

Introduction

- Our RNA-DRIVEr[™] platform identified MERTK as a key driver of metastatic progression¹
- MERTK is a member of the TAM family of receptor tyrosine kinases
- MERTK Is overexpressed in various solid and hematologic cancers and expression is associated with worse outcomes²⁻¹¹
- Binding of its ligands (Gas6, protein S) drives cancer cell proliferation, survival, invasion and drug resistance^{4, 12}
- MERTK is expressed on M2 tumor-associated macrophages (TAMs), playing a role in the production of immunosuppressive cytokines and efferocytosis¹²
- MERTK expression within retinal pigment epithelium (RPE) has resulted in retinal degeneration in animal models exposed to MERTK complete antagonists¹³

Background

RGX-019

- Humanized mAb with high affinity and specificity to human MERTK (Kd = 3.0 nM) with partial agonist activity¹
- Efficiently internalizes MERTK leading to degradation of surface MERTK
- As a partial agonist, blocks binding of Gas6 ligand and diminishes signal transduction
- Induces expression of pro-inflammatory cytokines in M2 macrophages





Figure 1. (A) RGX-019 Internalization. RGX-019 and IgG control were labeled with a pH-sensitive dye, pHrodo, that becomes fluorescent in low pH (eg. lysosomes).SK-Mel5 melanoma cells were treated with 6.7 nM pHrodo-labeled RGX-019 or IgG control for 24 hr before imaging with Image-Stream flow cytometer. (B) Human cytokine array analysis. M2 macrophages were differentiated from human monocytes by culturing with M-CSF for 7 days, and treated with 6.7 nM RGX-019 or IgG for 48hr. n=4; mean +/- SEM.

RGX-019-MMAE

RGX-019 MMAE-conjugated ADC

- MMAE payload: direct killing of mitotic cells & bystander efficacy
- Site-specific conjugation
- DAR 4
- Cleavable linker
- Additional spacer moiety to increase solubility

2 for 1 approach





2 for 1: Targeting tumor-associated macrophages and cancer cells with a novel MERTK-targeting antibody-drug conjugate (ADC)

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Results

MERTK expression is observed in human solid cancers and AML

Expression of MERTK is seen in the majority of ovarian, urothelial, pancreas, and non-small cell lung cancers (NSCLC)





Figure 2. MERTK immunohistochemistry (IHC) in human tumors. MERTK expression in human tumors across seven indications was analyzed by IHC and quantified as Tumor Proportion Score (TPS). Scale bar 100 µm.

Higher MERTK expression is found in M3 and M5 subtypes of AML as well as AML with t(9;11) translocations or PTPN11 mutations



RGX-019-MMAE demonstrates efficient internalization and target-dependent in vitro cytotoxicity



Davs elapsed (Months)



Figure 4. (A) RGX-019-MMAE internalization. SK-Mel5 and RPMI-8226 cells were treated with 6.7 nM pHrodo-labeled ADCs for 24 hr, and analyzed by flow cytometry. pHrodo MFI was normalized by DOL (Degree of Labeling). (B) Evaluation of in vitro cytotoxicity. SK-Mel5 or MERTK CRISPR-knock-out SK-Mel5 cells were incubated with indicated antibodies/ADCs for 7 days. Cell viability was measured using CellTiter Glo.



RGX-019-MMAE exhibits robust in vivo anti-tumor efficacy MERTK-hiał MERTK-intermediate MERTK-low MDA-MB-231 NCI-H1581 SK-Mel5 (TNBC) (NSCLC) (melanoma) Vehicle control Vehicle cont Vehicle contro RGX-019 RGX-019-MMAE 2 ma/ka RGX-019 - RGX-019-MMAE - RGX-019-MMAE RGX-019-MMAE 5 ma/k Treatment (5 mg/kg) Treatment (3 mg/kg 600 Davs after first treatmer Davs after first treatment

Figure 5. In vivo anti-tumor efficacy in xenograft models. SK-Mel5 or NCI-H1581 cells were injected subcutaneously into NSG or athymic mice, respectively. MDA-MB-231 cells were injected into the mammary fat pad of athymic nude mice. n=8-10, mean +/- SEM. Images represent MERTK IHC staining of control tumors. Scale bar 100 µm.

RGX-019-MMAE shows bystander activity via M2 macrophages



Figure 6. (A) MERTK expression in human immune cells. In vitro differentiated M2 macrophages and other immune cells isolated from human PBMCs were analyzed by anti-MERTK Western blot. (B) Evaluation of bystander-killing by M2 macrophages. M2 macrophages were co-cultured with CFSE-labeled MERTK-KO SK-Mel5 cells in the presence of RGX-019-MMAE for 6 days. Cells were stained with anti-CD14 and anti-MERTK antibodies, and numbers of MERTK-KO SK-Mel5 cells were quantified by flow cytometry. n=3-4; mean +/- SD.

RGX-019-MMAE exhibits anti-tumor immune modulation

To address the contribution to tumor inhibition from the host immune system, we injected MMAE resistant tumor cells (MC38-HM) into either wild type mice or mice that express a humanized version of MERTK

Tumor growth was suppressed only in animals expressing humanized MERTK • RGX-019-MMAE depletes tumoral M2 macrophages



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RGX-019-MMAE treatment is not associated with retinal degeneration in mice

Partial agonist activity of RGX-019 prevents complete inhibition of RPE phagocytosis, which is needed for maintenance of retinal function, as compared to a full MERTK inhibitor





Figure 9. Assessment of retinal degeneration in mice. huMERTK-KI mice were treated with 5 mg/kg IgG1, RGX-019, or RGX-019-MMAE by weekly IV administration for 28 days and eyes were collected for H&E staining. As a positive control, an alkylating agent MMS (methyl methanesulfonate) was administered at 75 mg/kg by a single IP. The thickness of ONL (outer nuclear layer) and IS+OS (photoreceptor inner segment, outer segment) were quantified as measures of retinal degeneration (n=8-10). Scale bar = 50 µm.

Summary

- MERTK is overexpressed in multiple solid and hematologic cancers with expression in healthy tissues generally limited to the retinal pigment epithelium and immune cells, mainly macrophages
- RGX-019 triggers MERTK receptor internalization and partial agonist activity enabling selective delivery of toxic payload to MERTK expressing cancer cells
- Inspirna has developed an RGX-019 based MMAE ADC to enhance cancer cell killing and activate anti-tumor immunity
- Robust in vitro and in vivo efficacy of RGX-019-MMAE is seen in solid and hematologic cancers
- Lack of retinal degeneration observed in cynomolgus monkey¹⁴ and mice expressing huMERTK, after treatment with RGX-019 and RGX-019-MMAE, differentiates RGX-019-MMAE from complete antagonist inhibitors of MERTK

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