SUPPLEMENTAL MATERIAL

Supplementary Table 1: qPCR primers	
18S F	CGCGGTTCTATTTTGTTGGT
18S R	AGTCGGCATCGTTTATGGTC
β actin F	TCATCACTATTGGCAACGAGCGGTTC
β actin R	TACCACCAGACAGCACTGTGTTGGCA
CD19 F	ACCAGTACGGGAATGTGCTC
cD19 R	TTCATAGGCCTCCCCTTCTT
Gadd45α F	CCGAAAGGATGGACACGGTG
Gadd45α R	TTATCGGGGTCTACGTTGAGC
Hprt F	CTGGTGAAAAGGACCTCTCG
Hprt R	TGAAGTACTCATTGTAGTCAAGGGCA
Lck F	CCTTCGGGATCTTGCTTACA
Lck R	GTTGTCAGGTCTCACCATGC
p21 F	GACATTCAGAGCCACAGGCAC
p21 R	GTCAAAGTTCCACCGTTCTCG
Rag1 F	TGGGAATCGTTTCAAGAGTGAC
Rag1 R	CATCTGCCTTCACGTCGATCC
Rag2 F	ACACCAAACAATGAGCTTTCCG
Rag2 R	CCGTATCTGGGTTCAGGGAC
Wasp F	GCTCCCTCCTACTCCAGTGTC
Wasp R	AGGGCACCTACTAGGCCTTC



SUPPLEMENTAL FIGURE 1. Pro-B and pro-T cells suppress *Rag2* expression in response to ionizing radiation. (**A-B**) qRT-PCR quantification of Rag2 and p21 mRNA in BM (**A**) or thymuses (**B**) from non-irradiated *Rag1*- mice or *Rag1*- mice at indicated times after exposure to 10 Gy IR. Data in each graph are from 3 or more independent experiments with a total of 4-6 mice per timepoint. Data averages are shown with error bars indicating SEM. p-values calculated using Dunnett's post-test after ANOVA. *p<0.05, **p<0.01, ***p<0.001.



SUPPLEMENTAL FIGURE 2. Neither p53 activity nor downregulation of Gadd45 α mRNA levels is required for suppression of *Rag1* and *Rag2* transcription in response to IR. (A) qRT-PCR quantification of Gadd45 α mRNA in non-irradiated *EµBCL2* pre-B cells or irradiated *EµBCL2* pre-B cells at indicated times after exposure to 4 Gy IR. Data are from 11 independent experiments. Data are normalized to 1.0 for non-irradiated cells. For irradiated cells, data averages are shown with error bars indicating SEM. pvalues calculated using Dunnett's post-test after ANOVA. ***p<0.001. (B) qRT-PCR quantification of EU-labeled Gadd45 α mRNA levels relative to EU-labeled Hprt mRNA levels in non-irradiated EuBCL2 pre-B cells and irradiated $E\mu BCL2$ pre-B cells at indicated times after addition of EU and exposure to 4 Gy of IR. Data are presented as the ratio of relative levels of each mRNA in irradiated cells compared to non-irradiated cells. The dotted line represents a value of 1, which would indicate that IR had no effect on the transcription rate of an assayed gene. Data are from 4 independent experiments. p-values were determined using one-tailed T test with Bonferroni's correction for multiple testing. $***p \le 0.001$. (C) qRT-PCR quantification of Rag1 and Rag2 mRNA in non-irradiated and irradiated $Rag1^{D708A}$ Abl pre-B cells transduced with an empty retroviral vector or the same retroviral vector containing an ER-Gadd45 α cDNA. For 2 hours before harvesting non-irradiated cells or exposing to IR the irradiated cells, all cells were treated with tamoxifen to induce nuclear translocation of ER-Gadd45 α . Data are from one representative experiment. (D) Western blot showing ER-Gadd45 α and actin expression in Rag1^{D708A} Abl pre-B cells transduced with an empty retroviral vector or the same retroviral vector containing an ER-Gadd45 α cDNA. Data are from one representative experiment. (E) qRT-PCR quantification of Rag1 and Rag2 mRNA in non-irradiated $VavCre^+Tp53^{flox/flox}$ pre-B cells or irradiated $VavCre^+Tp53^{flox/flox}$ pre-B cells at indicated times after exposure to 4 Gy IR. Data are from 2 independent experiments. Data are normalized to 1.0 for non-irradiated cells harvested one hour following IR exposure of irradiated samples. For other samples, data averages are shown with error bars indicating SEM.



SUPPLEMENTAL FIGURE 3. Bleomycin treatment inhibits recombination of endogenous $Ig\kappa$ loci. (A) qRT-PCR quantification of Rag1, Rag2, and p21 mRNA in *Artemis*- $E\mu BCL2$ Abl cells untreated or treated with STI571 or STI571 and bleomycin for the indicated amounts of time. Data are from 6 independent experiments. (B) Representative Southern blot analysis and graphical quantification of $J\kappa$ cleavage in *Artemis*- $E\mu BCL2$ Abl cells untreated or treated with STI571 or STI571 and bleomycin for the indicated amounts of time. Data are from 6 independent experiments. (B) Representative Southern blot analysis and graphical quantification of $J\kappa$ cleavage in *Artemis*- $E\mu BCL2$ Abl cells untreated or treated with STI571 or STI571 and bleomycin for the indicated amounts of time. Data are from 3 independent experiments.