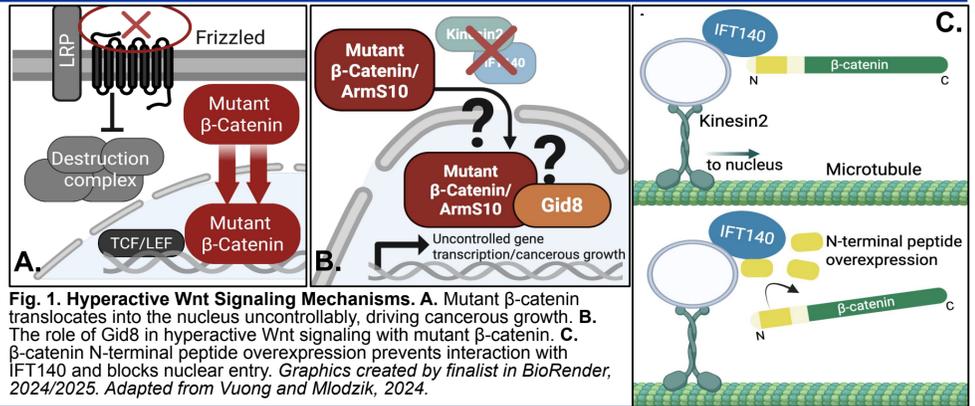


SELECTIVE DUAL INHIBITION OF β -CATENIN IN WNT-DRIVEN CANCERS VIA GID8-ASSISTED TRANSLOCATION AND A PRECISION PEPTIDE-BASED GENE THERAPY

INTRODUCTION

- Hyperactive Wnt signaling drives ~90% of hepatocellular carcinoma (HCC) cases, yet remains "undruggable"^[1-3] (Fig. 1A)
 - The hallmark mechanism, mutant β -catenin (ArmS10 in *Drosophila*) nuclear translocation, remains unclear^[4, 5]
- Gid8 has been shown to affect normal Wnt signaling, but its role in cancerous, hyperactive Wnt signaling is unstudied^[6] (Fig. 1B)
- What role does Gid8 play in regulating mutant β -catenin nuclear translocation in cancers?**
- In liver cells, β -catenin acts at the **membrane to maintain adherens junctions/polarity** and in the **nucleus to drive proliferation/cancer**^[7]
- A β -catenin²⁴⁻⁷⁹ peptide has been shown to suppress Wnt signaling^[8] (Fig. 1C)
- The full-length peptide is too large for selective targeting of nuclear β -catenin while preserving wild-type β -catenin, a major roadblock in cancer therapy^[8]
- Does truncating the β -catenin²⁴⁻⁷⁹ peptide enable selective inhibition?**



METHODOLOGY

Phase 1:

- Gal4/UAS *Drosophila* crosses were performed (4 experimental groups)
- Offspring's adult wings, wing discs, and salivary glands were dissected
- Immunofluorescence staining was conducted
- Co-IP western blot between Gid8 and Arm/ArmS10 was performed^[9]
- HEK293 cells were transfected with Gid8 and Gid8* (with C-terminal 3 mutations)

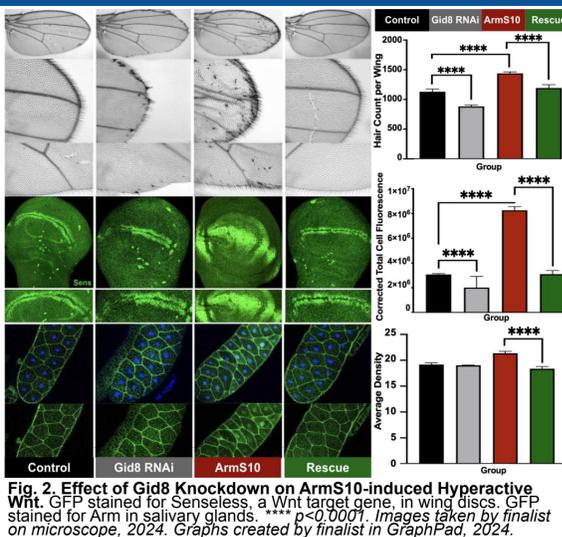
Phase 2:

- β -catenin²⁴⁻⁷⁹ peptide and 3 truncated fragments within the peptide were cloned
- Huh7, HepG2, and Hep3B cells were cultured and transfected with each peptide construct separately^[10]
- MTT assays, immunofluorescence staining, and Western blots were performed to quantify hyperactive Wnt activity

RESULTS AND DISCUSSION

Gid8 REGULATES HYPERACTIVE WNT SIGNALING

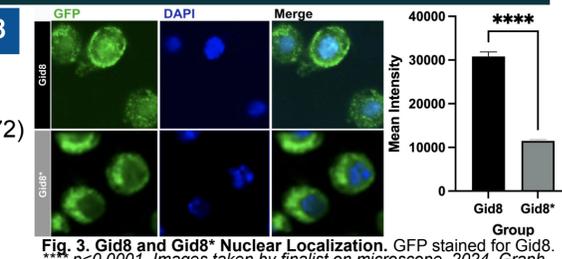
- Adult wings: Knockdown or underexpression of Gid8 **rescued** ArmS10-induced excessive hair growth, an indicator of hyperactive Wnt ($p < 0.0001$; Fig. 2)
- Gid8 amplifies hyperactive Wnt**
- Wing discs: Knockdown Gid8 + ArmS10 showed less Senseless expression than ArmS10 alone ($p < 0.0001$)
- Gid8 promotes Wnt hyperactivation, linking Gid8 to cell proliferation not degradation**
- Salivary glands: Knockdown Gid8 + ArmS10 had less Arm in the nucleus (less Wnt; $p < 0.0001$)
- First evidence linking Gid8 to hyperactive Wnt signaling**



Identifies Gid8 as a novel regulator of hyperactive Wnt with implications for cancer regulation

TARGETABLE REGION IN GID8

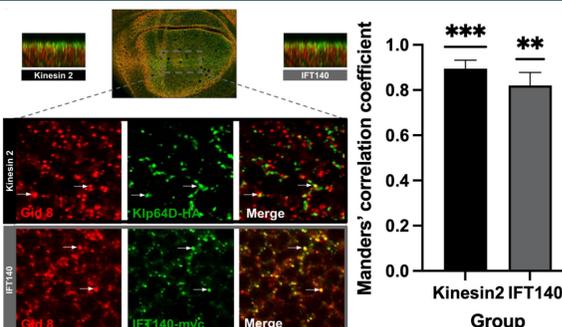
- In HEK293 cells with active Wnt, expression of Gid8* (contains mutated residues 167, 168, and 172) nearly abolished Gid8's nuclear localization ($p < 0.0001$; Fig. 3)
- Identifies these 3 residues as critical for Gid8 nuclear entry**



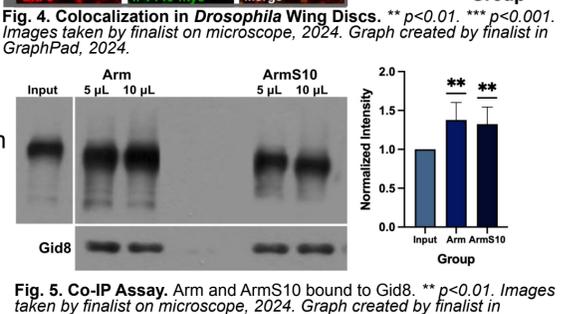
Defines precise mutations in Gid8's nuclear localization sequence (C-terminal) that can be targeted to limit hyperactive Wnt signaling

GID8 GUIDES NUCLEAR TRANSLOCATION

- Wing discs: Gid8 colocalized with Kinesin2 and IFT140, proteins that guide wild type β -catenin to the nucleus ($p < 0.01$; 0.001; Fig. 4)
 - Links Gid8 to intracellular transport
- Mutant β -catenin nuclear entry is mediated by Gid8-dependent engagement of Kinesin2/IFT140 transport pathways
- Co-IP assays confirmed interactions between Gid8 and mutant ArmS10 ($p < 0.01$; Fig. 5)
 - Transport-dependent mechanism of Gid8 controlling nuclear entry during hyperactive Wnt signaling



Mutant β -catenin enters the nucleus with Gid8, a previously unclear mechanism

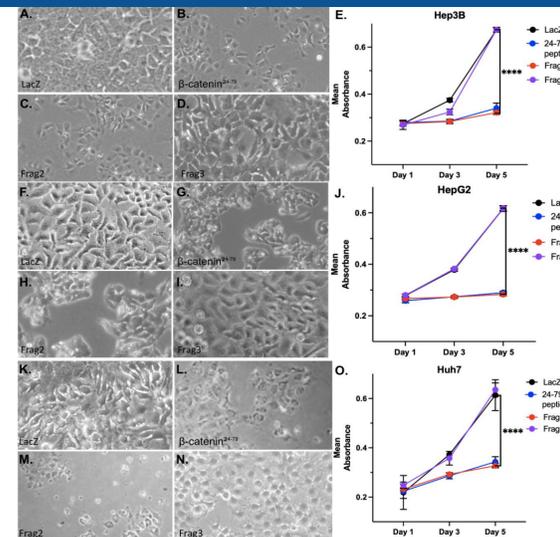


FRAGMENT 2 MITIGATES HCC PROLIFERATION

- Overexpression of residues **41-58 in β -catenin (fragment 2)** consistently produced the strongest reduction in HCC proliferation across Huh7, Hep3B, and HepG2 cells ($p < 0.0001$; Fig. 6)
- Fragment 2 defines the essential inhibitory region within the β -catenin²⁴⁻⁷⁹ peptide

Fragment 2 overcomes delivery and stability limitations of the full-length β -catenin peptide by defining an 18-amino-acid inhibitory region

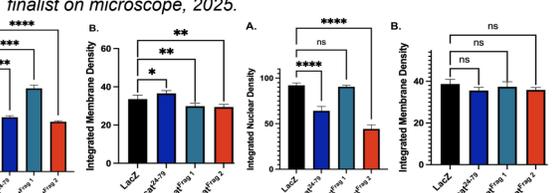
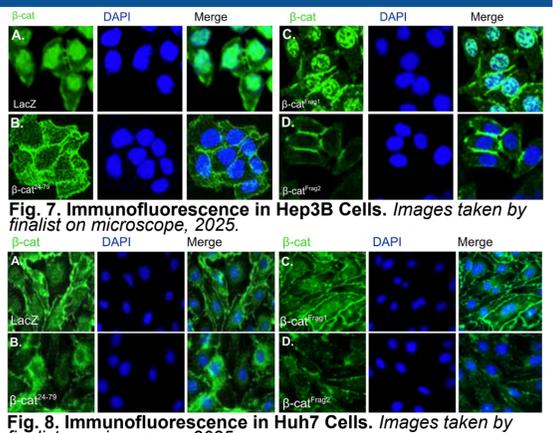
- Makes selective β -catenin inhibition clinically achievable due to small size



FRAGMENT 2 ACHIEVES SELECTIVE INHIBITION

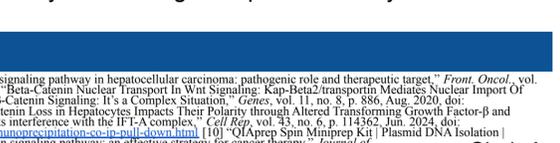
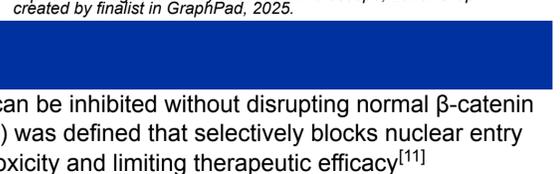
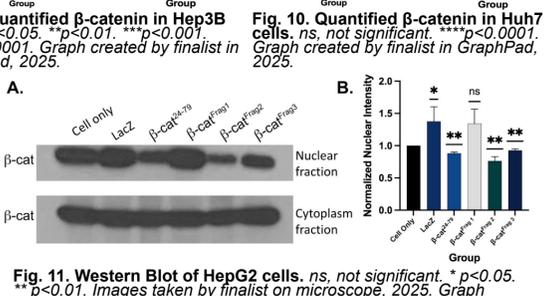
- In Hep3B and Huh7 cells, Fragment 2 produced the largest reduction in cancerous, nuclear β -catenin ($p < 0.0001$; Fig. 7D, 8D, 9A, 10A)
- Membrane β -catenin showed smaller changes, indicating retention at adherens junctions and preservation of normal β -catenin functions (ns; $p < 0.01$; Fig. 9B, 10B)
- Previously unachievable to block nuclear entry without disrupting membrane β -catenin

Fragment 2 resolves the central Wnt targeting problem by separating β -catenin's cancerous nuclear function from its essential membrane role



- In HepG2 cells, Fragment 2 significantly reduced nuclear β -catenin ($p < 0.01$; Fig. 11)
- Cytoplasmic levels of β -catenin remained unchanged

Shows that restricting nuclear access alone is sufficient to suppress hyperactive Wnt signaling without disrupting cytoplasmic pools



CONCLUSION

- Resolves **two key gaps in Wnt-driven cancers**: how mutant β -catenin enters the nucleus and how cancerous Wnt signaling can be inhibited without disrupting normal β -catenin
- Gid8 was identified as a transport-specific regulator that controls mutant translocation, and an inhibitory region (residues 41-58) was defined that selectively blocks nuclear entry
 - Upstream and downstream Wnt inhibitors failed: they disrupt essential β -catenin membrane functions, causing on-target toxicity and limiting therapeutic efficacy^[11]
- Initiates a novel baseline for precision peptide-based gene therapy**

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