Introduction to HT-KIT an mRNA Frame **Shifting Therapeutic** Significantly Inhibits **Tumor Growth in Mast Cell-Derived Cancers** 





# **HOTH**THERAPEUTICS



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# HT-KIT



HT-KIT is a novel antisense oligonucleotide under development for the treatment of cancers resulting from aberrant KIT signaling. The HT-KIT drug is designed to more specifically target the receptor tyrosine kinase KIT, which is required for the proliferation, survival and differentiation of bone marrow-derived hematopoietic stem cells. Mutations in the KIT pathway have been associated with several human cancers, such as mast cell-derived cancers (aggressive systemic mastocytosis (ASM), systemic mastocytosis with associated hematological neoplasm (SM-AHN), or mast cell leukemia (MCL)), gastrointestinal stromal tumors, and acute myeloid leukemia.

HT-KIT is currently in the pre-clinical stage of development.



# Targeting KIT by frameshifting mRNA transcripts as a therapeutic strategy for aggressive mast cell neoplasms

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Fig. 1. ESO-mediated alternative splicing of exon 4 in *c-KIT* pre-mRNA. (A) HT-KIT ESO was designed to target the donor splice site of exon 4, which led to exclusion of exon 4 by the spliceosome. This is predicted to introduce a premature stop codon due to a frameshift in the open reading frame of the mRNA transcript. Yellow boxes represent exons; thick black bar represents introns. FL-c-KIT = full-length c-KIT, t-c-KIT = truncated c-KIT. (B) RT-PCR demonstrating splice-switching of *c-KIT* by HT-KIT ESO in comparison to control oligo in HMC-1.2 cells. Black arrow = full-length *c-KIT*, orange arrow = alternatively spliced truncated *c-KIT* mRNA. (C) qRT- PCR of KIT transcripts in HMC-1.2 cells demonstrates a significant reduction in KIT mRNA transcripts with HT-KIT relative to control cells after correction against  $\beta$ -actin.





Fig. 2. HT-KIT ESO reduces KIT expression and induces apoptosis of neoplastic human mast cells. (A) Mean flow cytometry data for total KIT expression calculated from the geometric MFI. (B) Combined data from flow cytometry for Annexin V expressed as the geometric MFI. (C) Combined geometric MFI of LIVE/DEAD staining in HMC-1.2 cells at each time point. (D) Viable cell counts of HMC-1.2 cells in culture. Data are the mean  $\pm$  SEM from 3 independent experiments. \*p<0.05, \*\*p<0.001, \*\*\*p<0.001, ANOVA with Sidak's post-test.



## Figure 3



### Systemic delivery HT-Fig. 4. of human KIT ESO inhibits tumor growth in a humanized xenograft mast cell neoplasia model.

the xenograft model used. (A) Examples of tumors in mice during treatment. The red arrows indicate the tumor and the green arrows show a secondary growth caused by drag of the needle during inoculation. (B) Measurements of tumor volume over time during treatment. Arrows indicate days that HT-KIT or vehicle control was administered. (C) Photograph of tumors after mice were euthanized at day 14. Measurements are in centimeters. (D) Weight of excised tumors after mice were euthanized on day 14. (E) Spleen weight at the conclusion of the experiment. (F) Liver weight at the conclusion of the experiment. (G) Mice weight was monitored over the course of the experiment. No significant differences were observed over the course of the experiment. Data are the mean ± SEM from 5 mice per group. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001. ANOVA with Dunnett's post-test (B), or unpaired t-test (**D** & **E**).

The top schematic representation of the protocol for





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P815 cells were transduced to make intracellular luciferase. (A) DBA/2J female mice protocol for the isograft model. Vehicle control or HT-KIT ESO was administered every 3 days. (B) Bioluminescence from P815 neoplastic cells was significantly reduced in intraturmoral (IT) (p=0.0007) and intravenous (IV) (p=0.0220) HT-KIT ESO treatment groups compared to controls. (C-E) IVIS images overlay primary site tumor where bioluminescent P815neopl astic cells were detected. Equalsized ROIs over the primary tumor site standardized area for photon flux (total counts) detected from all mice on study (representative images shown). (F-I) Tumor weight (F), volume (G), and depth (H) collectively represent decreased primary site tumor burden (F-G) and penetration into underlying structures (H) for IT and IV HT-KIT treatment groups compared to controls. (F) Tumor mass was reduced for IT (p=0.0132) and IV (p=0.0674) HT-KIT ESO treatment group compared to controls. (G) Tumor volume was reduced for IT (p=0.0188) and IV (p=0.0530) HT-KIT ESO treatment group compared to controls. (H) Tumor depth was significantly reduced for IT (p < 0.0001) and IV (p=0.0008) HT-KIT ESO treatment group compared to controls. (G) Tumor necrosis was increased for IT (p=0.0434) and IV (p=0.0588) HT-KIT ESO treatment group compared to controls. \*p≤0.05, \*\*p≤0.01, \*\*\*p≤0.001, \*\*\*\*p≤0.0001.

# Fig. 5. HT-KIT ESO reduces tumor growth in an isograft model of MC neoplasia.





Fig. 6. HT-KIT ESO reduced cardinal signs of mast cell leukemia in an isograft model of mutant mast cell neoplasia. (A) Percent of bone marrow infiltrated by P815 neoplastic cells. (B) Representative micrograph of bone marrow from vehicle controls (B) compared to IT (C), and IV (D) HT-KIT ESO. (E) Displacement of hematopoietic bone marrow tissue (myelophthisis) by neoplastic mast cells. (F) Packed red blood cell volume (hematocrit). (G) Bone marrow hematopoiesis as measured by myeloid progenitor to erythroid progenitor (M:E ratio). (H) Circulating neoplastic mast cells on differential counts. (I) Blood smear of neoplastic mast cells with Romanowsky stain (representative image shown from vehicle control animal blood). (J) H&E histopathology assessment of neoplastic emboli and circulating neoplastic cells in organs. (K-L) Emboli of neoplastic mast cells in pulmonary vessels from H&E histopathology sections from control (K) and intravenous HT-KIT ESO (L) treatment groups. \*p≤0.05, \*\*p≤0.01, \*\*\*p≤0.001, \*\*\*\*p≤0.0001.

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### Rationale

Activating mutations in the proto-oncogene c-KIT are associated with the mast cell (MC) clonal disorders cutaneous mastocytosis, systemic mastocytosis and its variants including aggressive systemic mastocytosis and mast cell leukemia, and MC sarcoma. Currently, therapies inhibiting KIT signaling are a leading strategy to treat MC proliferative disorders. However, these approaches may have off-target effects and in some patients, complete remission or improved survival time cannot be achieved. These limitations led us to develop an approach using chemically stable exon skipping oligonucleotides (ESO) that induce exon skipping of pre-mRNA to alter gene splicing and introduce a frameshift into mature KIT mRNA transcripts.

### Hypothesis

Exon-skipping *c-KIT* to remove exon 4 will force a frameshift in mature mRNA introducing an immediate STOP codon triggering nonsense-mediated mRNA decay. If this pathway is saturated, production of a severely truncated nonfunctional KIT protein would result. Both of these mechanisms would deplete functional KIT, inhibit proliferation and induce apoptosis of neoplastic MCs.

### Summary

This alternate approach to MC neoplasia results in marked downregulation of KIT expression, diminished KIT signaling, inhibition of MC proliferation and rapid induction of apoptosis in neoplastic HMC-1.2 MCs. This downregulation of KIT expression is achieved by combining two independent mechanisms involving nonsense-mediated mRNA decay and the production of alternatively spliced non-functional proteins. We further demonstrate that in vivo administration of KIT targeting ESOs significantly inhibits tumor growth and systemic organ infiltration using both an allograft mastocytosis model and a humanized xenograft MC tumor model. We propose that our innovative approach, which employs well-tolerated, chemically stable oligonucleotides to target KIT expression through unconventional pathways, has potential as a KIT-targeted therapeutic alone, or in combination with agents that target KIT signaling, in the treatment of KIT-associated malignancies.

### Figure 1



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## Acknowledgements

Funding: Department of Molecular Biomedical Sciences, College of Veterinary Medicine, start-up funds (G.C.), NCSU Center for Human Health and the Environment grant #ES025128, NIH T32OD011130 and the Division of Intramural Research of NIAID within the National Institutes of Health. Conflict of Interest disclosure: Dr. Cruse has filed a patent application related to the research reported in this study. Dr. Cruse has research support from Hoth Therapeutics and also serves on their Scientific Advisory Board. The terms of this arrangement have been reviewed and approved by NC State University in accordance with its policy on objectivity in research.

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