

Mesenchymal stromal cell diversity and regulation of immune cell development



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Background

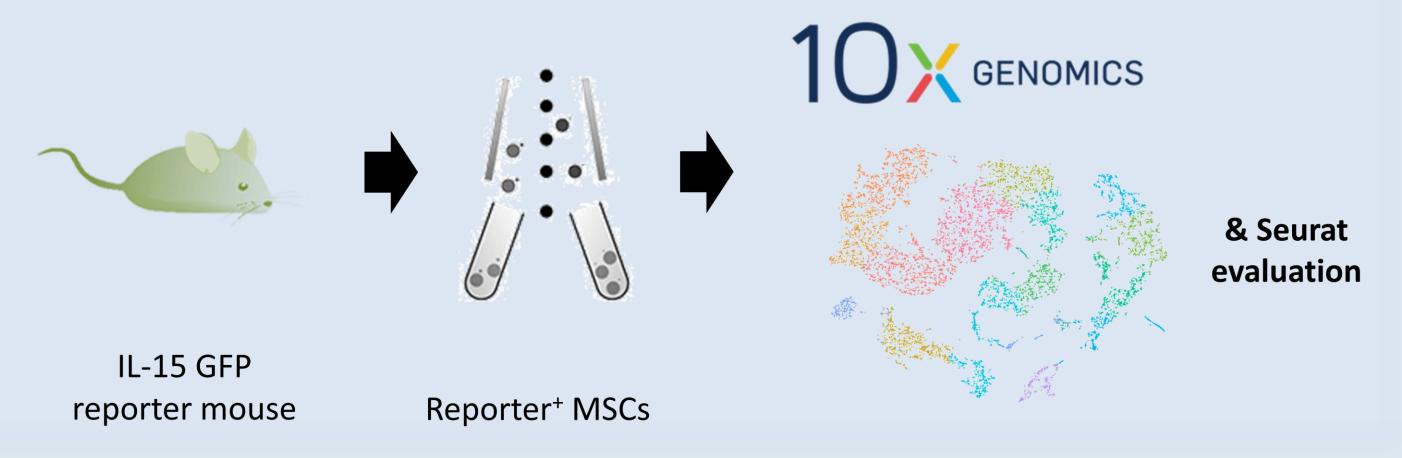
Bone marrow (BM) mesenchymal stroma cells (MSCs) provide distinct **lineage-instructive differentiation and survival signals** for hematopoietic stem and progenitor cells, such as CXCL12 and IL-15 (1). Hematopoietic stem cells (HSCs) for example reside and self-renew in CXCL12-rich perivascular MSC niches. While macrophages and dendritic cells are major IL-15 expressors, studies have shown that they are not sufficient to support maintenance of IL-15 dependent immune cell subsets such as NK, ILC1 and memory T cells in the BM (2), **indicating the presence of other IL-15 sources**.

Hypothesis

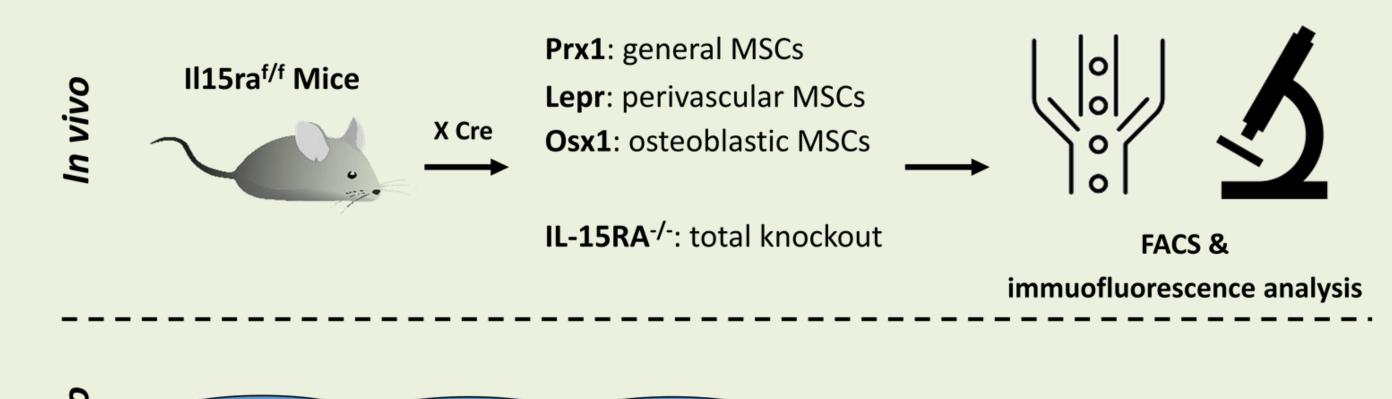
Heterogeneity among cytokine-/chemokine-expressing BM MSCs governs functional specialization, such as expression of distinct lineage-instructive differentiation signals for immune cell development ("developmental niches") or distinct survival/adhesion signals for (memory) immune cells ("survival niches").

Aims

Aim 1: Single-cell profiling of IL-15+ BM MSC subsets



Aim 2: Functional relevance of IL-15+ BM MSC subsets

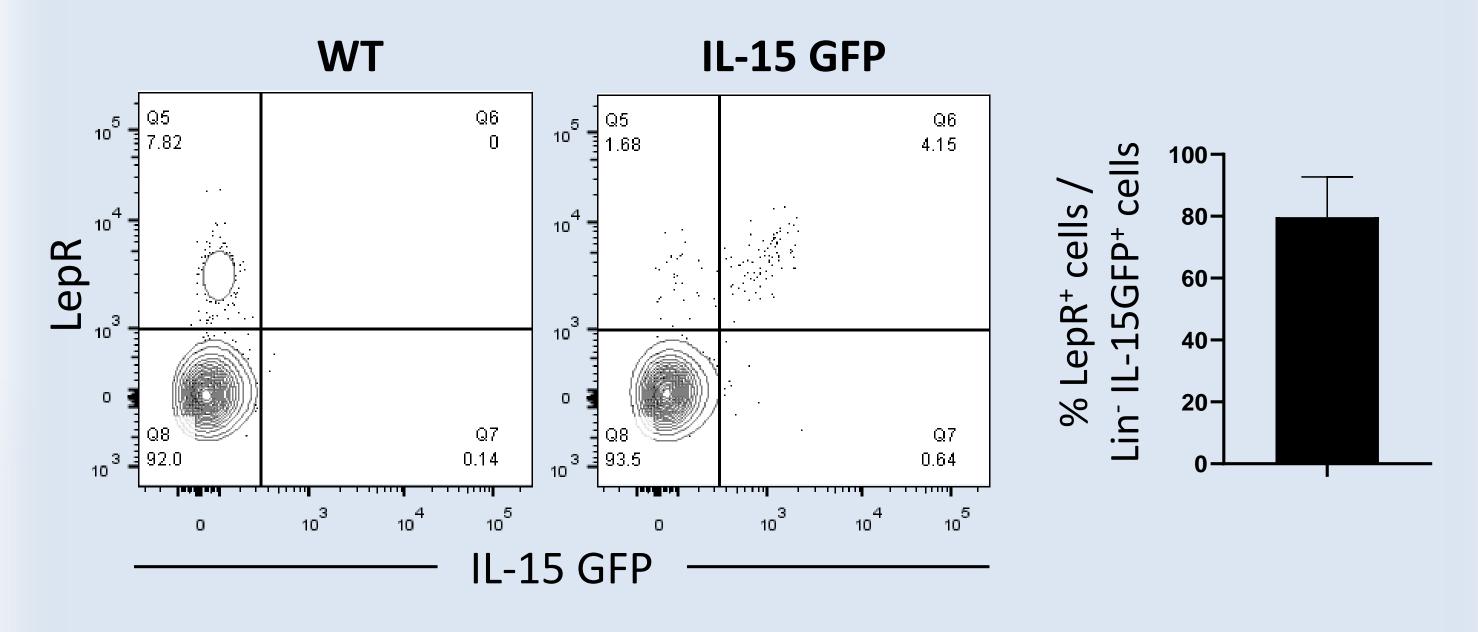


Co-culture of MSC subsets with IL-15 dependant immune cell subsets

Results

Our conditional KO experiments indicate a physiological relevance of IL-15+ MSCs for CD8+ memory T cell maintenance and NK cell development / maintenance. Nevertheless, the LepR expressing subset, which represents stem cell-like MSCs does not seem to be the causative subpopulation. The high co-expression of LepR on IL-15 GFP+ MSCs indicates a potential role of a rare, more mature LepR- IL-15+ MSC subset.

IL-15 GFP+ MSCs express LepR and locate close to hematopoietic cells



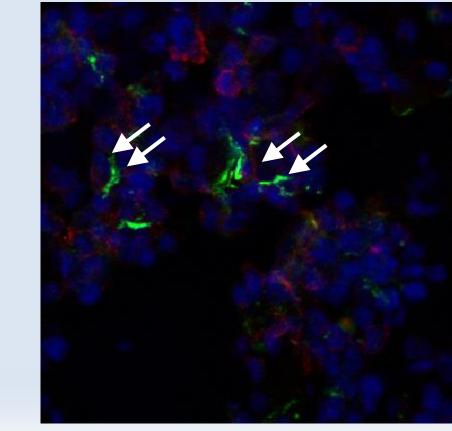


Fig.1: Identification and quantification of IL-15+ MSCs in the BM

Up: Quantification of LepR expression on IL-15+ MSCs

Down: Immunofluorescence image of IL-15 (green),

CD45 (red) and DAPI (blue)

Deletion of IL15RA in MSCs impairs NK cell but not ILC1 development

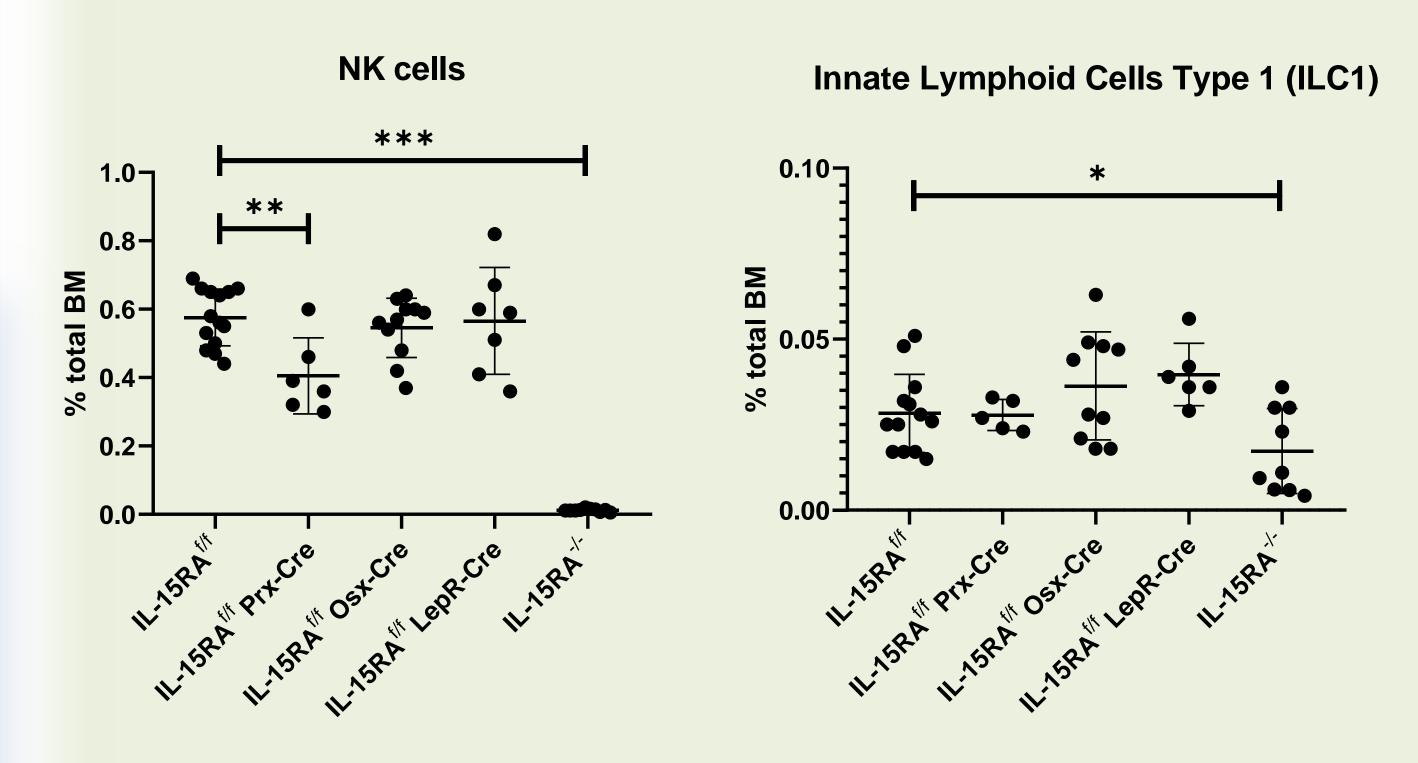
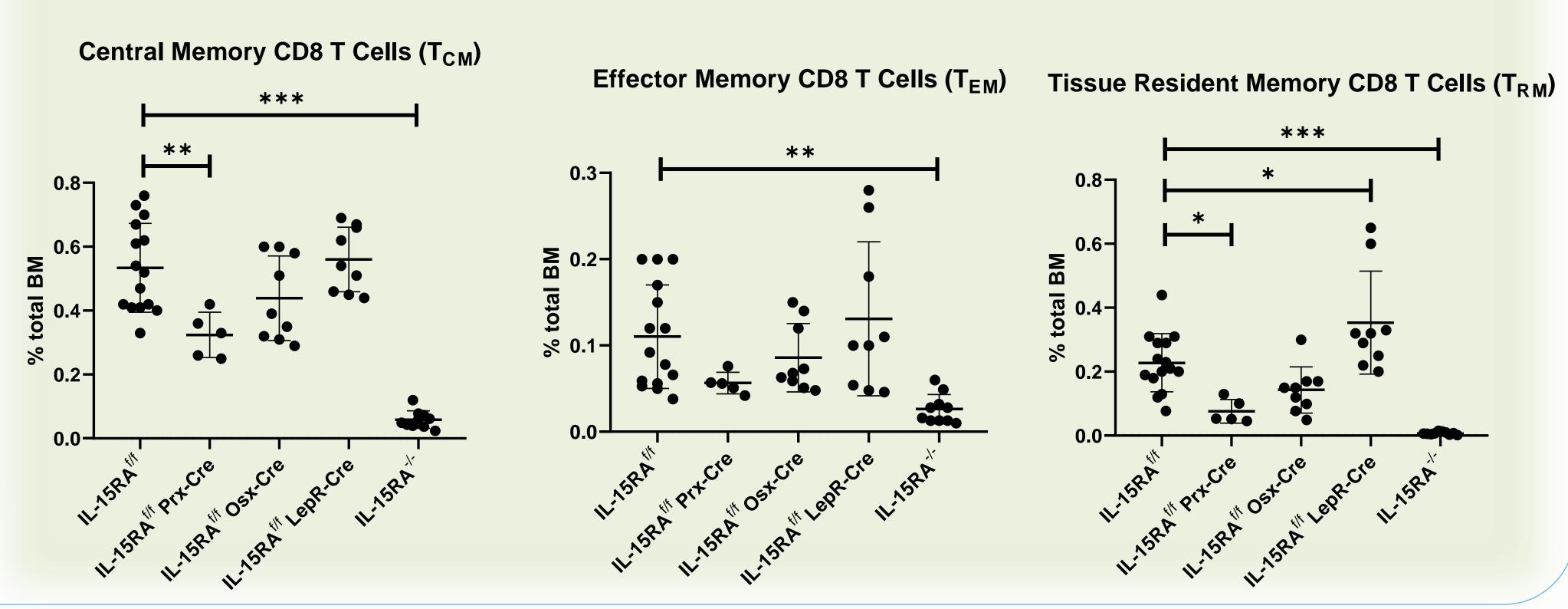


Fig.2: Percentage of group 1 ILCs and memory CD8+ T cells following conditional KO of all MSCs (Prx), osteoblastic MSCs (Osx) and LepR+ MSCs vs controls in the BM. . Mean ± s.e.m. are shown. Mean ± s.e.m. are shown. ** P < 0.01 and *** P < 0.001 (unpaired two-tailed Student's t-test).

Deletion of IL15RA in MSCs impairs survival of memory CD8+ T cell subsets



Conclusion

Our results indicate a physiological role of non-stem-like IL-15+ MSCs in development and maintenance of IL-15 dependent immune cell subsets. scRNAseq data will give further information about their identity as well as crosstalk mechanisms. The output will be validated via co-culture experiments.

With these investigations we will increase the understanding of stromal cells as regulators of immunity and immunological memory.

References

- (1) Cordeiro Gomes, A. et al. Hematopoietic Stem Cell Niches Produce Lineage-Instructive Signals to Control Multipotent Progenitor Differentiation. *Immunity* 45, 1219-1231, doi:10.1016/j.immuni.2016.11.004 (2016).
- (2) Mortier, E., et al. Macrophage- and dendritic-cell-derived interleukin-15 receptor alpha supports homeostasis of distinct CD8+ T cell subsets. Immunity 31, 811-822, doi:10.1016/j.immuni.2009.09.017 (2009).

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