RESEARCH ARTICLE

Ferroptosis Induced Following Iron Sucrose Injection into Ascites of Hepatocellular Carcinoma

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PUBLISHED

30 April 2025

CITATION

Yu, B., Zhang, H., et al., 2025. Ferroptosis Induced Following Iron Sucrose Injection into Ascites of Hepatocellular Carcinoma. Medical Research Archives, [online] 13(4).

https://doi.org/10.18103/mra.v13i4.6472

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DOI

https://doi.org/10.18103/mra.v13i4.6472

ISSN

2375-1924

ABSTRACT

Most of advantage stages of HCC is going into cachexia with asictes which is lacking an effective treatment. Ferroptosis-Enhanced Immunotherapy is used for treating ascites of HCC by an chitosan hydrochloride and oxidized dextran (CH-OD) designed to load sulfasalazine (SSZ). Since iron sucrose (IS) commonly used for treating iron deficiency by IV injection and iron is a critical component of many cellular functions. However, excess iron can be highly toxic, resulting in oxidative DNA damage to induce cancer cell death which is called a ferroptosis. Our current study has indicated that the interaction between mesothelial cells, MPs, neutrophils, TandNK and B cells are induced by IS local fusion treatment ascites of HCC, which promotes the expression of a large number of genes and leads to upregulation of immune response and feroptosis. Our evidence of study has approved local delivery IS can be used as ferroptosis inducer for ferronan immune adjuvant in treatment of cancer ascites of HCC.

Keywords: iron sucrose; ferroptosis; immunotherapy with iron sucrose; local immune inducer; local immune booster.

Introduction

Ascites in advanced hepatocellular carcinoma (HCC) is a complex clinical problem and lacks effective treatment.^{1,2} Their weight and immunity decline, and sometimes lose the ability to take care of themselves, in this condition, any chemotherapy treatment will lead to worsening of the disease, and even lead to death. Therefore, the exploration of non-chemotherapy drugs is useful for take care of ascites.

Iron sucrose (IS) is commonly IV injected for treatment of iron deficiency.3 Iron is a critical component of any cellular functions including DNA replication and repair, and it is essential for cell vitality. However, excess iron can be highly toxic, resulting in oxidative DNA damage to induce cancer cell death.⁴ Ferroptosis-Enhanced Immunotherapy is used for treating ascites of HCC by a chitosan hydrochloride and oxidized dextran (CH-OD) designed to load sulfasalazine (SSZ).5 To enable growth of cancer cells, they have an increased demand for iron compared to normal non-cancer cells. This dependence on iron can make cancer cells more vulnerable to ironcatalyzed necrosis, known as ferroptosis. identification of an FDA-approved ferroptosis inducer has raised high expectations for ferroptosis as a promising new way to kill treatment-resistant cancers.6 There is not report about relation between IS and ferroptosis.

In this study, IS was used for explore of treatment ascites of HCC by local injection replaced IV injection. Tolerance of a single dose for adults and seniors is 10ml (200mg iron) of this product by injection for at least 10 minutes.3 Safety and efficacy of iron sucrose in patients sensitive to iron dextran: North American clinical trial have showed that it can be administered without a test dose by IV push or infusion; and they concluded that iron sucrose injection administered as 1,000 mg in 10 divided doses by IV push without a prior test dose is safe and effective for the treatment of iron deficiency in patients with dialysisassociated anemia.^{7,8} Doctors have reason to believe that IS is safe and common when used locally, usually offlabel. IS was injected with single drug of 10ml (200mg) into ascites of three patients, samples of ascites were taken before and day 1 and day 5 after IS injection. We used single-cell RNA sequencing (scRNA-Seq) to obtain transcriptome profiles of a total of 30909 cells including Mesothelial Cells (UPK3B, WT1, KRT19, MPs ((CSF3R,S1-CD14,LYZ,C1QC), **Neutrophils** 9A,FCGR3B), TandNK (CD3D, CD3E, CD3G) and B cells (CD79A, MS4A1, CD19), and the proportion of changes in immune cells before and after treatment.

Materials and Methods

ETHICAL STATEMENT

All procedures and protocols in the study have been reviewed and approved by the Ethical Committee of the Shandong Baofa Cancer Institute (TMBF 0010, 2021). All informed consent forms for three patients have been signed prior to the start of the study. This experimental treatment was approved by the hospital ethics committee.

CLINICAL SPECIMENS

200 mg of iron sucrose was injected into ascites of HCC,

and no adverse reactions such and bleeding were observed. Ascites samples (25-30 ml) were taken before the treatment as control and after treatment at day 1 and day 5 for scRNA-Seq analysis.

TISSUE DISASSOCIATION AND CELLS COLLECTION

The fresh cells samples were immediately stored in the sCelLiVE® Tissue Preservation Solution (Singleron) on ice. Finally, the single cell suspension was collected after resuspension with PBS, and trypan blue (Sigma) staining was used to calculate cell activity and cell count under a microscope.9

SINGLE-CELL RNA SEQUENCING

Briefly, the scRNA-seq library was constructed using the GEXSCOPE® Single Cell RNA Library Kits (Singleron). The library was lastly sequenced with 150 bp that was diluted to 4nM and paired-end reads on the IlluminaHiSeq X platform following an established protocol. Sequencing data processing and quality control was performed as described in previous publications. 10, 11

DATA PROCESSING AND ANALYSIS

To identify differentially expressed genes (DEGs), genes expressed in more than 10% of the cells were selected in both groups of cells and with an average log (fold changes) value greater than 1 as DEGs. The InferCNV package was used to detect the CNAs in malignant cells. Non-malignant cells (T and NK cells) were used as control references to estimate the CNVs of malignant cells. Genes expressed in more than 20 cells were sorted based on their loci on each chromosome. To investigate the potential functions of DEGs between clusters, the Gene Ontology (GO) and Kyoto Encyclopedia of Genes and Genomes (KEGG) analysis were used with the "clusterProfiler" R package 3.16.1.12

Cell-cell interaction (CCI) between B cells, Epithelial cells, Fibroblasts, Mononuclear phagocytes, Mast cells, Neutrophils, T and NK cells were predicted based on known ligand—receptor pairs by Cellphone DB v2.1.0.13

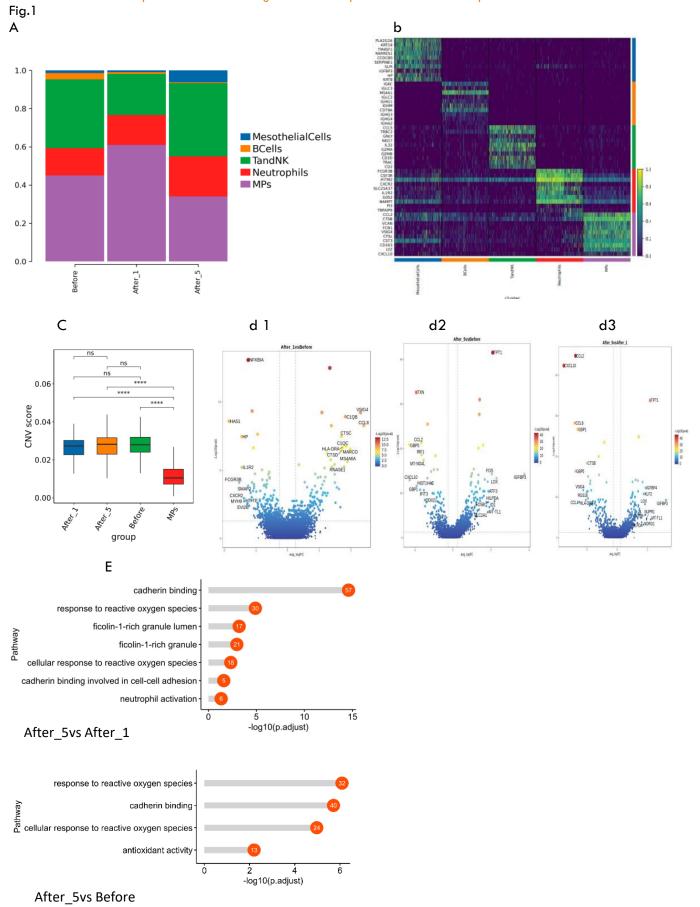
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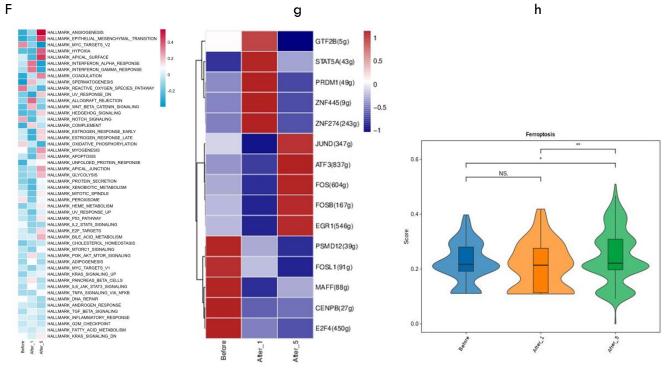
CLINICAL BENEFIT

After IS treatment, patient felt a relive from incomfortable, ascites of HCC is stable for a long time without pumping out of ascites, no more side effect like chemotherapy and extend her life for 6 months, a low-grade fever is the only sensation and may be related to an immune response.

LANDSCAPE OF SINGLE CELL TRANSCRIPTOME SEQUENCING BEFORE AND AFTER TREATMENT

The three samples had a total of 30,909 cells, annotated as 5 cell types, mesothelialCells (UPK3B, WT1, KRT19), MPs (CD14, LYZ, C1QC), neutrophils (CSF3R, S1-9A, FCGR3B), TandNK (CD3D, CD3E, CD3G) and B, respectively cells (CD79A, MS4A1, CD19) (Fig. 1a/b). The proportion of TandNK decreased at 1 day after treatment, but increased at 5 days after treatment. The proportion of neutrophils increased 5 days after treatment. MPs increased 1 day after treatment and decreased 5 days after treatment (Fig. 1 b).





- A. Histogram of the proportion of cell types from different sample sources;
- B. Heat-map display of Top10 differential genes in different cell types;
- C. CNVscore between mesothelial cell groups;
- D. D1Differential gene volcano map compared with 1d after treatment and before treatment;
 - D2: Differential gene volcano map compared with 1d after treatment and before treatment;
 - D3: Differential gene volcano map 5 days after treatment compared with 1 day after treatment;
- E. The upregulated gene enrichment results of mesothelial cells 5 days after treatment;
- F. Inter-mesothelial cell GSVA Analysis (Hallmark gene set);
- G. Intergroup transcription factor analysis of mesothelial cells;
- H. Ucell analysis of iron death gene set in mesothelial cells.

CHANGES OF MESOTHELIAL CELLS BEFORE AND AFTER TREATMENT: IRON DEATH GENE SCORE

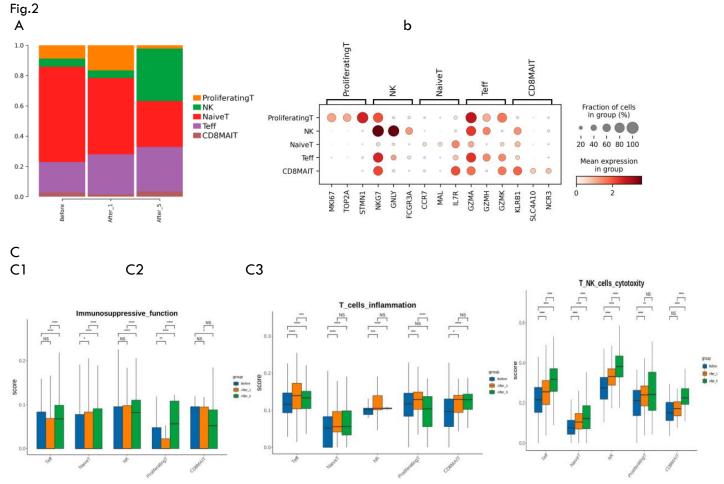
The inferCNV analysis of mesothelial Cells against MPs showed deletion on chromosomes 1, 6, 19 and 21, and significant amplification on chromosomes 2, 3, 4, 7, 8, 12, 16 and 18 (Fig. 1c). The results of CNVscore showed that significant changes occurred before treatment atlday and 5day, but there was no significant difference among the three patients (Fig. 1d). The expression of immunerelated genes such as CCL8, VSIG4, MARCO, C1QC, C1QB and CTSD was up-regulated 1 day after treatment compared with that before treatment (Fig. 1d). The expressions of IGFBP3, LOX, HILPDA, MT-TL1, TPT1, FOS, ID1, ATF3, FOSB, SLC2A1 and other genes were up-regulated 5 days after treatment compared with those before treatment (Fig. 1d) and mesenchymal cell enrichment analysis after 5 days of drug treatment revealed activation of ROS related biological processes and molecular functions (such as reactive oxygen species response, cadherin binding, and antioxidant activity pathways) (Fig. 1e).

The function of each cluster was identified based on GSVA analysis (hallmark database). The After_1 group mesenchymal cells enriched IFN- α and IFN- γ signaling pathways, and the After_5 group mesenchymal cells enriched epithelial mesenchymal transformation and heme metabolism (Fig. 1 f).

Pyscenic analysis was used to evaluate the significantly active regulators in the TF regulatory network of mesothelial cells in different groups, and the results showed that: The transcriptional activity of ATF3 and EGR1 transcription factors in mesodermal cells was enhanced after 5 days of drug treatment, and activation of transcription factor 3 (ATF3) (Fig.1g). Intergroup iron death gene set analysis was performed on mesothelial cells based on Ucell analysis, and it was found that the iron death gene set score 5 days after drug treatment was significantly higher than that before treatment and 1 day after treatment (Fig.1h).

CHANGES OF TANDNK CELLS AFTER TREATMENT

There are 8587 TandNK cells in total, which are proliferating T, NK, NaiveT, Teff, CD8MAIT respectively after marker gene annotation(Fig. 2a). The proportion of Teff cells increased successively. The proportion of NK cells at 5day increased significantly after treatment. Proliferating T ratio increases on 1 day after treatment, and declines on 5 day after treatment (Fig. 2a). T cell receptor signaling pathway expression increases in Proliferating T, NK, NaiveT, Teff, and CD8MAIT 5 days after treatment, and Th1, Th17, and Th2 cells differentiate in NK cells and decline 5 days after treatment (Fig. 2b). The inflammatory effect of T cells was increased at 1 and 5 days after NK, NaiveT, Teff and CD8MAIT treatment. Toxic effects in Teff, NaiveT, NK, Proliferating T, and CD8MAIT increase sequently 1 day after treatment and 5 days after treatment (Fig. 2c).



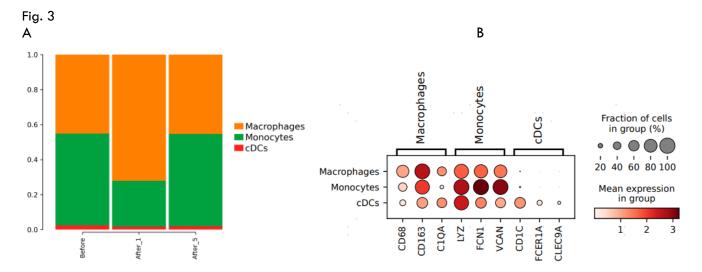
- A. TandNK subpopulation cell proportion histogram;
- B. marker gene bubble map of TandNK subgroup;
- C. C1: Marker gene bubble map of TandNK subgroup;
 - C2: T cell inflammation gene set score;
 - C3: T cell toxicity gene set score.

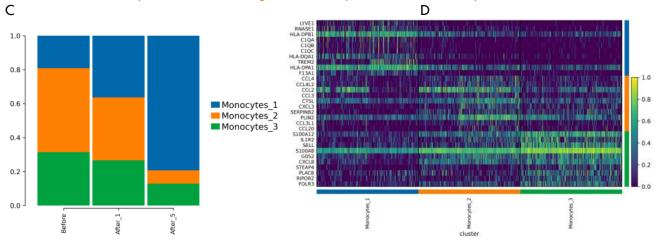
CHANGES OF MONOCYTES AFTER TREATMENT

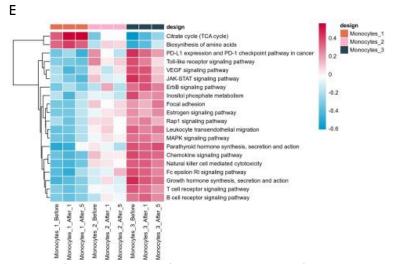
MPs cells are divided into Macrophages, Monocytes, and cDCs. Macrophages increased in 1 day and decreased in 5 day post treatment. The proportion of monocytes decreased 1 day after treatment and increased 5 days after treatment (Fig. 3.a/b).

A total of 5109 monocytes were divided into 3 subpopulations after unsupervised clustering, Monocytes_1, Monocytes_2, and Monocytes_3 (Fig. 3.c-e). The cell proportion of Monocytes_1 showed an

increasing trend after treatment, while the cell proportion of Monocytes_2 and Monocytes_3 showed a decreasing trend after treatment (Fig.3 c). Monocytes_1 highly expressed C1QA, C1QB, C1QC and other immune-related genes; Monocytes_2 highly expressed CCL2, CCL3, CCL4 and other chemokine-related genes; Monocytes_3 highly expressed FOLR3 and S100A12 extracellular adhesion molecules (Fig. 3 d). Monocytes_3 mainly has immune-related functions but no significant changes after treatment (Fig. 3 e).







A: Bars of cell proportion of each subpopulation of MPs

B: Marker gene of MPs grouping;

C: Cell proportion bar chart of monocytes;

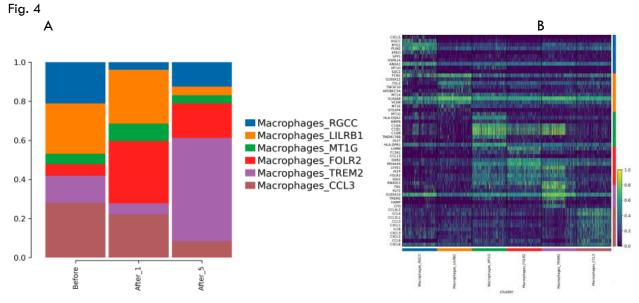
D: Cell proportion bar chart of monocytes;

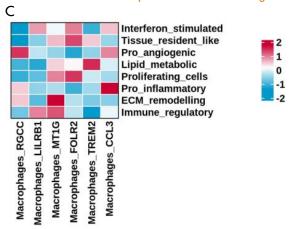
E: Intergroup GSVA analysis of different subpopulations

CHANGES OF MACROPHAGES AFTER TREATMENT

Macrophages consist of 7776 cells divided into 6 subpopulations Macrophages_CCL3, new Macrophages_FOLR2, new ges_lilrb1, new ges_mt1g, new ges_rgcc and new ges_TREM2 (Fig. 4 a/b). Macrophages_RGCC is decreased 1d and increased 5 days after treatment. Macrophages_LILRB1 has no significant change in 1d posttreatment, but significantly

decreased 5 day post-treatment. Macrophages_FOLR2 increased in 1 day, but decreased in 5 day. The proportion of Macrophages_TREM2 increases significantly after 5 days, while the subgroup of Macrophages_CCL3 gradually decreases (Fig. 4 b). Macrophages_CCL3 is a proinflammatory macrophage (Fig. 4 c).



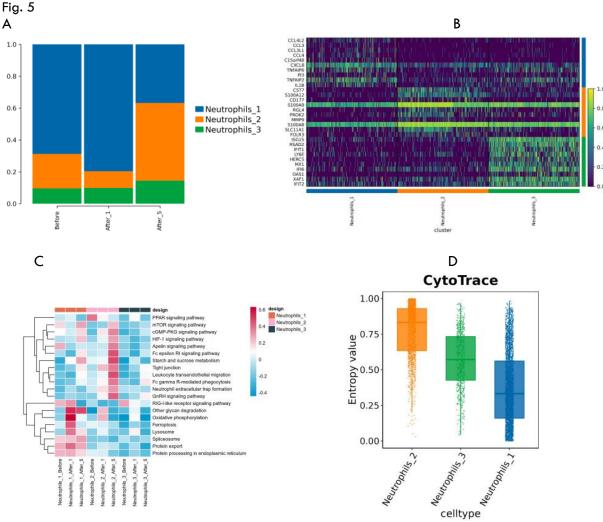


- A: The cell proportion after macrophage cluste;
- B: Differential gene heat map after clustering;
- C: Macrophage subpopulation function assessment gene set scoring heat map.

CHANGES IN NEUTROPHILS AFTER TREATMENT

There were 5117 Neutrophils in total, which were divided into 3 subpopulations by unsupervised clustering, namely Neutrophils_1, Neutrophils_2, and Neutrophils_3 (Fig. 5 a). The proportion of Neutrophils_1 increased on day 1 and decreased on 5 days after treatment. The proportion of Neutrophils_2 decreased at 1 day and increased at 5 day. However, Neutrophils_3 did not change significantly at 1 day and 5 days after treatment (Fig. 5 a).

Neutrophils_3 highly expressed IFIT1 and IFIT2, indicating that this group was related to interferon response (Fig. 5 b). Glycolysis, oxidative phosphorylation, and iron death increased 1 day after Neutrophils_1 treatment compared with before treatment, but decreased 5 days after treatment. Immune-related pathways were elevated at 1 and 5 days after Neutrophils_2 treatment (Fig. 5 c).



- A. The cell proportion of each subgroup of neutrophils;
- B. Heat maps showing the top10 differential genes of each subgroup of neutrophils
- C. GSVA analysis;
- D. CytoTrace analysis of neutrophils

Discussion

Iron sucrose (IS) was commonly used for treatment of iron deficiency and iron is important for cancer patient's nutrition.³ Ferroptosis-Enhanced Immunotherapy is reported for treating ascites of HCC by a chitosan hydrochloride and oxidized dextran (CH-OD) designed to load sulfasalazine (SSZ) and induces higher levels of immunogenic ferroptosis, then combined with anti-PD-1 immunotherapy to control ascites of HCC.⁵

In this study, we used IS local injection in to ascite of HCC to explore if treat ascites of HCC and induce ferroptosis related with immune response including MesothelialCells (UPK3B, WT1, KRT19), MPs (CD14, LYZ, C1QC), Neutrophils (CSF3R, S1-9A, FCGR3B), TandNK (CD3D, CD3E, CD3G) and B cells, respectively cells (CD79A, MS4A1, CD19) (Fig.1a/b).

Changes of mesothelial Cells before and after treatment has indicated iron death gene score, The inferCNV analysis of Mesothelial Cells against MPs showed deletion on chromosomes 1, 6, 19 and 21, and significant amplification on chromosomes 2, 3, 4, 7, 8, 12, 16 and 18 (Fig. 1c/d). The results of CNVscore showed that significant changes occurred before treatment, 1 day after treatment and 5 day after treatment. Induction of HCC cells TGF - beta 1 / BMP - 7 pathways imbalances by increasing differentiation inhibitor 1 (ID1) expression significantly promote HCC cells and dye.13 The function of each cluster was identified based on GSVA analysis (hallmark database). The After_1 group mesenchymal cells enriched IFN-α and IFN-y signaling pathways, and the After_5 group mesenchymal cells enriched epithelial mesenchymal transformation and heme metabolism, among which: Has a tendency to literature reported death iron exists a strong correlation between EMT and death may have a promoting effect on EMT of iron.14 A high propensity for iron death and EMT may result in poor prognosis and non-response to immunotherapy. Have other literature reports after large amounts of iron is input mitochondria, used for synthesis of hemoglobin and hemoglobin disorders has recently been shown to induce cell death of iron (Fig. 1 f,g,h).15 local high dosage of IS has still induced cell death of iron and increased TandNK highly expressed in proliferating cells, and the expression of amino acid biosynthesis, citric acid cycle, thermogenesis and other pathways decreased at 5 days after treatment. According to the characteristic gene set score of TandNK cells, immunosuppressor effect will significantly increase on ProliferatingT, NK, NaiveT, and Teff 5 days after treatment, and will also significantly increase on proliferatingt, NK, Naivet, and TEFF 5 days after treatment compared with 1 day after treatment (Fig. 2).

MPs cells are divided into Macrophages, Monocytes, and cDCs. Macrophages increased in 1 day and decreased in 5 day post treatment (Fig. 3). Multiple tumor studies have reported that tumor macrophages expressed by SPP1 and MMP9 are associated with tumor progression, metastasis, and poor prognosis in the remodeled ECM. 16,17 These macrophages play an immune surveillance and pro-inflammatory role and may actively recruit and regulate immune cells during tumor-related inflammatory responses. 18

In summary, our current study indicated that the interaction between mesothelial cells, MPs, neutrophils, TandNK and B cells are induced by IS treatment ascites of HCC, which promotes the expression of a large number of genes and leads to upregulation of immune response and feroptosis. There is still need to explore this study in large number of ascites of HCC for providing an strong evidence: IS can used for ferroptosis and immune adjuvant in treatment ascites of HCC and regulate the immune cell function.

Credit author statement

Baofa, Yu, MD conceived the concept and provided overall supervision of all experiments and write.
Hongxi Zhang, help to write English
Feng Gao, MD helped the clinical sample collection
Peng Jing, MD, He helped the clinical data collection.
Guoqin Zheng, MS conducted and communicated with single cell company and data analysis.

Conflict of Interest Statement: All of authors do not have any conflict interest for the study.

Translational relevance to the manuscript

This research is valuable for medical practice. The study provides evidence to support possible precision immunotherapy with iron sucrose, a old drug for treating iron deficiency, as off label use for inducing ferroptosis and local immune booster that can potentially be applied to treatment of ascites fluid of hepatocellular carcinoma.

Ethical Statement: All procedures and protocols in the study have been reviewed and approved by the Ethical Committee of the Beijing Baofa Cancer Hospital (TMBF 0010, 2015). All informed consent forms from patients have been signed prior to the start of the study. Ethical Approval: approved by the Ethical Committee of the Beijing Baofa Cancer Hospital (TMBF 0010, 2015).

Data Availability Statement: The data that support the findings of this study are available from [third party name] but restrictions apply to the availability of these data, which were used under license for the current study, and so are not publicly available. Data are however available from the authors upon reasonable request and with permission of [third party name].

Funding: There is not outside funding for this research.

Financial support: This study was sponsored in part by Tai Mei Baofa Cancer Hospital, Dongping, Shandong Province, China 271500.

Clinical trial number: Not applicable.

Translational relevance to the manuscript:

This research is valuable for medical practice. The study provides evidence to support possible precision immunotherapy with iron sucrose, an old drug for treating iron deficiency, as off label use for inducing ferroptosis and local immune booster that can potentially be applied to treatment of ascites fluid of hepatocellular carcinoma.

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