# Immune microenvironment dysfunctions enable malignification at the onset of MDS



Ganan-Gomez I, Ma F, Chien KS, Yang H, Montalban-Bravo G, Wildeman BE, Kumar B, Kim YJ, Daher M, Takahashi K, Garcia-Manero G and Colla S

### Background

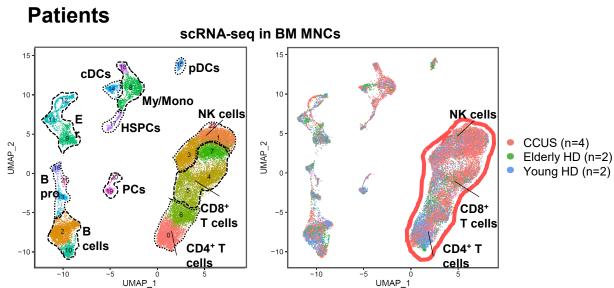
- Myelodysplastic syndromes (MDS): clonal stem cell malignancies
- Standard therapy not curative, transient responses, poor prognosis
- Are prevention or early intervention possible in MDS?
- Clonal cytopenias of undetermined significance (CCUS): aging-related premalignant state, low-grade clonal hematologic disorders at high risk of progression to MDS and leukemia
- HSC-intrinsic alterations and extrinsic inflammatory factors cooperate to induce abnormal differentiation in CCUS (Ganan-Gomez et al. ASH Meeting 2021).

### Aim

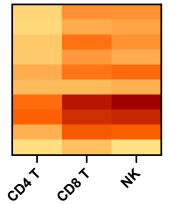
· Characterize early alterations in the bone marrow (BM) immune microenvironment that lead to the expansion of the MDS clone

## **Methods**

- BM mononuclear cells (MNCs) from patients with CCUS and young and elderly healthy donors (HDs)
- Single-cell transcriptomics (scRNA-seq)
- In vitro functional assays
- Single-cell genomics/antigen expression analysis (Tapestri)



#### -Log<sub>10</sub>[P val] 10 20 30 40 50



#### Pathway analysis of significantly upregulated genes in CCUS

Formation of ATP by chemiosmotic coupling Interferon alpha/beta signaling Neutrophil degranulation Mitochondrial translation initiation Cytokine Signaling in Immune system Membrane Trafficking **Eukaryotic Translation Elongation** Respiratory electron transport, ATP synthesis by chemiosmotic coupling Adaptive Immune System Parasite infection

# 1. BM Innate Immune Cells Are Activated in CCUS

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# 2. Inhibitory Crosstalk Predicted Between NK and CD8<sup>+</sup> T Cells in CCUS

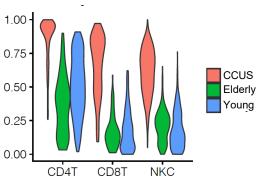
#### **CellPhoneDB analysis of DEGs** Ligand-expressing cells Myelomonocytic 20 NK cell cDC 15 CD8 T cell HSPC 10 pDC 5 B cell **B** Precursor CD4 T cell Plasma cell **RBC** Precursor NK cell сDC pDC CD8 T cell B cell HSPC Myelomonocytic CD4 T cell Plasma cell B Precursor **RBC** Precursor **Receptor-expressing cells**

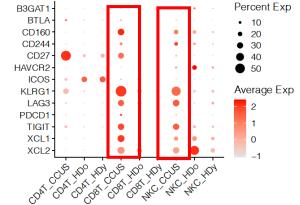
NK cell activation by CD8<sup>+</sup> T cells HLA-E:CD94/NKG2C CD58:CD2 FASLG:TNFRSF1A NK cell CD8<sup>+</sup> T Immunosuppresion of

CD8<sup>+</sup> T cells by NK cells CD48:CD244 TGFB1:TGFBR1

# 3. CCUS NK/T Cells Are Terminally Differentiated toward Exhaustion

scRNA-seg in sorted CD3<sup>+</sup> T and CD56<sup>+</sup> NK cells Young HD (n=2) Elderly HD (n=2) CCUS (n=3) 3 3- $\sim$ UMAP\_2 NK UMAP\_ CD4 T CD8 -3 -3. 10 10 -5 0 -5 0 5 UMAP 1 UMAP 1 CytoTRACE differentiation scores **Exhaustion marker expression** 





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## 4. NK Cells from CCUS Patients Are Irreversibly Dysfunctional

## 5. NK Cells Are Part of the Mutant CCUS Clone

Joint scDNA-seq and surface protein analysis (Tapestri)

