

Inhibiting Ehmt2/Ezh2 histone methyltransferases enhances immune microenvironment in a *Trp53*^{-/-} murine ovarian cancer model

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Background

- Ovarian high grade serous carcinoma (HGSC) is a poor prognosis disease with mutations in *TP53* found in around c.100% of all cases¹.
- Ovarian cancer prognosis is strongly associated with development of an anti-tumour immune response². However, tumours can employ epigenetic mechanisms to silence immunostimulatory genes, such as chemokines³.
- We investigated whether a novel dual inhibitor of the histone methyltransferases Ehmt2/Ezh2 was able to derepress expression of critical chemokines and augment the immune response in a murine ovarian cancer model.

Methods

- Trp53*^{-/-} ID8 cells were previously generated by CRISPR-Cas9 technique⁴.
- An 84-chemokine RT-qPCR array was used to analyse *in vitro* transcript changes following treatment with an Ehmt2/Ezh2 inhibitor.
- C57BL/6 female mice were inoculated with *Trp53*^{-/-} ID8 cells intraperitoneally (IP). Following this, tumours and haemorrhagic ascites develops approximately 40-45 days post-inoculation. Treatments were delivered IP.
- Following tumour dissociation and digestion, cells were stained for flow cytometry and analysed using Cytex Aurora flow cytometer.

Result 1: Chemokine gene upregulation by Ehmt2/Ezh2 inhibition

- In vitro*, combination Ehmt2/Ezh2 treatment significantly ($p < .05$) upregulates the expression of lymphocyte-attractant chemokines, such as *CXCL10* (3-fold), *CXCL9* (22-fold) and *CCL5* (14-fold), after stimulation with IFN γ (fig. 1)

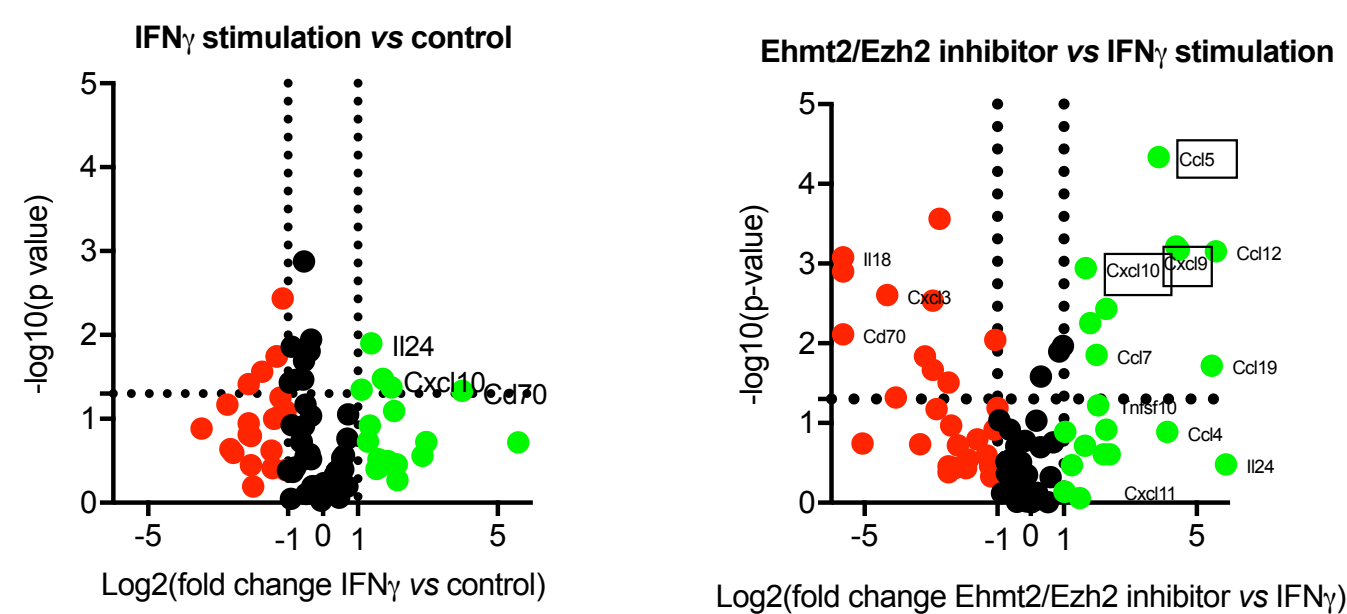


Figure 1: *Trp53*^{-/-} ID8 cells a) stimulated with 1ng/ml IFN γ alone or b) stimulated with FN γ and treated with 6uM of dual Ehmt2/Ezh2 inhibitor. Green colour denotes increase, red colour decrease and black colour no change.

Result 2: Dual Ehmt2/Ezh2 blockade confers a survival advantage

- In vivo*, dual Ehmt2/Ezh2 inhibition confers a survival advantage over control (52d vs 45d, $p < .0001$) (fig 2a). Tumour weight and ascites volume were significantly lower in the group treated with the dual inhibitor (mean 120mg vs 178mg, $p = .006$; 3.7ml vs 5.6ml, $p = .003$) compared to control (fig 2b).

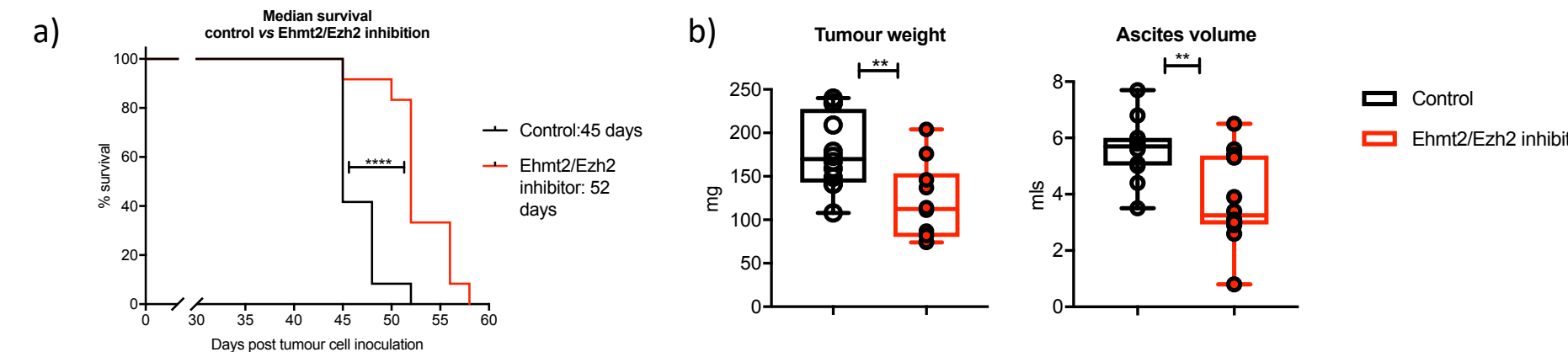


Figure 2: Female C57BL/6 mice were injected IP with *Trp53*^{-/-} ID8 cells on day 0 and treated with a dual Ehmt2/Ezh2 inhibitor 20mg/kg BD (n=12) or vehicle (0.9% NaCl, 1% Tween and 3.5% DMSO, n=12) for 14 days via IP injection. Subsequently, mice were allowed to reach humane endpoint. At endpoint, weight and ascites volume were measured.

Result 3: Ehmt2/Ezh2 inhibition shapes the immune microenvironment within tumour deposits

- Ehmt2/Ezh2 inhibition results in an increase in effector CD8⁺ T cells (71.2% vs 54.4% $p = .03$) with a simultaneous decrease in naïve CD8⁺ T cells (3.44% vs 0.68%, $p = .02$) in the TME (fig 3a). There are significant increases in both mean fluorescence intensity (MFI) of CXCR3, the receptor involved in CXCL9/CXCL10 axis (MFI 3959 vs 1862, $p < .001$) and granzyme-B producing CD8⁺ cells (65.1% vs 27.2% $p < .0001$) following Ehmt2/Ezh2 inhibition (fig 3b)

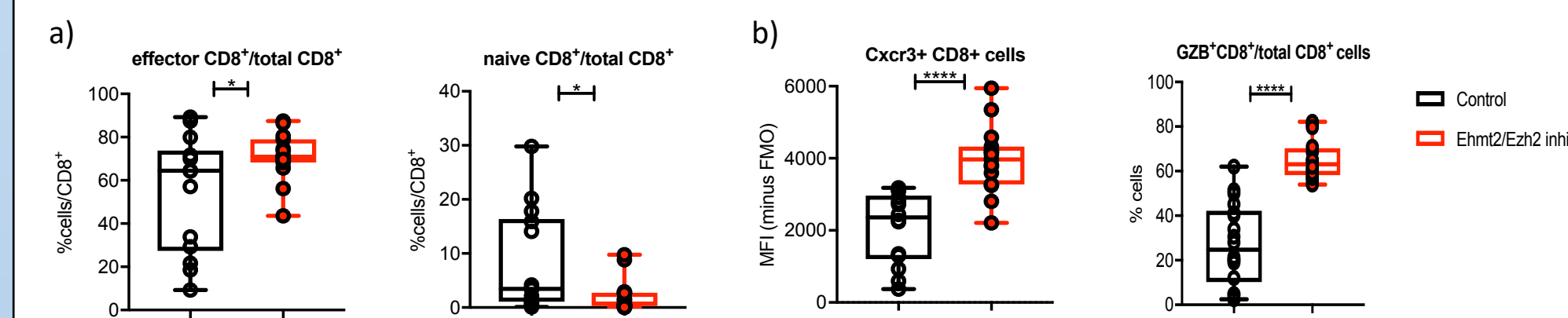


Figure 3: a) Quantitative analysis of flow cytometry performed on mouse tumours, gating on (a) CD8⁺ T cells stained with CD44 and CD62L markers for identification of effector (CD44⁺CD62L⁻) and naïve (CD44⁺CD62L⁺) populations, (b) mean fluorescence intensity of CXCR3 receptor on CD8⁺ cells and CD8⁺ T cells stained with granzyme B antibody following stimulation with PMA/ionomycin, followed by brefeldin A/monensin treatment.

- Treatment with Ehmt2/Ezh2 inhibitor results in an increase in both Natural Killer (NK) cells (5.2×10^6 vs 2.8×10^6 cells/g, $p = .007$) and their CXCR3 receptor expression (MFI 1507 vs 440, $p < .001$) (fig 4).

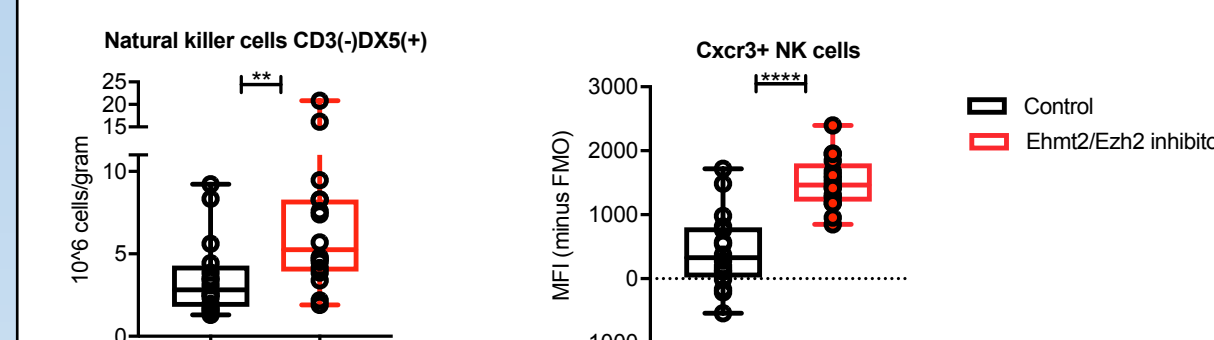


Figure 4: Quantitative analysis of flow cytometry gating on CD3⁺DX5⁺ for identification of natural killer population (NK cells) and MFI of CXCR3 receptor on NK cells.

- Ehmt2/Ezh2 inhibition reduces the immunosuppressive population of FoxP3⁺CD4⁺ Treg cells (0.9×10^6 vs 2.2×10^6 cells/g, $p = .02$).

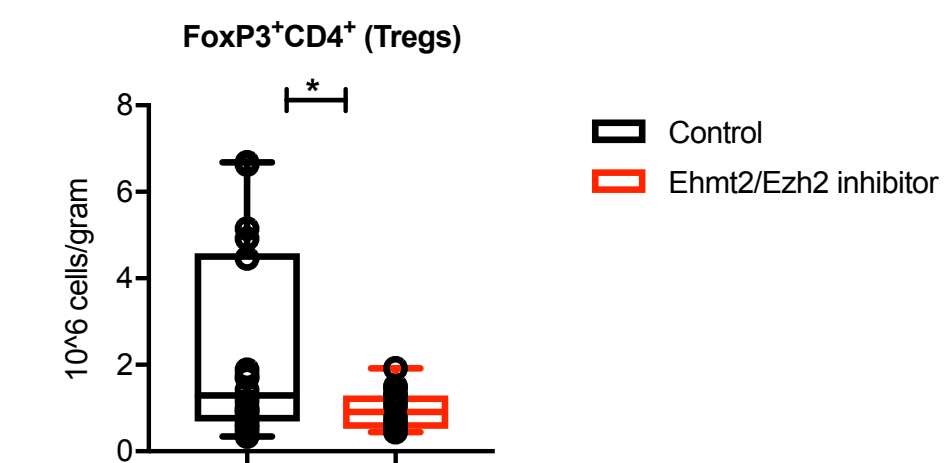


Figure 5: Quantitative analysis of flow cytometry gating on CD4⁺ T cells stained with FoxP3 marker.

- Treatment with Ehmt2/Ezh2 inhibitor increases the number of dendritic cells (DCs) (3.3×10^6 vs 1.3×10^6 cells/mg, $p = .0003$). These DCs have higher expression levels of the activating receptor CD86, compared to control (mean fluorescence intensity 22365 vs 16181, $p = 0.009$) (fig 6).

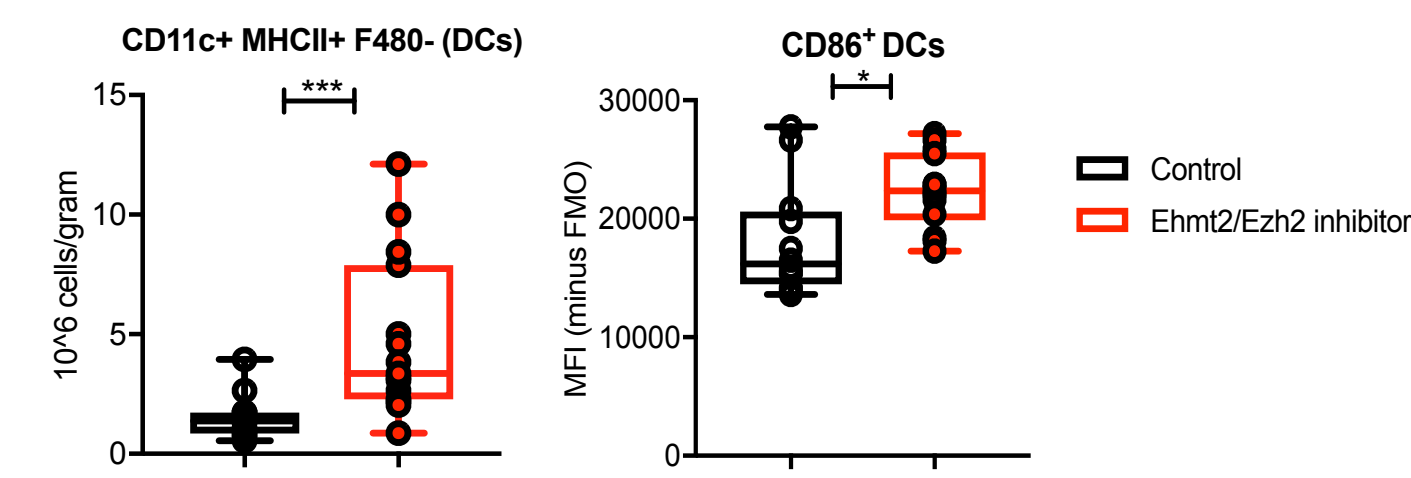


Figure 6: Quantitative analysis of flow cytometry gating on CD11b⁺Ly6G⁻SiglecF⁻F4/80⁻MHCII⁺CD11c⁺ cells (DCs) and quantification of CD86 MFI activation marker on DCs.

Result 4: Ehmt2/Ezh2 inhibition augments activation of splenic CD8⁺ cells

- CD8⁺ cells derived from mice after treatment with Ehmt2/Ezh2 inhibitor demonstrate higher level of perforin (10.2% vs 5.9%, $p = .004$) and IFN γ (22% vs 18%, $p = .01$), compared to control.

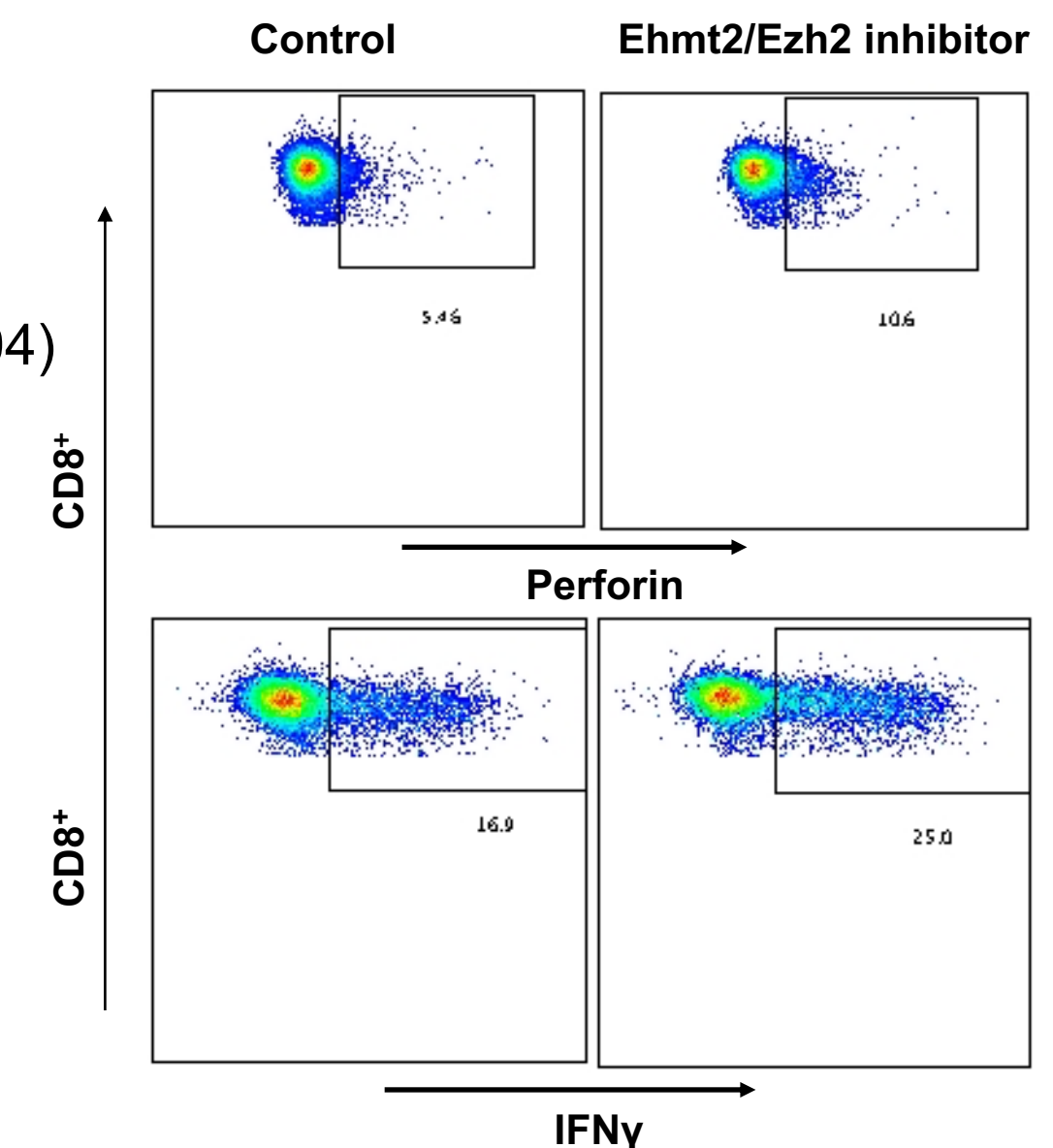


Figure 7: Mouse splenocytes were stimulated with PMA/ionomycin, treated with brefeldin A/monensin and then subjected to flow analysis after stained with intracellular markers against Perforin and IFN γ . Plots are showing a representative sample closest to the mean and numbers represent frequencies respective to entire CD8⁺ population.

Conclusion

Inhibition of Ehmt2/Ezh2 stimulates expression of chemokines involved in T cell, NK and DC recruitment. *In vivo*, Ehmt2/Ezh2 inhibition alters the immune microenvironment and confers a survival benefit. This suggests that Ehmt2/Ezh2 inhibition could augment the anti-tumour immune response in ovarian cancer.