Pharmacodynamic Biomarker Characterization of ALX148, a CD47 Blocker, in Combination with **Established Anticancer Antibodies in Patients with Advanced Malignancy**

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Background

- CD47-SIRPα signaling is a myeloid checkpoint mechanism that signals the macrophage to ignore the cell on which CD47 is expressed. Tumors upregulate CD47 to evade the immune response
- ALX148 is a fusion protein with a high affinity CD47 blocker linked to an inactive human immunoglobulin Fc region (Figure 1). It blocks the CD47-mediated 'don't eat me' signal and enhances anti-tumor immunity.
- AT148001, a first-in-human phase 1 study, evaluates ALX148 administered as a single agent (Part 1) and in combination with established anti-cancer antibodies (Part 2).24

Figure 1. ALX148: A Unique High Affinity SIRPa Fusion Protein¹



Molecular weight half the size of typical antibody. • Fc domain enables antibody-like PK

 ${\sf SIRP}\alpha$ with picomolar binding affinity

Figure 2. ALX148 Bridges Innate and Adaptive Immunity¹



ALX148 Designed to Safely Maximize Anti-Cancer Antibody Activity

Selectively binds CD47 to block its interaction with

Fc domain mutated to eliminate Fcy receptor binding



ALX148 Increases Ratio of Inflammatory M1 to Suppressive M2 in Tumor



ALX148 Activates Dendritic Cells and Enhances Cross-Priming of T Cells

References

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Methods

AT148001 Study Design

• Part 1 (single agent): No MTD reached, maximum administered dose 30 mg/kg QoW • Part 2 (combination): ALX148 (10 mg/kg QW) combined with standard regimens of pembrolizumab (200 mg IV Q3W) or trastuzumab (8 mg/kg IV + 6 mg/kg Q3W)

Table 1. ALX148 Combination Solid Tumor Cohorts



- Patients with HER2-positive advanced malignancy (n=30), including gastric/GEJ cancers progressed on or after trastuzumab therapy, received ALX148 + trastuzumab. Patients with advanced malignancy (n=52,) including HNSCC progressed on platinum-based chemotherapy, as well as NSCLC [progressed on prior checkpoint inhibitor (CPI) or tumor proportion score (TPS) <50%] received ALX148 + pembrolizumab.
- Archival tumor tissue, on-treatment biopsies and whole blood samples were collected. CD8, CD68, CD163, PD-L1 and HER2 expression in tumor tissue were neasured by immunohistochemistry (IHC) assays. Percent positive values for CD8, CD68 and CD163 were derived by image analysis. PD-L1 (Clone 22C3) TPS and combined positive score (CPS) were obtained by central pathologist review. HER2 levels were determined using HercepTest™.
- RNA expressions from paired tumor biopsies were assessed using NanoString IO360 Panel. Cell type abundance and pathway profiling analyses using pre-defined gene signatures were performed using NanoString nSolver Analysis Software.
- Peripheral CD47 Target Occupancy (TO) and immunophenotyping were measured by flow cytometry

Results

Note: Data Cutoff 18Apr2019

Patient Baseline Characteristics

• 82 patients with advanced solid tumor malignancies have been enrolled into Part 2 combination cohorts (Table 2)

Table 2. Baseline Characteristics

	Pembrolizumab N=52	Trastuzumab N=30
Median Age, Years (range)	61 (32-81)	58 (45-79)
Sex, n		
F	23	9
Μ	29	21
Race, n		
White	34	13
Black	3	1
Native American	1	-
Asian	11	14
Other	3	2
ECOG PS, n		
0	16	8
1	36	22
Primary Disease, n		
NSCLC	26	-
HNSCC	20	-
Gastric/GEJ/Esophageal	-	25
Breast	-	2
Colorectal	2	-
Ovarian	2	1
Pancreatic	-	1
Peritoneal	1	-
Appendiceal	1	-
Urothelial	-	1

Clinical Results

• ALX148 in combination with trastuzumab and pembrolizumab was well tolerated with no maximum tolerated dose reached. Treatment related adverse events (TRAE) were mostly of low grade and frequency and have been previously reported.^{2.4}

Figure 3. ALX148 + Trastuzumab Clinical Activity in HER2 positive Gastric/GEJ Cancer Expansion Cohor



2019 Patients who received at least one dose of AI X148 in the expansion phase, had a baseli assessment, and at least one post-baseline disease assessment. One patient with clinical progression plots; ND = Not Done; HER2 Score retrospectively assessed using archival tissue by a central IHC lab.

Figure 4. Al X148 + Pembrolizumab Clinical Activity in Advanced HNSCC Expansion Cohorts



Notes: Data Cutoff 23Sept2019. Patients who received at least one dose of ALX148 in the expansion phase, had a baseline assessment, and at least one post-baseline disease assessment; One patient with progressive disease n included in plots.

Figure 5. ALX148 + Pembrolizumab Clinical Activity in Advanced NSCLC Expansion Cohort



Note: Data Cutoff 23Sept2019. Patients who received at least one dose of ALX148 in the expansion phase, had a baseline assessment, and at least one post-baseline disease assessm



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Figure 6. Complete CD47 Target Occupancy by ALX148



Figure 7. No Changes of Circulating Immune Cell Profiles Following ALX148 Combinations





- ALX148 + Trastuzumab (N=10) - ALX148 + Pembrolizumab (N=12)

Figure 8. Increased Infiltrating Macrophages or Lymphocytes in Intratumoral Areas as essed by IHC Following ALX148 Combination





Figure 10. Effects of ALX148 Combinations on Immune Pathways in Tumor Biopsies As Assessed by RNA Expression









Figure 13. IHC and RNA Expression Results from Patient 3 Gastric Cancer HER2 (3+)



Note: Mast cells signature for this sample was determined based on a partial gene list

Conclusions

Intended for combination, ALX148 is designed to block the CD47-SIRPa myeloid checkpoint pathway, safely maximizing innate and adaptive immune response to cancer.

- ALX148 in combination with standard regimens of trastuzumab or pembrolizumab is well tolerated with a favorable hematologic safety profile.2-4
- ALX148 demonstrates emerging anti-cancer activity that compares favorably with anticipated activity of trastuzumab alone in patients with advanced gastric/GEJ cancer that has progressed on prior HER2-targeted therapies; ALX148 also demonstrates emerging anti-cancer activity that compares favorably with single agent pembrolizumab in patients with HNSCC that has progressed on prior systemic therapy.
- ALX148 at 10 mg/kg QW achieves complete CD47 target occupancy with no impact on circulating immune cells in combination with trastuzumab and pembrolizumab.
- Assessment of tumor infiltrating immune cells and molecular signatures in paired tumor biopsies reveals differences in pharmacodynamic changes following different ALX148 combinations, providing insights to ALX148's mechanism as a myeloid checkpoint inhibitor. Further studies are needed to confirm the relationship betweer pharmacodynamic changes and clinical responses
- ALX148 in combination with pembrolizumab showed increased immune infiltration of both innate and adaptive immune cells as well as increased inflammation signatures.
- ALX148 in combination with trastuzumab showed increased macrophage infiltration without increased inflammation signatures

Patients in combination cohorts continue to be followed (NCT03013218)