



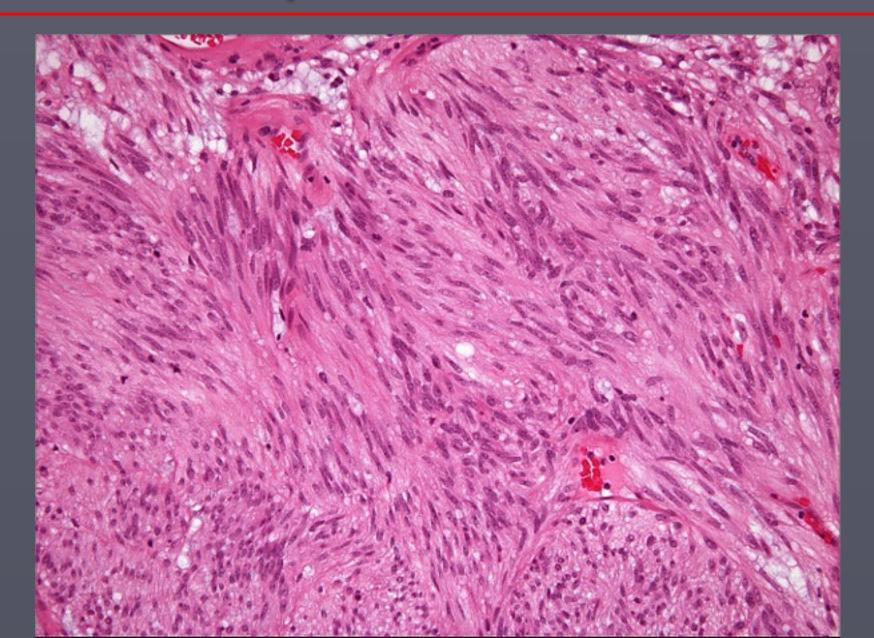
Brigham and Women's Hospital, Harvard Medical School

Biology of GIST Translating Cancer Research into Targeted Therapeutics

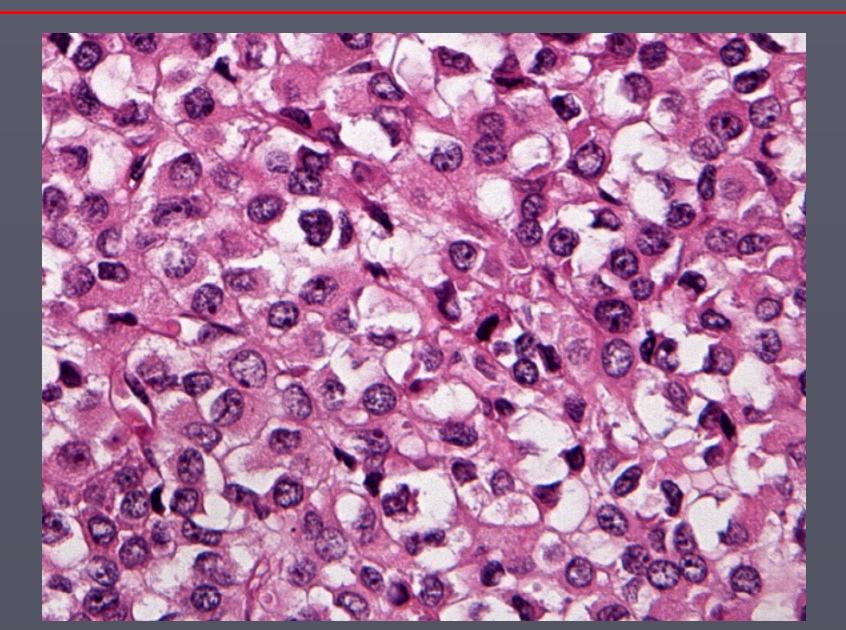
Jonathan A. Fletcher, M.D. Depts of Pathology & Pediatrics Brigham and Women's Hospital Dana-Farber Cancer Institute Harvard Medical School

Dana-Farber/Harvard Cancer Center

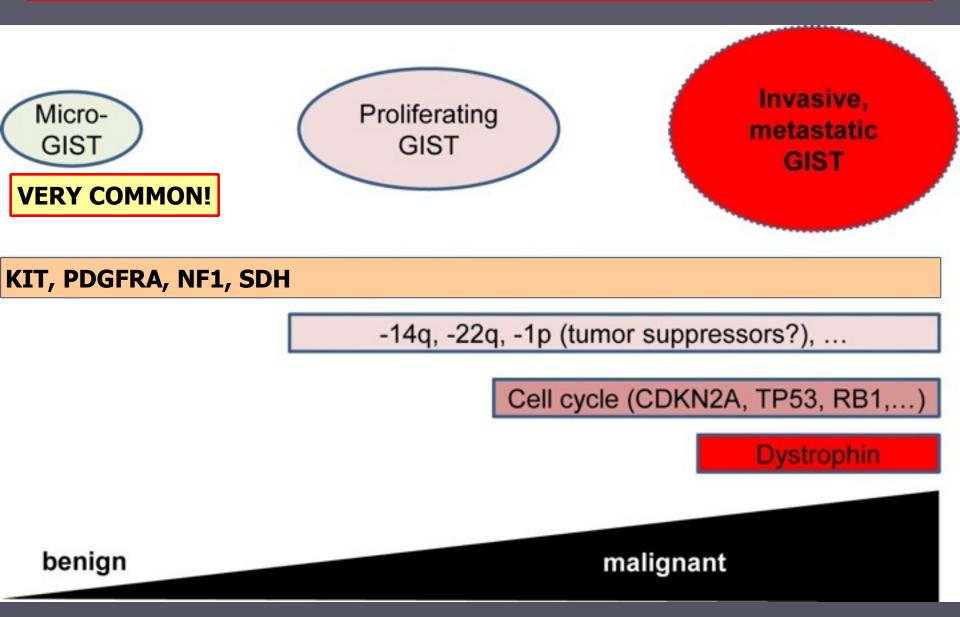
Spindle-cell GIST



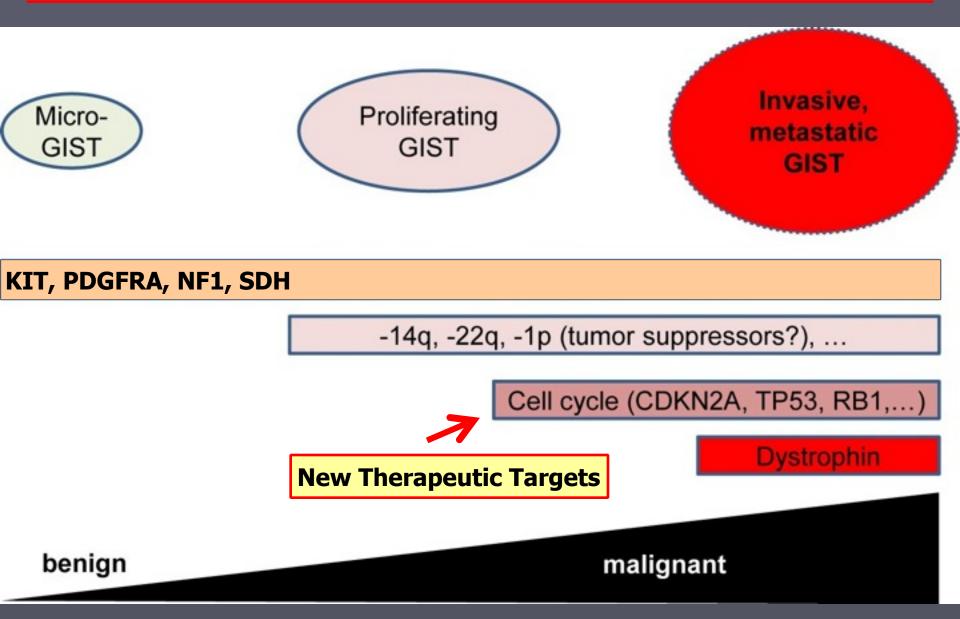
Epithelioid GIST



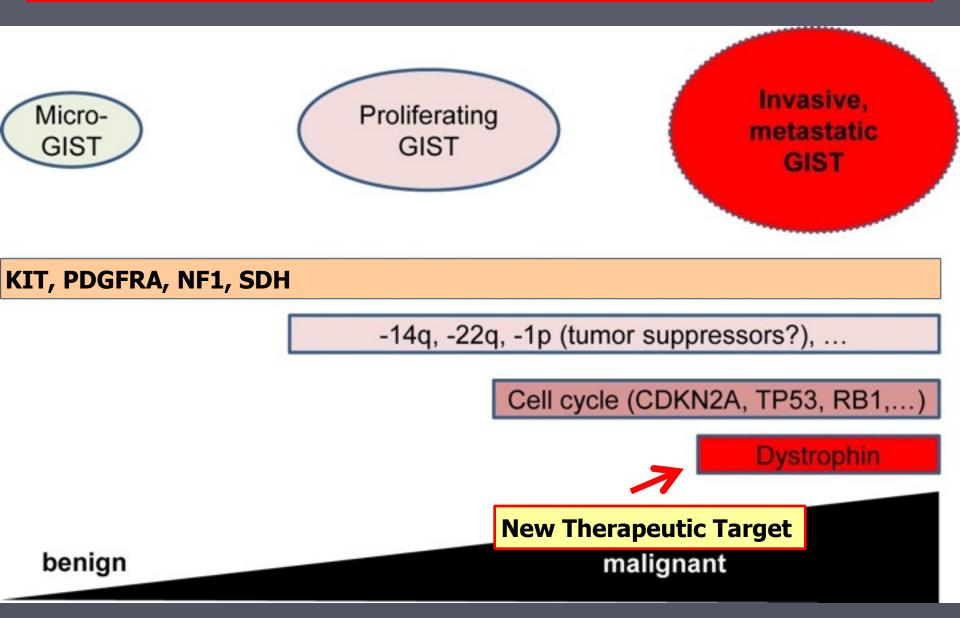
GIST Biologic Progression



GIST Biologic Progression



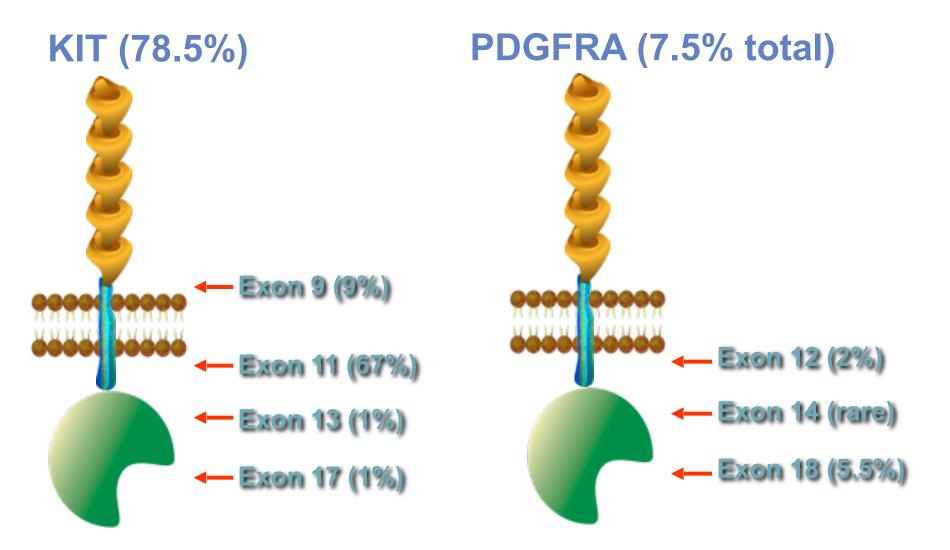
GIST Biologic Progression



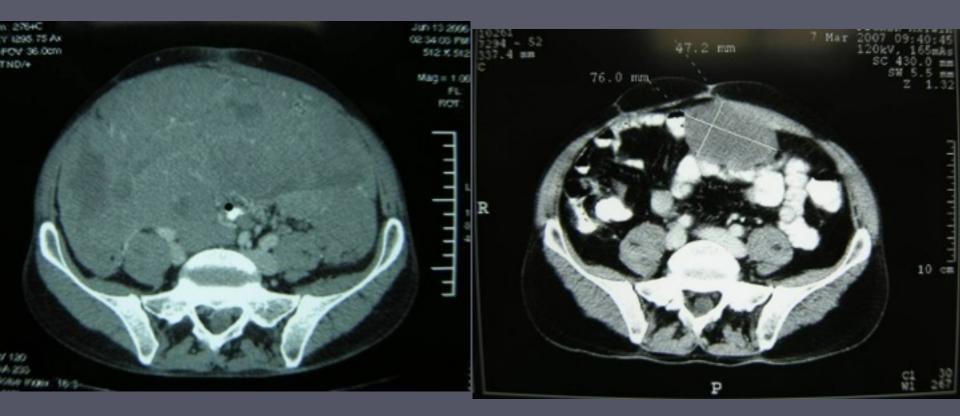


KIT and PDGFRA Mutations in >2000 GISTs (Heinrich-Corless)

Overall Mutation Frequency: 86%

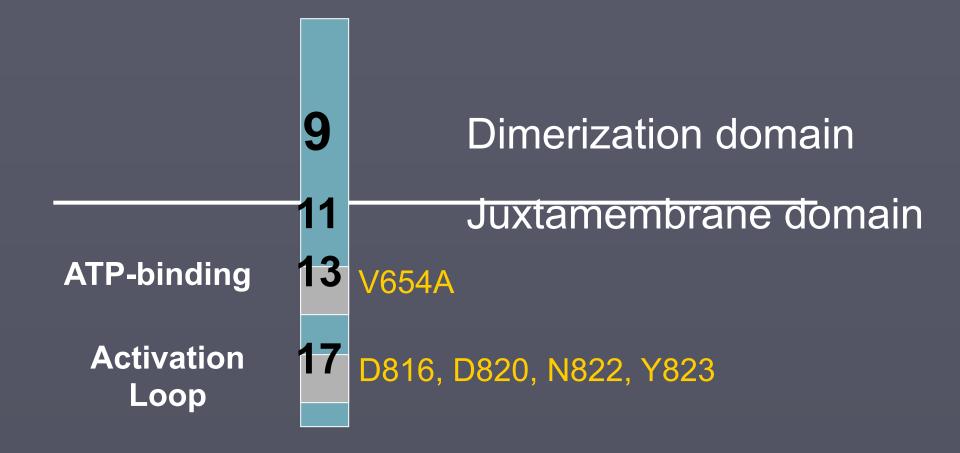


Metastatic GIST Major response after 6 months of Imatinib

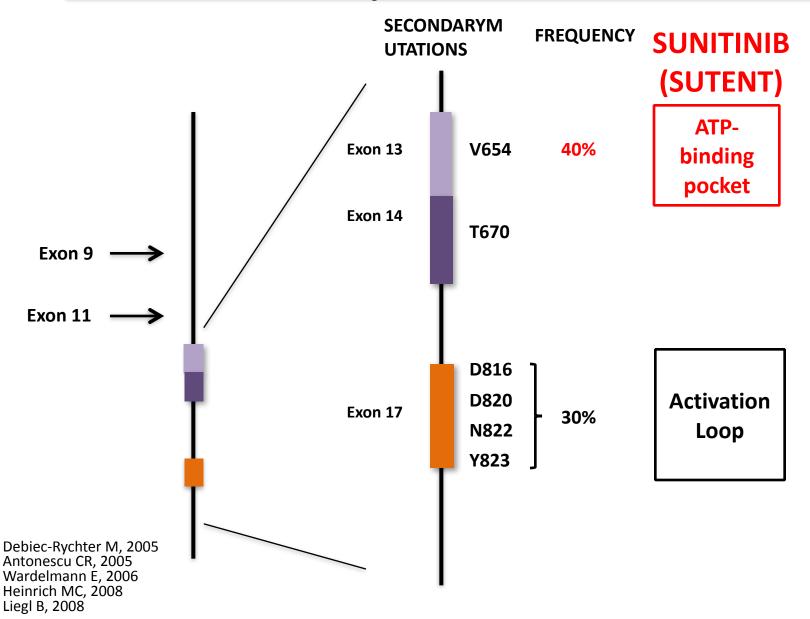


KIT exon 11 mutation

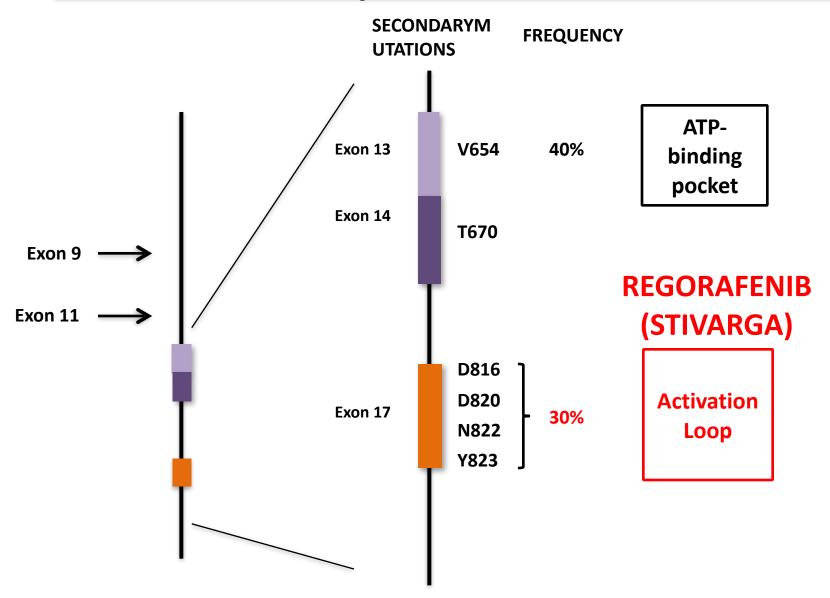
Secondary imatinib-resistance KIT mutations in GIST



Secondary resistance in GIST



Secondary resistance in GIST



Rapid alternation regimen

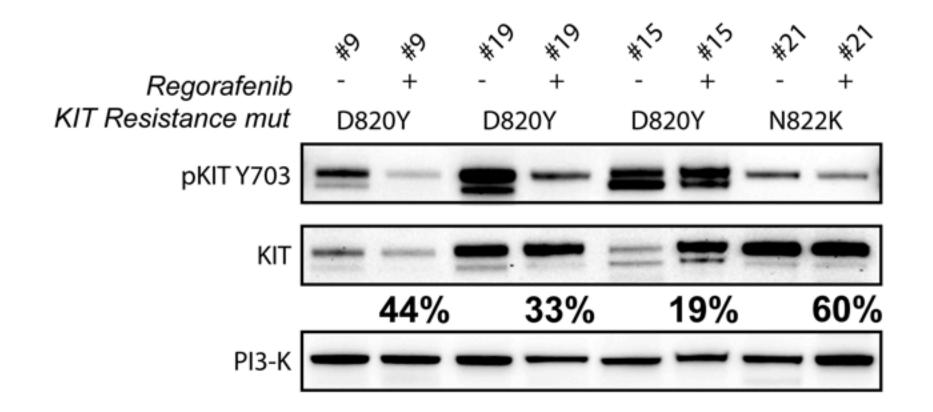


✓Rapid alternation regimen might minimize toxic effects.

✓Alternation of complementary drugs increases the spectrum of effective inhibition of IM-resistant clones.

✓SuRe Trial – (Drs. Serrano, George and colleagues)





Mutations <u>activate</u> KIT/PDGFRA, causing GIST cells to grow and survive



KIT baseline activation Normal GIST precursor cell

Increased KIT/PDGFRA activation causes GIST to grow Imatinib and other kinase inhibitors down-regulate KIT/PDGFRA activation to levels that no longer support cell growth



KIT or PDGFRA mutant GIST

KIT/PDGFRA activation restored to low levels

KIT and PDGFRA imatinib-resistance mutations are life-savers for GIST but at same time they STRESS the cells

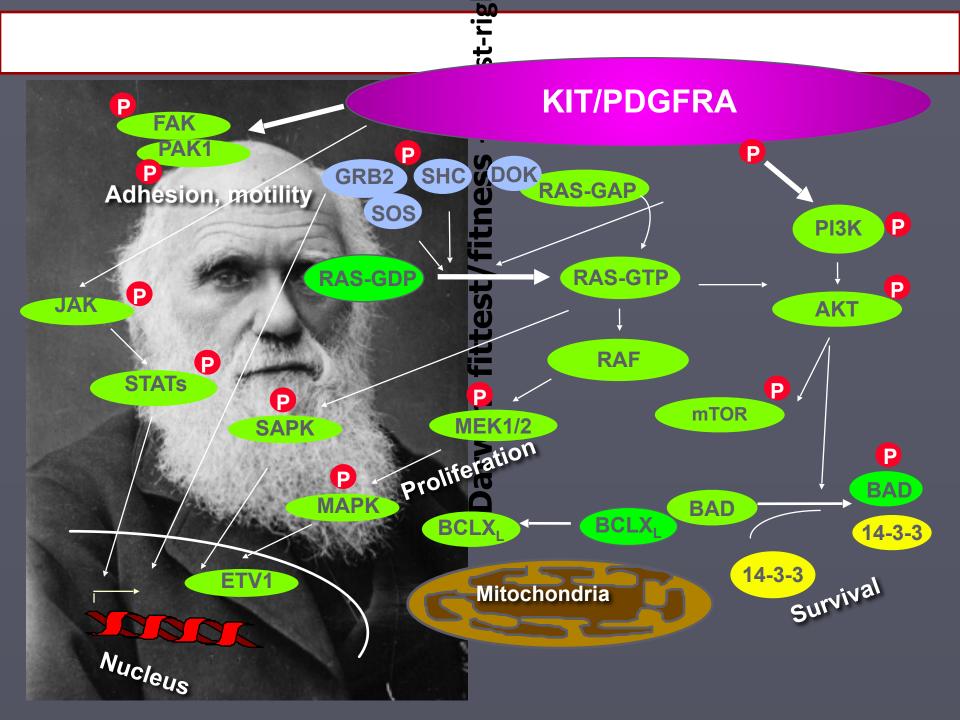


2nd Mutation Imatinibresistance

Optimal energy-level for GIST



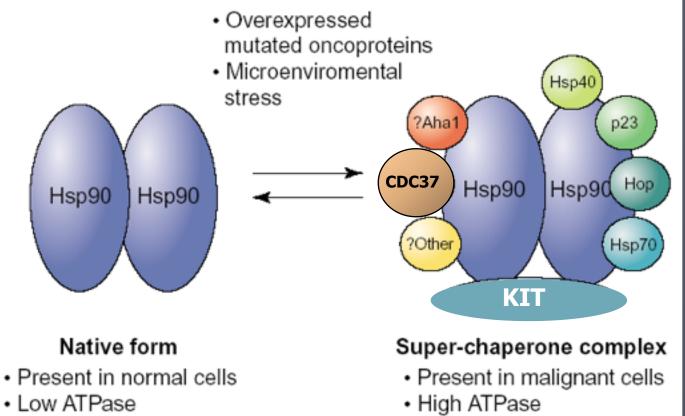
Destroy GIST by selectively STRESSING the most Gleevec-resistant cells



HSP90: Key KIT oncoprotein chaperone in GIST

Chaperone family:

- protein folding
- translocation
- stabilization



Low 17AAG affinity

High 17AAG affinity

Paul Workman

Screen <u>11,000 genes</u> to determine which the GIST cells need most

CDC37 = most essential gene among 11,000 genes screened!

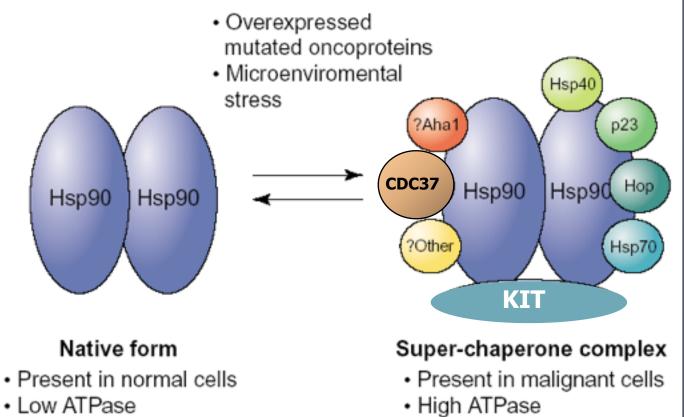
NAME	RIGER_RANK	RIGER_SCORE	RIGER_LEV
CDC37:NM_007065	1	-1.82	0.8
VCP:NM_007126	3	-1.81	0.6
PSMC4:NM_006503	2	-1.81	0.8
ZNF206:NM_032805	4	-1.78	0.4
PARN:NM_002582	5	-1.77	0.6
DES:NM_001927	6	-1.77	0.8
EIF5B:NM_015904	7	-1.76	0.75
ZNF207:NM_003457	8	-1.76	0.6
PDHA1:NM_000284	10	-1.75	0.6

Adrian Marino-Enriquez – Oncogene, 2014

CDC37 targeting 10-fold more selective than HSP90 targeting

Chaperone family:

- protein folding
- translocation
- stabilization

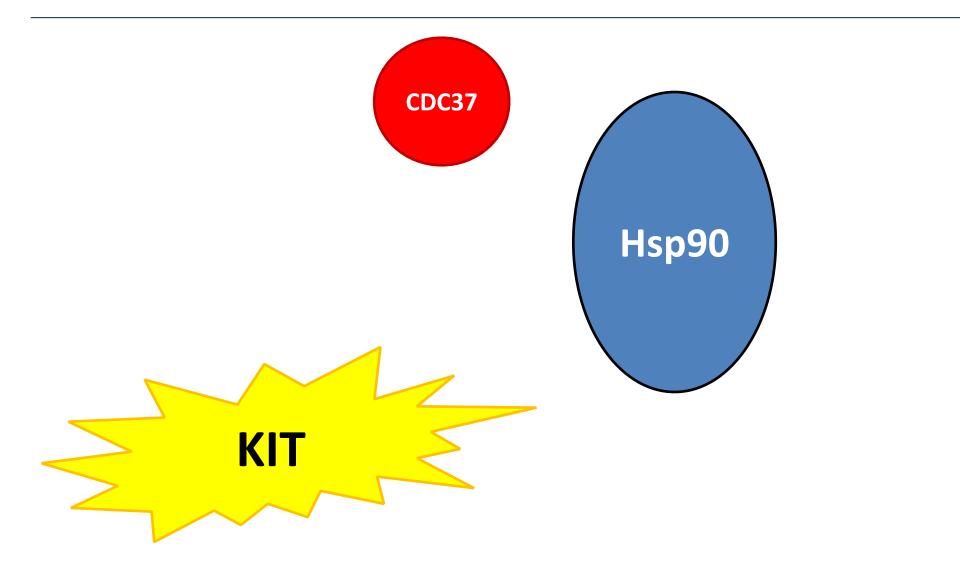


Low 17AAG affinity

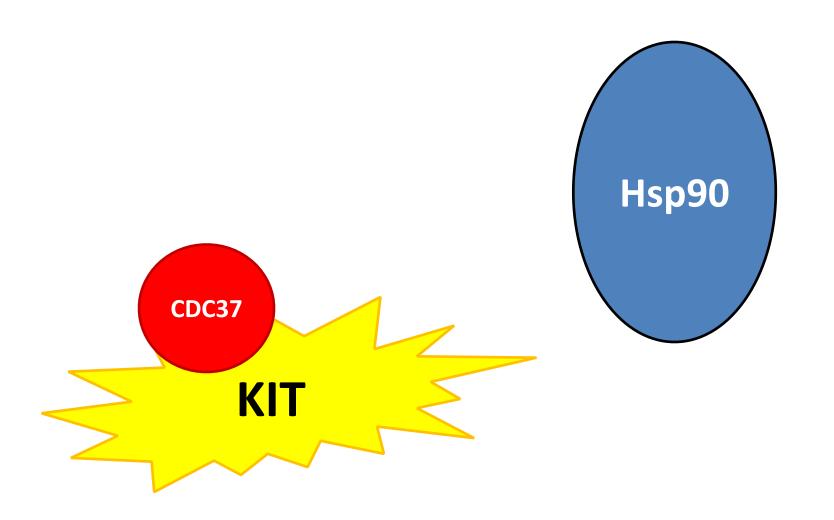
High 17AAG affinity

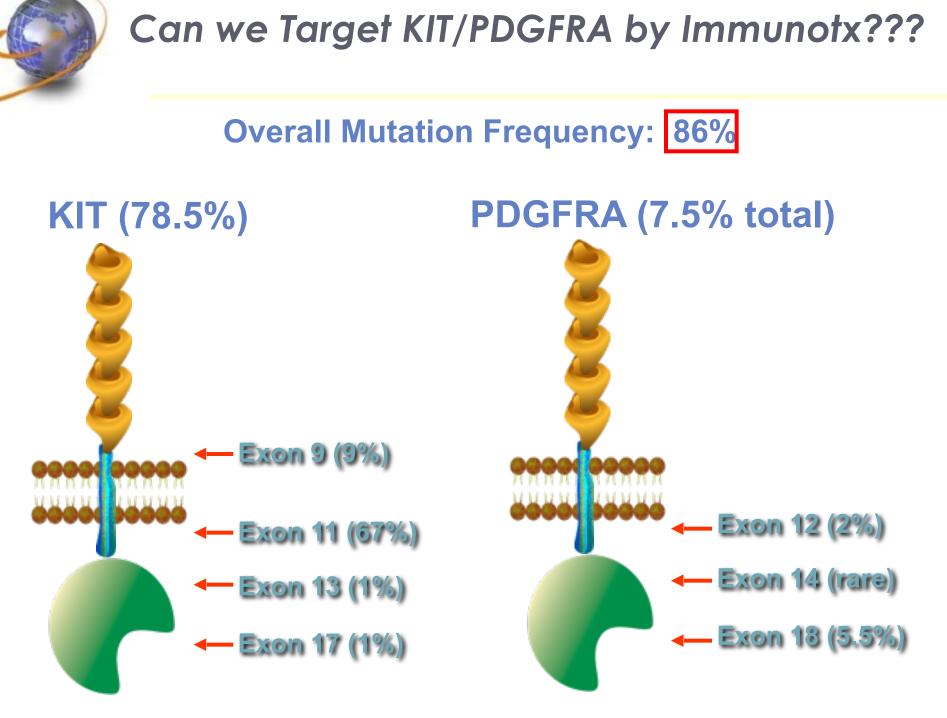
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shRNA pooled library screen in GIST



shRNA pooled library screen in GIST

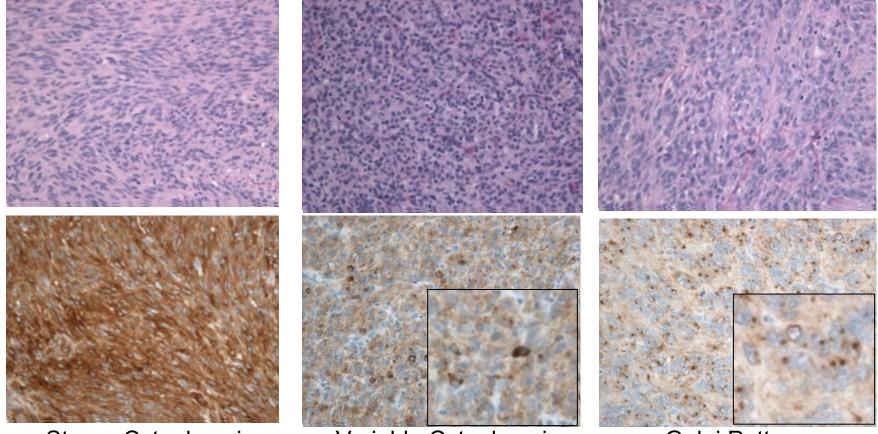




3 TKI-resistant GIST metastases (same patient): Varied cytoplasmic <u>NOT MEMBRANE</u> oncoprotein target expression

SPINDLE CELL KIT: DIFFUSE-STRONG

EPITHELIOID KIT: WEAK SHORT SPINDLE CELLS KIT: GOLGI PATTERN



Strong Cytoplasmic

Variable Cytoplasmic

Golgi Pattern

Models: Defining GIST Biology

Cell Lines

- Cell cultures created from GIST surgical specimens
- Studies are quick & inexpensive
- Cells can lose dependence on key targets
- Xenografts
 - GIST surgical specimens implanted into mice
 - Nuanced evaluation of complex biology
- Genetically-engineered models
 - GIST developing in a mouse, eg due to a KIT mutation in the mouse
 - Potentially most nuanced, although might not represent the true biology of human GIST
 - Most expensive