

Supplementary Figures and Tables

Supplementary Figure 1

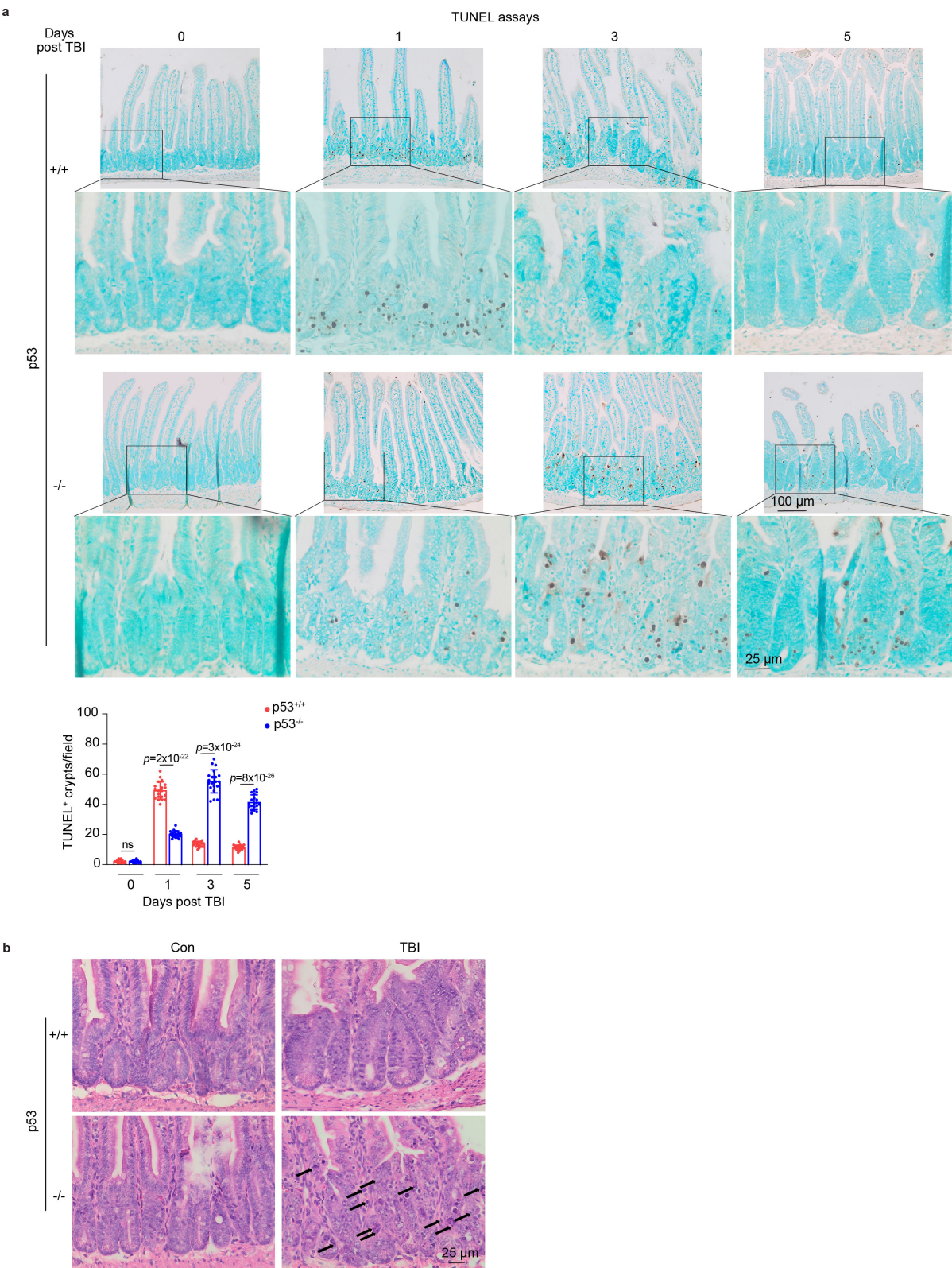


Figure S1. p53 deficiency results in severe cell death in the SI of mice post-TBI.

a. Representative images from at least 3 independent mice (upper panels) and quantification (bottom panel) of TUNEL assays in the SI of p53^{+/+} and p53^{-/-} mice at different time points post 12 Gy TBI. $n = 20$ fields from at least 3 mice/group. **b.** Representative images from at least 3 independent mice showing mitotic catastrophe in the SI of p53^{+/+} and p53^{-/-} mice at 3 days post 12 Gy TBI. Data are presented as mean \pm SD from at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test. Source data are provided as a Source Data file.

Supplementary Figure 2

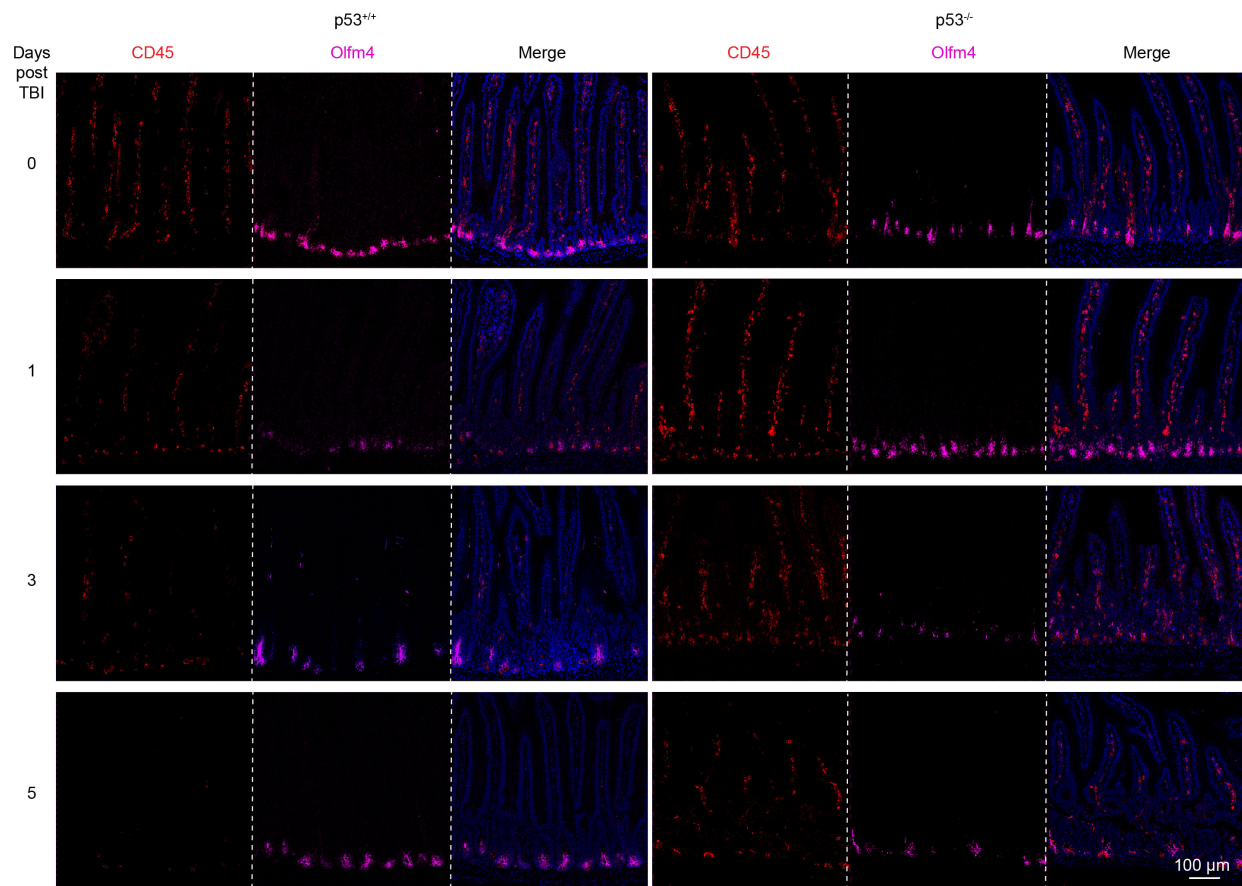


Figure S2. p53 deficiency enhances the inflammatory immune response and ISC damage in the SI of mice post-TBI.

Representative images from at least 3 independent mice showing IF staining of CD45 and Olfm4 in the SI of $p53^{+/+}$ and $p53^{-/-}$ mice at different days post 12 Gy TBI.

Supplementary Figure 3

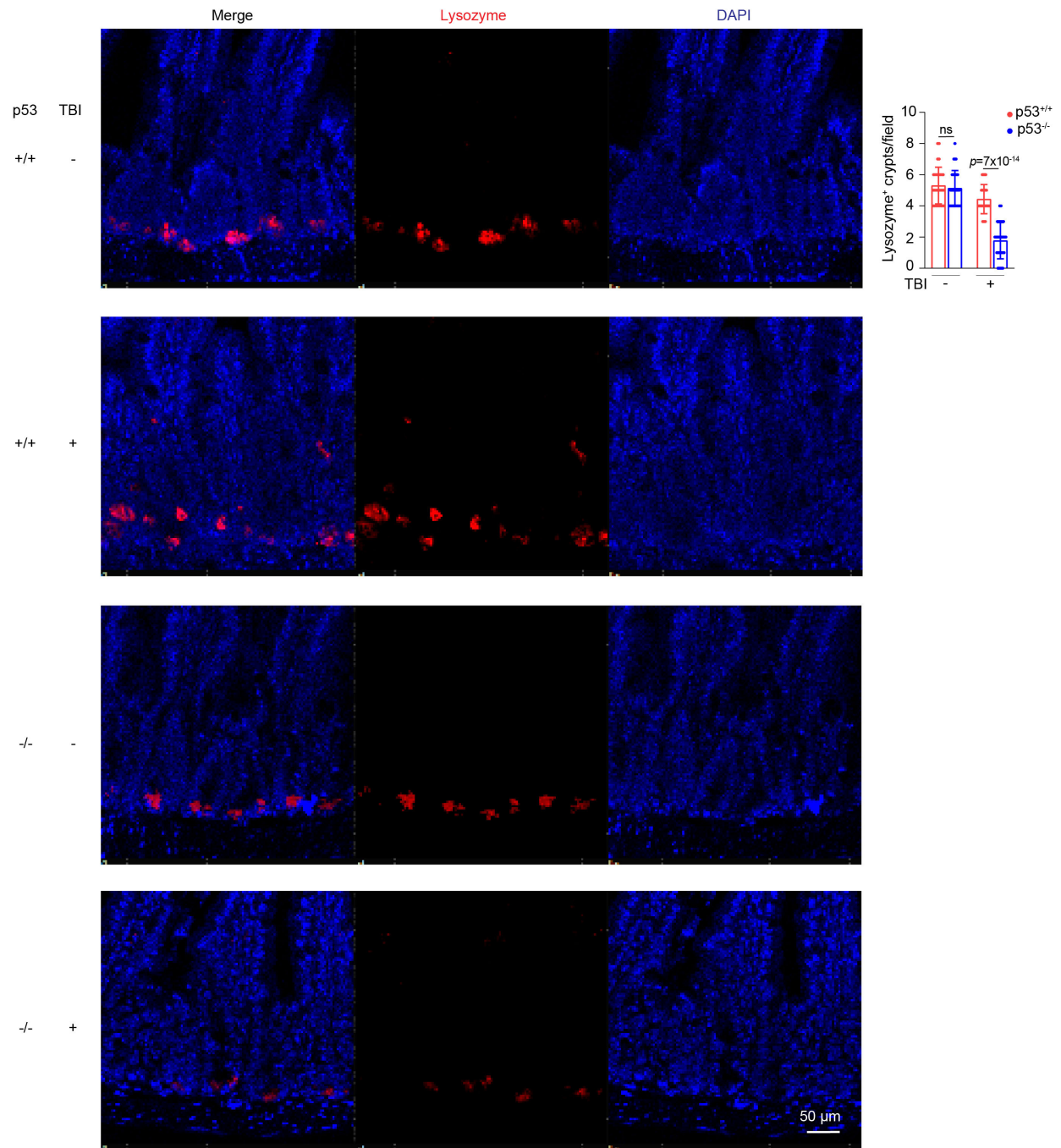


Figure S3. p53 deficiency enhances the damage of Paneth cells in the SI of mice post-TBI. Representative images from at least 3 independent mice (left panels) and quantification (right panel) of IF staining for lysozyme, a Paneth cell marker, in the SI of p53^{+/+} and p53^{-/-} mice at 3 days post-TBI. n = 30 crypts from at least 3 mice/group. Data are presented as mean ± SD from

at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test. Source data are provided as a Source Data file.

Supplementary Figure 4

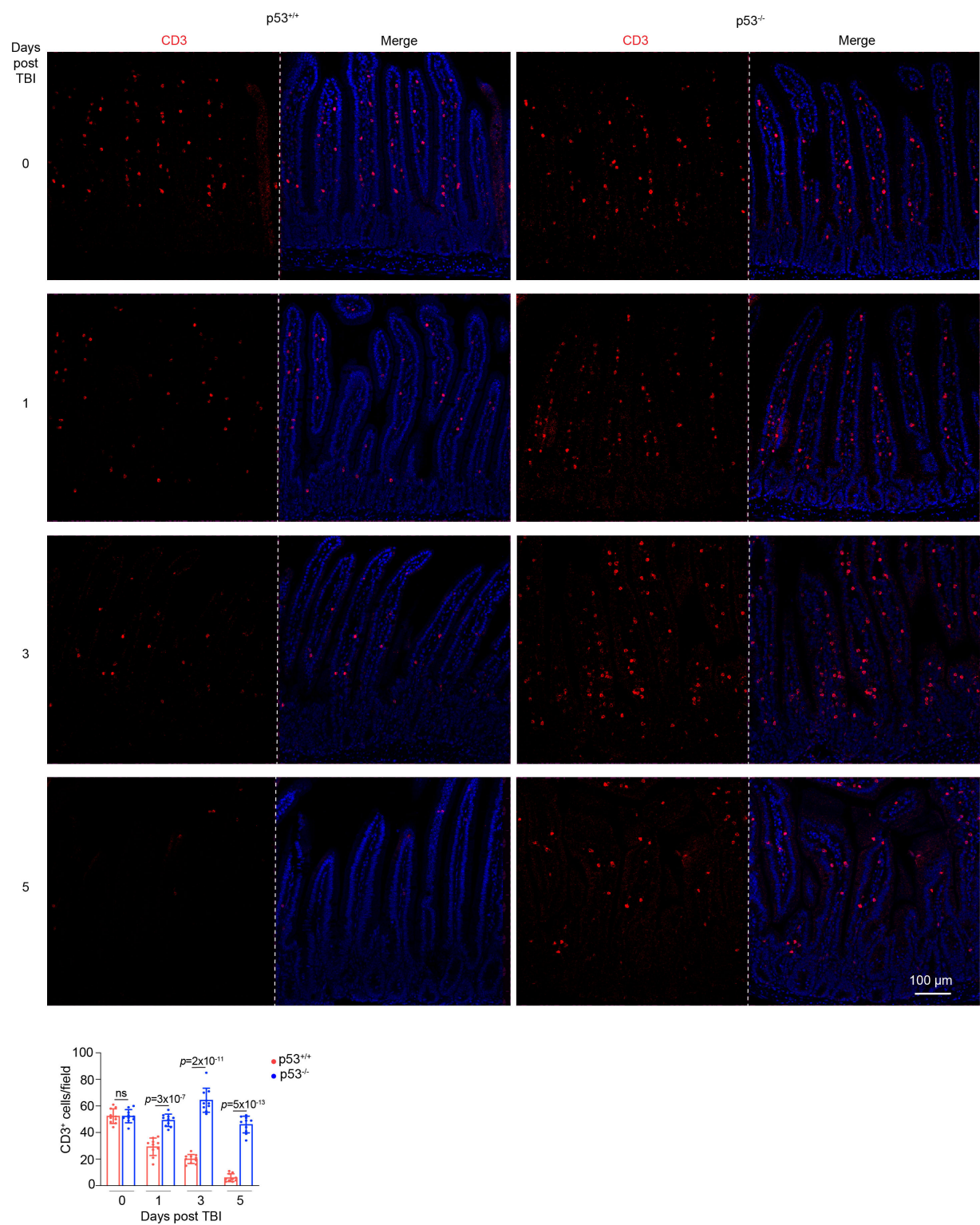


Figure S4. p53 deficiency enhances the number of CD3⁺ T cells in the SI of mice post-TBI.

Representative images from at least 3 independent mice (upper panels) and quantification (bottom panel) of IF staining of CD3 in the SI of p53^{+/+} and p53^{-/-} mice at different days post-TBI. n = 10 fields from at least 3 mice/group. Data are presented as mean \pm SD from at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test. Source data are provided as a Source Data file.

Supplementary Figure 5

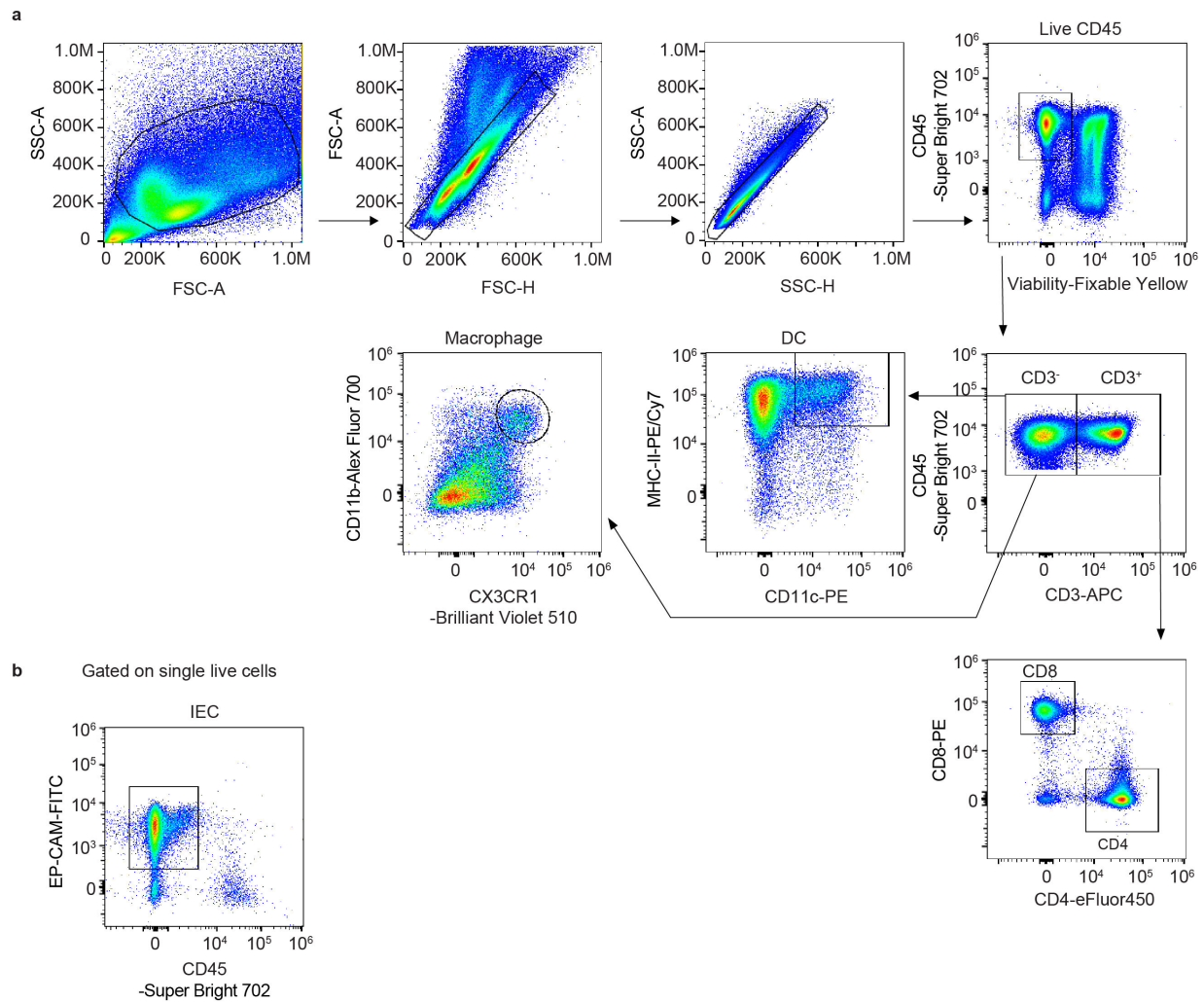


Figure S5. Gating strategies for different immune cell populations and IEC population.

a. Gating strategies for macrophages ($CD45^+CD3^-CX3CR1^+CD11b^+$), DCs ($CD45^+CD3^-MHC-II^+CD11c^+$), $CD8^+$ T cells ($CD45^+CD3^+CD8^+$), and $CD4^+$ T cells ($CD45^+CD3^+CD4^+$). **b.** Gating strategies for IECs ($CD45^+EpCAM^+$).

Supplementary Figure 6

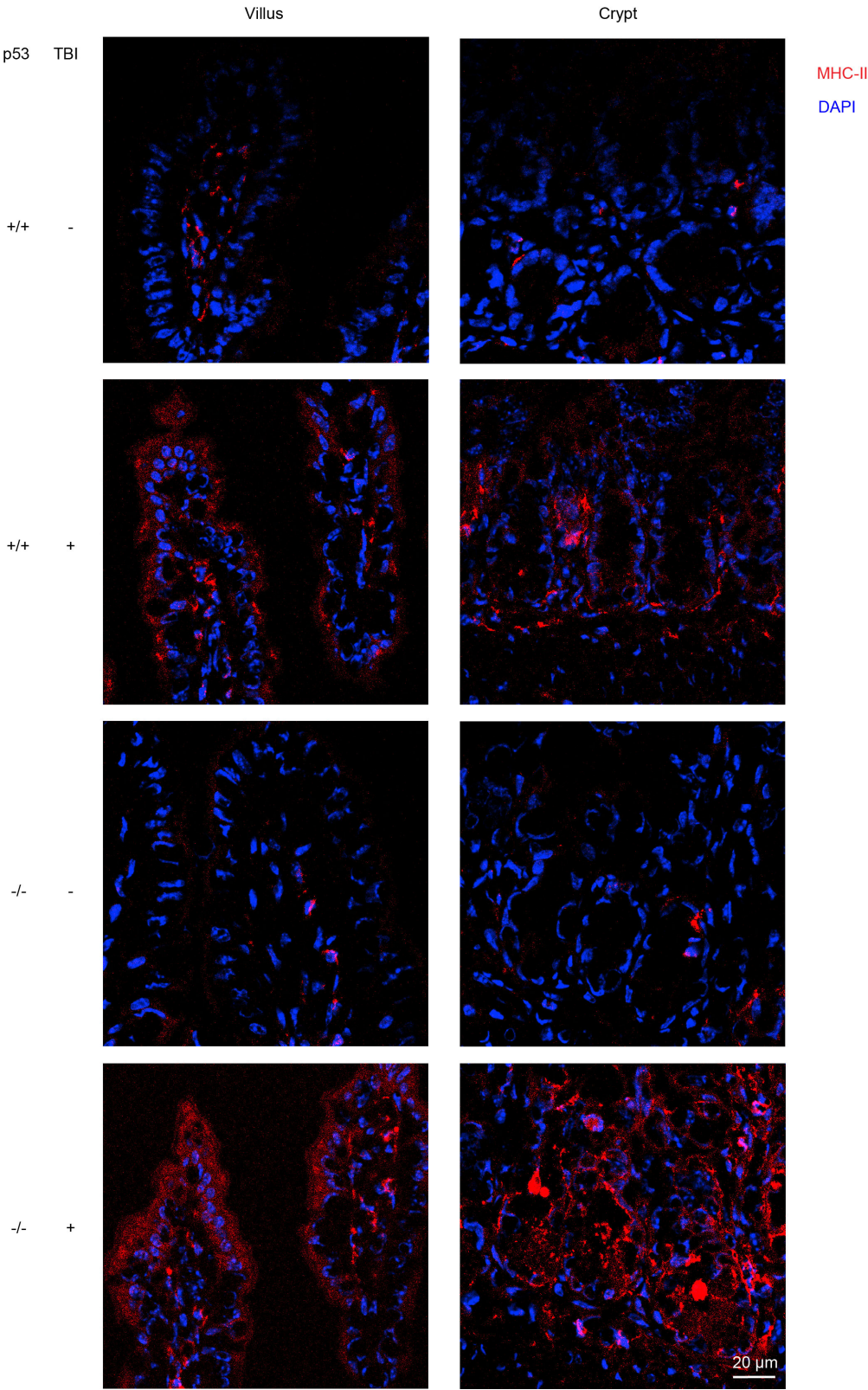


Figure S6. p53 deficiency induces MHC-II expression on both villus and crypt cells in the SI of mice post-TBI.

Representative images from at least 3 independent mice showing IF staining of MHC-II in the SI of p53^{+/+} and p53^{-/-} mice at 3 days post-TBI.

Supplementary Figure 7

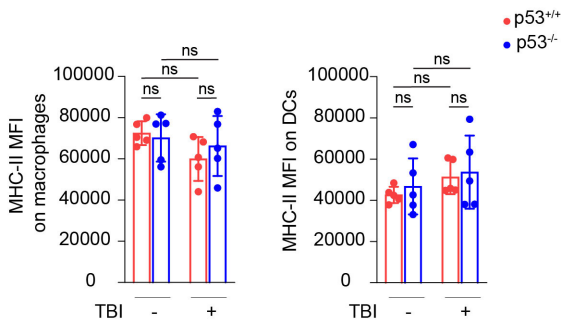


Figure S7. p53 deficiency has no obvious effect on MHC-II expression on macrophages and DCs in the LP of mice post-TBI.

Quantifications of MFI of MHC-II on macrophages (left panel) and DCs (right panel) in the LP of mice at 3 days post-TBI as determined by flow cytometric assays. $n = 5$ mice/group. Data are presented as mean \pm SD from at least 3 independent experiments. ns: not significant, two-tailed Student's t -test followed by Bonferroni correction. Source data are provided as a Source Data file.

Supplementary Figure 8

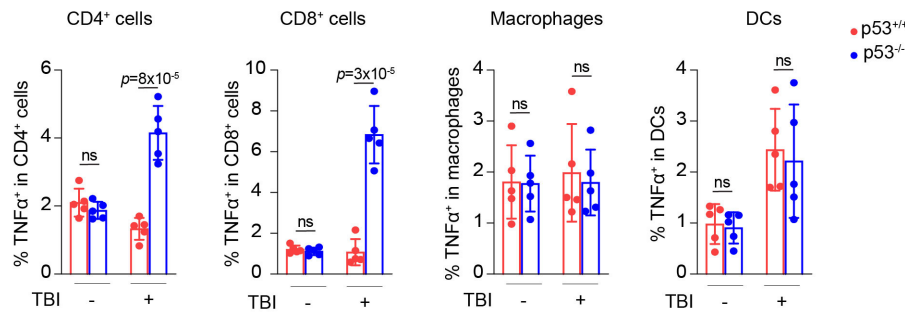


Figure S8. p53 deficiency promotes TNFα secretion from T cells in the LP of mice post-TBI.

Quantification of TNFα secretion from CD4⁺, CD8⁺, macrophages and DCs in the LP of mice at 3 days post-TBI as determined by flow cytometric assays. LP cells isolated from mice were stimulated with eBioscience™ Cell Stimulation Cocktail containing PMA, ionomycin, brefeldin A and monensin for 6 hours at 37°C. After cell surface staining, cells were fixed and permeabilized, followed by intracellular staining of TNFα. n = 5 mice/group. Data are presented as mean ± SD from at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test. Source data are provided as a Source Data file.

Supplementary Figure 9

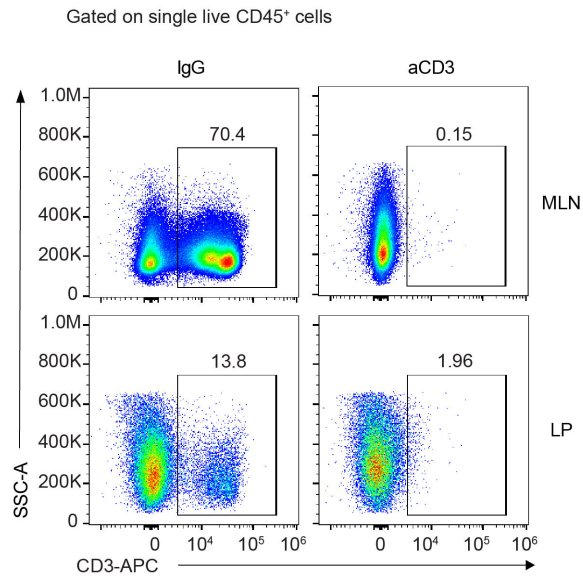


Figure S9. The depletion of CD3⁺ T cells in mice upon the aCD3 antibody administration.

Representative flow cytometric images from at least 3 independent mice showing the depletion of CD3⁺ T cells in the MLN and LP of p53^{+/+} mice at 6 days upon IgG or the aCD3 antibody (200 µg/mouse, once) treatment.

Supplementary Figure 10

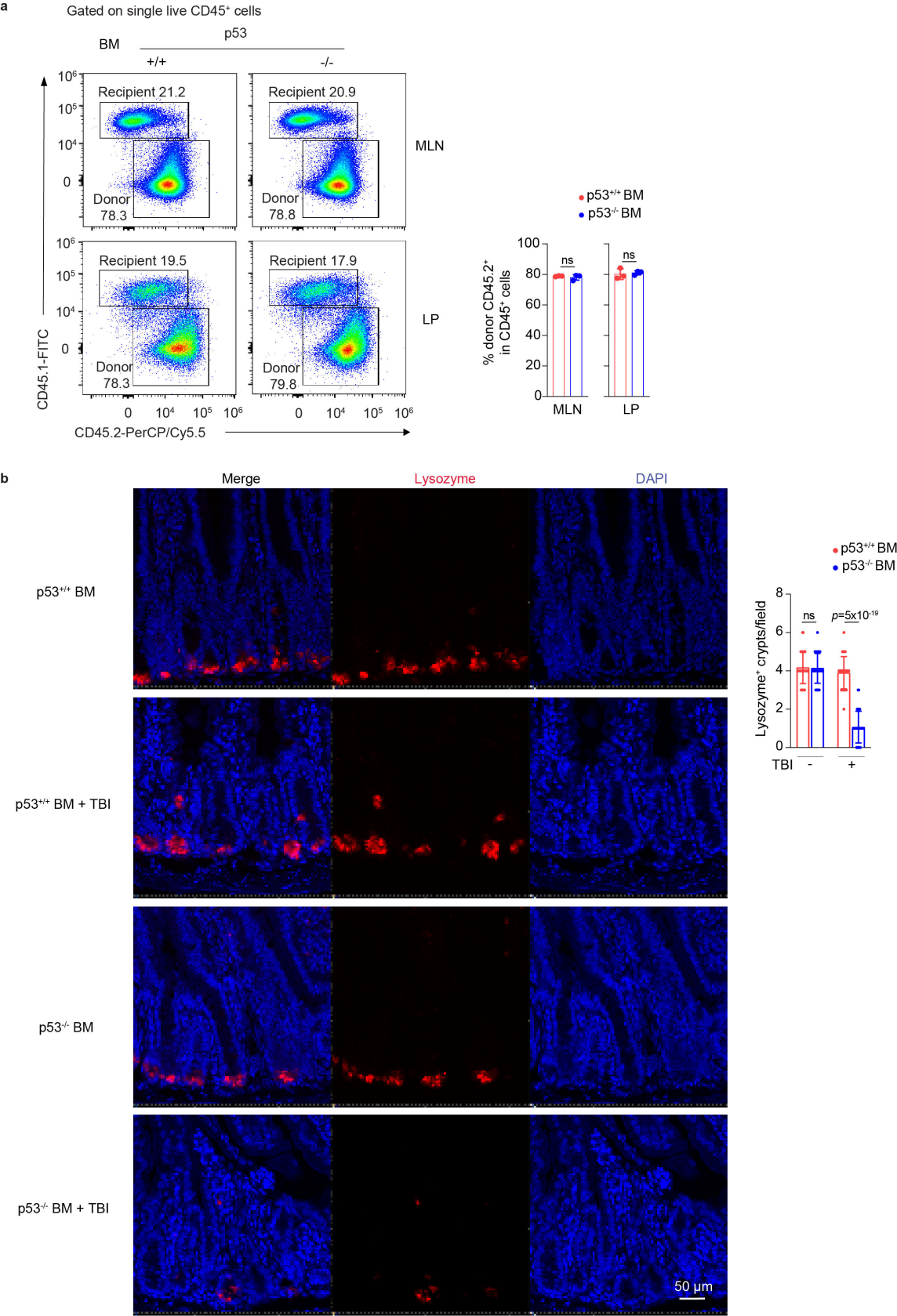


Figure S10. Adoptive transfer of the BM derived from p53^{-/-} mice into p53^{+/+} mice exacerbates the damage of Paneth cells post-TBI.

a. CD45.1⁺ mice were adoptively transferred with BM cells isolated from CD45.2⁺ p53^{+/+} and p53^{-/-} mice. The reconstitution of donor CD45.2⁺ CD45⁺ immune cells in the MLN and LP was examined at 28 days after adoptive transfer by flow cytometric assays. *n* = 3 mice/group. **b.** Representative images from at least 3 independent mice (left panels) and quantification (right panel) of IF staining of the lysozyme in the SI of mice at 5 days post 12 Gy TBI. *n* = 30 crypts from at least 3 mice/group. Data are presented as mean ± SD from at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test. Source data are provided as a Source Data file.

Supplementary Figure 11

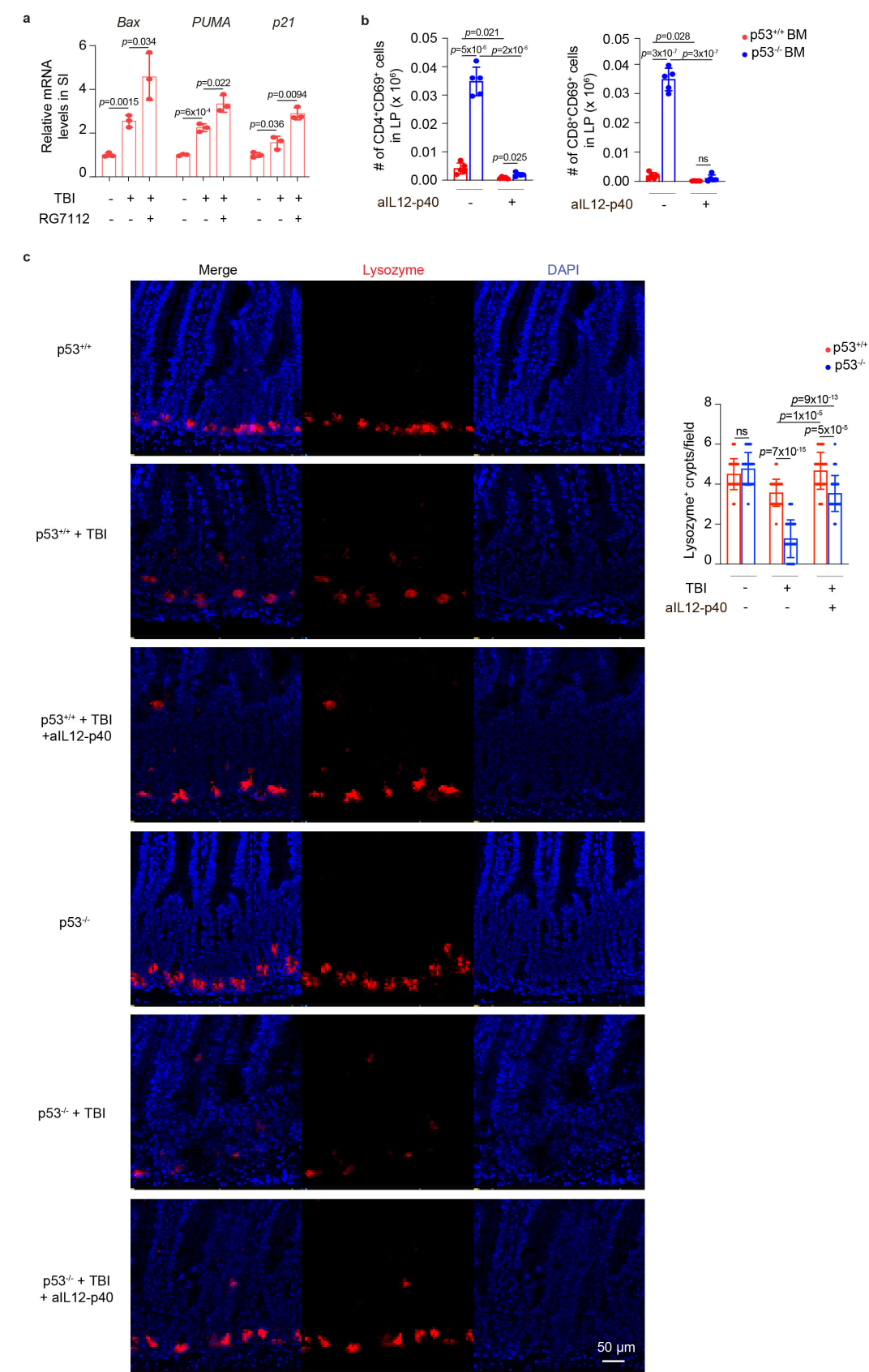


Figure S11. aIL12-p40 antibody reduces the damage of Paneth cells in the SI of mice post-TBI.

a. The mRNA levels of *BAX*, *PUMA* and *p21* in the SI of p53^{+/+} mice at 6 hours post 12 Gy TBI along with or without RG7112 treatment (*i.p.*, 50 µg/g body weight, twice, 8 hours before and 1 hour post-TBI) determined by qPCR assays. n = 3 mice/group. **b.** The number of CD69⁺ activated CD4⁺ (left panel) and CD8⁺ (right panel) T cells in the LP of mice at 3 days post-TBI determined by flow cytometric assays. n = 5 mice/group. **c.** Representative images from at least 3 independent mice (left panels) and quantification (right panel) of IF staining of lysozyme in the SI of mice at 3 days post-TBI. n = 30 crypts from at least 3 mice/group. Data are presented as mean ± SD from at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test followed by Bonferroni correction. Source data are provided as a Source Data file.

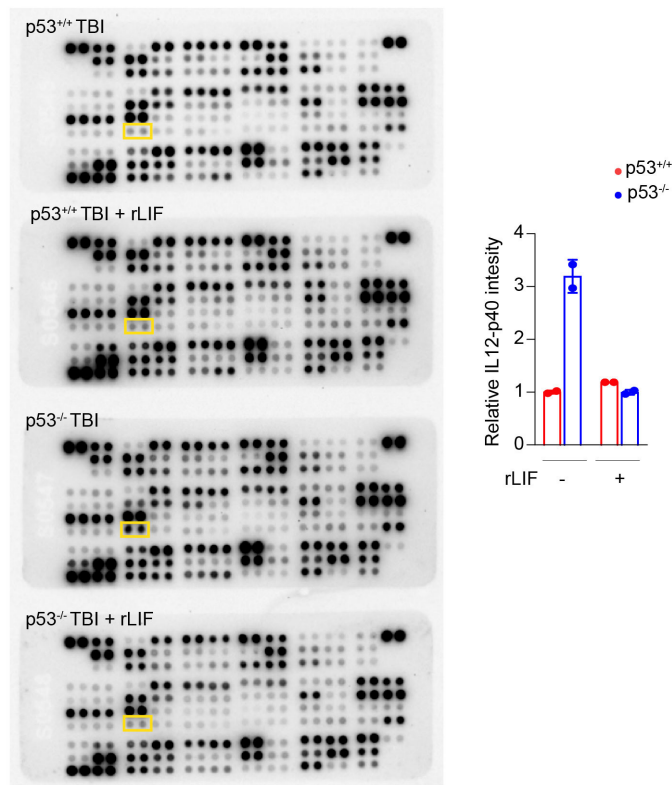


Figure S12. rLIF administration inhibits the IL12-p40 production in $p53^{-/-}$ mice post-TBI.

Left: representative images of cytokine levels in the serum from $p53^{+/+}$ and $p53^{-/-}$ mice with or without rLIF administration at 3 days post 12 Gy TBI detected by using the Proteome Profiler Mouse XL Cytokine Array. Right: quantification of IL12-p40 levels in Cytokine Array. Each array represents a pool of 3 mice with the same treatment. Dots in yellow box represent IL12-p40. Source data are provided as a Source Data file.

Supplementary Figure 13

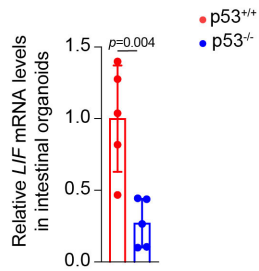


Figure S13. *p53* deficiency reduces *LIF* expression levels in intestinal organoids.

The mRNA levels of *LIF* in intestinal organoids derived from *p53*^{+/+} and *p53*^{-/-} mice determined by qPCR assays. $n = 5/\text{group}$. Data are presented as mean \pm SD from at least 3 independent experiments. Two-tailed Student's *t*-test. Source data are provided as a Source Data file.

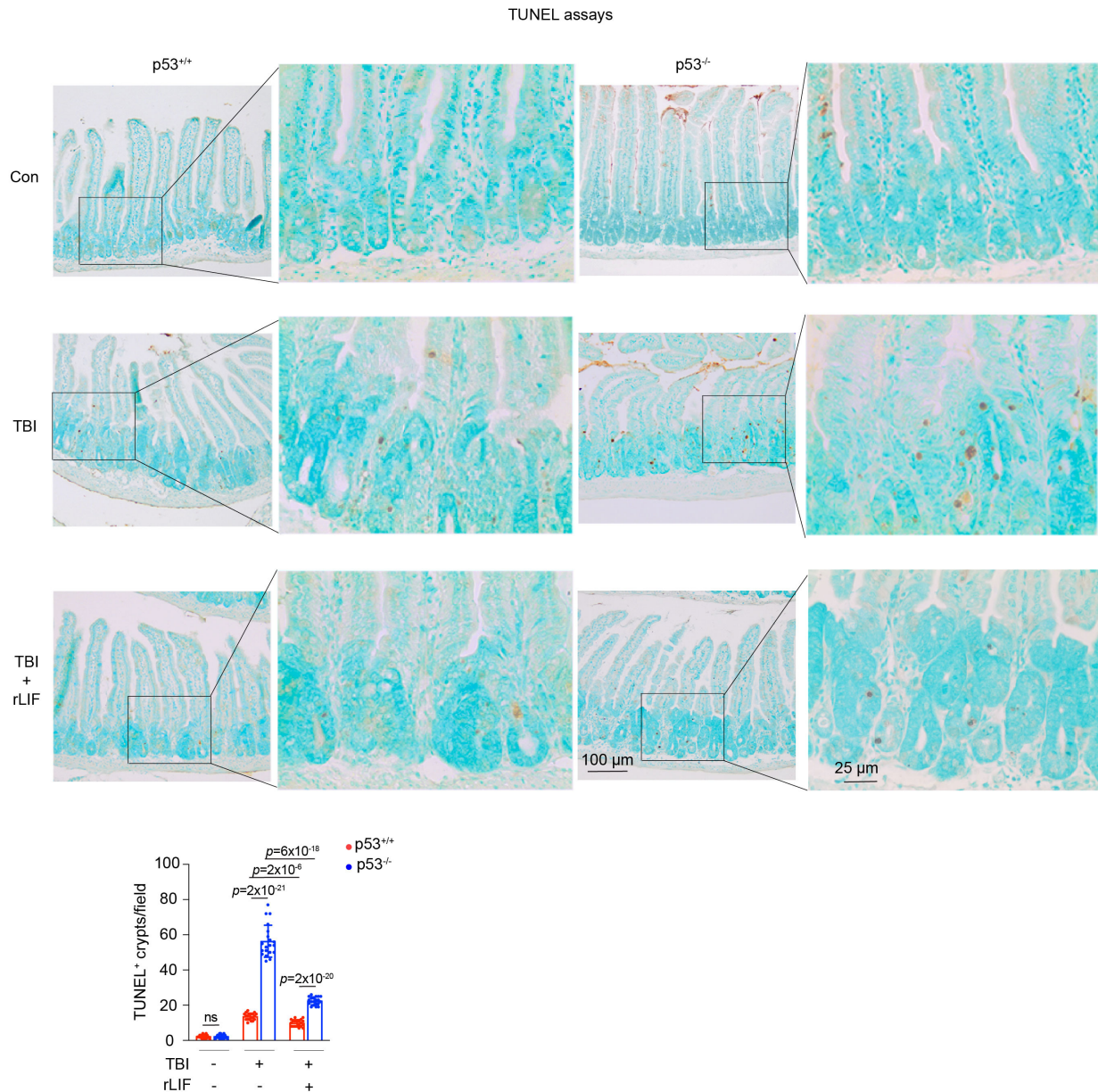


Figure S14. rLIF administration reduces cell death in the SI in $p53^{-/-}$ mice post-TBI.

Representative images from at least 3 independent mice (upper panels) and quantification (bottom panel) of TUNEL assays in the SI of $p53^{+/+}$ and $p53^{-/-}$ mice with or without rLIF administration at 3 days post 12 Gy TBI. $n = 20$ fields from at least 3 mice/group. Data are presented as mean \pm SD from at least 3 independent experiments. ns: not significant, two-tailed Student's t -test followed by Bonferroni correction. Source data are provided as a Source Data file.

Supplementary Figure 15

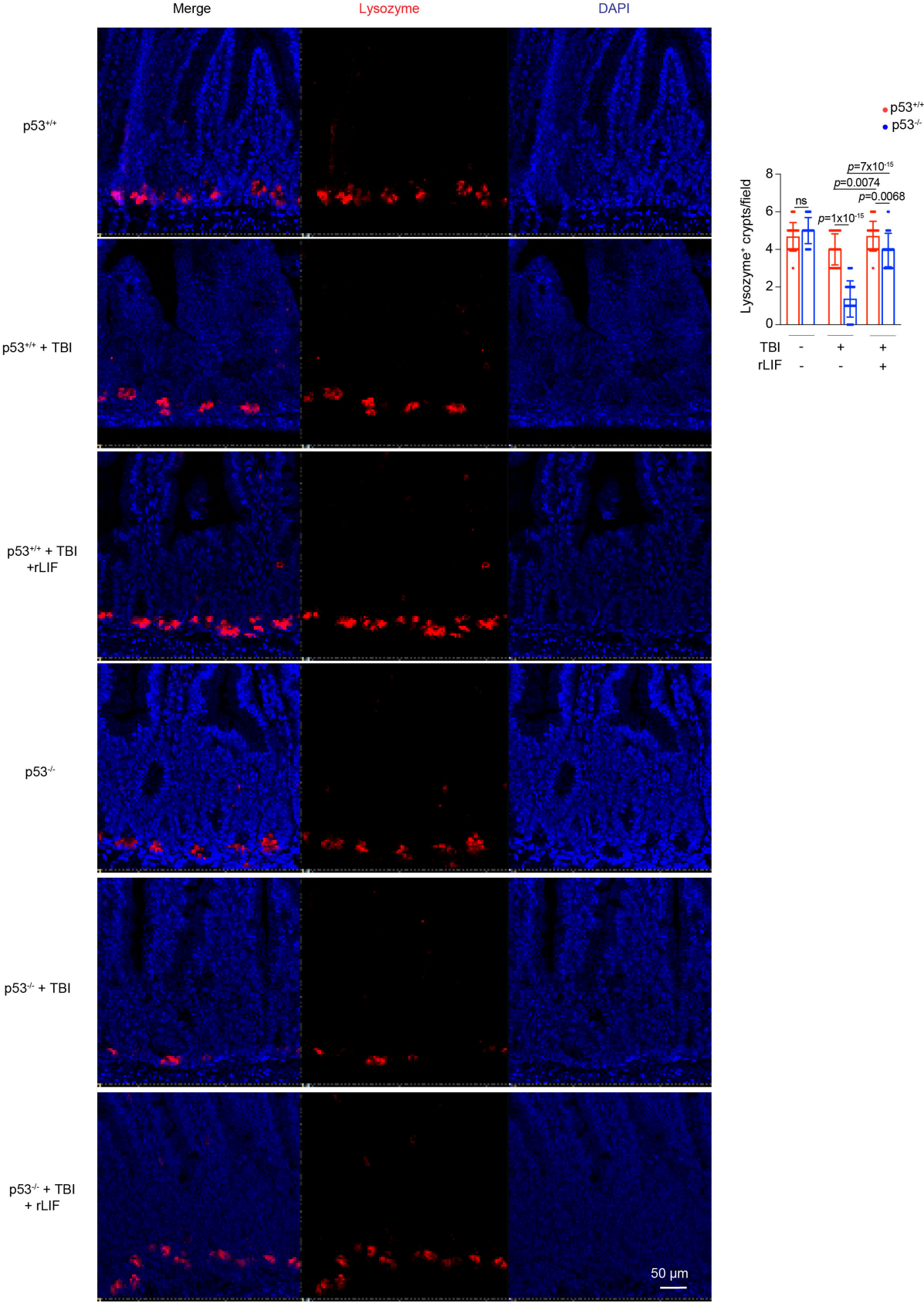


Figure S15. rLIF administration reduces the damage of Paneth cells in p53^{-/-} mice post-TBI. Representative images from at least 3 independent mice (left panels) and quantification (right panel) of IF staining of lysozyme in the SI of mice at 3 days post-TBI. n = 30 crypts from at least 3 mice/group. Data are presented as mean \pm SD from at least 3 independent experiments. ns: not significant, two-tailed Student's *t*-test followed by Bonferroni correction. Source data are provided as a Source Data file.

Supplementary Table S1. Information of flow antibodies used in this study

| Flow antibodies and reagents | | | |
|--|-------------------|------------|-------------|
| REAGENT or RESOURCE | SOURCE | Cata# | CLONE |
| eFluor450 anti-mouse CD4 | eBioscience | 48-0042-82 | RM4-5 |
| Brilliant Violet 510™ anti-mouse CX3CR1 | Biolegend | 149025 | SA011F11 |
| Super Bright 702 anti-mouse CD45 | Thermo Fisher | 67-0451-82 | 30-F11 |
| FITC anti-mouse Ep-CAM | Biolegend | 118208 | G8.8 |
| FITC anti-mouse CD45.1 | Biolegend | 110706 | A20 |
| Per/Cy5.5 anti-mouse CD45.2 | Biolegend | 109828 | 104 |
| PE anti-mouse CD11c | Biolegend | 117308 | N418 |
| PE anti-mouse CD8 | Biolegend | 100708 | 53-6.7 |
| PE/Cy7 anti-mouse MHC-II | Biolegend | 107630 | M5/114.15.2 |
| APC anti-mouse CD3 | Biolegend | 100236 | 17A2 |
| PE/Cy7 anti-mouse CD25 | BD | 552880 | PC61 |
| Alex Fluor 700 anti-mouse CD69 | Biolegend | 104539 | H1.2F3 |
| Alex Fluor 700 anti-mouse CD11b | Biolegend | 101222 | M1/70 |
| Alex Fluor 700 anti-mouse TNF α | Biolegend | 506338 | MP6-XT22 |
| Cell Stimulation Cocktail | Thermo Scientific | 00-4975-03 | N/A |
| BD cytofix/cytoperm Fixation/Permeabilization Solution Kit | BD | 554714 | N/A |
| Ultra-LEAF™ Purified anti-mouse CD16/32 | Biolegend | 101339 | 93 |
| LIVE/DEAD™ Fixable Yellow Dead Cell Stain Kit | Thermo Fisher | L34968 | N/A |
| Abc Total Antibody Compensation Bead Kit | Thermo Fisher | A10497 | N/A |

Supplementary Table S2. Sequences of qPCR primers

| Gene | Forward Primer | Reverse Primer |
|----------------|--------------------------------------|--------------------------------|
| β -actin | 5' GAACCCTAAGGCCAACCGTGAAAAGATGAC 3' | 5' GCAGGATGGCGTGAGGGAGAGCA 3' |
| TNF α | 5' CTGAACTTCGGGGTGATCGG 3' | 5' GGCTTGTCACCTCGAATTTTGAGA 3' |
| BAX | 5' TGAAGACAGGGGCCTTTTGTG 3' | 5' AATTCGCCGGAGACACTCG 3' |
| PUMA | 5' GTCGCTACCGTCGTGACTTC 3' | 5' CAGACATGCACCTACCCAGC 3' |
| p21 | 5' CCTGGTGATGTCCGACCTG 3' | 5' CCATGAGCGCATCGCAATC 3' |