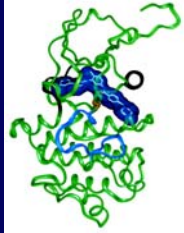


### Imatinib Resistance and Other Diseases Targeted by Imatinib

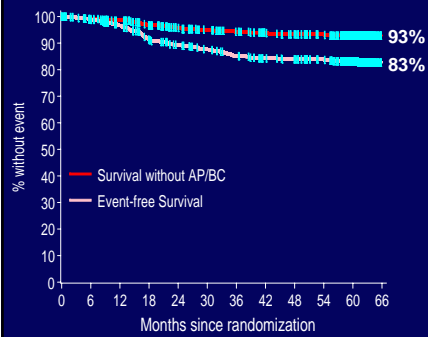


Brian J. Druker, MD

### Summary of CML Clinical Trials

- Imatinib yields high response rates with minimal toxicity in all phases of CML
- Durable responses are achieved in chronic phase patients
- Resistance in advanced phase patients is common

### Relapses and Disease Progression



### Relapse Rate (4 Years)

- Chronic (IFN failure) 26%
- Accelerated 73%
- Blast 95%

### Why Do Some Patients Relapse?

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Is BCR-ABL kinase inhibited?

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Yes

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Yes

- Additional mutations

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No

Yes

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### Why Do Some Patients Relapse?

Is BCR-ABL kinase inhibited?

No

Yes

- Drug efflux
- Bcr-Abl amplification
- Kinase mutations
- Drug metabolism
- Others
- Additional mutations

### BCR-ABL Substrates

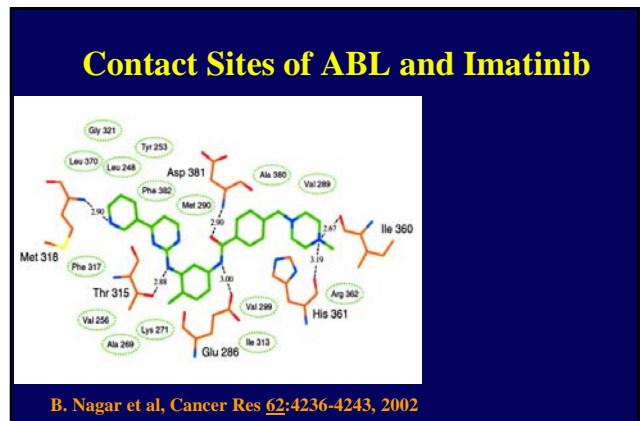
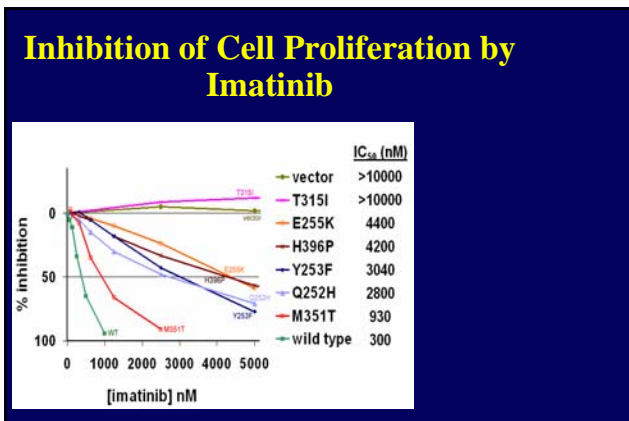
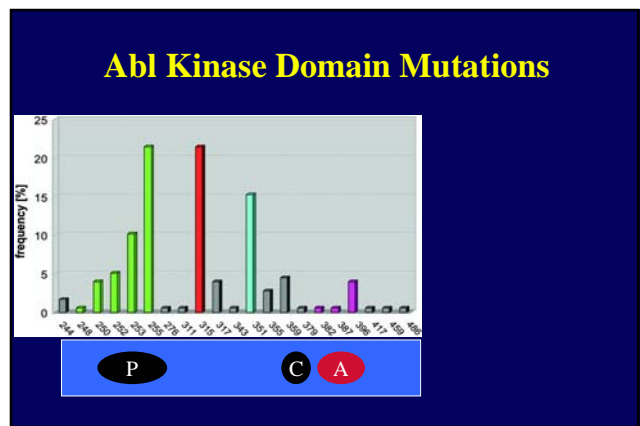
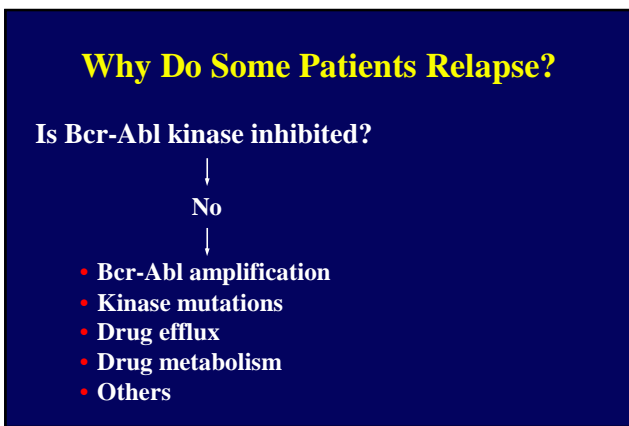
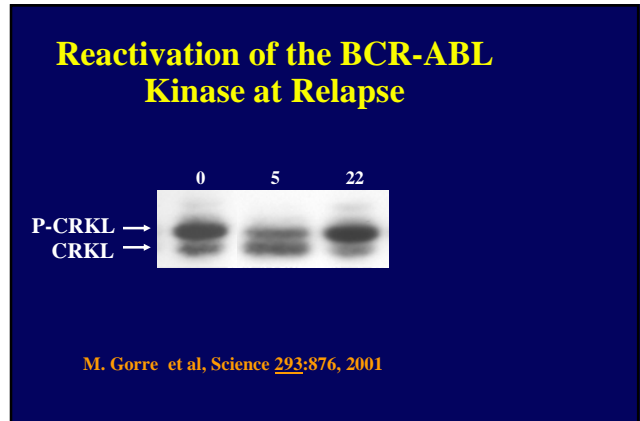
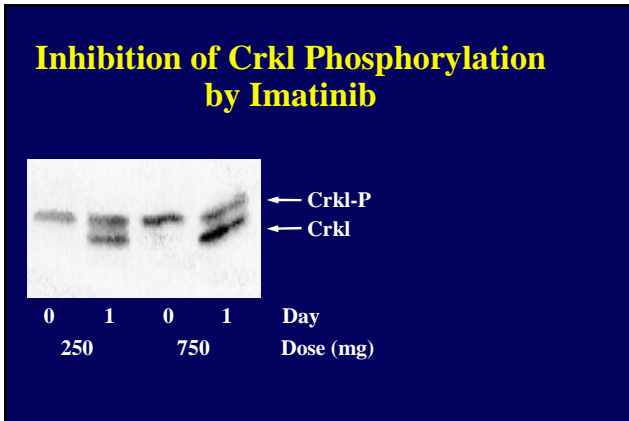
Tyrosine Phosphorylated Proteins  
in CML Patient Samples

- Bcr-Abl
- CrkL
- p62Dok
- STAT5

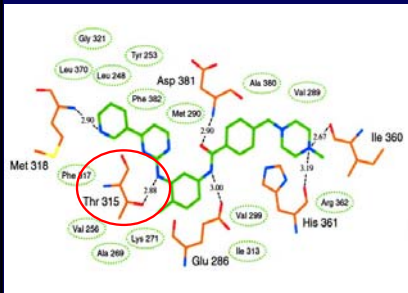
### CrkL



- Most heavily tyrosine phosphorylated protein in CML cells
- Direct substrate of BCR-ABL
- Required for BCR-ABL transformation

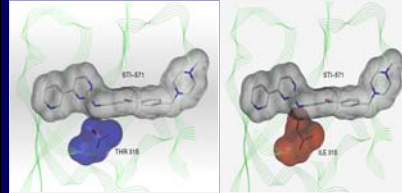


### Contact Sites of ABL and Imatinib



B. Nagar et al, Cancer Res 62:4236-4243, 2002

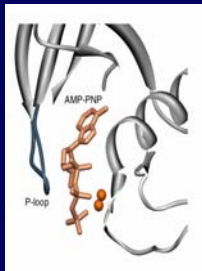
### T315I



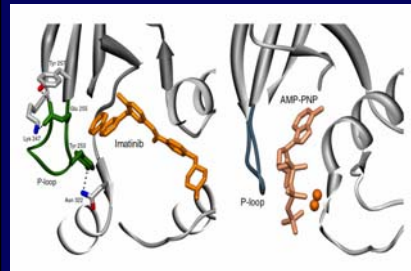
WILD-TYPE

T315I MUTANT  
(MODEL)

### P-loop Mutants



### P-loop Mutants



### Structure of the ABL Kinase Domain

ABL kinase with imatinib      ABL kinase with PD173955



B. Nagar et al, Cancer Res 62:4236-4243, 2002

### Novel ABL Inhibitors

- Nilotinib (Tasigna, AMN107)
  - Modification of imatinib structure to allow tighter binding

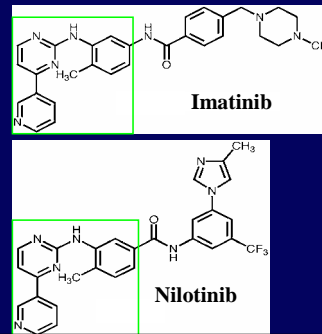
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  - Structural distinct SRC/ABL inhibitor
  - Fewer structural constraints to binding
  - Inhibits more kinases than imatinib

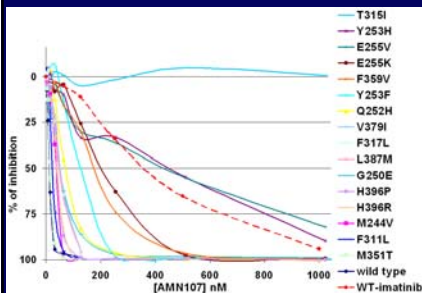
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- Both more potent ABL inhibitors than imatinib

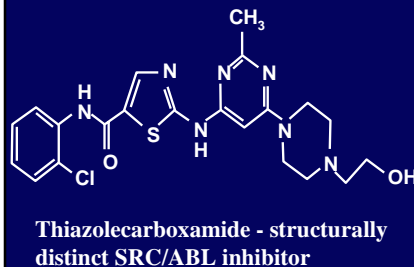
### Imatinib vs Nilotinib

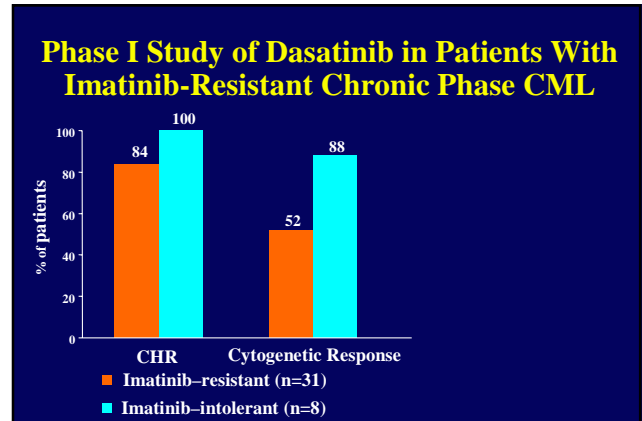
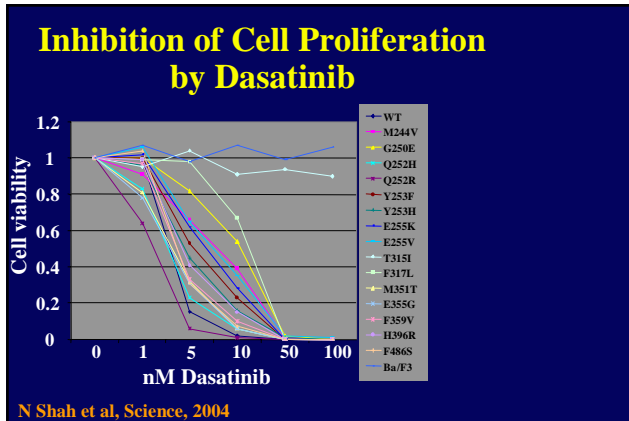


### Inhibition of Cell Proliferation by Nilotinib



### Dasatinib





### Summary of Dasatinib and Nilotinib Trials

- Significant activity in imatinib-resistant patients
- Activity observed against all imatinib-resistant mutants except T315I
- Relapses common in advanced phase
  - T315I
  - Other causes

### Summary of Imatinib Resistance

- Relapses mostly due to kinase domain mutations
  - Novel ABL inhibitors have significant activity

### Summary of Imatinib Resistance

- Relapses mostly due to kinase domain mutations
  - Novel ABL inhibitors have significant activity
- T315I remains insensitive to all current inhibitors
  - Numerous preclinical compounds

### Other Diseases Targeted by Imatinib

### Other Diseases Targeted by Imatinib

- Gastrointestinal stromal tumor
  - Driven by KIT mutations
- Hypereosinophilic syndrome
  - Driven by PDGF receptor rearrangement
- A few other rare diseases

### Imatinib and Gastrointestinal Stromal Tumor (GIST)

### Gastrointestinal Stromal Tumor

- GIST: intestinal sarcoma (formerly intestinal leiomyosarcoma) – KIT positive
- US annual incidence: ~5,000 cases

### Gastrointestinal Stromal Tumor

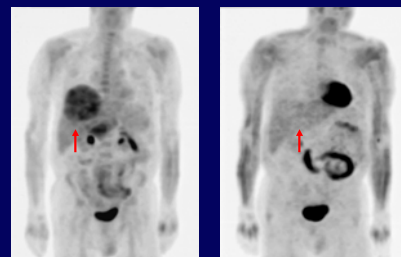
- Response rates to chemotherapy <5%
- Activating KIT mutations are present in the majority of patients

### Imatinib Response Data - GIST

	n= 147
Partial response (%)	54%
Stable disease (%)	28%
Progression (%)	14%

G. Demetri, et al, N Engl J Med 347:472-480, 2002

### PET Scan - GIST



Pre - 12/7/00

1/9/01

G. Demetri, et al

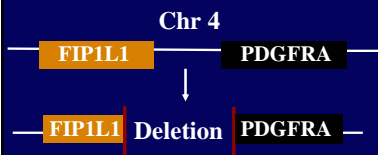
### Idiopathic Hypereosinophilic Syndrome (HES)

- Prolonged eosinophilia
  - >1,500/ul in blood
- Exclusion of other etiologies
  - Parasitic infection, allergy, etc.
- End organ damage
  - Heart, lungs, CNS, skin

### HES and Imatinib

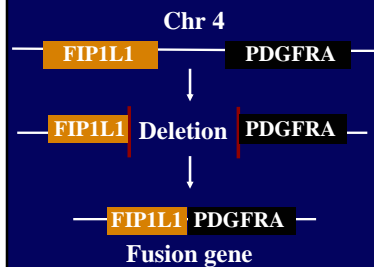
- Anecdotal reports of dramatic responses to imatinib
- Suggests HES is caused by a kinase that is inhibited by imatinib
  - ABL, ARG, KIT, PDGFRA, PDGFRB

### Molecular Pathogenesis of HES



J. Cools et al, N Engl J Med 348:1201-14, 2003

### Molecular Pathogenesis of HES



J. Cools et al, N Engl J Med 348:1201-14, 2003

An understanding of the molecular targets of a drug can lead to insights into molecular pathogenesis