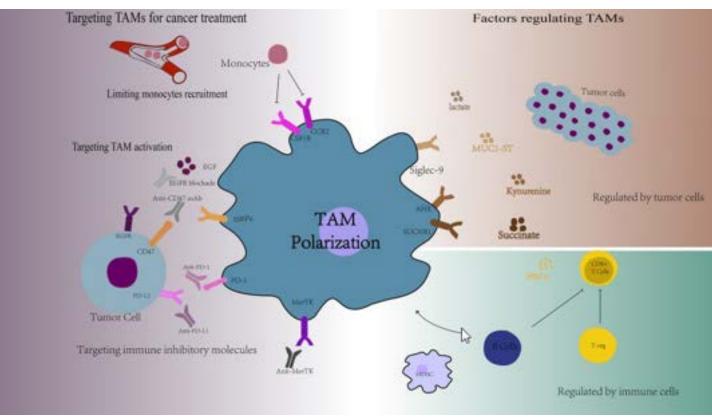
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Introduction

Tumor-Associated Macrophages (TAMs) play an important role in tumor development modulation of neoangiogenesis, immune suppression, and metastasis. A high infiltration of macrophages in the tumor is also often correlated with a poor prognosis in several types of cancer. Therefore, they became an attractive target for cancer immunotherapies. Several macrophage-targeting approaches in anticancer therapy are under development, including TAM depletion, inhibition of new TAM differentiation, or re-education of TAM activation for cancer cell phagocytosis.



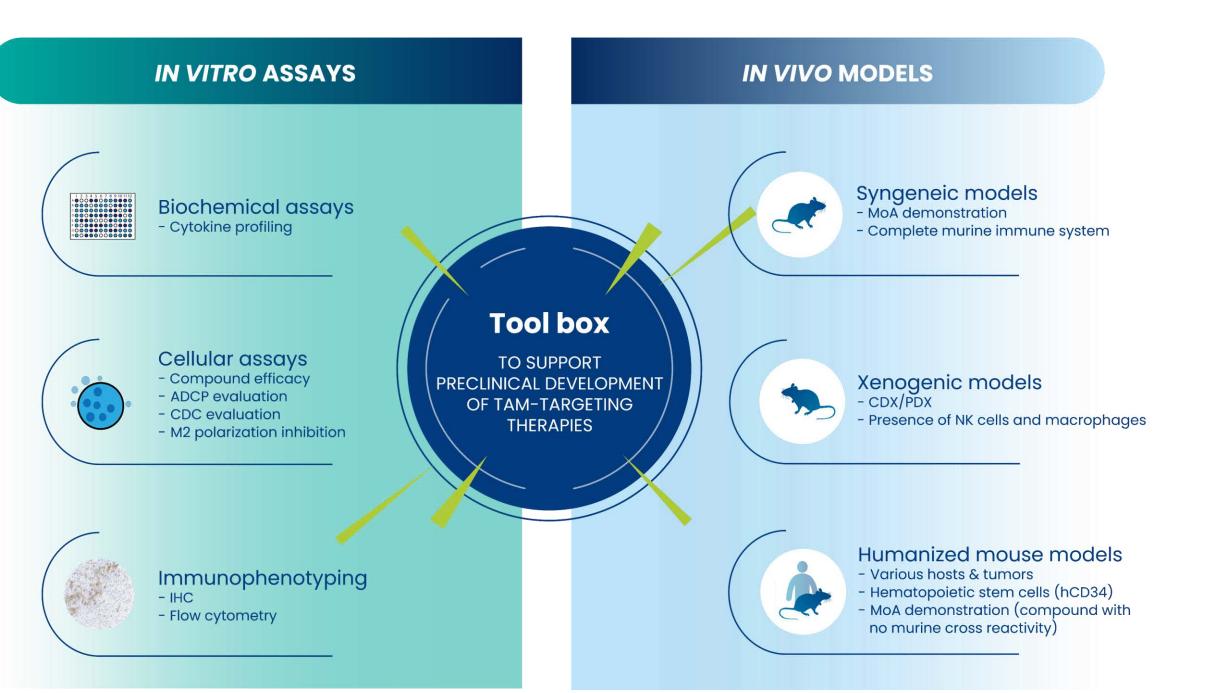
From Pan et al., Front. Immunol., 2020

In order to better evaluate and support the pre-clinical development of novel TAM-targeting strategies, we implemented a large panel of *in vitro* assays and *in vivo* models:

- Antibody-dependent cellular phagocytosis (ADCP) has been used to demonstrate a crucial mechanism of action of several antibody (Ab)-based therapies targeting macrophages.
- Some of these Ab (including anti-CSF1R) have been tested in syngeneic in vivo tumor models
- We demonstrated that the site of tumor implantation in mice (PAN-02, subcutaneous vs orthotopic) could impact the macrophage polarization (M1 vs M2). Differences in the ratio of M1 and M2 subtypes infiltrating the PAN-02 murine pancreatic tumors were observed
- Anti-CSF1R antibodies induced a slight reduction in the tumor mass of Renca murine kidney tumor by eliminating TAMs, but not in other syngeneic tumor models like MC38
- In an orthotopic Hepa1-6 murine liver cancer model, we showed high antitumor efficacy of compounds targeting the STAT6 pathway by reprogramming immunosuppressive TAMs into an M1 phenotype that promotes the induction of a cytotoxic immune response.
- For compounds displaying no cross-reactivity with murine target, we developed models and characterized the TAMs in breast, colon, melanoma and head & neck PDX and CDX tumors in different huCD34-engrafted mouse models.

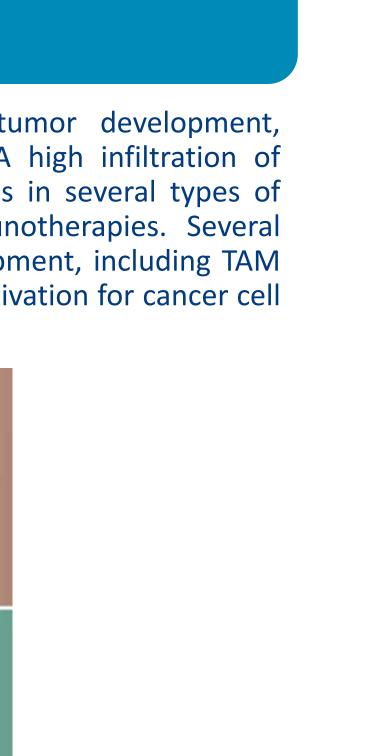
Conclusion

We designed a comprehensive preclinical platform to support the development of TAM-targeting therapies.

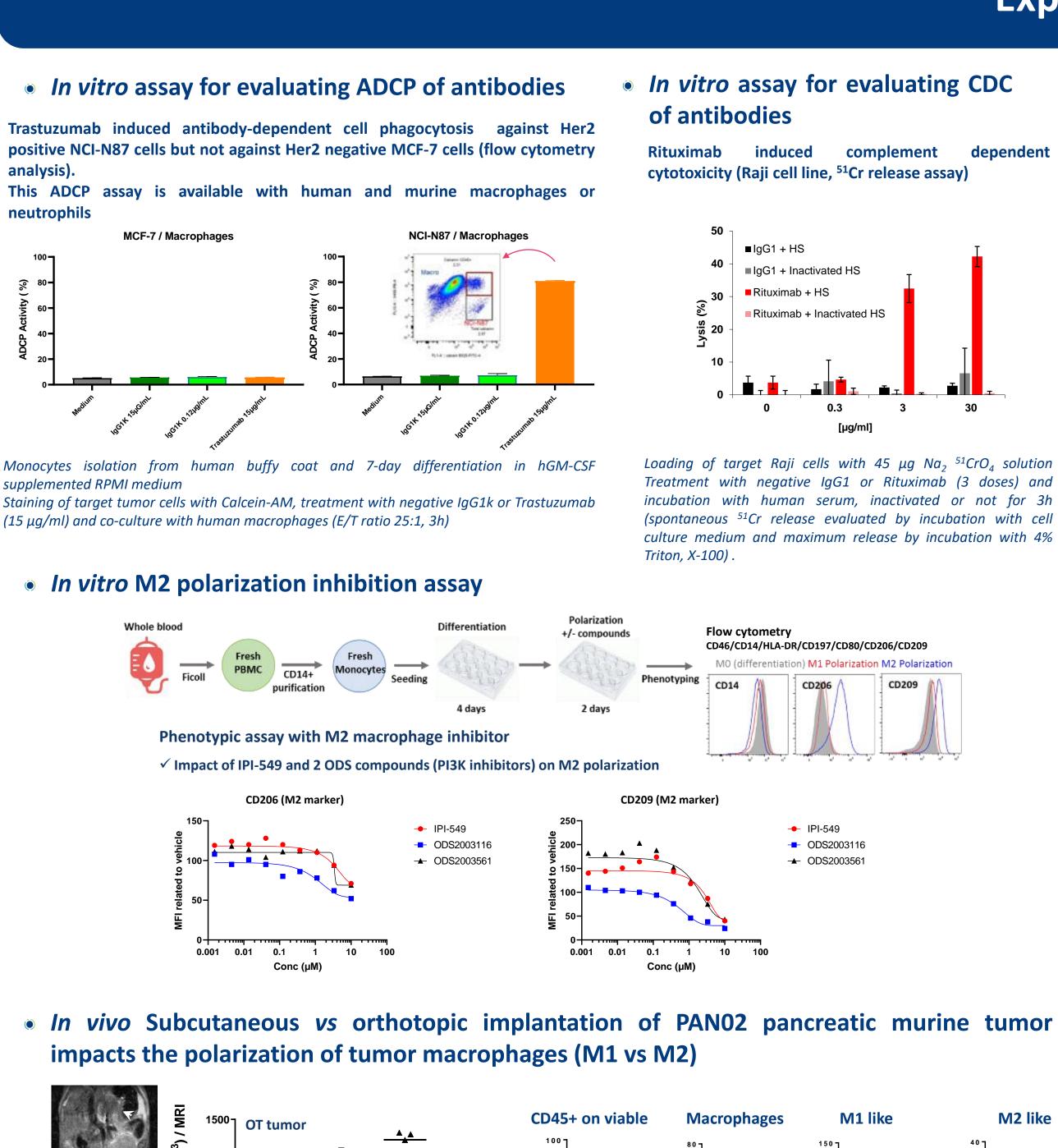


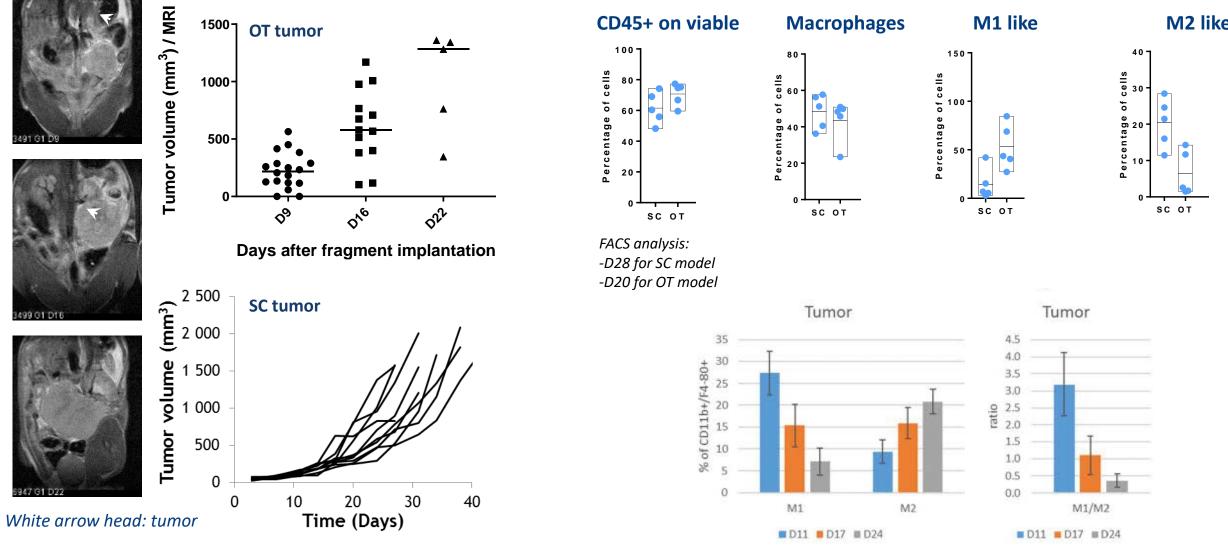
OncoTAM, a comprehensive preclinical platform to explore Macrophages as key drivers of cancer progression and develop new therapies against Tumor-Associated-Macrophages

Caroline Mignard, Damien France, Francis Bichat, Nicolas Legrand, Olivier Duchamp Oncodesign Services, 21000 Dijon - France

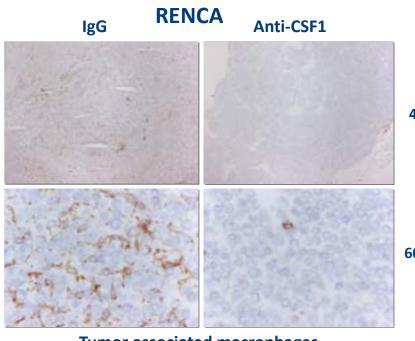






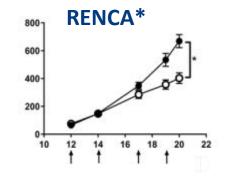


• TAMs elimination by *in vivo* anti-CSF1R treatment in Renca murine kidney tumors or MC38 colon murine tumors



Tumor associated macrophages F4/80 IHC in OT Renca tumor samples

any efficacy in MC38 tumor model despite significant TAM depletion

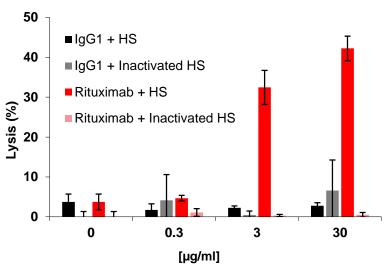


*From O'Brien S. et al, Cancer Immunol, 2021 Only RENCA aCSF1R-treated tumors had a significant increase in neutrophil cell density, promoting CAF to secrete chemokines involved in neutrophil recruitment. Not observed across all tumor models, potentially due to differences in fibroblast content

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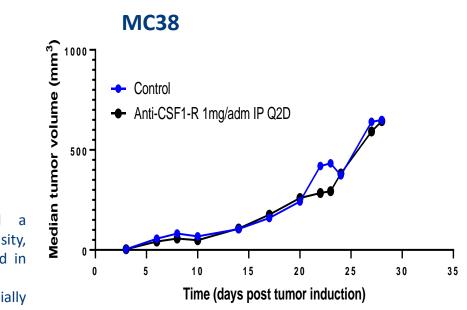
Experiments

In vitro assay for evaluating CDC

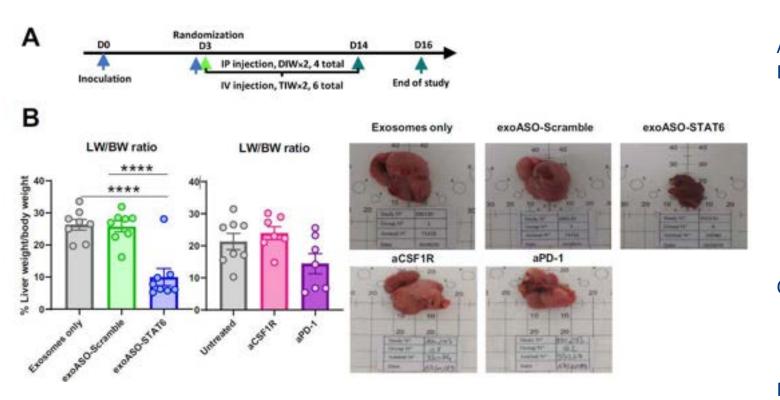


(spontaneous ⁵¹Cr release evaluated by incubation with cell culture medium and maximum release by incubation with 4%

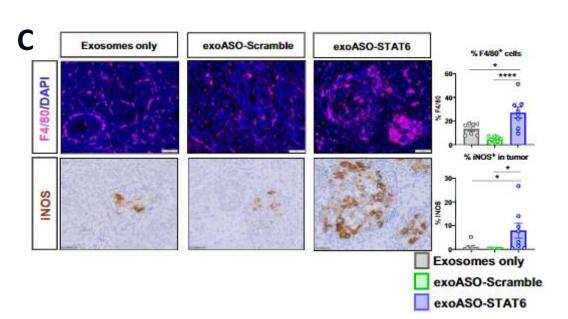
Anti-CSF1R treatment show a slight anti-tumor efficacy in Renca tumors but did not show



From Kamerkar S. et al., Sci Adv, 2022, in vivo work done at Oncodesign Services



exoASOSTAT6 results in effective re-programming to an M1 phenotype that promotes the induction of a cytotoxic immune response and an antitumoral TME

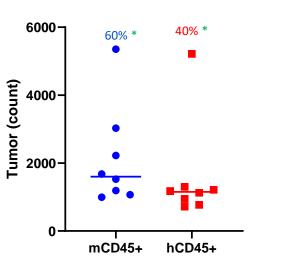


• Immune-humanized mice to monitor the MOA of compounds on human macrophages Myeloid lineage engraftment is enhanced by the use of immunodeficient mice expressing specific human cytokines such as NOG-EXL and **BRGSF** mice

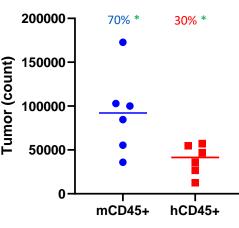
Mouse host	Hu. Method	Premium features
NOG-EXL	hHSC	 Express human GM-CSF and human IL-3 Increased myeloid populations Long term stable engraftment, no GvHD
BRGSF	hHSC	 hDC boost on demand by hFLT3L injection Activity of complement (C5) Suitable for infection & vaccination studies
NOG-FcγR ^{-/-}	hHSC	 Lack expression of functional mouse FcγRs Lack activity of endogenous ADCC Suitable for assessment of antibody-based therapies

Immunophenotypical characterization of 2 xenograft models in hCD34-engrafted NOG-EXL or BRGSF mice

A. Breast xenograft/HuNOG-EXL

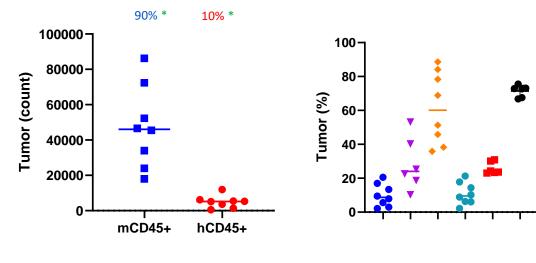


B. Lung xenograft/HuNOG-EXL

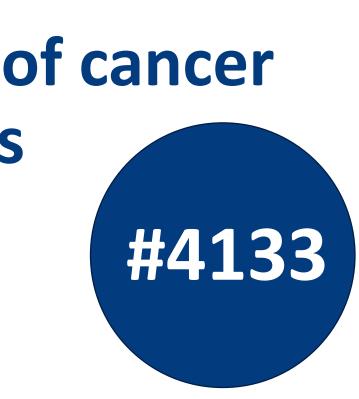


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C. Breast xenograft/HuBRGSF



All animal procedures were approved by the Animal Care and Use Committee of Oncodesign Services (Oncomet - CNREEA agreement N° 91)



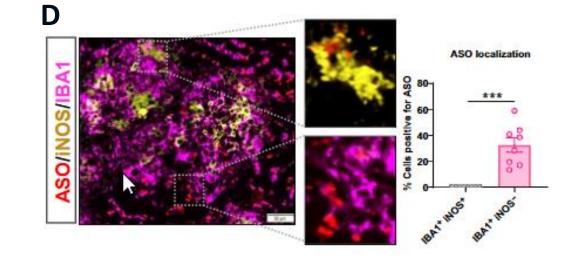


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• C57BL/6 mice bearing orthotopic Hepa1-6 tumors in the liver, treated with systemic administration of exoASO-STAT6 result in a potent monotherapy anti-tumor response

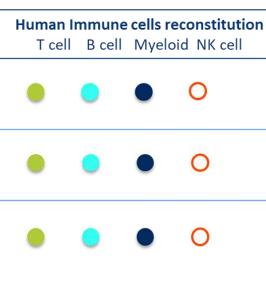
A. Experimental study design

- Antitumoral efficacy as represented by (12mg)(TIW.) (PTGFRN++) (12mg)(TIW, CSF1R (10mg/kg)(DIW) anti-PD-2 (10 mg/kg)(BIW)
- Quantification of F4/80 (macrophage) and iNOS expression, performed by immunofluorescence and IHC (immunohistochemistry) respectively, in Hepa1-6 tumor sections
- D. Representative images and quantification of ASO localization in iNOS-positive, IBA1+ (M1), and iNOS-negative, IBA1+ (M2), macrophages.



• Optimal level of presence

O Potential level of presence



- human leukocytes (% of m+hCD45+)
- ▲ human myeloid cells (% of hCD45+)
- human macrophages (% of hCD45+) human T cells (% of hCD45+)
- human CD8 T cells (% of hCD3+)
- human M1 macrophages (% of macrophages)
- human M2 macrophages (% of macrophages * Expressed as % of h+m hCD45+ cells

- human leukocytes (% of m + hCD45+ cells) ▲ human myeloid cells (% of hCD45+)
- human macrophages (% of hCD45+ cells)
- human T cells (% of hCD45+ cells)
- human CD8 T cells (% of hCD3+ cells)
- * Expressed as % of h+m hCD45+ cells
- noderately high frequency macrophages within the tumor.
- \rightarrow Immunophenotypical profile is tumor and host-dependent
- \rightarrow Mouse immune cells, which are strickly from the myeloid lineage, represent the dominant myeloid population.
- Others internal data indicate that immune cell repertoire i driven by the tumor
- human leukocytes (% of m + hCD45+ cells)
- human macrophages (% of hCD45+ cells) human T cells (% of hCD45+ cells)
- human CD8 T cells (% of hCD3+ cells)
- human M1 macrophages (% of macrophages)
- human M2 macrophages (% of macrophages) * Expressed as % of h+m hCD45+ cells



