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Targeting Macrophages to Reduce Colorectal Cancer Metastasis: Diminished Effect in the Alcohol-injured Liver

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Summer Undergraduate Alcohol Research Program



Background and Aims

The dissemination of colon cancer to the liver is a major cause of morbidity and mortality in colorectal cancer patients. Alcohol use is a risk factor for colorectal liver metastasis (CRLM) yet contributing mechanisms are undefined.

Macrophages play a crucial role in alcohol-associated liver disease (AALD) as well as in CRLM. Studies indicate that treatment with gadolinium chloride (GdCl3) leads to the inactivation of Kupffer cells (KCs), the resident liver macrophage population. Also, it is known that carcinoembryonic antigen (CEA) from colorectal cancer cells is internalized by KCs leading to protumor cytokine and chemokine production and the promotion of liver metastasis.

However, it is not known whether targeting macrophages with GdCl₃ to reduce CRLM would be effective considering the flux of macrophage phenotypes in the liver. Furthermore, it is not known how preexisting alcohol-mediated liver injury will affect GdCl₃ inactivation of macrophages and hepatic colorectal cancer tumor burden.

The aim of this study was to determine the effect of GdCl₃ inactivation of KCs after CRLM establishment in healthy and alcohol-injured livers.

Methods

Preclinical model of AALD: C57BL/6 mice were fed control (C) or 5% ethanol (E) Lieber-DeCarli diets for 4 weeks to establish alcohol-associated liver injury.

Alcohol-CRLM model and treatments: The C/E-fed mice were intravenously injected with GdCl₃ (20mg/kg) or saline 9 and 12 days after the intrasplenic delivery of 1×10^{6} MC38+CEA colorectal cancer (CRC) cells.

Assessments: Animals were sacrificed 13 days following CRC cell injections and continued C/E diets. Tumor burden, AALD, macrophage phenotype, and cytokine expression were assessed in serum and liver tissue. Assays performed by W.L:

PCR:

- RNA was isolated from frozen liver tissue using the PureLink RNA Mini Kit (ThermoFisher).
- Quantification of isolated RNA was conducted using the Ribogreen Assay.
- cDNA was generated from RNA using the High-Capacity cDNA Reverse **Transcription Kit**
- TaqMan Real-Time PCR was conducted on cDNA to detect cytokine/chemokines: TNF- α , Nos2, TGF- β , IL-10, CD163, and HO-1.

Immunohistochemistry:

- Formalin-fixed, paraffin-embedded liver sections were dewaxed and subject to antigen unmasking using citric acid-based solution.
- The sections were incubated overnight at 4° with primary antibodies specific for smooth muscle α -actin (SMA, Abcam 5694), caspase-3 (R&D AF835), and heme oxygenase 1 (HO-1, Abcam 189491).
- Secondary antibody incubation was done using anti-rabbit HRP (Cell Signaling).
- Immunofluorescent detection was performed by tyramide signal amplification using AlexaFluor 647, 568, and 488 tyramide (ThermoFisher).
- Fluorescent micrographs were captured using a Nikon Eclipse 80i microscope.

Targeting Macrophages to Reduce Colorectal Cancer Metastasis: Diminished Effect in the Alcohol-injured Liver

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