LRP8-mediated selenocysteine uptake is a targetable vulnerability in MYCN-amplified neuroblastoma.

José Friedmann Angeli, Hamed Alborzinia, Marcus Conrad, Bernhard Michalke, Christoph Bartenhagen, Werner Schmitz, Umut Yildiz, Elisa Espinet, Andreas Trumpp, Martin Eilers, Matthias Fischer, Andres Florez, Nesrine Aroua, Giulio Supert-Furga, Gabriele Büchel, zhiyi chen, florencio freitas, Felix Vogel, Julianna Varga, Jasmin batani, Svenja Meierjohann, Enrico Girardi, Ancely dos Santos, Thamara Xavier da Silva, Adriana przybylla, Petra Zeisberger, Almut Schulze, Ulrich Schweizer, Jiashuo Zheng, Tasneem Cheytan, Julie Haenlin, Marietta Fabiano, and Lisa Schlicker **DOI: 10.15252/emmm.202318014**

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25th May 2023

25th May 2023

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***** Reviewer's comments *****

Referee #1 (Remarks for Author):

Ferroptosis is a regulated form of cell death characterized by the accumulation of oxidatively damaged phospholipids. Recent

research suggests that inducing ferroptosis could be beneficial in treating forms of cancer that are resistant to conventional therapies. For example, studies report that MYCN-amplified neuroblastomas are exceptionally sensitive to ferroptosis triggered by GPX4 inhibition. However, the precise mechanisms behind why MYCN-amplified neuroblastomas are particularly sensitive to ferroptosis remain poorly understood.

In the current manuscript, genome-wide CRISPRa screens reveal an exceptional requirement for LRP8 as a ferroptosis suppressor in MYCN-amplified neuroblastoma. Notably, while previous studies have found that LRP8 loss sensitizes cancer cells to ferroptosis induction, this research convincingly establishes that the loss of LRP8 alone is sufficient to trigger ferroptosis in MYCN-amplified neuroblastoma. A key strength of this study lies in the characterization of the underlying mechanism behind the LRP8 dependency. Their investigations demonstrate that the LRP8 dependency is due to the low expression of system Xc-and, consequently, the limited uptake of selenium. This pivotal insight not only reaffirms the significance of LRP8 but also sheds light on the critical role of selenium metabolism as a key therapeutic vulnerability in MYCN-amplified neuroblastoma.

The experimental design and execution presented in this manuscript are exemplary, reflecting rigorous methodology and thoughtful analysis. The authors effectively address previous reviewers' comments, further strengthening the overall quality of their work. Its implications for targeted therapeutic approaches in this devastating form of cancer make it an important addition to the scientific literature. Based on the substantial contribution of these findings to our understanding of ferroptosis regulation in MYCN-amplified neuroblastoma, I recommend immediate publication of this manuscript.

-James Olzmann

19th Jun 2023

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Materials and methods, Human samples: please indicate the details of authority granting ethics approval, provide reference number for approval. Please include a statement confirming that informed consent was obtained from all subjects and that the experiments conformed to the principles set out in the WMA Declaration of Helsinki and the Department of Health and Human Services Belmont Report. Please correct the checklist accordingly (it is currently stated "please check with Matthias Fischer").
 Acknowledgements: The funding information provided in the manuscript should be the same as the one provided in the submission system. Currently, SFB873, "RiskY-AML", the "Integrate-TN" Consortium funded by the Deutsche Krebshilfe, the Dietmar Hopp Foundation, Humboldt Postdoctoral Fellowship, FOR2314, SFB 1399, Förderverein für krebskranke Kinder e.V.

Köln Austrian Academy of Sciences, a Marie Sklodowska-Curie fellowship (MSCA-IF-2014-661491) are missing from the submission system.

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Problem Ferroptosis, a distinct form of cell death, has emerged as a potential approach for eliminating solid tumors that are refractory to treatment. However, inhibiting the selenoprotein glutathione peroxidase 4 (GPX4), which suppresses ferroptosis, remains challenging due to the absence of suitable inhibitors and to the potential systemic toxicity. Our previous findings have demonstrated a notable dependence on GPX4 in high-risk MYCN-amplified neuroblastomas, yet the precise factors contributing to this phenomenon remain incompletely understood.

Results

Our study reveals that high-risk MYCN-amplified neuroblastomas exhibit a significant dependency on LRP8. Through genomewide and single-cell transcriptomics CRISPR-activation screens, we identify the low-density lipoprotein receptor-related protein 8 (LRP8) as a critical factor in selenium/selenocysteine metabolism in MYCN-amplified cancers. We find that MYCN-amplified cells ineficiently activate alternative selenium/selenocysteine pathways, resulting in vulnerability to LRP8 inhibition. These metabolic vulnerabilities provide a unique opportunity to target LRP8 to induce ferroptosis selectively and safely. Impact

In light of the limited success in repurposing adult oncology drugs for neuroblastoma treatment, our research presents a significant advance by introducing innovative strategies based on ferroptosis. By specifically focusing on targeting LRP8, we have identified a potential breakthrough that not only offers novel therapeutic approaches for neuroblastoma but also holds promise for a range of pediatric malignancies and MYCN-driven cancers.

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- Ferroptosis, a cell death modality, is gaining interest as a therapeutic approach against challenging tumors.
- GPX4 is crucial for suppressing ferroptosis, but suitable in vivo inhibitors are lacking, limiting translation to cancer therapies.
- Genome-wide and single-cell CRISPR-activation screens reveal LRP8 as a critical ferroptosis suppressor in MYCN-amplified neuroblastoma.
- Blocking selenium/selenocysteine uptake via LRP8 disrupts GPX4 function and selectively induces ferroptotic cell death.
- LRP8 dependency emerges as the result of the low system Xc- activity suggesting that targeting LRP8 could be explore in other entities such as AML and lymphoma."

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I look forward to receiving your revised manuscript as soon as possible,

With kind regards,

Lise

Lise Roth, PhD Senior Editor EMBO Molecular Medicine The authors addressed the remaining minor editorial issues.

2nd Revision - Editorial Decision

23rd Jun 2023

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Thank you for providing your revised files. I am pleased to inform you that your manuscript is now accepted for publication in EMBO Molecular Medicine pending one minor change:

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With my best wishes,

Lise

Lise Roth, Ph.D Senior Editor EMBO Molecular Medicine

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Journal Submitted to: EMBO Molecular Medicine	
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- The data shown in figures should satisfy the following conditions: the data were obtained and processed according to the field's best practice and are presented to reflect the results of the experiments in an accurate and unbiased manner.

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- → a specification of the experimental system investigated (eg cell line, species name).
- the assay(s) and method(s) used to carry out the reported observations and measurements
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- are there adjustments for multiple comparisons?
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Short novel DNA or RNA including primers, probes: provide the sequences.	Yes	sequences of this focused libary can be found in Supplementary Table 2. Sequence information of individual gRNA-cloning are listed in the Methods -
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Cell lines: Provide species information, strain. Provide accession number in repository OR supplier name, catalog number, clone number, and/OR RRID.	Yes	Human neuroblastoma cell lines: SK-N-FI, SK-N-DZ, cells were purchased from ATCC.
Primary cultures: Provide species, strain, sex of origin, genetic modification status.	Not Applicable	
Report if the cell lines were recently authenticated (e.g., by STR profiling) and tested for mycoplasma contamination.	Yes	Materials and Methods - Cell culture
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Laboratory animals or Model organisms: Provide species, strain, sex, age, genetic modification status. Provide accession number in repository OR supplier name, catalog number, clone number, OR RRID. Animal observed in or captured from the field: Provide species, sex, and age where possible. Please detail housing and husbandry conditions. Plants and microbes Plants: provide species and strain, ecotype and cultivar where relevant, unique accession number if available, and source (including location for collected wild specienes). Microbes: provide species and strain, unique accession number if available, and source.	the manuscript? Yes Not Applicable Yes Information included in the manuscript? Not Applicable Yes Information included in	(Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section) Materials and Methods Materials and Methods In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section) Methods: Bacteria used for amplification of plasmid DNA In which section is the information available?
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Include a statement about sample size estimate even if no statistical methods were used.	Yes	Materials and Methods
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Could your study fall under dual use research restrictions? Please check biosecurity documents and list of select agents and toxins (CDC): https://www.selectagents.gov/sat/list.htm	Not Applicable	
If you used a select agent, is the security level of the lab appropriate and reported in the manuscript?	Not Applicable	
If a study is subject to dual use research of concern regulations, is the name of the authority granting approval and reference number for the regulatory approval provided in the manuscript?	Not Applicable	

Reporting
The MDAR framework recommends adoption of discipline-specific guidelines, established and endorsed through community initiatives. Journals have their own policy about requiring specific guidelines and recommendations to complement MDAR.

Adherence to community standards	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
State if relevant guidelines or checklists (e.g., ICMJE, MIBBI, ARRIVE, PRISMA) have been followed or provided.	Not Applicable	
For tumor marker prognostic studies, we recommend that you follow the REMARK reporting guidelines (see link list at top right). See author guidelines, under 'Reporting Guidelines'. Please confirm you have followed these guidelines.	Not Applicable	
For phase II and III randomized controlled trials, please refer to the CONSORT flow diagram (see link list at top right) and submit the CONSORT checklist (see link list at top right) with your submission. See author guidelines, under Reporting Guidelines'. Please confirm you have submitted this list.	Not Applicable	

Data Availability

Data availability	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
Have primary datasets been deposited according to the journal's guidelines (see 'Data Deposition' section) and the respective accession numbers provided in the Data Availability Section?	Yes	Methods - Data availability
Were human clinical and genomic datasets deposited in a public access- controlled repository in accordance to ethical obligations to the patients and to the applicable consent agreement?		Comparison of RNA-seq and microarray-based models for clinical endpoint prediction. Genome Biol 16, 133 (2015) - described in Materials and Methods
Are computational models that are central and integral to a study available without restrictions in a machine-readable form? Were the relevant accession numbers or links provided?	Not Applicable	
If publicly available data were reused, provide the respective data citations in the reference list.	Yes	Results and Material and Methods