Integrated biomarker trials to evaluate myeloid and lymphoid composition of HNSCC and solid tumors treated with pepinemab and combinations with checkpoint inhibitors

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Summary

Anti-semaphorin 4D (SEMA4D, CD100) blocking antibody promotes immune infiltration, Recruitment of immunosuppressive myeloid cells into the tumor microenvironment (TME) is a critical limitation to the efficacy of immune checkpoint inhibitors (ICIs) in patients with head and neck squamous cell carcinoma (HNSCC). In preclinical models, antibody blockade of Semaphorin 4D (SEMA4D, CD100) reduced function and recruitment of immunosuppressive myeloid cells within the TME. Importantly, combinations of anti-SEMA4D with ICIs enhanced T cell activity and tumor regression. Pepinemab, an IgG4 humanized monoclonal antibody targeting SEMA4D, is currently being evaluated in window of opportunity, integrated biomarker trials to characterize immunomodulatory effects of treatment.

SEMA4D exerts multi-faceted effects within the tumor microenvironment by creating a parrier at the tumor-stroma margin restricting immune cell infiltration and promoting mmunosuppressive activity of myeloid-derived cells. In preclinical in vitro and in vivo studies, blocking antibody to SEMA4D directly enhanced M1/M2 ratio and reduced both expression of chemokines that recruit MDSC and the ability of MDSC to suppress T cell activity. SEMA4D antibody treatment simultaneously restored the ability of dendritic cells and cytotoxic T cells to infiltrate the TME and increased ratio of Teffector to Tregulatory cells. This coordinated shift from immunosuppression to tumoricidal activity complemented effects of other immunotherapies in syngeneic tumor models

At present, three biomarker-driven window of opportunity trials are recruiting patients with four resectable indications to investigate novel combinations of pepinemab with ICIs; 1) HNSCC (NCT03690986, n=36), 2) pancreatic ductal adenocarcinoma and colorectal cancer with resectable liver metastases (NCT03373188, n=32), and 3) metastatic melanoma (NCT03769155, n=36). Presurgical treatment cohorts include combinations of pepinemab with nivolumab and /or ipilimumab, single agents, or no treatment. Three to seven weeks later, surgically resected tumors are collected under the guidance of a pathologist for comparison of tumor infiltrates across treatment groups and with a pre-dose tissue biopsy. Blood is collected for PK, PD, and additional correlative biomarker assessments. Correlative multiplex flow cytometric and immunohistochemistry panels have been established to phenotype cells in the TME and periphery; preliminary biomarker analysis will be presented.



RPMI 8226 Multiple Myeloma M1 = CD14⁻CD16⁺ M2 = CD14⁺CD16⁺ MDSC Recruitment: In vitro treatment of MOC1 tumor cells In vivo treatment: MOC1 TME CXCL CXCL1 CXCL2 _ CXCL1 CXCL2 CXCL5 CTRL αS4D CTRL αS4D CTRI aS4D

SEMA4D blockade increases ratio of M1/M2 when exposed to SEMA4D+ tumors. (A) Human PBMC were cultured with conditioned tumor media from co-culture of multiple myeloma RPMI 8226 vith human bone marrow stroma (mock), HNSCC tumors HN6 or Cal27, and incubated with Control Ig or anti-SEMA4D/2503 (α S4D) for 24 hours. M1 and M2 were determined by flow cytometry. SEMA4D directly promotes function of myeloid derived suppressor cells (MDSC); suppression

is reversed by antibody blockade. (B) MDSC were isolated from HNSCC MOC1 in vivo tumors and co-cultured ex vivo with rSEMA4D (10 µg/ml) or antibodies, in presence of naïve T cells labeled with CFSE in a T cell suppression assay (right panel). gMDSC were assessed for ARG1 expression via qRT-PCR after *ex vivo* exposure to recombinant protein or antibody for 3 hours (n=5/group). Arginase production in TME suppresses T cell function (right panel)

SEMA4D Mab inhibits cytokines that recruit MDSC (C) MOC1 tumor cells cultured in vitro were exposed to Sema4D mAb (10 μg/mL) or isotype for 24 hours and analyzed for myeloid chemokine expression by qRT-PCR. (D) Mice bearing MOC1 tumors were treated *in vivo* with isotype control or anti-Sema4D Ab (αS4D) (n=5/group). Whole tumor digests were analyzed for myeloid chemokine expression via gRT-PCR.

PRECLINICAL: Anti-SEMA4D Mab neutralizes SEMA4D barrier at tumor margin and shifts the balance of tumor immunity B. Increased infiltration of pro-inflammatory C. Shift in balance of tumor immunity in TME gradien APC and CD8+ T cells Chemokine Teff:Treg



SEMA4D is strongly expressed at the invasive margin of tumors. Antibody blockade of SEMA4D facilitates migration of APCs and T cells into the tumor and a shift toward anti-tumor immune activity. A) SEMA4D expression at invasive margin of murine Colon26 tumors restricts infiltration of PLXNB1+ DC into TME. Brackets indicate area of SEMA4D gradient. B) Anti-SEMA4D MAb promotes infiltration of pro-inflammatory CD11c+/F4-80+ antigen presenting cells, which recruit and activate CD8+ T cells within TME. Colon26 tumor-bearing mice were treated with Control Ig or anti-SEMA4D/MAb67 antibodies (50 mg/kg, weekly IP) and tumors were harvested on day 27 and FFPE sections were stained by IHC or C) tumors were dissociated and assessed for immune cell markers by flow cytometry. Leukocytes were enriched from whole tumor digests using lympholyte-M and cultured for 2-days and supernatants were assessed for chemokine/cytokine secretion using cytometric bead array, or for T cell activity by ELISPOT, n=8-12 mice/group.

SEMA4D and PLXNB1 are over-expressed in HNSCC tumor and myeloid cells compared to non-malignant tissue



PRECLINICAL: Combination Immunotherapy



Immunomodulatory effects of SEMA4D blockade can enhance other immunotherapies. A) MOC1 HNSCC (5x10⁶ cells) were subcutaneously implanted into C57BI/6 mice, that were then treated with aSEMA4D/MAb67 (10 mg/kg, weekly IP), aCTLA-4 / MAb 9H-10 (5 mg/kg, q5D); n=10. B-D) Colon26 (500,000 cells) were subcutaneously implanted into Balb/c mice, that were then treated with αSEMA4D / MAb67 (10 mg/kg, weekly IP X4), αLAG3/C9B7W (10 mg/kg 2x/week X4; n=20); αCTLA-4 / MAb UC10-4F10 (100/50/50 μg, q3 days; n=20), αPD-1 / MAb RMP1-14 (10 mg/kg, twice/week, starting on day10, n=20).



SEMA4D+ immune cells reside in the stroma but are excluded tumor. B) SEMA4D and Plexin receptors are over-expressed in tumor tissue. of SEMA4D and Plexin receptors were assessed by flow cytometr







tolerability and efficacy of pepinemab in combination with avelumab in 62 subjects with advanced (IIIB/IV) NSCLC.





Integrated Biomarker Window of Opportunity Study: HSNCC, MSS CRC with resectable liver metastases, PANC, Melanoma





The key observations relate to distribution of T cells and MDSC. Surgical resections were analyzed from one CRC patient/treatmer arm following 3-5 weeks of treatment with pepinemab, pepinemab + ipilimumab, and one patient who did not receive antibody treatment No conclusion can be drawn regarding tumor necrosis because patients received neoadjuvant chemotherapy prior to surgery A) 5 micron FFPE sections were stained sequentially for each marker and scanned at 40X. Scans were co-registered for each stain in Invasive margin: CD8+ T cells (red) overlays on cytokeratin stain (green) at tumor/normal liver margin are shown (3.3x) (Bottom) Center of tumor: with S100A9+/CD33+ MDSC (blue) overlays (10x). B) Total number of S100A9+/CD33+ cells were quantified d area, normalized by area of tumor bed using Visiopharm software, and 2 sections/patient were averaged in bar graphs, C) Percent of total tumor bed area for each component was quantified and averaged from 2 sections/patient, D) Target saturation ative data from melanoma patient 06A dosed on Days 1 and 21 with pepinemab (15mg/kg) and nivolumab (240mg); rested 15-28 days later. TIL were isolated and CD3+ T cells were assayed for saturation of the cellular SEMA4D target. Isotype, Anti Hu IgG4, Anti Hu IgG4 + pepinemab Similar data were observed in TIL isolated from HNSCC patients