

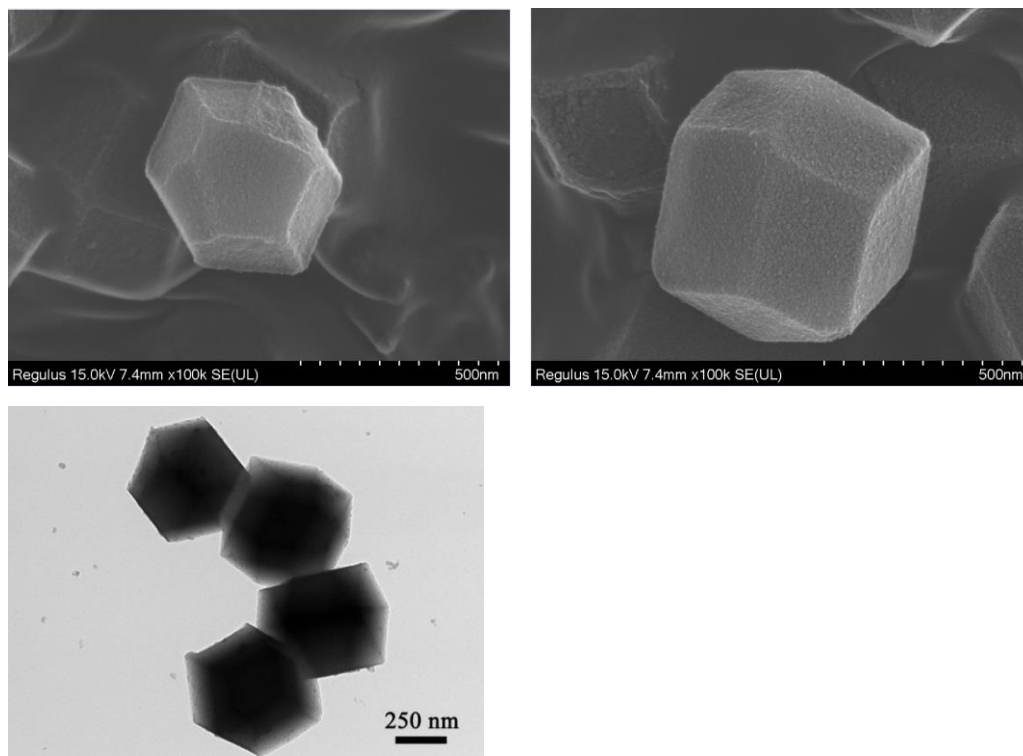
## Supporting Information

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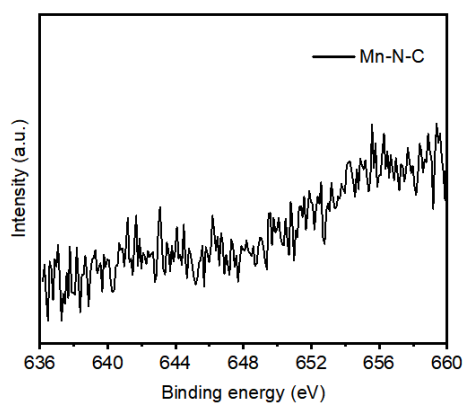
A Single-Atom Manganese Nanozyme Mn-N/C Promotes Anti-Tumor Immune Response via Eliciting Type I Interferon Signaling

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A



B



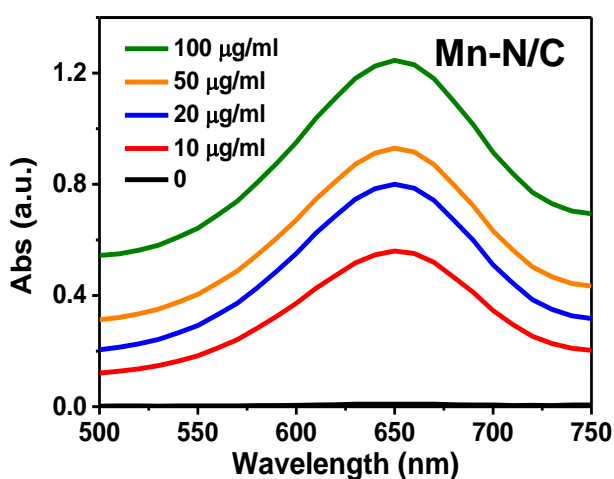
**Figure S1. Characterization of single-atom manganese nanozyme Mn-N/C.**

**A.** High-resolution SEM and TEM images of Mn-N/C.

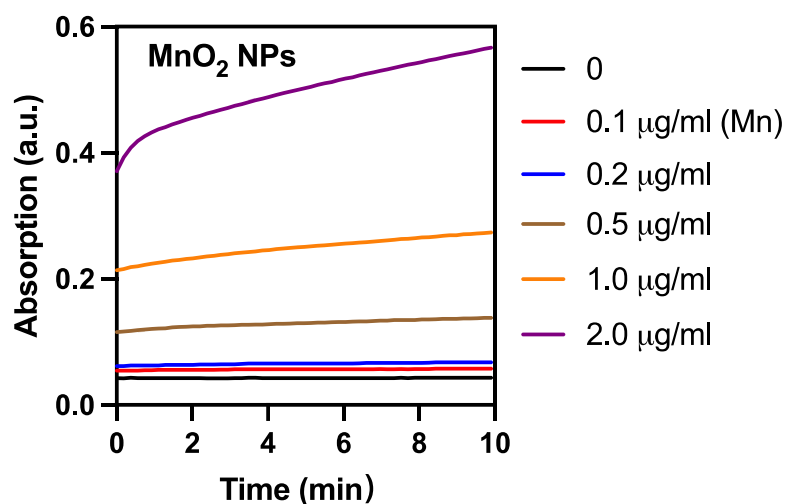
**B.** XRD data of Mn-N/C, showing their amorphous feature.

**Figure S2**

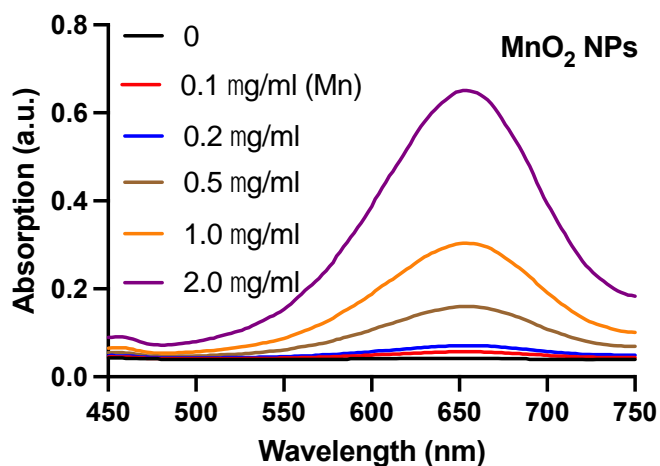
**A**



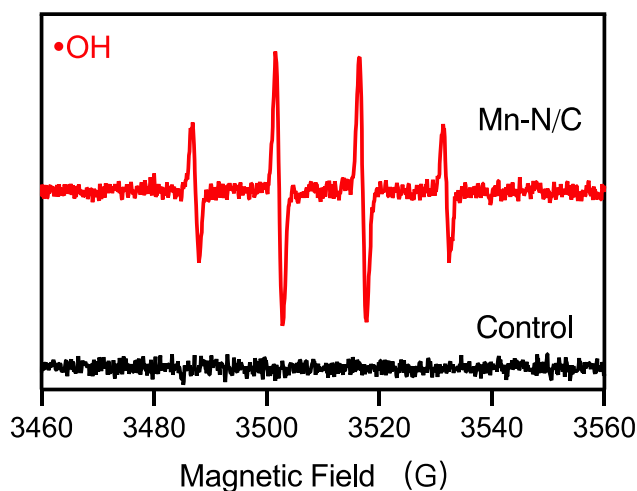
**B**



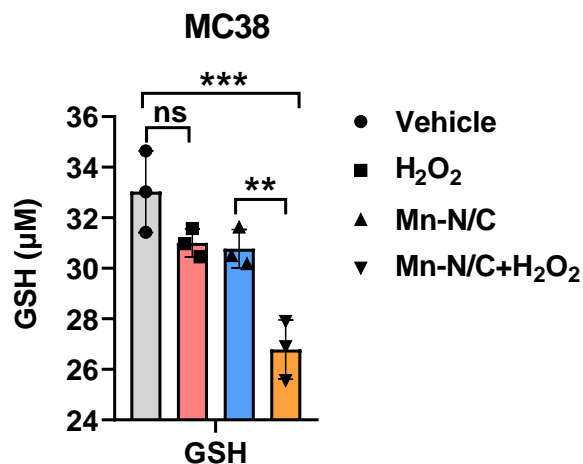
**C**



**D**



**E**



**Figure S2. The peroxidase-like activity of Mn-N/C and ROS generation.**

**A.** The absorbance at different wavelengths in the absence (black) and presence of different concentrations of Mn-N/C.

**B.** The time-dependent absorbance changes of TMB (652 nm) in the absence (black) or presence of different concentrations of MnO<sub>2</sub>.

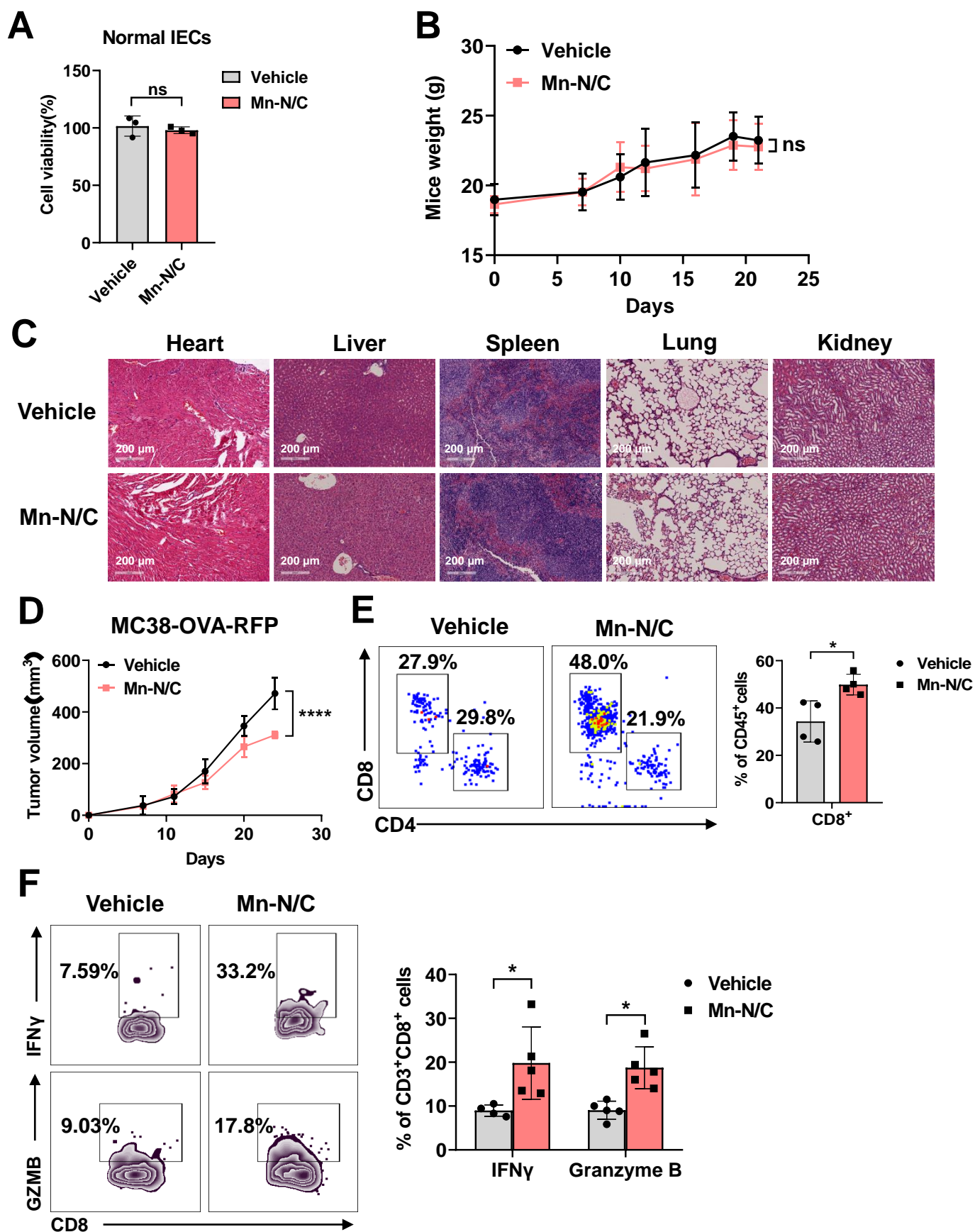
**C.** The UV-Vis spectrum of TMB in the absence or presence of different concentrations of MnO<sub>2</sub>.

**D.** EPR spectra for detection of •OH by DMPO.

**E.** GSH levels in MC38 tumors treated with H<sub>2</sub>O<sub>2</sub> or Mn-N/C (100 µg/mL) for 24 hr. Data are shown as mean ± SEM (n=3). Two-tailed unpaired t-test was performed for the comparisons between two groups.

\*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; ns, no significance.

**Figure S3**



**Figure S3. Mn-N/C suppresses tumor growth without tissue damage.**

**A.** The cell viability of normal intestinal epithelial cells (IECs) treated with vehicle or Mn-N/C for 24 hr by CCK8 assays. Quantitative data are shown as mean $\pm$ SEM (n=3). Two-tailed unpaired t-test was performed for the comparison between two groups.

**B.** Body weight curve of CT26 tumor-bearing mice treated with vehicle or Mn-N/C. Data are shown as mean $\pm$ SEM (n=4-5 mice per group). Two-way ANOVA (mixed model) and Sidak's multiple comparisons test were performed.

**C.** Representative IHC images of heart, liver, spleen, lung and kidney of CT26 tumor-bearing mice treated with vehicle or Mn-N/C.

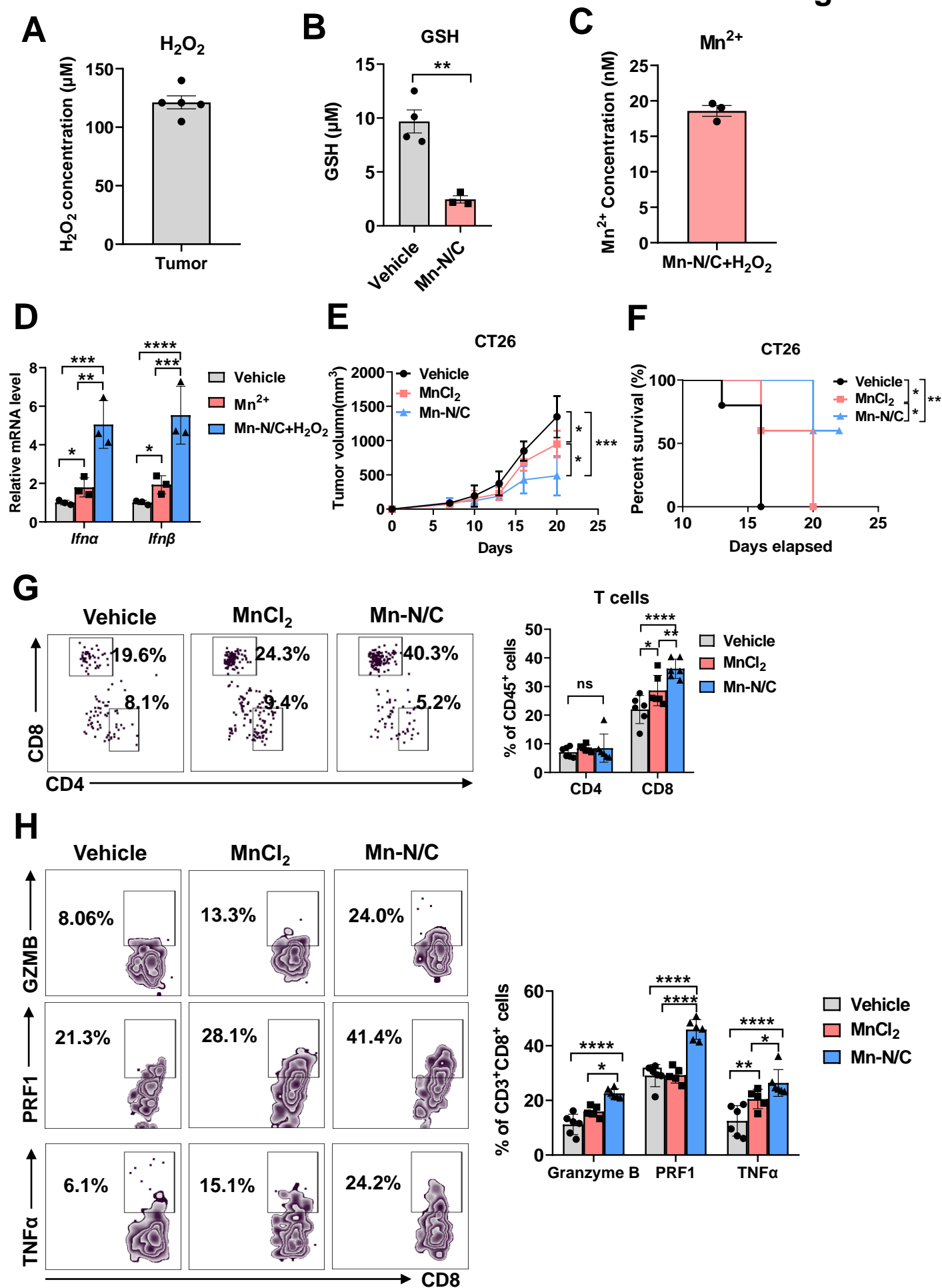
**D.** Tumor growth curve of MC38-OVA-RFP in the WT mice treated with vehicle or Mn-N/C. Data are shown as mean $\pm$ SEM (n=5 mice per group). Two-way ANOVA (mixed model) and Sidak's multiple comparisons test were performed.

**E.** FACS analysis (left) and quantification (right) of tumor-infiltrating CD8<sup>+</sup> T cells in the mice treated with vehicle or Mn-N/C as **D** described. Data are shown as mean $\pm$ SEM (n=4 mice per group). Two-tailed unpaired t-test was performed for the comparisons between two groups.

**F.** FACS analysis (left) and quantification (right) of IFN $\gamma$ <sup>+</sup> and Granzyme B<sup>+</sup> of tumor-infiltrating CD8<sup>+</sup> T cells in the mice treated with vehicle or Mn-N/C as **D** described. Data are shown as mean $\pm$ SEM (n=4-5 mice per group). Two-way ANOVA and Sidak's multiple comparisons test were performed.

\*,  $P<0.05$ ; \*\*\*\*,  $P<0.0001$ ; ns, no significance.

Figure S4



**Figure S4. Mn-N/C displayed superior tumor suppression compared to MnCl<sub>2</sub>.**

**A.** H<sub>2</sub>O<sub>2</sub> concentration in CT26 tumor tissues. Data are shown as mean±SEM (n= 5 mice).

**B.** GSH levels in CT26 tumor tissue following vehicle or Mn-N/C treatment. Data are shown as mean±SEM (n=3-4 mice per group). Two-tailed unpaired t-test was performed for the comparisons between two groups.

**C.** Mn<sup>2+</sup> concentration in the supernatant of MC38 tumor cells treated with Mn-N/C and H<sub>2</sub>O<sub>2</sub> for 24 hr. After treatment, supernatants were subjected to high-speed centrifugation to remove residue Mn-N/C. The content of dissociated Mn<sup>2+</sup> in the supernatant was determined by ICP-MS. Data are shown as mean±SEM (n=3).

**D.** Real-time PCR analysis of *Ifnα* and *Ifnβ* gene expression levels in MC38 tumor cells treated with Mn<sup>2+</sup> (20 nM) or Mn-N/C combined with H<sub>2</sub>O<sub>2</sub> for 24 hr. Data are shown as mean±SEM (n=3). Two-way ANOVA and Sidak's multiple comparisons test were performed.

**E.** Tumor growth curve of CT26 in the WT mice intratumorally injected with vehicle, MnCl<sub>2</sub> or Mn-N/C. Data are shown as mean±SEM (n=5 mice per group). Two-way ANOVA (mixed model) and Sidak's multiple comparisons test were performed.

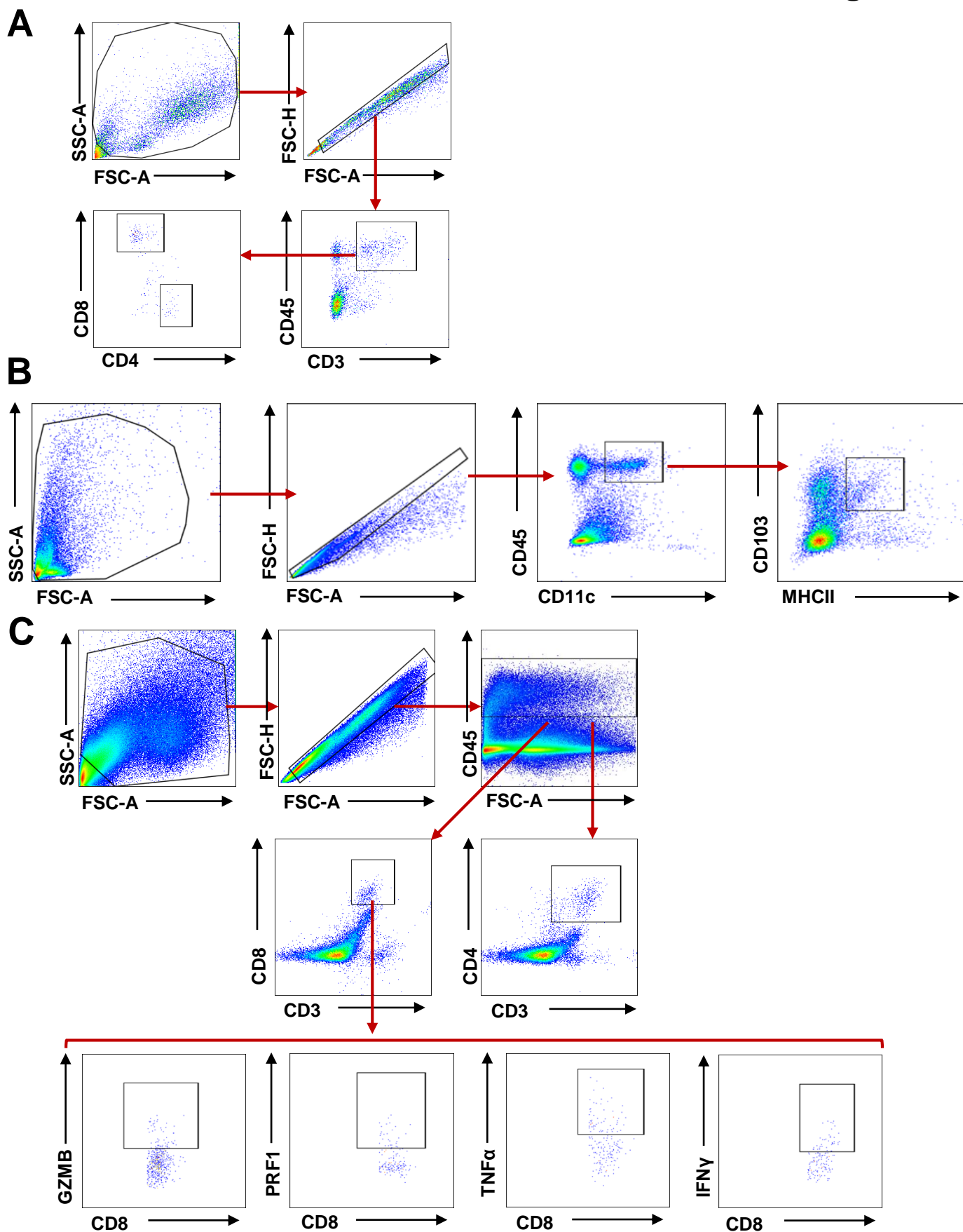
**F.** Kaplan–Meier analysis of mice survival after vehicle, MnCl<sub>2</sub> or Mn-N/C treatment as described in **E** (n=5 mice per group).

**G.** FACS analysis (left) and quantification (right) of tumor-infiltrating CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the mice treated with vehicle, MnCl<sub>2</sub>, or Mn-N/C. Data are shown as mean±SEM (n=6 mice per group). Two-way ANOVA and Sidak's multiple comparisons test were performed.

**H.** FACS analysis (left) and quantification (right) of Granzyme B<sup>+</sup>, PRF1<sup>+</sup>, TNFα<sup>+</sup> of tumor-infiltrating CD8<sup>+</sup> T cells in the mice treated with vehicle, MnCl<sub>2</sub>, or Mn-N/C. Data are shown as mean±SEM (n=6 mice per group). Two-way ANOVA and Sidak's multiple comparisons test were performed.

\*,  $P<0.05$ ; \*\*,  $P<0.01$ ; \*\*\*,  $P<0.001$ ; \*\*\*\*,  $P<0.0001$





**Figure S5. The gating strategies for flow cytometry.**

**A.** Gating strategy for tumor-infiltrating CD4<sup>+</sup> and CD8<sup>+</sup>T lymphocytes.

**B.** Gating strategy for DCs.

**C.** Gating strategy for cytotoxic molecules of tumor-infiltrating CD8<sup>+</sup>T cells.

**Table S1****Primers used in RT-PCR**

<b>Primer name</b>	<b>Sequence (5'-3')</b>
Murine <i>Gapdh</i> -F	AGCTTGTCATCAACGGGAAG
Murine <i>Gapdh</i> -R	TTTGATGTTAGTGGGGTCTCG
Murine <i>Ifna</i> -F	ATTTCCCCTGACCCAGGAAGATG
Murine <i>Ifna</i> -R	CCCAGCACATTGGCAGAGG
Murine <i>Ifnb</i> -F	AGCTCCAAGAAAGGACGAACA
Murine <i>Ifnb</i> -R	GCCCTGTAGGTGAGGTTGAT
Murine <i>Cxcl10</i> -F	CCAAGTGCTGCCGTCATTTTC
Murine <i>Cxcl10</i> -R	GGCTCGCAGGGATGATTTCAA
Murine <i>Hmgb1</i> -F	GGCGAGCATCCTGGCTTATC
Murine <i>Hmgb1</i> -R	GGCTGCTTGTCATCTGCTG
Murine <i>Ddit3</i> -F	CTGGAAGCCTGGTATGAGGAT
Murine <i>Ddit3</i> -R	CAGGGTCAAGAGTAGTGAAGGT
Murine <i>Irf7</i> -F	TACCATCTACCTGGGCTTCG
Murine <i>Irf7</i> -R	GCTCCATAAGGAAGCACTCG
Murine <i>Pdl1</i> -F	GCTCCAAAGGACTTGTACGTG
Murine <i>Pdl1</i> -R	TGATCTGAAGGGCAGCATTTTC
Murine <i>Isg15</i> -F	GGAACGAAAGGGGCCACAGCA
Murine <i>Isg15</i> -R	CCTCCATGGGCCTTCCCTCGA
Murine gDNA <i>Tert</i> -F	CTAGCTCATGTGTCAAGACCCTCTT
Murine gDNA <i>Tert</i> -R	GCCAGCACGTTTCTCTCGTT
Murine gDNA <i>Polg1</i> -F	GATGAATGGGCCTACCTTGA
Murine gDNA <i>Polg1</i> -R	TGGGGTCCTGTTTCTACAGC
Murine mtDNA <i>Dloop 1</i> -F	CCCTTCCCCATTTGGTCT
Murine mtDNA <i>Dloop 1</i> -R	TGGTTTCACGGAGGATGG
Murine mtDNA <i>Dloop 2</i> -F	CCCTTCCCCATTTGGTCT
Murine mtDNA <i>Dloop 2</i> -R	TGGTTTCACGGAGGATGG