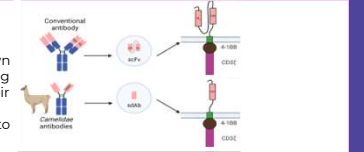


Mégane Jassin¹, Chloé Onkelinx¹, Bianca E Silva¹, Valentina Bocuzzi¹, Alix Block¹, Guillaume Marcion¹, Frédéric Baron^{1,2}, Emmanuel Di Valentin³, Grégory Ehx¹, Jo Caers^{1,2} and Tham Nguyen¹
¹Laboratory of Hematology, GIGA-13, University of Liège, Liège, Belgium
²Department of Hematology, CHU de Liège, Liège, Belgium
³Viral Vectors, GIGA, University of Liège, Liège, Belgium

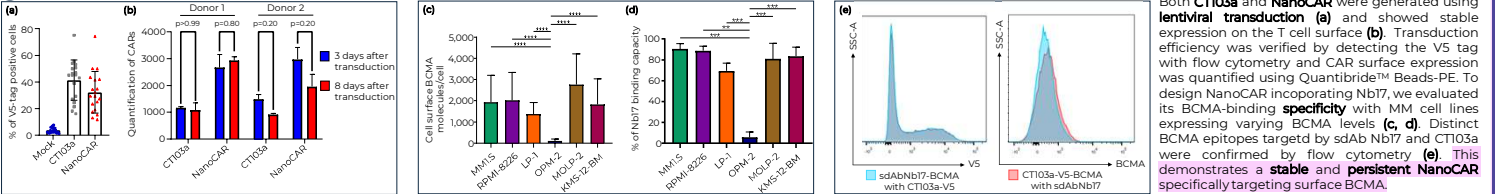
Introduction & Objective

- Multiple myeloma (MM) is an incurable hematologic malignancy of plasma cells. Chimeric antigen receptor T lymphocyte (CAR-T) immunotherapy has shown remarkable results in relapse patients. Single-domain antibody (sdAb) offers an excellent alternative to scFv due to the advantage of having a small and stable folding structure, therefore avoiding tonic signaling. Despite the development of numerous nanoCAR-Ts, there is limited data demonstrating the direct differences and their mechanisms of action.
- This project aims to characterize and compare the activity of a nanoCAR-T containing the sdAb Nb17 to the ScFv CAR-T CT103a, an FDA-approved CAR-T, to try to elucidate differences of mechanisms between them.



Methods & Results

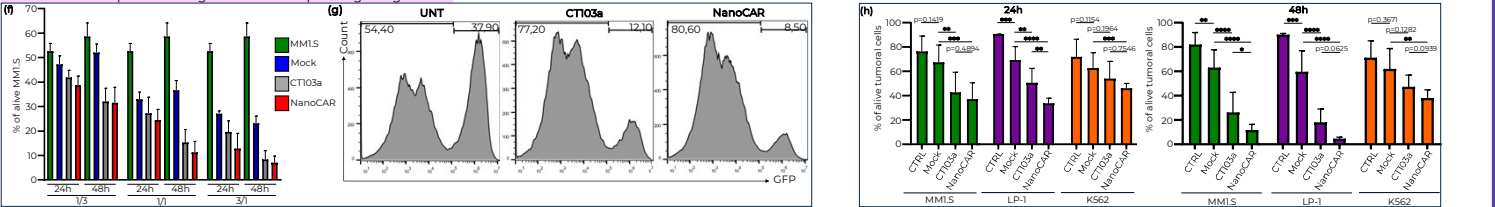
I Construction of a NanoCAR sequence containing sdAb Nb17



Both CT103a and NanoCAR were generated using lentiviral transduction (a) and showed stable expression on the T cell surface (b). Transduction efficiency was verified by detecting the V5 tag with flow cytometry and CAR surface expression was quantified using QuantiBrite™ Beads-PE. To design NanoCAR incorporating Nb17, we evaluated its BCMA-binding specificity with MM cell lines expressing varying BCMA levels (c, d). Distinct BCMA epitopes targeted by sdAb Nb17 and CT103a were confirmed by flow cytometry (e). This demonstrates a stable and persistent NanoCAR specifically targeting surface BCMA.

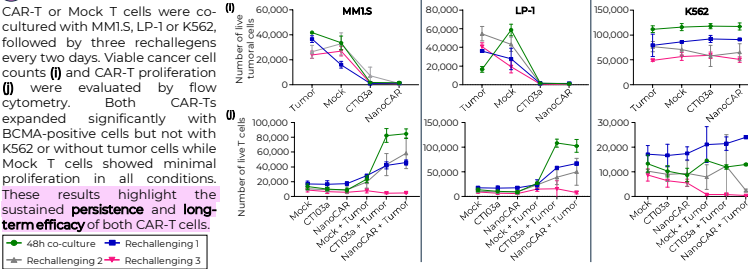
II In vitro efficacy of CT103a and NanoCAR in killing BCMA-positive MM cell lines

The cytotoxic capacity of NanoCAR, CT103a and Mock T cells was evaluated by flow cytometry after co-culture with MM1S for 24 to 48 hours at effector-to-target (E/T) ratios of 1/3, 1/1 and 3/1. Killing efficiency improved with higher ratios and longer incubation (f, g). A 1/1 ratio killing assay confirmed specific targeting of BCMA-positive cells (MM1S, LP-1) while sparing BCMA-negative K562 cells (h). These results demonstrate the efficacy and selectivity of CT103a and NanoCAR as therapeutic strategies for BCMA-expressing malignancies.



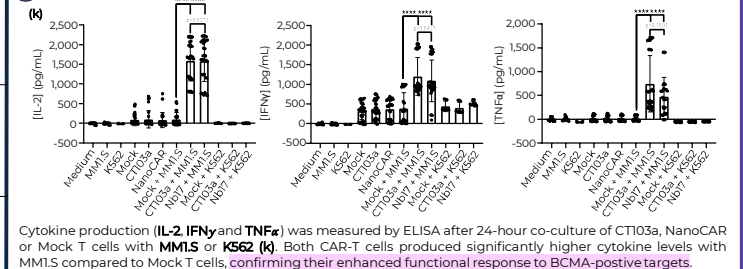
III Persistence of CAR-T killing ability following repeated antigen challenges

CAR-T or Mock T cells were co-cultured with MM1S, LP-1 or K562, followed by three rechallenges every two days. Viable cancer cell counts (i) and CAR-T proliferation (j) were evaluated by flow cytometry. Both CAR-Ts expanded significantly with BCMA-positive cells but not with K562 or without tumor cells while Mock T cells showed minimal proliferation in all conditions. These results highlight the sustained persistence and long-term efficacy of both CAR-T cells.



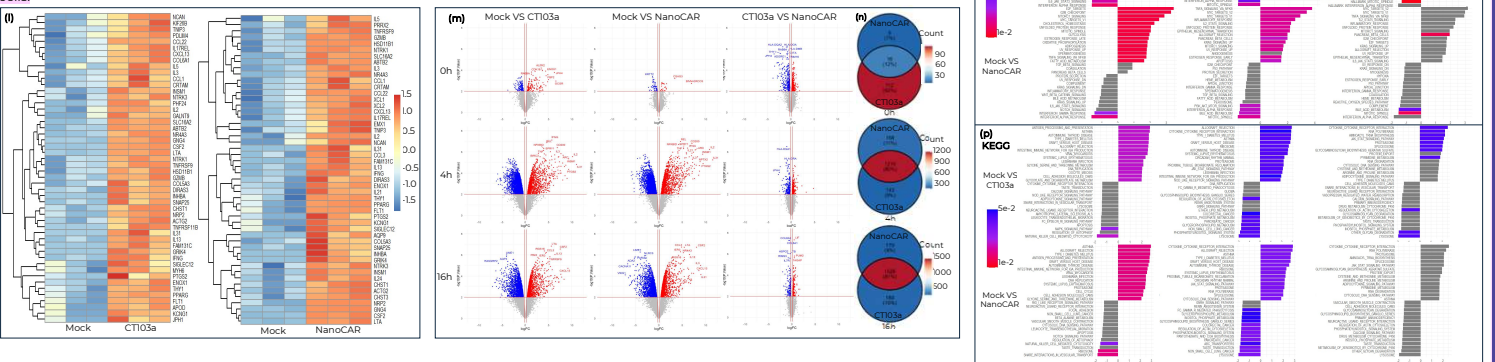
IV CT103a and NanoCAR produce cytokines when co-cultured with MM

Cytokine production (IL-2, IFNγ and TNFα) was measured by ELISA after 24-hour co-culture of CT103a, NanoCAR or Mock T cells with MM1S or K562 (k). Both CAR-T cells produced significantly higher cytokine levels with MM1S compared to Mock T cells, confirming their enhanced functional response to BCMA-positive targets.



V CT103a and NanoCAR expressed similar set of genes or pathways when co-cultured with MM1S

Bulk RNA seq compared transcriptional profiles of CT103a, NanoCAR and Mock T cells at 0h, 4h and 16h post-incubation with MM1S. Mock T cells served as a negative control and CT103a as the comparator. Heatmaps of the top 50 genes at 16h (l) showed upregulation of activation, proliferation and effector function genes in both CAR-T cells. Volcano plots highlighted minimal differences between CT103a and NanoCAR with many upregulated and downregulated genes shared (m, n). GSEA (Hallmark and KEGG) at 4h revealed enriched cytokine production, cytokine interactions and proliferation pathways in both CARs compared to Mock (o, p). However, these differences disappeared by 16h due to basal cytokine production by pre-activated Mock T cells. Despite similar transcriptional profiles, both CT103a and NanoCAR showed tumor-specific activation mechanisms distinct from Mock T cells.



Conclusion(s)

To conclude, we first designed a persistent and stable NanoCAR containing the sdAb Nb17. The specificity of CT103a and the NanoCAR CAR-T cells for BCMA-positive targets while sparing BCMA-negative cells, such as the K562 leukemia cell line, confirmed the potential of CT103a and NanoCAR as effective and selective therapeutic strategies for targeting BCMA-expressing malignancies. Moreover, both CT103a and the NanoCAR demonstrated sustained persistence and long-term efficacy, suggesting their potential for prolonged tumor control. In addition, the cytokine production highlighted the robust cytotoxic potential and functional activity of CT103a and NanoCAR, confirming their functional readiness to engage and kill target. Also, according to overall bulk RNA seq data, while CT103a and NanoCAR demonstrated similar transcriptional profiles, their distinct gene signatures relative to Mock revealed shared mechanisms of tumor-specific activation. Thus, our nanoCAR-T demonstrates excellent anticancer efficacy as CT103a *in vitro*. Further investigation *in vivo* in a mice model should finally validated *in vitro* results.

Acknowledgements

Members of the Laboratory of Hematology
 Viral Vector platform (GIGA) – Flow Cytometry platform (GIGA) – RNA seq platform (GIGA)

Contact Information

Mégane JASSIN, PhD Student at the Laboratory of Hematology, University of Liège – GIGA, Belgium
 megane.jassin@uliege.be