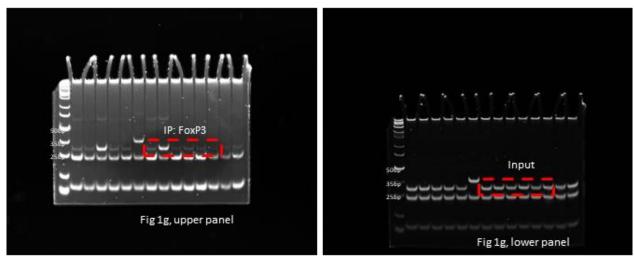




Used in Figure 1c. Right panel.



Used in Figure 1f and 1h

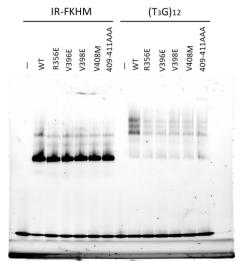


Used in Figure 1g

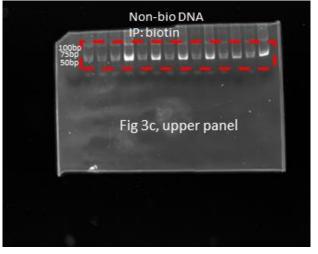
#### For Figure 2 and Figure 3



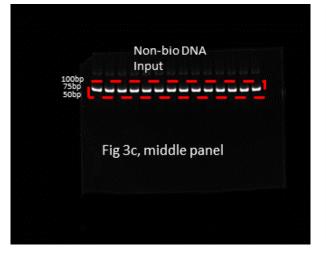
Used in Figure 2e

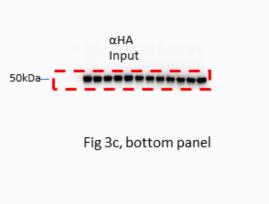


Used in Figure 3b



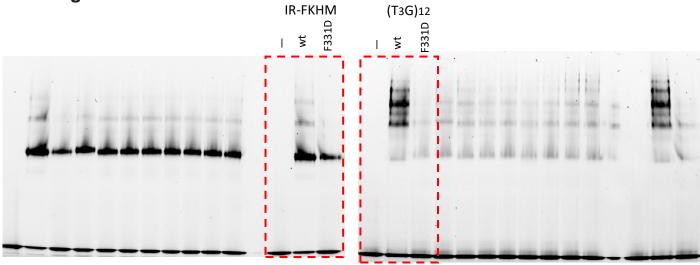
Used in Figure 3c, upper panel



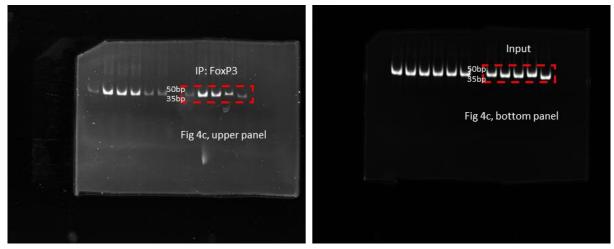


Used in Figure 3c, middle panel

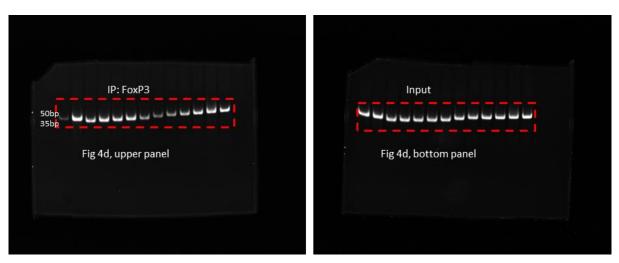
#### Used in Figure 3c, bottom panel



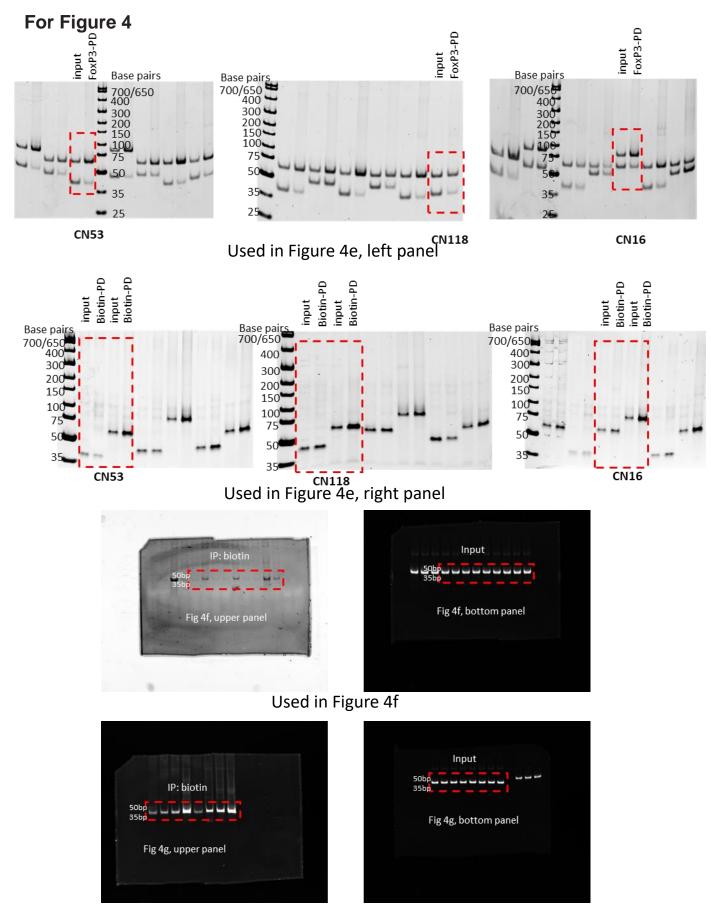
Used in Figure 4b



Used in Figure 4c

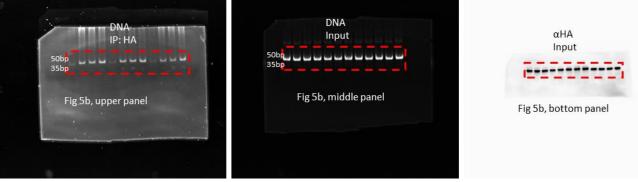


Used in Figure 4d

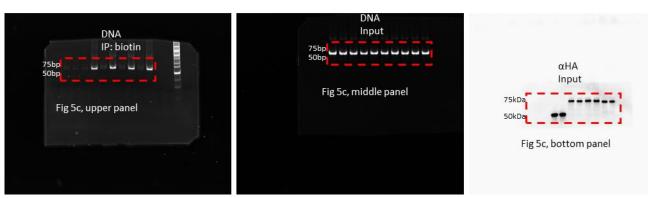


Used in Figure 4g

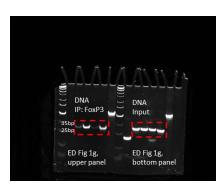
### For Figure 5 and ED Figure 1



#### Used in Figure 5b



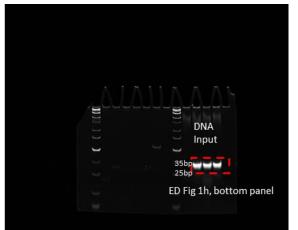
Used in Figure 5c



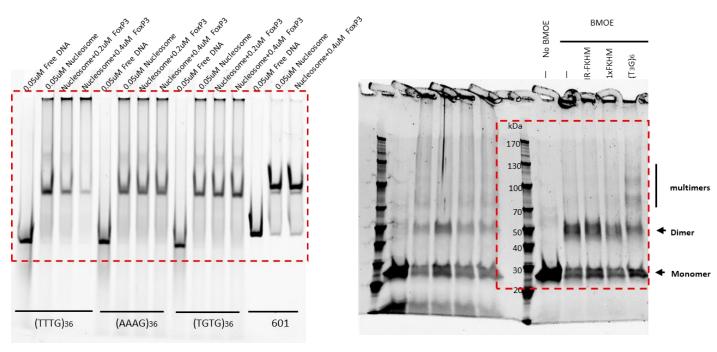
#### Used in ED Figure 1g



ED Fig 1h, upper panel



#### Used in ED Figure 1h



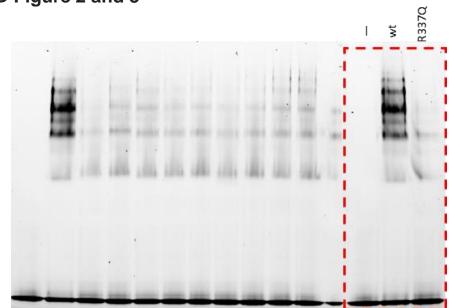
Used in ED Figure 1i

Used in ED Figure 1j

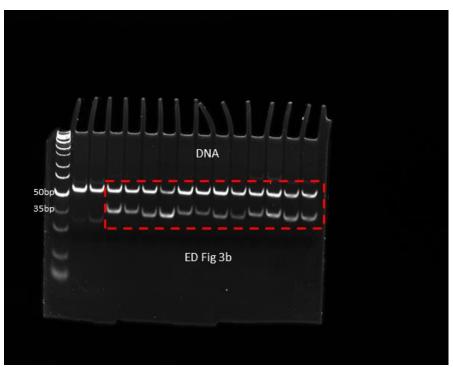


Used in ED Figure 1k

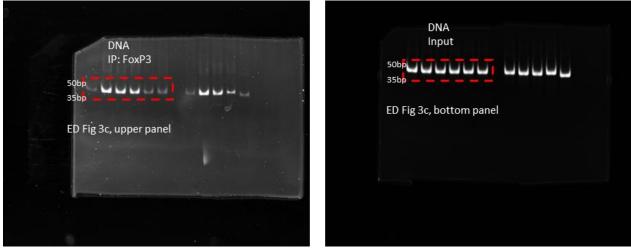
For ED Figure 2 and 3



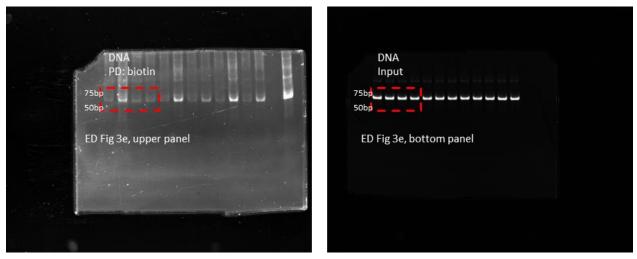
Used in ED Figure 2h



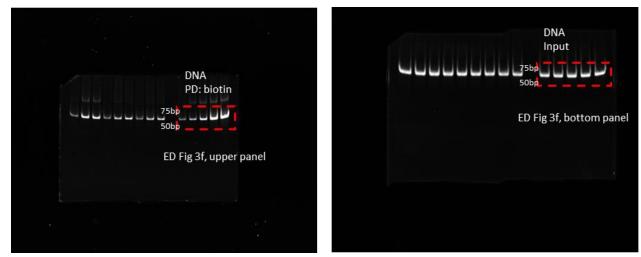
Used in ED Figure 3b



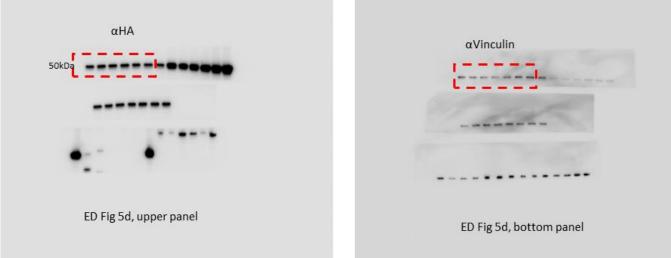
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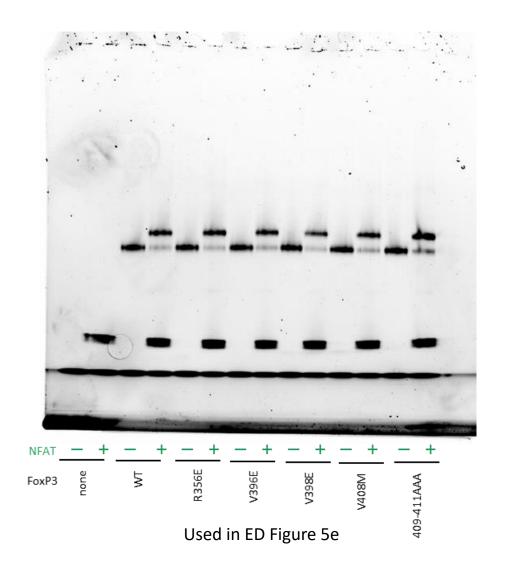
Used in ED Figure 3e

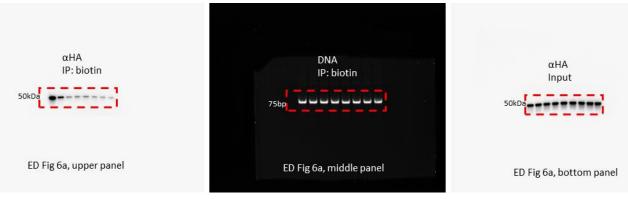


Used in ED Figure 3f

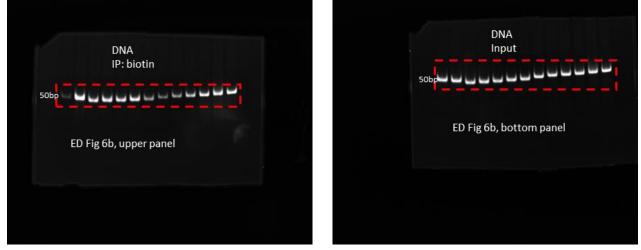




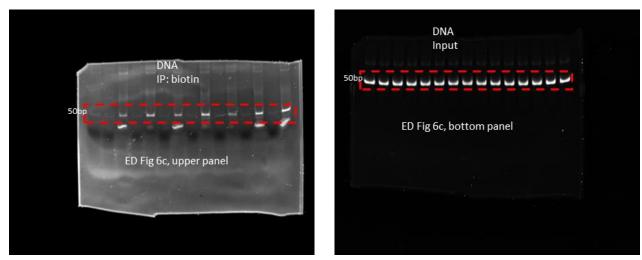




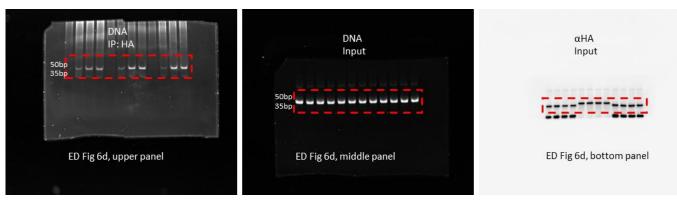
Used in ED Figure 6a



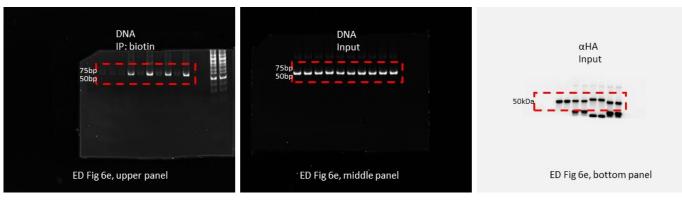
Used in ED Figure 6b



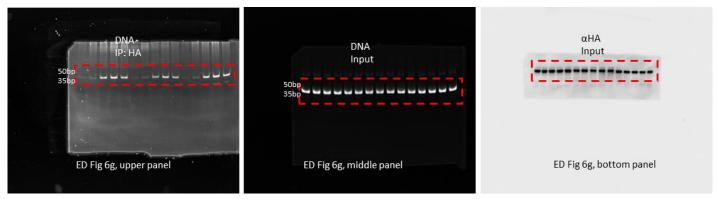
Used in ED Figure 6c



#### Used in ED Figure 6d



#### Used in ED Figure 6e



#### Used in ED Figure 6g

## **Supplementary Tables**

**Supplementary Table 1.** Summary of FoxP3 binding motifs. 1a, FoxP3 binding motif *in vitro*. 1b, FoxP3 binding motifs *in vivo*. 1c, other FKH TFs' binding motifs.

**Supplementary Table 2.** List of DNA oligos used in this study. 2a, allelic bias sites information. 2b, DNA oligo sequences.

**Supplementary Table 3.** Chromatin contacts at FoxP3-bound anchors. 3a, summary of the contacts with frequency>5 in WT Treg HiC and connected by two TnG anchors. 3b-3f, with an increasing FoxP3 PLAC-seq count threshold (>5, 10, 25, 50, 75 respectively), all the anchor information is displayed. 3g, genomic annotation for the CNR peaks overlapping with PLAC-seq counts above the threshold of 75.