Neoadjuvant CD40 agonism remodels the tumor immune microenvironment in locally advanced esophageal/gastroesophageal junction cancer

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BACKGROUND AND RATIONALE

Neoadjuvant chemoradiation (CRT) followed by surgical resection is the standard of care for patients with locally advanced esophageal/ gastroesophageal junction (E/GEJ) cancer. A pathologic complete response (pCR) at surgery is associated with improved survival outcomes. Sotigalimab (sotiga) is a potent CD40 agonist mAb capable of inducing and expanding anti-tumor immune responses [1].

In a phase II clinical trial of sotiga combined with neoadjuvant CRT in patients with locally advanced E/GEJ cancer, we saw pathologic complete responses (pCR) in 38% of patients [2]. Here, deep immune profiling was performed on samples from the circulation and tumor microenvironment (TME) from a subset of patients to gain insight into the mechanism of action of sotiga.

RESULTS





RESULTS CONTINUED



Figure 4. Single-cell immune repertoire analysis demonstrates induction of CD8⁺ T cell clones post-sotiga. A. Quantitative & B&C. network analysis of TCR by T cell subtype in all T cells (A – TME, B&C – PBMCs and TME) (A&B; Pre-existing – before sotiga - pink, Induced – only present after sotiga - green, Persistent - present pre and post - blue). B. Analysis by timepoint. C. Analysis by subtype (CD4⁺ T cellspink, CD8⁺ T cells- green, Tregs- blue).



STUDY PROTOCOL



This study reports on data from an initial cohort of patients from a larger study [2]. Blood samples (PBMC) were collected before (A) and after treatment (B), and after surgery (N=6). Paired tumor biopsies were collected before (A) and after (B) sotiga treatment (N=4). When available, tumor samples were collected from surgical specimens (C).

METHODS





Figure 1. Treatment with sotiga increases immune infiltration into the TME, including T cells and myeloid cells. **A.** Representative images of MIBI analyses of immune cells (CD45+, yellow), T cells (CD3+, orange) and myeloid cells (CD68+, magenta) of pre- and post-treatment biopsies from Patient 04. B. The density of each immune cell type per tumor area (n=4).



Figure 2. Tumor-infiltrating myeloid cells are activated postsotiga. A. Representative images of MIBI analyses of DCs (CD11b+, CD11c+, HLA-II+, white arrow; CD86+ activated DCs) of pre- and post-treatment biopsies from Patient 04. **B.** Quantification of DC and DC-subtype density. **C.** Expression of activation markers CD86 and HLA-II on DCs.





Figure 5. Sotiga induces effector function (x-axis) in association with downregulation of oxidative phosphorylation (y-axis) in monocytes/macrophages, DCs, CD8⁺ and non-Treg CD4⁺ T cells in the TME. scRNAseq analysis was used to calculate scores for metabolic and functional phenotypes based on gene expression in individual cells pre- and post-sotiga treatment. Scores for cells from the TME were then plotted for individual cells in dot-plot analysis.

CONCLUSION



Treatment with sotiga induced the activation of antigen presentation leading to the downstream generation of novel T cell clonotypes, enhanced T cell activation, and altered immune cell metabolism.

- **1.** Biopsy samples were analyzed by IonPath using multiplex ion beam imaging (MIBI).
- 2. Tumor samples were also dissociated and scRNAseq and TCR sequencing were performed using the 10X Genomics platform.
- **3.** PBMCs were isolated from whole blood, scRNAseq and TCR sequencing were performed using the 10X Genomics platform.



Figure 3. Sotiga induces an activated T cell infiltrate and reduces Tregs in tumors. A. Representative images of MIBI analyses of T cells (CD45RO-, naïve; CD45RO+, memory; granzyme B+, activated; FoxP3+, Treg) of pre- and posttreatment biopsies from Patient 04. B. Quantification of proportions of T cell subsets in the TME.

- This is the first demonstration that single-agent systemic sotiga can induce significant inflammatory responses in the TME.
- The conversion of "cold" tumors to "hot", a key mechanism of sotiga, is the foundation of immunotherapy.

REFERENCES

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