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Abstract

Tumor-associated myeloid cells inhibit anti-cancer immune responses systemically and in the tumor microenvironment (TME), thereby limiting the efficacy of T cell checkpoint inhibitors. However, the plasticity of myeloid cells may enable therapeutic intervention. The immune inhibitory receptor LILRB2 (also known as ILT4) is expressed primarily by myeloid cells (monocytes, macrophages, dendritic cells and neutrophils), and has emerged as a key myeloid checkpoint contributing to the tolerogenic activity of myeloid cells associated with cancer. LILRB2 has several ligands (classical and non-classical MHC-I, ANGPTL2/5, SEMA4A and CD1) and most of these are known to contribute to immune suppression in the TME. Using computational biology approaches applied to the RNA-seq dataset of TCGA, we found that *LILRB2* expression is associated with the gene expression "signature" of macrophages infiltrating solid tumors. Therefore, LILRB2 is a compelling target to overcome immune suppressive activity of cancer-associated myeloid cells.

IO-108 is a fully human IgG4 therapeutic antibody that binds to LILRB2 with high affinity and specificity. IO-108 binds to all myeloid cells in the solid TME and periphery. In vitro studies support that blockade of LILRB2 interaction with its various ligands is the anticipated mechanism of action of IO-108. The LILRB2 antagonist activity of IO-108 produces the desired pro-inflammatory (re)polarization of myeloid cells. IO-108 treatment results in increased pro-inflammatory responses and an enhanced antigen-presenting cell (APC) phenotype to multiple stimuli (e.g., T cell activators, STING and TLR agonists) in ex vivo assays. In addition, IO-108 precludes the antiinflammatory myeloid cell phenotype resulting from "tumor conditioning" and promotes the differentiation of monocytes and immature dendritic cells (DC) into pro-inflammatory DC. IO-108 enhances the effect of PD-1 blocking antibodies in allogeneic mixed leukocyte reactions of CD4⁺ T cells and macrophages. Moreover, IO-108 polarized primary myeloid cells isolated from solid tumor patient blood and ovarian cancer-associated ascites towards a pro-inflammatory phenotype and attenuated their suppressive effect on autologous T cell proliferation and production of pro-inflammatory cytokines. In vivo, IO-108 inhibits tumor growth in mouse models, which is associated with immune cell activation. Importantly, IO-108 presents a favorable pharmacokinetic and safety profile in preclinical

Collectively, the preclinical characterization of IO-108 enabled a comprehensive clinical biomarker plan and lends rationale to the clinical study (NCT05054348) of IO-108 as a novel immunotherapy for multiple solid tumor types, including those relapsed/refractory to standard of care therapies.



IO-108, a Fully Human Therapeutic Antibody Blocking the Myeloid Checkpoint LILRB2/ILT4, **Promotes Innate and Adaptive Anti-Cancer Immunity in Preclinical Studies**

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A. Flow cytometric analysis of IO-108 binding in tumor tissue samples from 3 different patients. CD11b is used as pan-myeloid cell marker and CD45 is used as pan-tumor-infiltrating leukocyte marker. B. Flow cytometric analysis of IO-108 binding (black-filled histogram) on myeloid cells in peripheral blood from patient #3. White filled histogram corresponds to IgG4 isotype control. C-F. The ex vivo activity of the antibody precursor to IO-108 was tested (at 10 µg/mL) in cells isolated from cancer patient-derived PBMC and ascites. IO-108 is a modified version of this antibody precursor optimized for developability while maintaining the same binding affinity, selectivity and potent biologic activity as its precursor (not shown). C-E. Each line represents results from a different patient. Monocytic (M)-MDSC and autologous T cells isolated from PBMC were co-cultured for 5 days. T cells were activated with soluble tetrameric anti-CD3/CD28 antibody complexes. C. Proliferation of T cells (alone and in co-culture with M-MDSC) measured by flow cytometry (CFSE dilution). ANOVA: ***p= 0.0002, **p= 0.0052. D. Flow cytometric analysis of M-MDSC phenotype at the end of co-culture with autologous T cells. E. Cytokine levels in culture media supernatants. F. CD14⁺ cells were isolated from an ovarian cancer ascites sample, incubated with anti-LILRB2 (IO-108 precursor) or isotype control antibodies for 7 days and their phenotype analyzed by flow cytometry. G. Proliferation of patient PBMC-derived T cells (alone and in co-culture with autologous PMN-MDSC) measured by flow cytometry (CFSE dilution). ANOVA: *p<0.05, n.s. = non-significant.

IO-108 decreases the immune suppressive activity and promotes a pro-inflammatory phenotype in cancer patient-derived MDSC ex vivo



A, C-F. Each symbol is result from one mouse. A. Flow cytometric analysis of blood leukocytes of LILRB2 transgenic mice showing that LILRB2 expression (mean ± SEM) is restricted to myeloid cells. B. IO-108 caused 41% (average) tumor growth inhibition (TGI) of LLC in syngeneic LILRB2 transgenic mice. C. Increased spleen size in IO-108-treated mice in B (study day 28) suggests enhanced immune activation. D-F. Immune cell profiling from mice in B (study day 28). D. IO-108 caused increased CD86 expression in tumor-infiltrating M-MDSC and MHCII^{Lo} TAM. E. IO-108 promoted activation, proliferation and cytotoxic activity in tumor-infiltrating T cells. F. IO-108 promoted T cell proliferation of peripheral (splenic) T cells.

	intration fittation		Pre-treatment	On-treatment vs pre-treatment
Low Mo in. High Mo in.			Myeloid cell phenotype	Receptor occupancy
Cancer type 1 (N=585) -	- 4		T cell activation markers	LILRB2 expression
Cancer type 2 (N=1073) –				- -
Cancer type 3 (N=299) -	3	Porinhoral		Myoloid coll phonotypo
Cancer type 4 (N=467) -		Peripiteral	Immune cell counts	· wyelold cell pliellotype
Cancer type 5 (N=494) -	2	biood		
Cancer type 6 (N=486) –	LILRB2		 Soluble HLA-G levels 	 T cell activation markers
Cancer type 7 (N=369) –	expression			
Cancer type 8 (N=504) –				 Immune cell counts
Cancer type 9 (N=360) -				
Cancer type 10 (N=281) -	0			Cytokine & chemokine levels
Breast cancer Colon cancer			LILRB2 protein expression	
LILRB2 CD163	LILRB2 CD163	Tumor	Gene expression studies	Gene expression studies
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			Patient	Confirmation of MoA and
			segmentation	therapeutic hypothesis
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IO-108 induces differentiation of monocytic myeloid cells

IO-108 inhibits tumor growth in a syngeneic mouse model by promoting innate and adaptive immunity

IO-108 clinical biomarker plan

A. Analysis of TCGA RNA-seq database shows that *LILRB2* mRNA expression in solid tumors is associated with a high macrophage infiltration gene expression "signature". B. IHC with a proprietary LILRB2 antibody confirms LILRB2 expression in tumor-associated macrophages (CD163+).