

Blockade of the myeloid CLEC-1 checkpoint enhances antitumor responses and tumor antigen cross-presentation

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Abstract

Myeloid cells represent the most abundant immune component of the tumor microenvironment, where they often assume immunosuppressive roles. We have identified CLEC-1, a member of the C-type lectin receptor (CLR) family as being expressed by myeloid cells, especially by conventional type one dendritic cells (cDC1), and by tumor-associated macrophages (TAM). Here, we investigated the effect of CLEC-1 blockade, either by genetic deletion or by antibody targeting, on myeloid function and anti-tumor response

First, we observed that CLEC-1 genetic deletion significantly increases the survival of syngeneic tumor-bearing mice in the Hepa1.6 hepatocarcinoma, as well as in the AK-7 mesothelioma, orthotopic models. Moreover, the synergy with chemotherapy treatment in the MC38 model of colon adenocarcinoma significantly enhanced complete responses. Next, we generated anti-human CLEC-1 monoclonal antibodies (mAbs) as well as CLEC-1 humanized mice. We found that CLEC-1 targeting through mAb treatment was able to prolong mouse survival as efficiently as by CLEC-1 genetic deletion in MC38 and Hepa1.6 models. CLEC-1 blockade profoundly impacted the tumor microenvironment: increase of the frequency of invigorated dendritic cells (DCs) and macrophages, activated and memory T cells, while frequencies of immunosuppressive myeloid cells and PD1expressing T cells largely decreased. Mechanistically, the in vivo phagocytosis of tumor cells (Hepa1.6 and MCA101) by macrophages was enhanced in CLEC-1 deficient animals compared to WT animals.

Altogether, our results demonstrate that CLEC-1 acts as a novel immune checkpoint in myeloid cells and highlight its high potential as a target for innovative immunotherapy in oncology.



CLEC-1 is expressed by tumor associated macrophages (TAM) and cross-presenting dendritic cells (cDC1)

CLEC-1 expression in mouse and human immune cells and M Φ /DCs subsets from different solid tumor microenvironment context or models was analyzed by RT-gPCR (A). CLEC1A expression was also characterized in human cancer (lung, breast, ovarian, colon cancers) tumor using single cell RNA sequencing public datasets (TME Blueprint) (**B**).





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in macrophages (F4/80+).

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macrophages in human

tumor cell phagocytosis

- CLEC-1 KO mouse:
 - Significant anti-tumor responses in monotherapy
 - Synergy with chemotherapy
 - Strong modification of TME (e.g. increased memory CD8 T cells) - Enhancement of *in vivo* phagocytosis of tumor cells

• CLEC-1 antagonist mAbs : chemotherapy in CRC

Chemotherapy resistances.



CLEC-1 inhibits tumor cell phagocytosis by mouse $M\Phi$

MCA101 cells (fibrosarcoma) or Hepa 1.6 cells (hepatoma) were stained with CellTracker (ThermoFisher) and intraperitoneally injected (1.10⁶) cells/mouse) in WT or KO mice. Exudates were then collected after 1H after a 5mL injection of PBS-EDTA to harvest all cells. Macrophages were stained using an anti-F4/80 mAb and phagocytosis of tumor cells was evaluated in macrophages positive for the CellTracker Green. (A) Gating strategy. (B) Percentage of harvested macrophages in exudates. (C) Phagocytosis of MCA in macrophages (F4/80+). (D) Phagocytosis of Hepa 1.6

Conclusion

- CLEC-1 is expressed by dendritic cells and tumor associated
- CLEC-1/CLEC-1L interaction inhibits macrophage

- Prolongs survival in HCC preclinical model and synergizes with
- Modifies the TME as observed in deficient mice
- High interest in immune desert and to fight **Radiotherapy and**

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