

# SDH-Deficient GIST-A Newly Identified GIST Subtype

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On behalf of the Consortium for Pediatric & wildtype GIST Research



**CPGR**  
Consortium for Pediatric  
& wildtype GIST Research

the NIH Pediatric & Wildtype GIST Clinic





## Wild-Type GIST

- 85% of GIST occurring in young population is lacking mutations in *KIT* or *PDGFRA*
- Stomach location, epithelioid histology
- May be syndromic (Carney Triad, Stratakis-Carney Syndrome)
- Tyrosine kinase inhibition is less effective compared to *KIT* or *PDGFRA* mutant tumors

# the NIH Pediatric & Wildtype GIST Clinic

**An international clinic twice a year  
115 patients have been seen in 10 clinics since 2008.**

## **Objectives of the Wildtype Clinic at NIH**

- To bring together healthcare providers who have the most experience treating and studying GIST**
- To obtain clinical history, response to prior treatments, histopathologic results, radiographic assessments and genetic/molecular analyses**
- Continue long-term follow-up for these patients**

# Report on First 78 Patients Characterized



CENTER FOR CANCER RESEARCH



- **12 PATIENTS-KINASE TYPE (Group A)**
- **22 PATIENTS-SDH DEFICIENT WITHOUT IDENTIFIED SDH MUTATION (Group B)**
- **44 PATIENTS-SDH DEFICIENT WITH SDH MUTATIONS, ALL BUT 3 GERMLINE (GROUP C)**

## Group A (n=12)

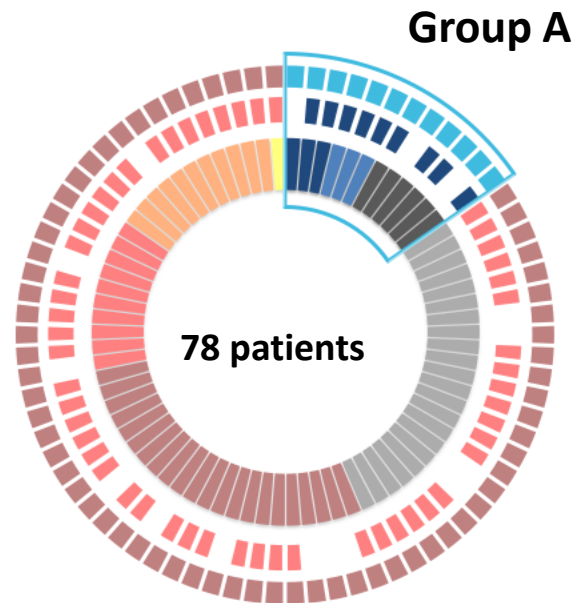
Positive SDHB IHC

Normal Methylation pattern

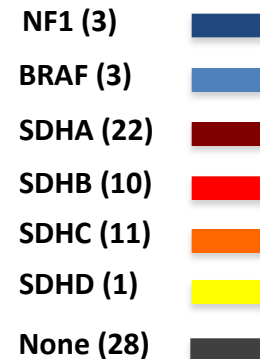
Mutations in NF1, BRAF or other unknown genes

These are Kinase-type GIST

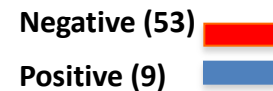
Remaining 66-SDH deficient



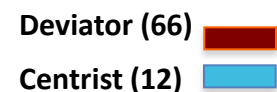
1<sup>st</sup> Circle  
Mutated  
Genes:



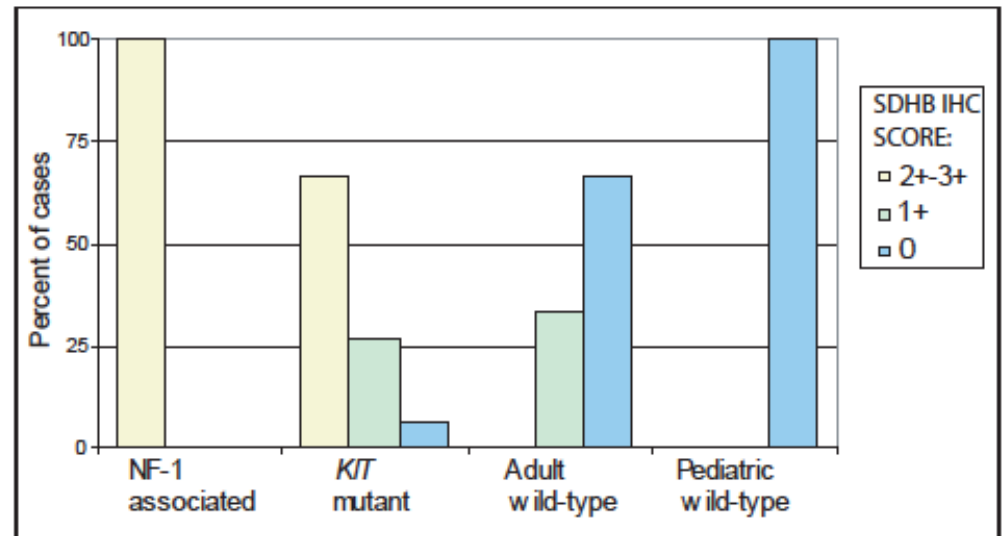
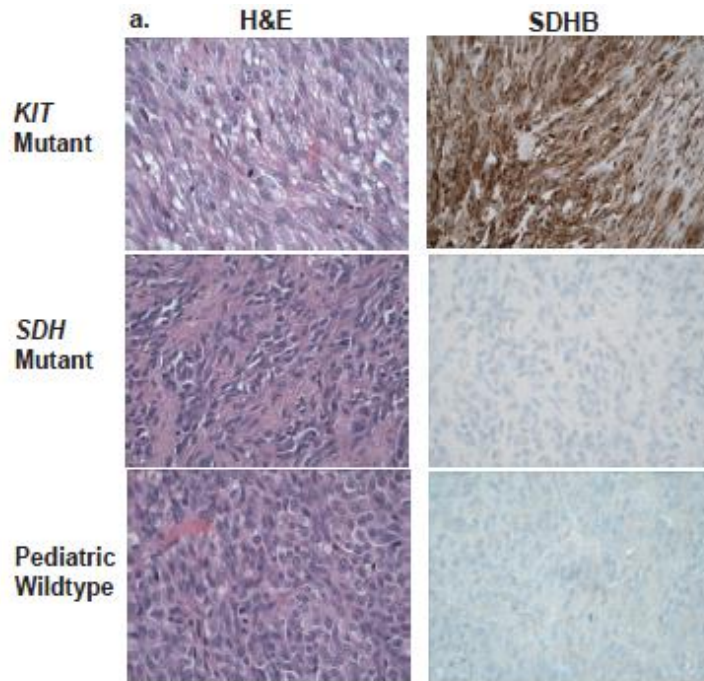
2<sup>nd</sup> Circle  
SDHB staining:



3<sup>rd</sup> Circle  
Methylation



# Loss of SDHB Protein Abundance

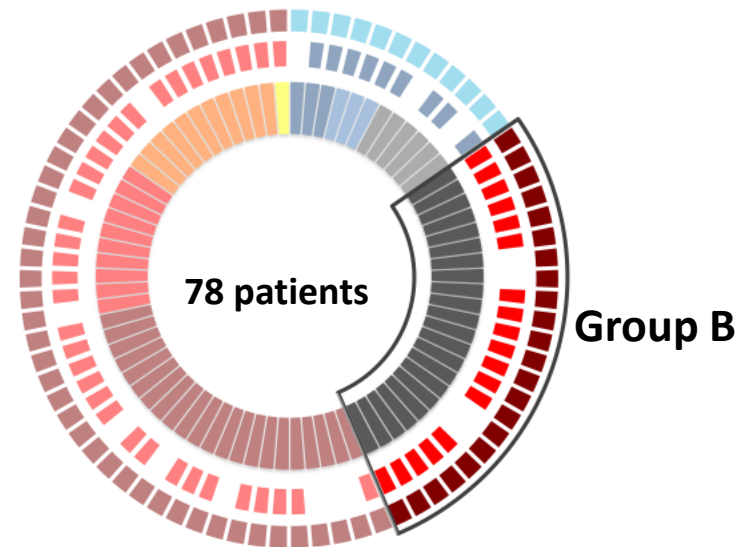


Janeway and Kim, et al. 2011 PNAS 108:314

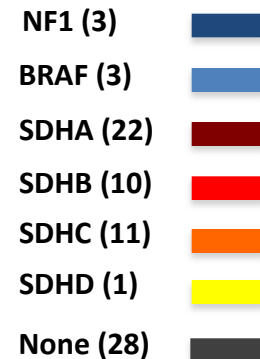
## Group B (n=22)

**Negative SDHB staining by IHC  
Hypermethylation (Deviator)**

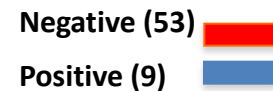
No identified mutations-early data  
suggest a mechanism of SDH  
deficiency



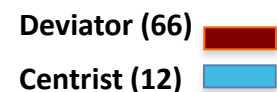
1<sup>st</sup> Circle  
Mutated  
Genes:



2<sup>nd</sup> Circle  
SDHB staining:



3<sup>rd</sup> Circle  
Methylation



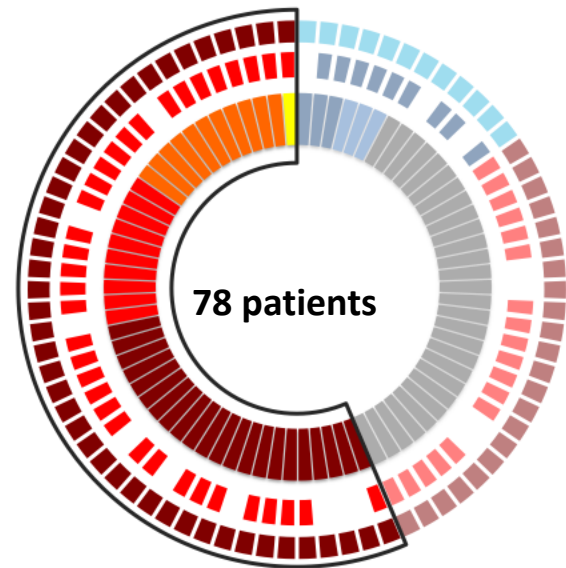
## Group C (n=44)

**Negative SDHB staining by IHC  
Hypermethylation (Deviator)**

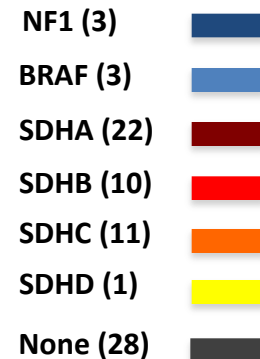
SDHA, B, C, D mutations

To date, all but 3 are germline  
mutations

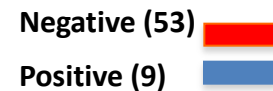
Group C



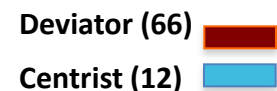
1<sup>st</sup> Circle  
Mutated  
Genes:



2<sup>nd</sup> Circle  
SDHB staining:

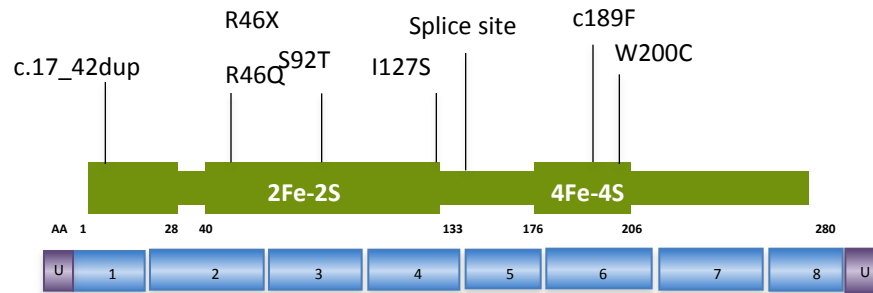
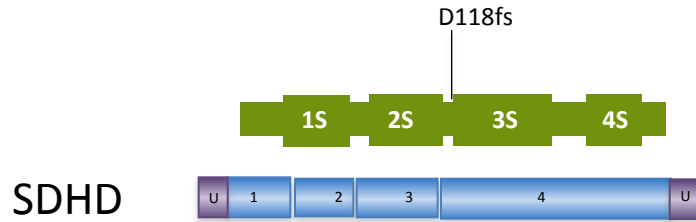


3<sup>rd</sup> Circle  
Methylation

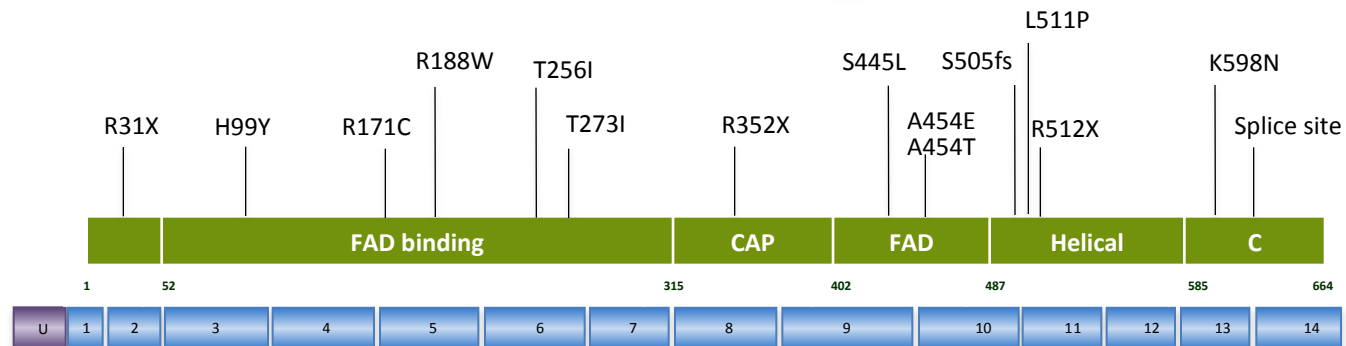




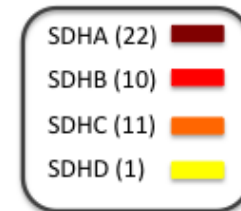
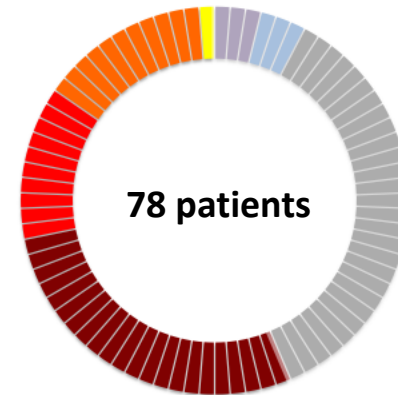
# Mutations Distributed across all exons-90% Germline



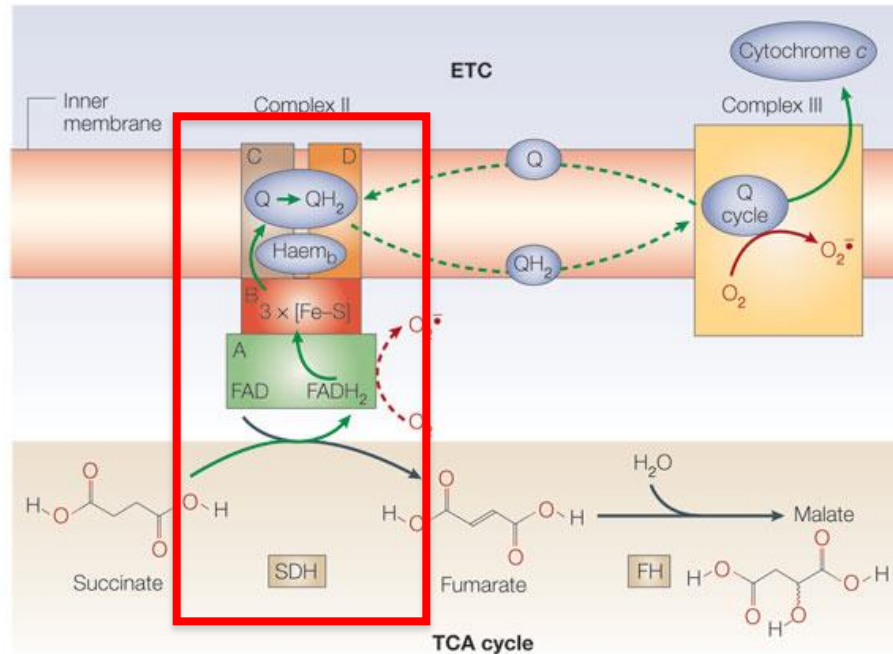
**SDHB**



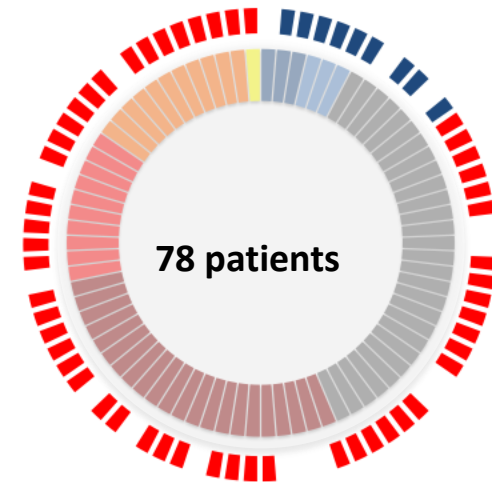
**SDHA**



- Tumors with SDHA, B, C, D mutations have always negative SDHB staining.



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Nature Reviews | Cancer



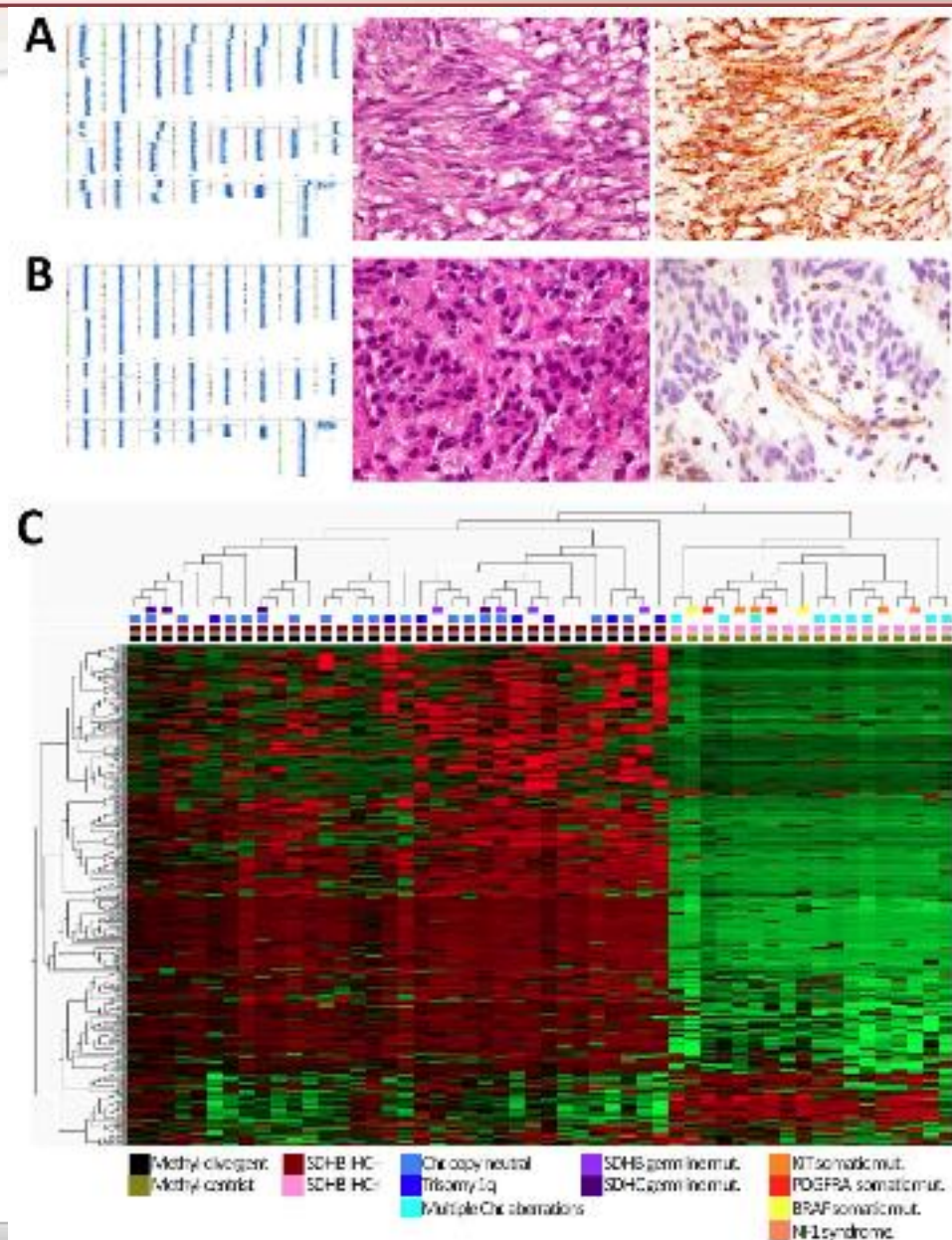
1<sup>st</sup> Circle  
Mutated  
Genes:

NF1 (3)	Dark Blue
BRAF (3)	Light Blue
SDHA (22)	Dark Red
SDHB (10)	Red
SDHC (11)	Orange
SDHD (1)	Yellow
None (28)	Grey

2<sup>nd</sup> Circle  
SDHB staining:

Negative (53)	Red
Positive (9)	Blue

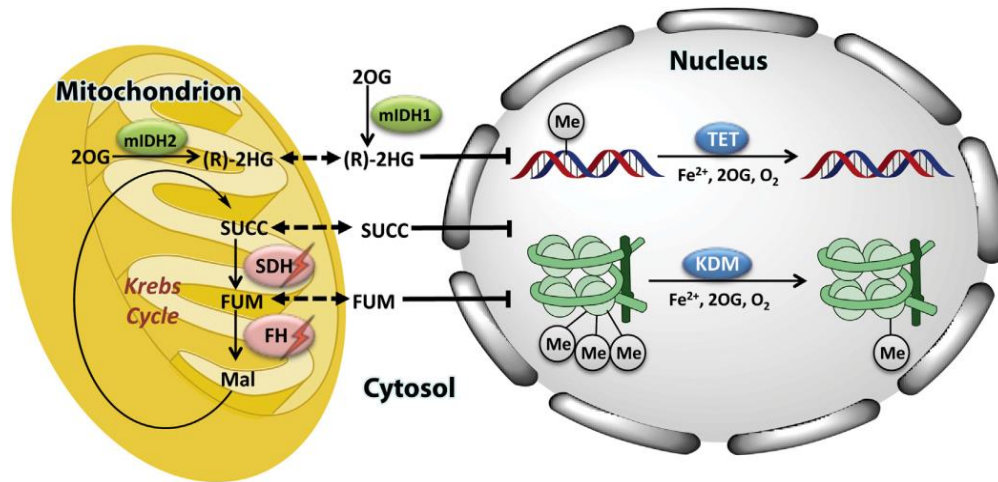
# SDH Deficient GIST Have Global Hypermethylation



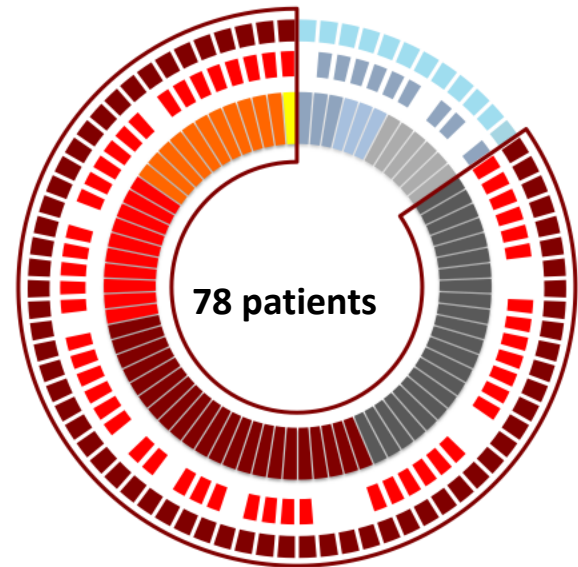
18 SDH mutations found

Killian K et al.  
Cancer Discovery 2013

# SDH loss leads to succinate inhibition of demethylases TET2 and KDM



Yang M, Pollard PJ Cancer Cell 2013



1<sup>st</sup> Circle  
Mutated  
Genes:

NF1 (3)	Dark Blue
BRAF (3)	Light Blue
SDHA (22)	Dark Red
SDHB (10)	Red
SDHC (11)	Orange
SDHD (1)	Yellow
None (28)	Grey

2<sup>nd</sup> Circle  
SDHB staining:

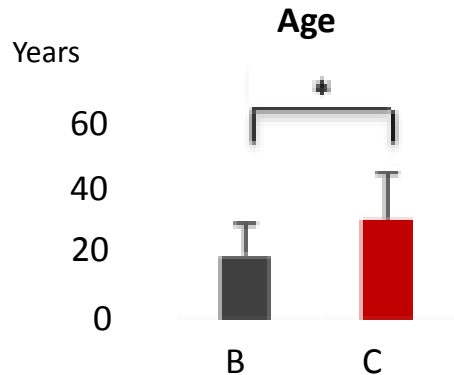
Negative (53)	Red
Positive (9)	Blue

3<sup>rd</sup> Circle  
Methylation

Deviator (66)	Dark Red
Centrist (12)	Light Blue

# Comparison of Group B and C

## 1. Group B=young age

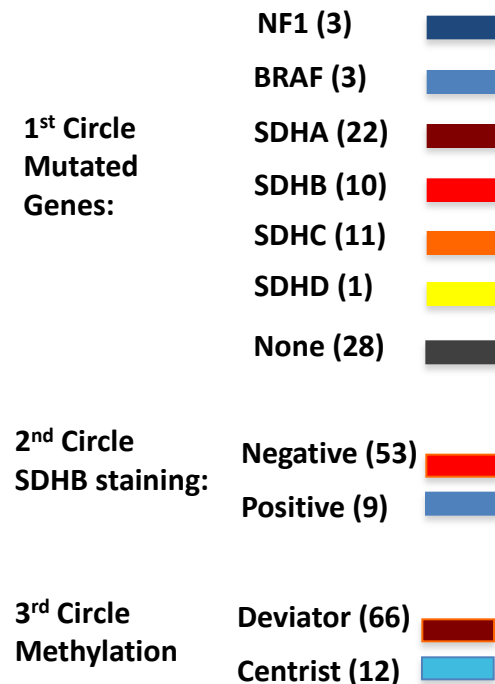
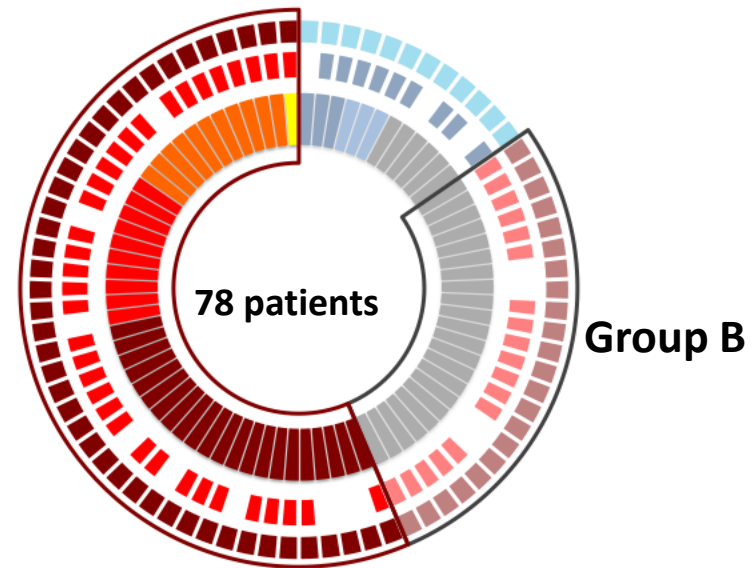


**2. All patients in Group B are females while in Group C 65% are females.**

**We have Carney Triad patients (chondroma, paraganglioma) or Carney-Stratakis Syndrome patients (paraganglioma) in both groups B and C.**

**At the moment we have no statistically significant differences in overall Survival, Recurrence free Survival.**

**Group C**



# Conclusions

## 3 distinct groups of Wildtype GIST

	Group A	Group B	Group C
Mutated genes	NF1, BRAF or other genes	Unknown genes	SDH genes
SDHB expression by ICH	normal	no	no
SDH Function	Normal	Impaired	Impaired
Methylation Pattern	Centrist	Deviator (hypermethylation)	Deviator (hypermethylation)
Gender	Both	<u>Females</u>	Female predominance
Age	Young adults, adults	<u>Pediatric</u> , young adults	Young adults, adults
Location	Gastric, small bowel	Gastric	Gastric



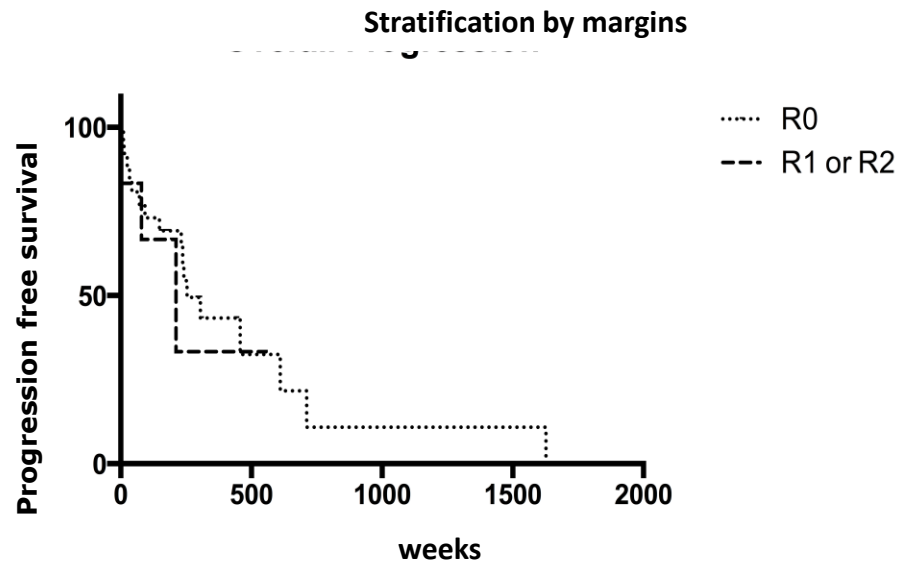
# Baseline patient and disease characteristics per group

		SDHB (+) N=9	SDHB (-) SDHx WT N=17	SDHB (-) SDHx mutant N=41
<b>Age, median years (range)</b>		44 (9-57)	16 (8-50)	24 (8-50)
<b>Sex, n (%)</b>	Male	3 (33.3%)	1 (5%)	14 (34%)
	Female	6 (66.6%)	16 (94%)	27 (65%)
<b>Location of primary tumor, n (%)</b>	Gastric	1 (11.1%)	17 (100%)	41 (100%)
	Duodenum	1 (11.1%)	0	0
	Small Bowel	6 (66.6%)	0	0
	Peritoneum	1 (11.1%)	0	0
<b>Size of primary tumor, median cm (range)</b>		10.5 (5.6-13.5)	5.5 (2-16)	5.8 (2-21)
<b>Multifocal Disease, n (%)</b>	Yes	2 (22.2%)	9 (52%)	14 (40%)
	No	7 (77.7%)	7 (41%)	21 (60%)
<b>Metastatic Disease, n (%)</b>	At presentation	0 (0%)	5 (29%)	11 (26%)
	Later	0 (0%)	7 (41%)	19 (46%)
	Never	0 (0%)	5 (29%)	11 (26%)
<b>Metastatic sites at presentation</b>	Lymph Nodes	0 (0%)	1 (5%)	4 (9%)
	Small Bowel	0 (0%)	0	0
	Liver	0 (0%)	4 (23%)	9 (21%)
	Peritoneum	0 (0%)	2 (11%)	4 (9%)
<b>Recurrence sites</b>	Stomach	0	5 (29%)	6 (14%)
	Small Bowel	1 (1,1 %)	0	0
	Liver	0	6 (35%)	12 (29%)
	Peritoneum	6 (66%)	4 (23%)	8 (19%)
<b>Other tumors</b>	Paraganglioma	0 (0%)	2 (11%)	5 (12%)
	Chondroma	0 (0%)	4 (23%)	2 (4%)

# SDH Deficient GISTs are Multi-focal



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# the NIH Pediatric & Wildtype GIST Clinic



Children's Hospital Boston

The first place for children



FOX CHASE  
CANCER CENTER



UNIVERSITY OF UTAH



National Human  
Genome Research  
Institute





**Our Thanks**

**To GIST support group members**

**To the patients and their families**

