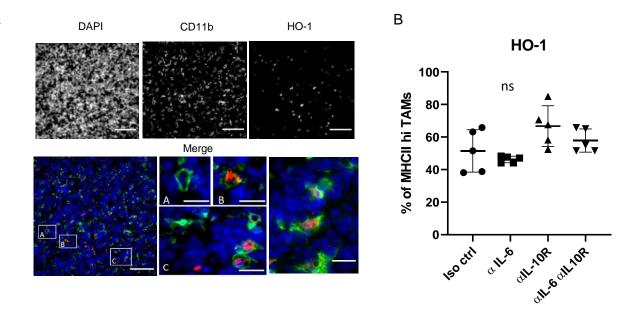
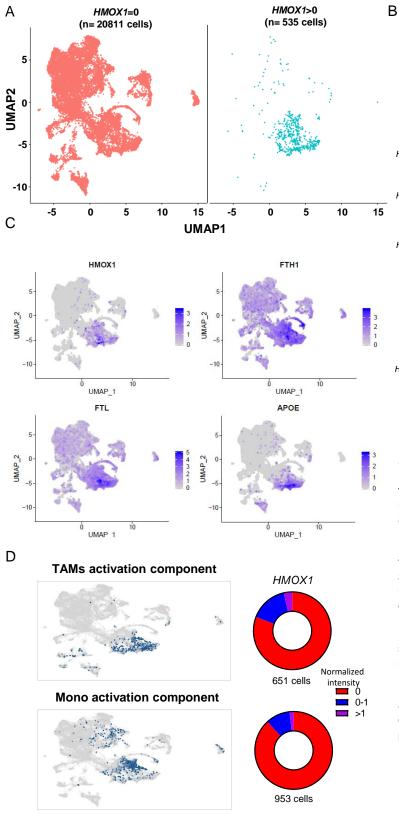
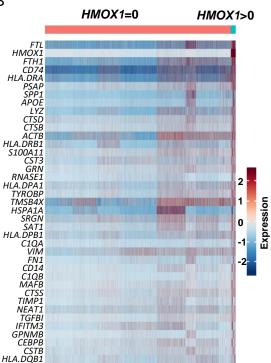
## Figure S1



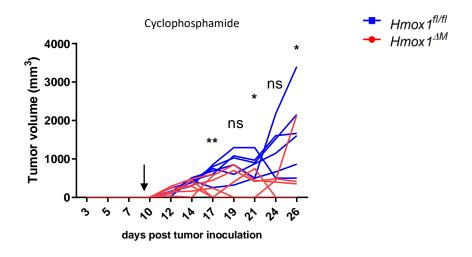
**Supplemental Figure 1. HO-1 expression by TAMs.** A) HO-1 staining (in red) combined with DAPI co-staining showing nuclei (in blue) are visualized in tumor slices by immunofluorescence in CD11b<sup>+</sup> myeloid cells (in green) in an EG7-OVA tumor 21 days after tumor inoculation in a wild type mouse. Scale bar = 5  $\mu$ m. B) HO-1 expression measured by flow cytometry among CD11b<sup>hi</sup>Ly6G<sup>-</sup>Ly6C<sup>lo</sup>CD64<sup>+</sup>MHCII<sup>+</sup> TAMs from EG7-OVA tumor 16 days after inoculation in wild type mice, after in vivo neutralization of IL-6, IL-10, or both (n=5). \*P<0,05; \*\*P<0,01; \*\*\*P<0,001; \*\*\*\*P<0,0001, ns: not significant. .Kruskall-Wallis and Dunn's multiple comparisons test.

## Figure S2

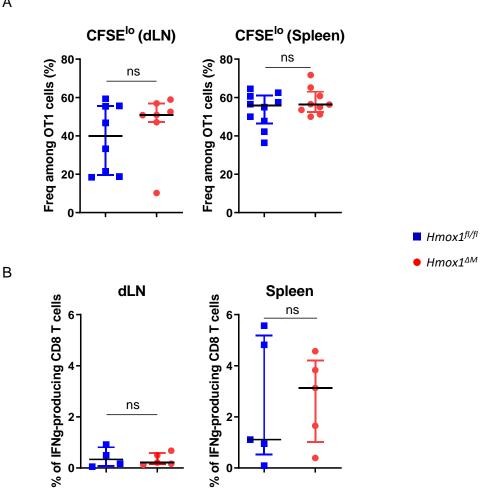




Supplemental Figure 2. Characterization of HMOX1-expressing cells in human breast tumors (A) HMOX1 positive (green) and negative (orange) cells were identified based on absolute count of HMOX1 in each cell and represented on UMAP plot. (B) Heatmap representing the expression of the genes that of were significantly upregulated in HMOX1+ cells as compared to HMOX1- cells. Gene expression is color-coded, using a scale based on z-score distribution (C) Normalized expression of representative genes that were associated with HMOX1 expression. Coordinates of cells are the same as in A. (D) The enrichment for TAMs and monocytes activation gene sets were plotted on UMAP with the same cells coordinates as in A using AUCell R package. Blue dots represent cells passing AUC threshold (0.2 for TAMs and 0.3 for Monocytes). Pie charts represent the proportion of HMOX1<sup>+</sup> cells among each subset.



**Supplemental Figure 3. Myeloid HO-1 inhibition improves the antitumor effect of cyclophosphamide treatment.** EG7-OVA tumor cells were inoculated intradermally at day 0 on the right flank of  $Hmox1^{\Delta M}$  mice (n = 6). All mice were injected with cyclophosphamide (i.p.) at day 10 post tumor inoculation. Their tumor volumes were compared to  $Hmox1^{fl/fl}$  littermates (n = 6) at regular intervals following implantation. Statistical analysis was performed with Mann-Whitney U test. \*P<0,05; \*\*P<0,01; \*\*\*P<0,001; \*\*\*\*P<0,0001.



Supplemental Figure 4. Myeloid HO-1 inhibition has no impact on T-cell priming in the secondary lymphoid organs. Intravenous adoptive transfer of CFSE labelled OT-1 cells (2x10<sup>6</sup> cells/mouse) was performed 10 days after tumor inoculation. This was followed by an immunization with subcutaneous injection of ovalbumin protein (50 µg/mouse) and poly(I:C) (50 µg/mouse) one hour later on the right flank of the animals. Two days later, EG7-OVA tumors were enzymatically and mechanistically digested and analyzed by flow cytometry. A) Tumor-infiltrating OT-1 cell proliferation assessed by CFSE dilution in the axillary and inguinal draining lymph nodes on the right flank and the spleen of Hmox1<sup>ΔM</sup> mice and Hmox1<sup>fl/fl</sup> littermates mice. dLN: n=8 (littermates) or n=7 (Hmox1<sup>ΔM</sup> mice). Spleen: n=10 (littermates) or n=9 (Hmox1<sup>ΔM</sup> mice). B) Production of IFNr was assessed by ex vivo stimulation overnight with OVA SIINFEKL peptide (and brefeldin A added 2 hours later). dLN: n=4 (littermates) or n=5 (Hmox1<sup>ΔM</sup> mice). Spleen: n=5 (littermates) or n=5 (Hmox1<sup>ΔM</sup> mice). Horizontal bars indicate median ± interquartile. Statistical analysis was performed with Mann-Whitney U test. \*P<0.05; \*\*P<0.01; \*\*\*\*P<0.001; \*\*\*\*P<0.0001.