

Figure S1

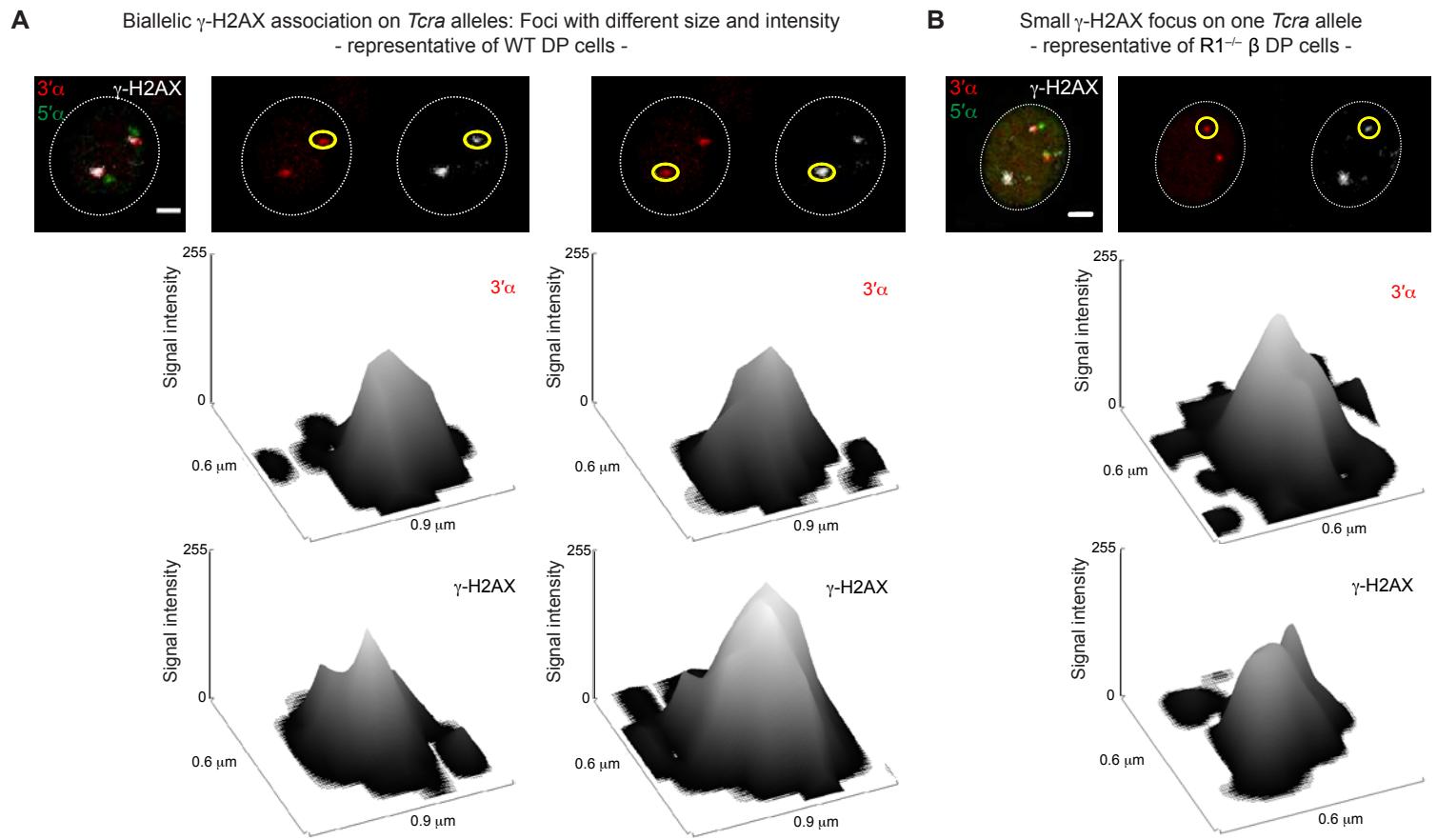
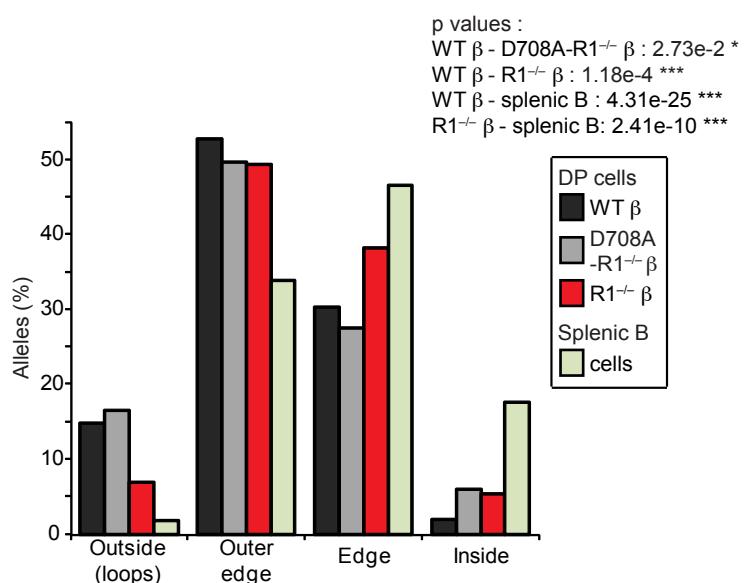
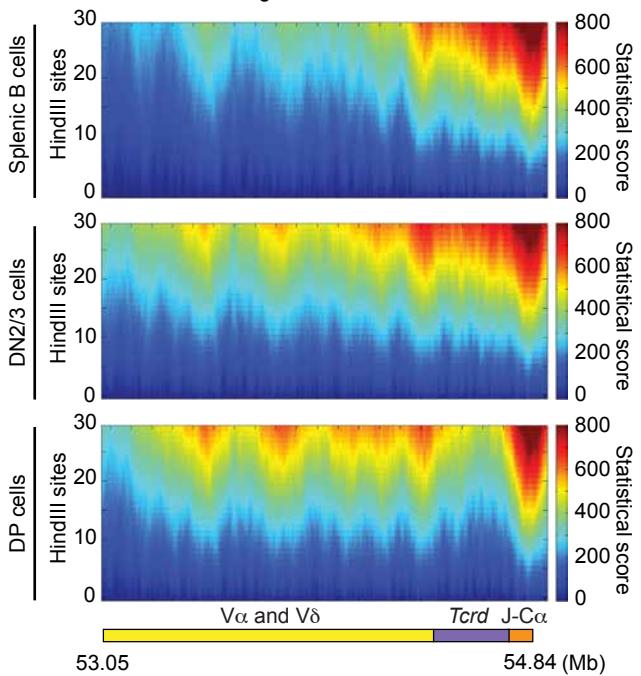


Figure S2

A Location of *Tcra* alleles relative to the chromosome 14 territory



B Domainograms of the *Tcra/d* locus



C

Monoallelic γ -H2AX association on *Tcra* in DP cells

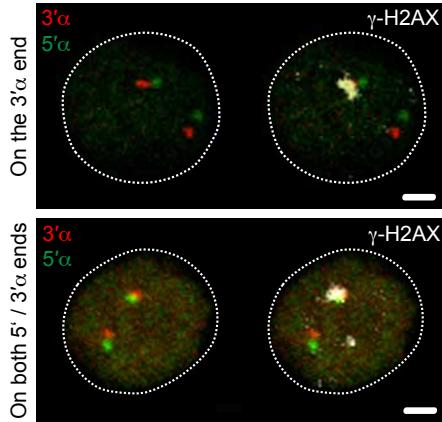
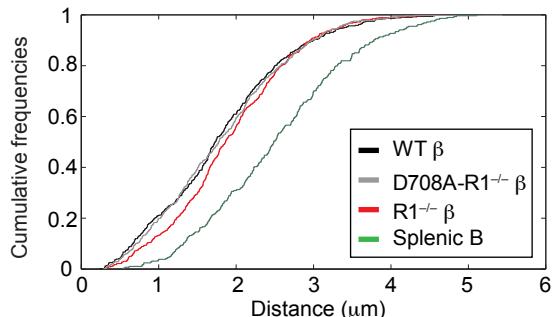


Figure S3

A Separation of *Tcra* alleles in WT, RAG1-mutant DP and splenic B cells



p values (entire distributions):

WT β-R1^{-/-} β : 3.17e-2 *

WT β-D708A-R1^{-/-} β : 8.60e-1 ns

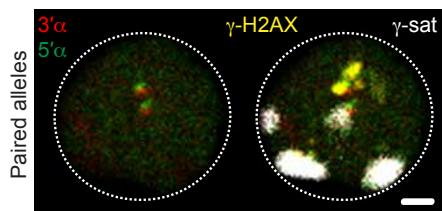
WT β-splenic B : 1.03e-17 ***

R1^{-/-} β-D708A-R1^{-/-} β : 1.13e-2 *

R1^{-/-} β-splenic B : 1.36e-13 ***

C

Nuclear location of paired *Tcra* alleles with the uncleaved allele at PCH (DP cells)



B

Location of paired *Tcra* alleles relative to the chromosome 14 territory

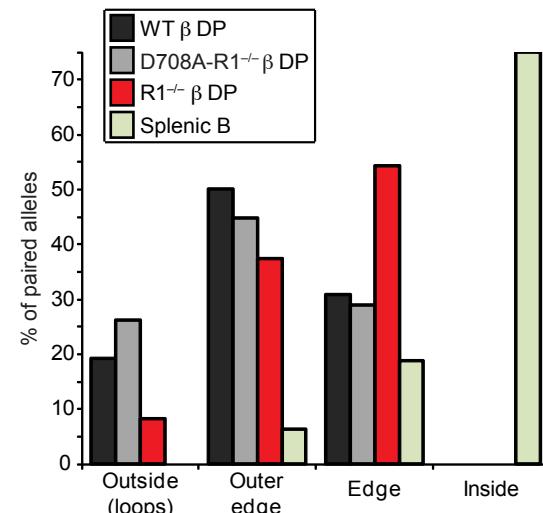
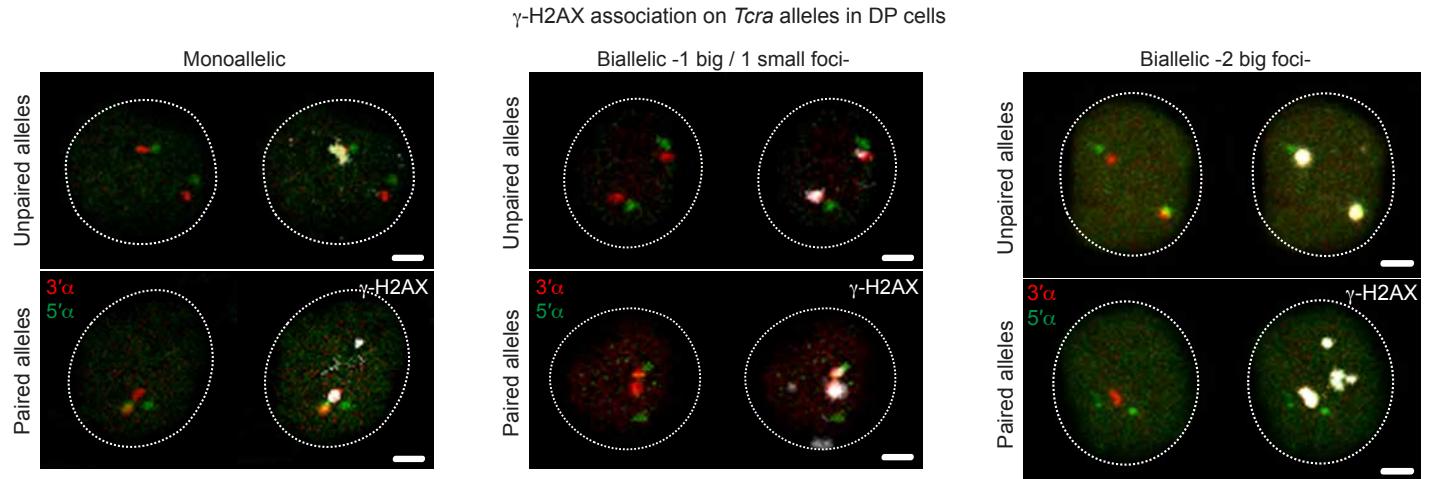
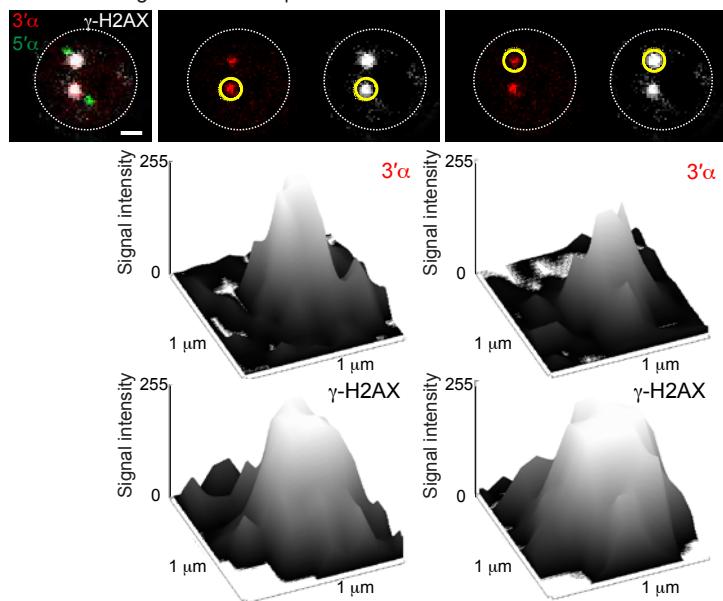


Figure S4

A



B Two large and intense γ -H2AX foci on *Tcra* alleles in *Atm*^{-/-} DP cells



C γ -H2AX association on *Tcra* alleles in *53bp1*^{-/-} DP cells

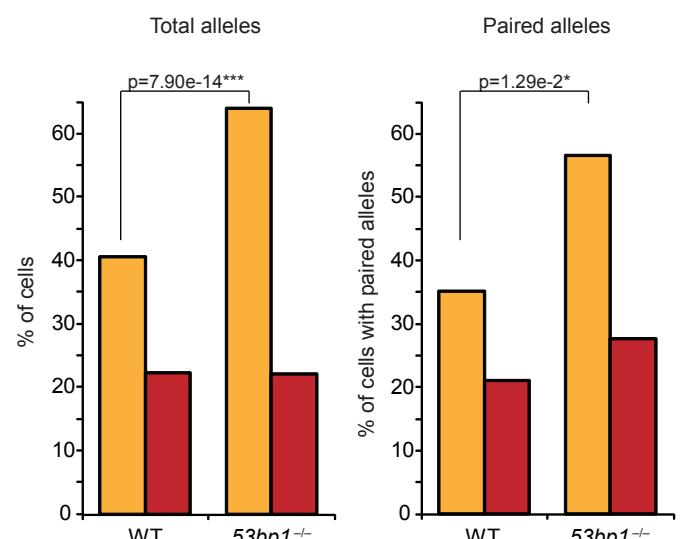


Figure S5

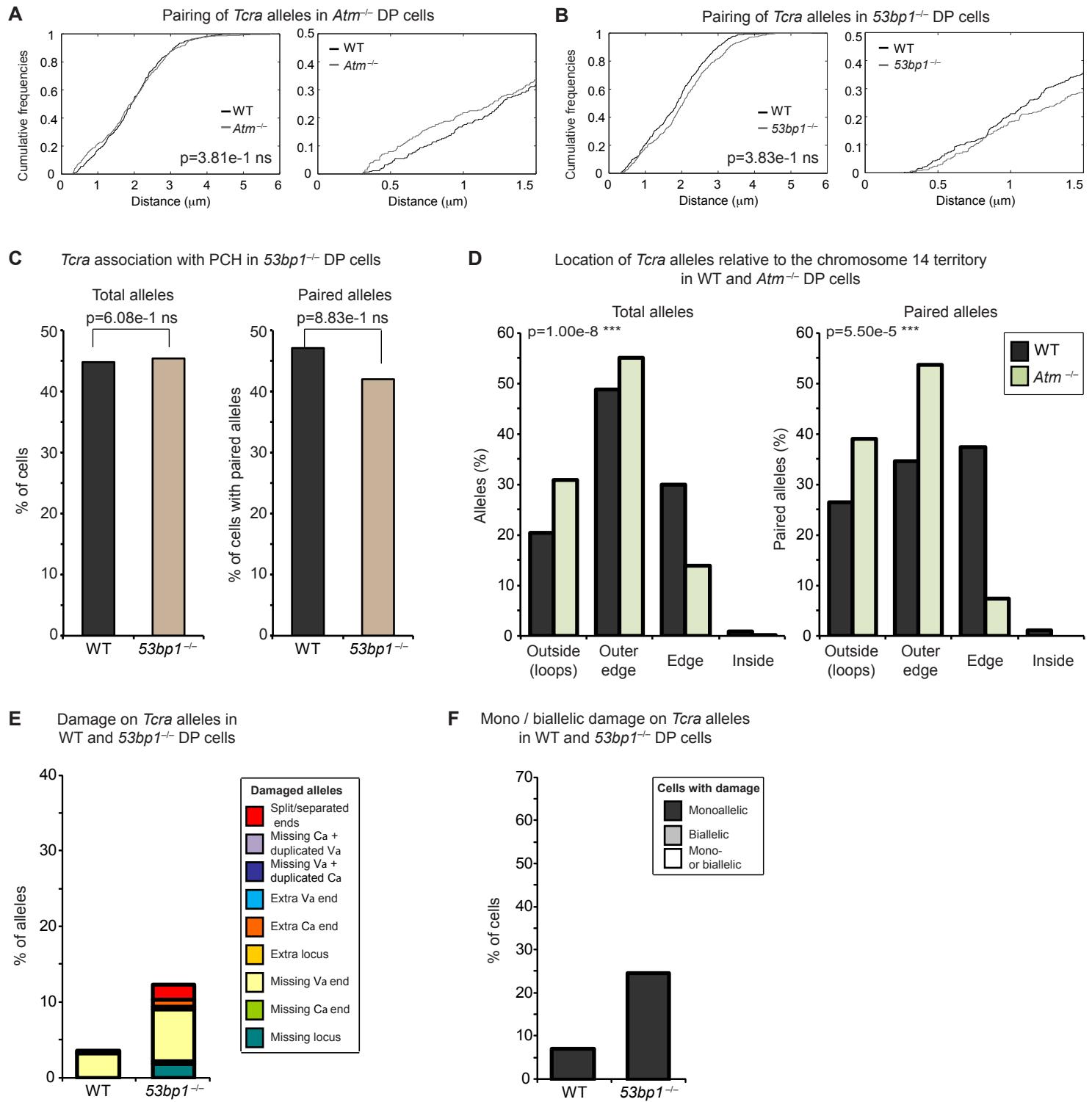


Figure S6

