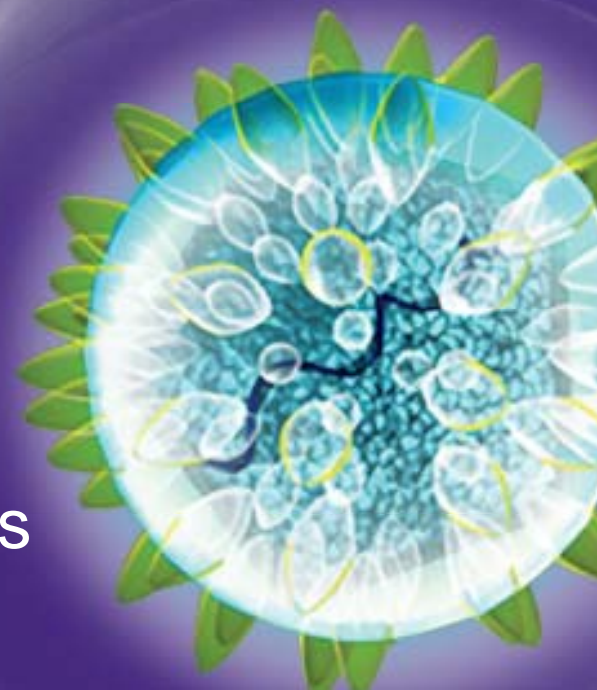


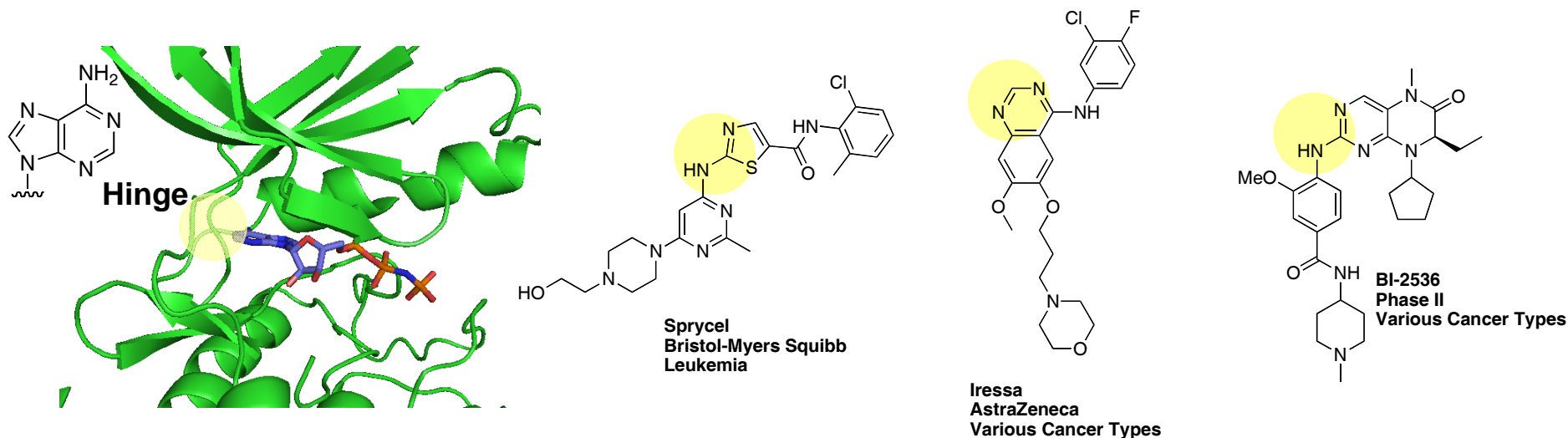
Design and synthesis of kinase inhibitors using novel heterocyclic systems



Dr Steven Durrant
Vertex Pharmaceuticals
23rd March 2011



Development of Kinase Inhibitors

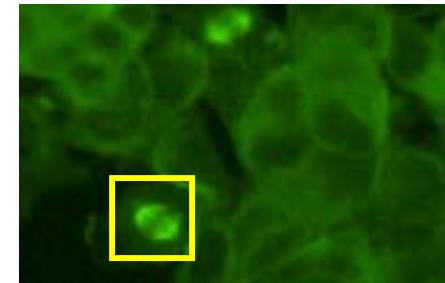


- 9 kinase inhibitors have reached the market
 - Many more have entered clinical trials
- ATP competitive inhibitors bind to “hinge” region using HBA/HBD motifs that mimic adenosine
- Mimicking this interaction leads to inherently flat aromatic systems
 - Poor physical properties result in need to append solubilising groups

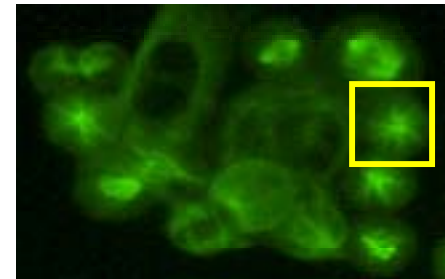
Plk1 is a promising oncology drug target

- Plk1 is overexpressed in multiple human tumours
 - Overexpression correlates with poor disease prognosis
- Plk family (1-4) are essential for accurate cell cycle progression
 - Plk1 plays a key role in the regulation of mitosis
 - Plks 2-4 have roles in tumour suppression / DNA damage response
- Small molecule inhibitors of Plk1 cause mitotic arrest and cell death (apoptosis) by activation of the mitotic checkpoint
- Development of ATP-competitive, Plk1-selective inhibitors could lead to a novel cancer therapy
 - Plk inhibitors are currently progressing through clinical trials

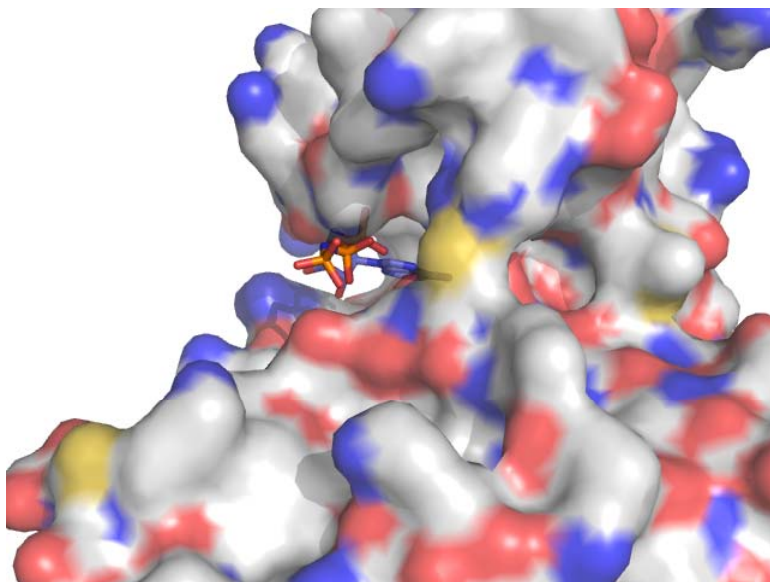
Control



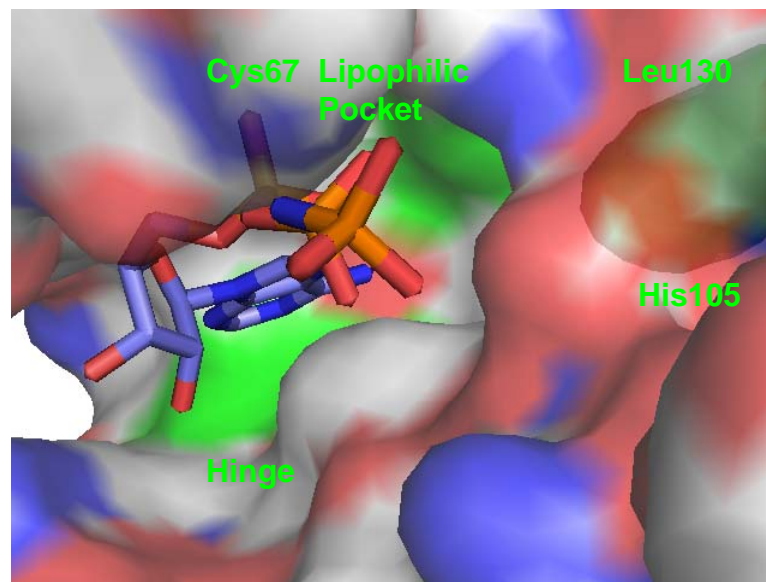
Plk1 inhibitor



Overview of the Plk active site



Overview of whole kinase

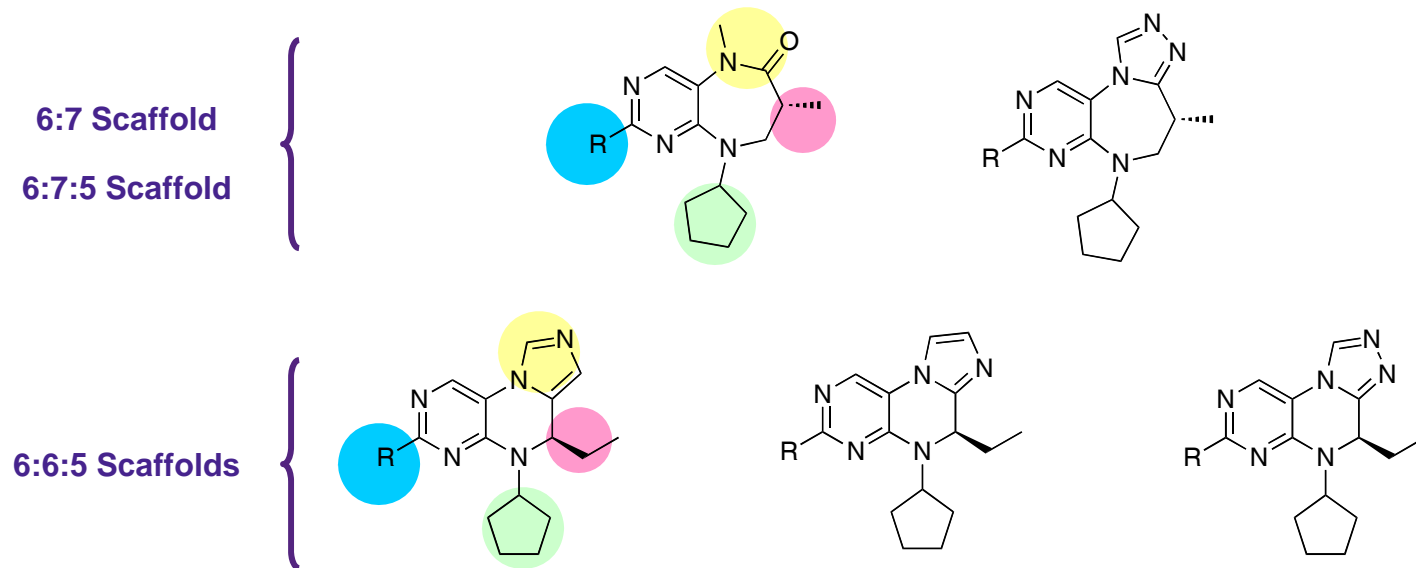


Key residues

- Overall structure typical of many kinases
 - Cleft where ATP binds
- Two areas believed to be important for potency and selectivity
 - Lipophilic pocket defined by Leu130/Cys67
 - His105

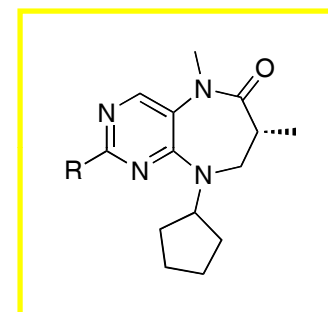
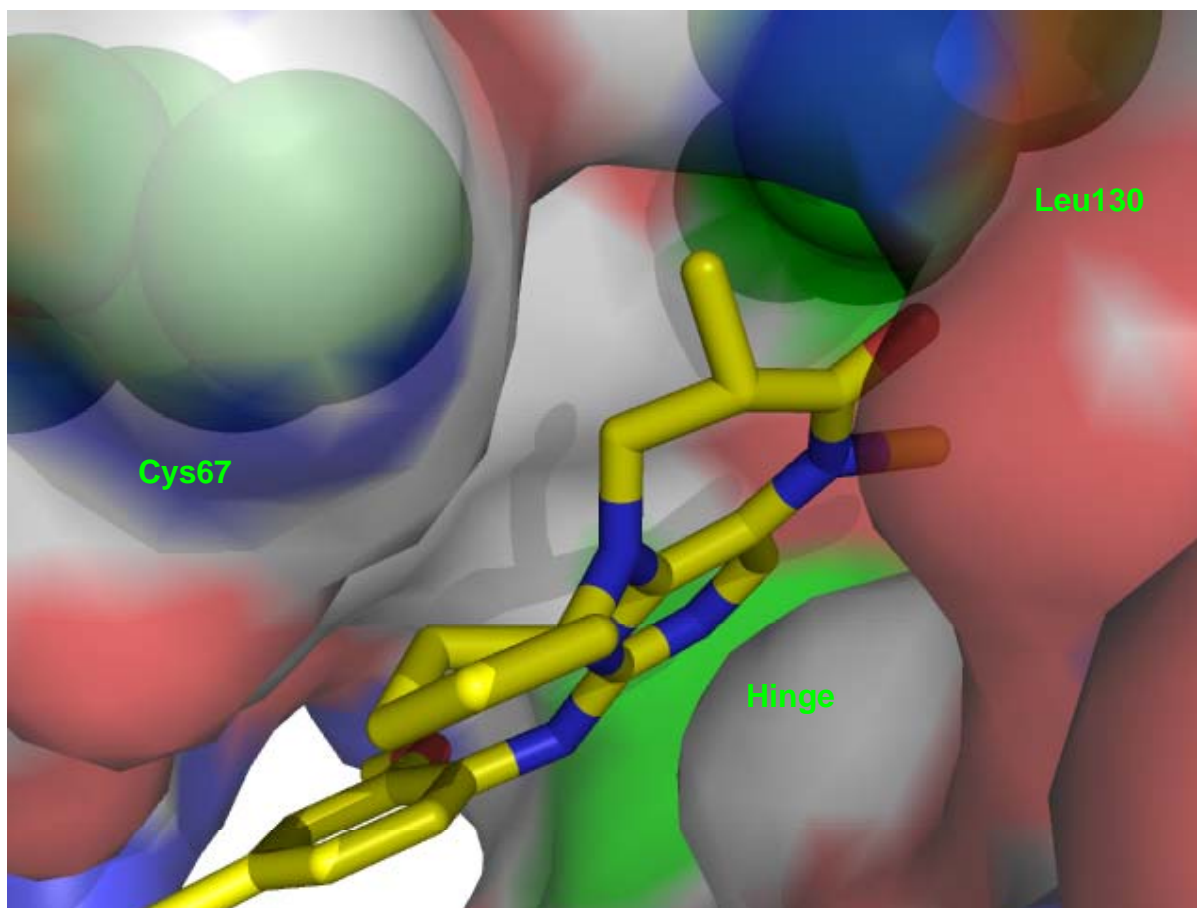


3-Dimensionality can be engineered to improve drug-likeness



- Array of novel related scaffolds offer a degree of 3-dimensionality
 - Expectation of good drug-like properties (e.g. solubility)
 - Opportunity to target 4 points of diversity

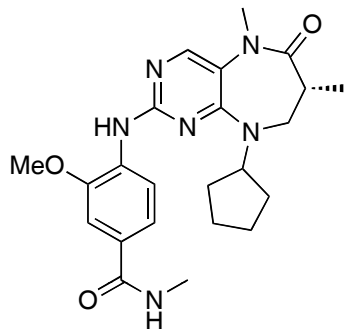
7-Membered Ring offers great potential for 3-D molecules



- Natural pucker of 7 membered ring allows good interaction with lipophilic pocket
 - Additional substitution enhances 3-dimensionality further



8,9-Dihydro-5H-pyrimido[4,5-*b*][1,4]diazepin-6(7*H*)-one scaffold (*a.k.a.* 6:7 scaffold)



Mw: 438.5 gmol⁻¹
cLogP: 2.73
PSA: 100 Å²

Compound 1

Plk1 K_i: 3 nM

Plk2 K_i: 162 nM

Plk3 K_i: 126 nM

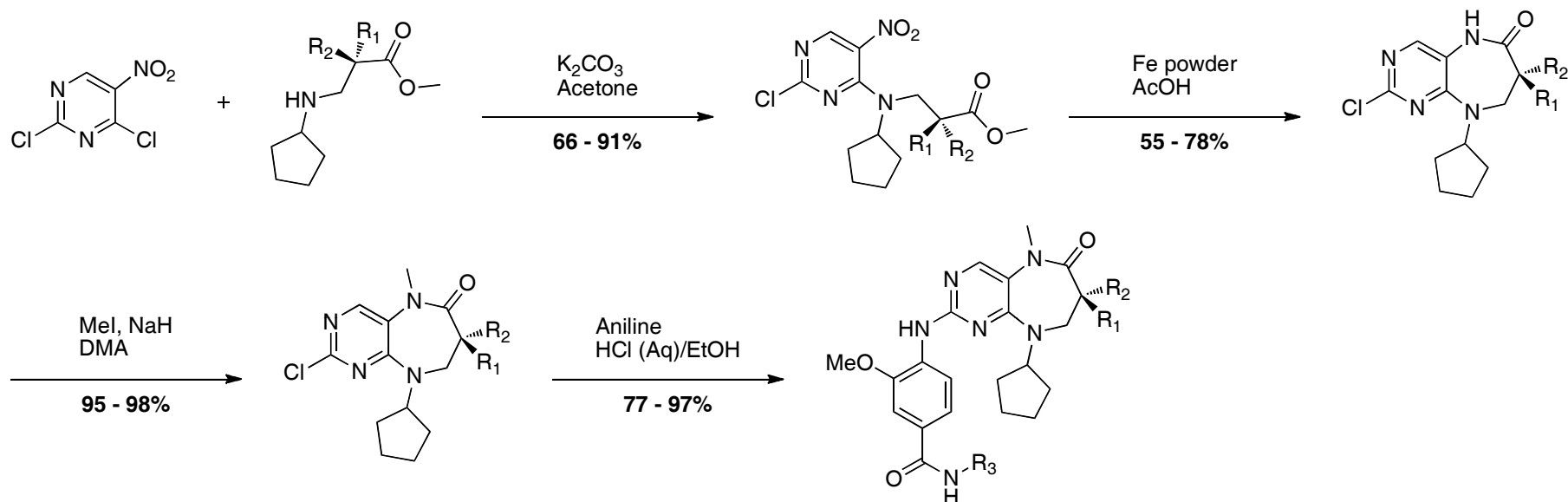
HCT-116 IC₅₀: 140 nM

Solubility at pH_{7.4}: > 200 μM

- 6:7 scaffold gives potency and selectivity consistent with drug candidate
- High solubility without need to append traditional solubilising groups

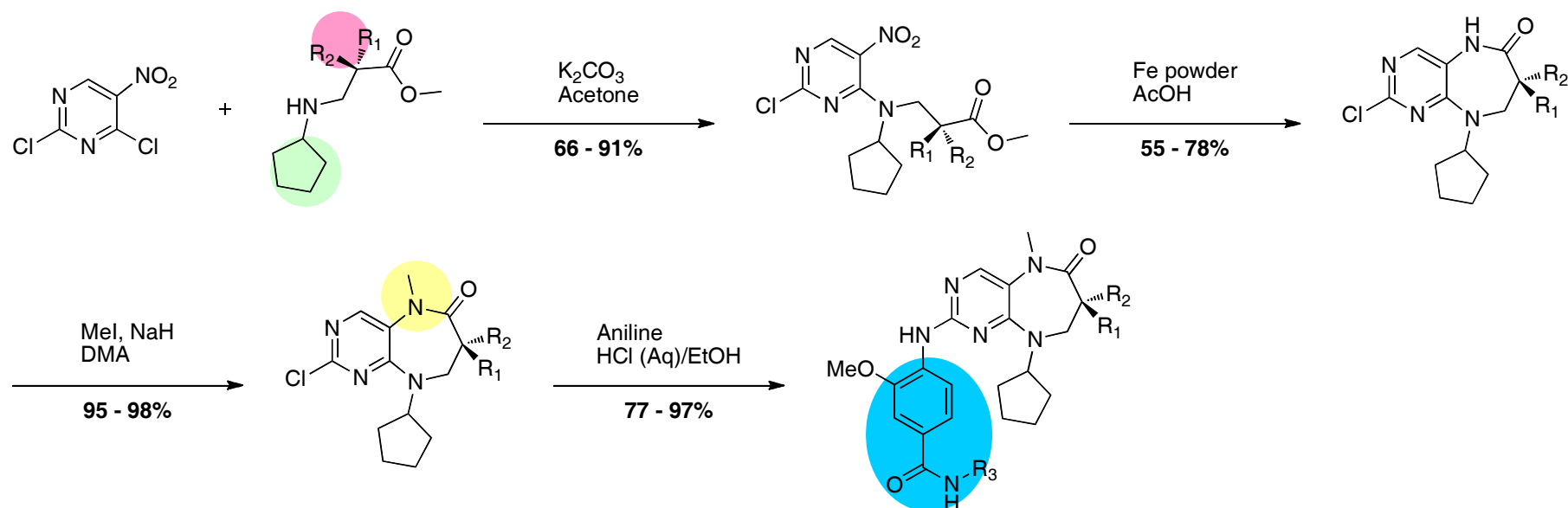


Synthesis of the 6:7 scaffold

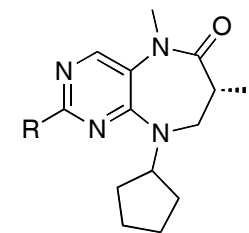


- Novel system can be accessed using optionally substituted β -amino acids
- Rapid, high yielding and versatile
 - 4 points of diversity to be manipulated

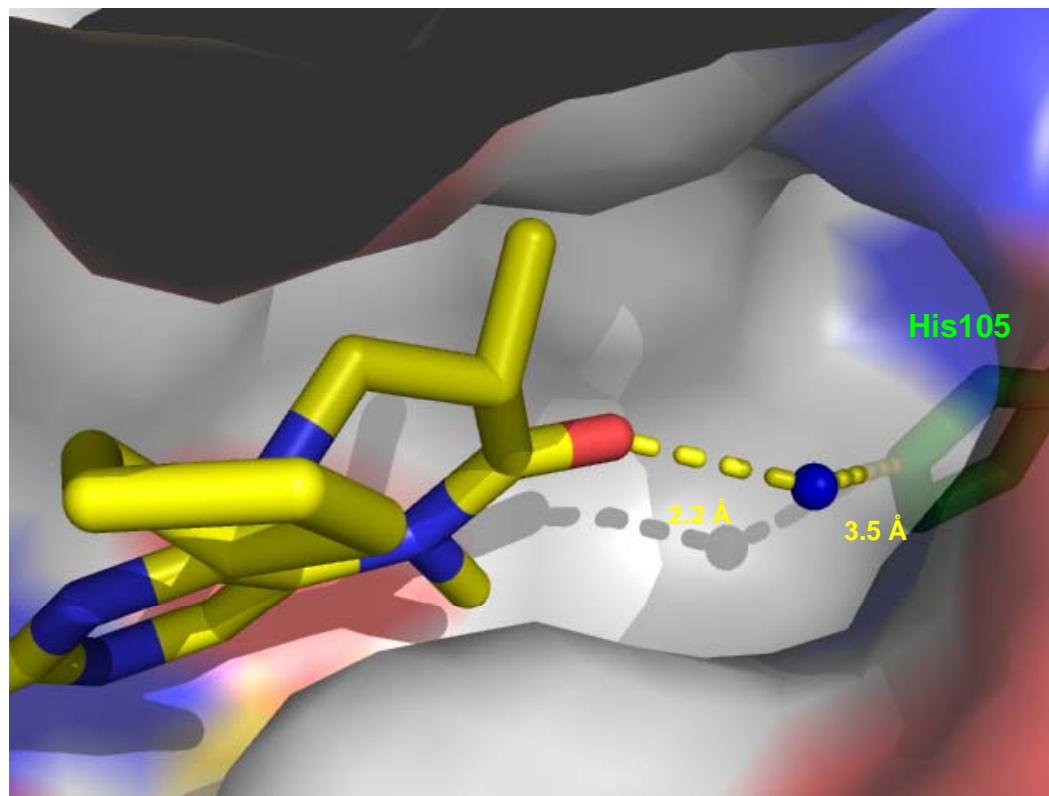
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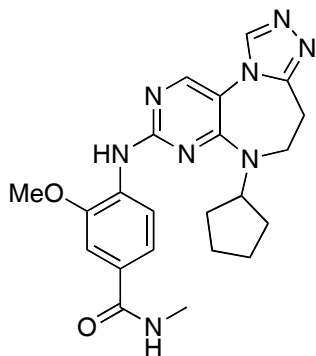


Interaction with Plk specific His105 Residue



- 6-Carbonyl makes water mediated interaction with specific His105 residue as shown through X-ray structures
- Objective: Make direct interaction with His105 as it is predicted to be energetically favourable

5,6-Dihydro-4*H*-pyrimido[4,5-*b*][1,2,4]triazolo[4,3-*d*][1,4]diazepine (a.k.a. 6:7:5 scaffold)



Mw: 434.5 gmol⁻¹ |
cLogP: 0.71
PSA: 110 Å²

Compound 2

PIk1 K_i: 19 nM

PIk2 K_i: 2850 nM

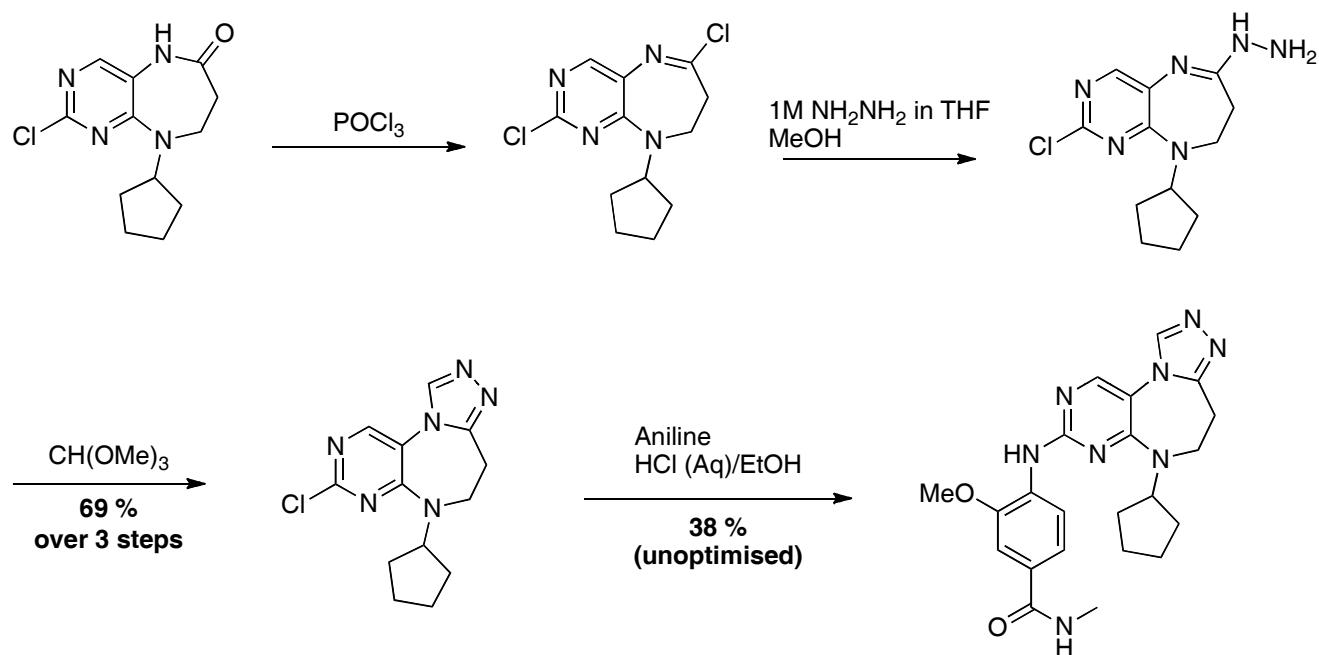
PIk3 K_i: 357 nM

HCT-116 IC₅₀: > 2000 nM

- Direct interaction with His105 with 6:7:5 scaffold suboptimal
- 7-ring no longer able to adopt appropriate conformation to interact in lipophilic pocket
- Small lipophilic substitutions failed to re-adjust binding complementarity

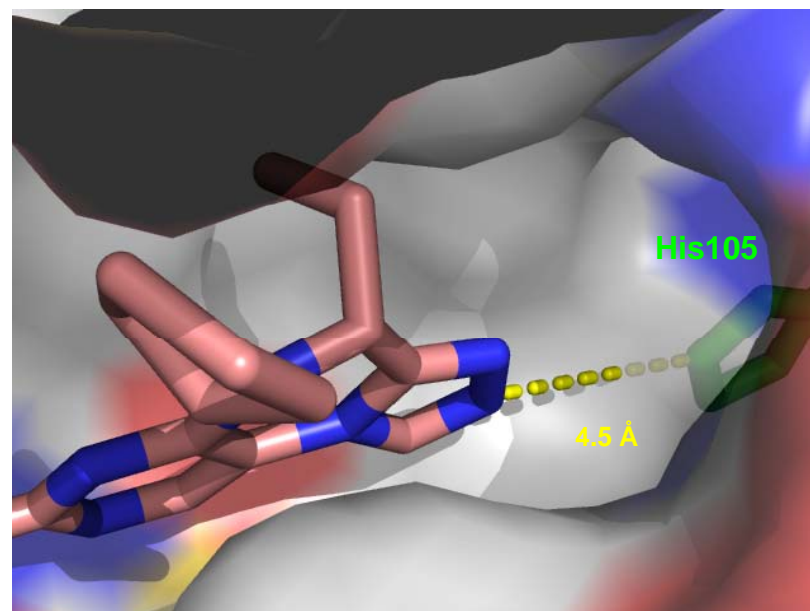
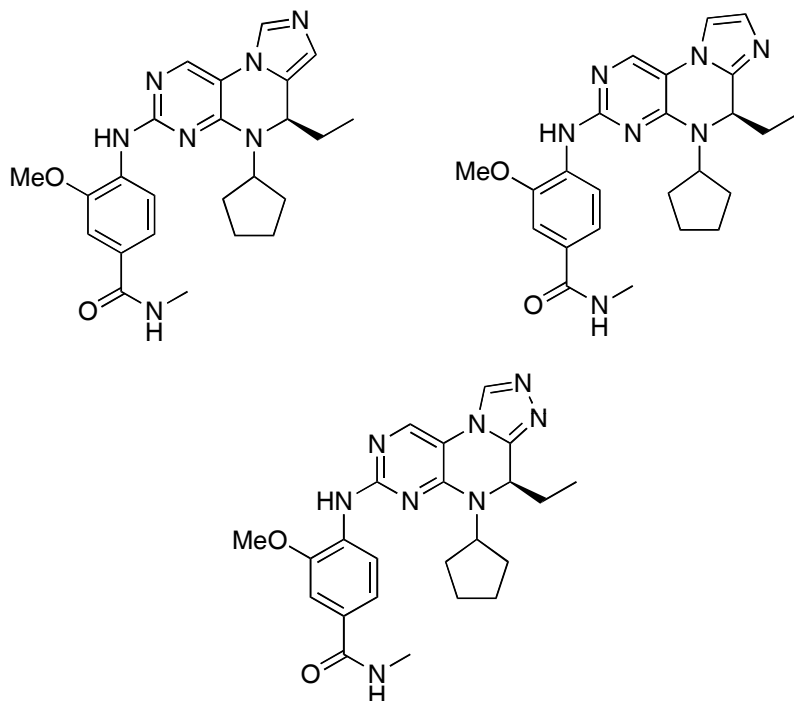


Synthesis of the 6:7:5 scaffold



- Novel tricyclic system accessed *via* same intermediate as bicyclic system
- Need to use non-aqueous hydrazine to obtain high yield
- Safety concerns: toxicity, explosive

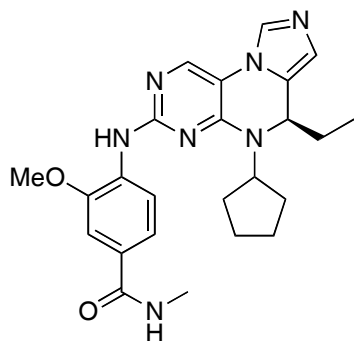
6:6:5 Scaffolds



- 3 potential scaffolds identified
 - All predicted to interact efficiently with His105 with no water mediated hydrogen bond
 - More rigid system enables substituents to adopt preferred conformation more easily



Optimising 5-membered ring interactions



Compound 3

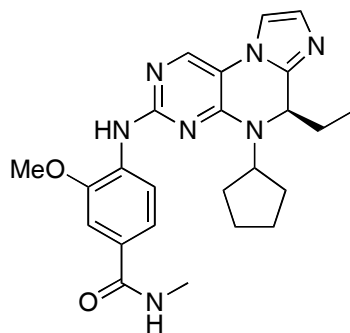
Plk1 K_i : 0.2 nM

Plk2 K_i : < 10 nM

Plk3 K_i : < 10 nM

HCT-116 IC_{50} : 45 nM

Solubility at $pH_{7.4}$: 24 μM



Compound 4

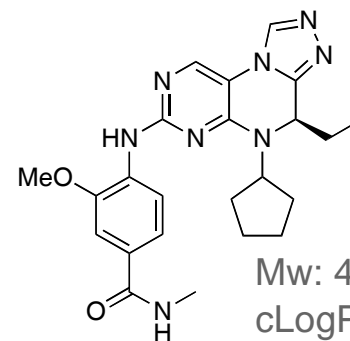
Plk1 K_i : 0.4 nM

Plk2 K_i : 19 nM

Plk3 K_i : 56 nM

HCT-116 IC_{50} : 179 nM

Solubility at $pH_{7.4}$: 31 μM



Compound 5

Plk1 K_i : 0.1 nM

Plk2 K_i : < 10 nM

Plk3 K_i : < 10 nM

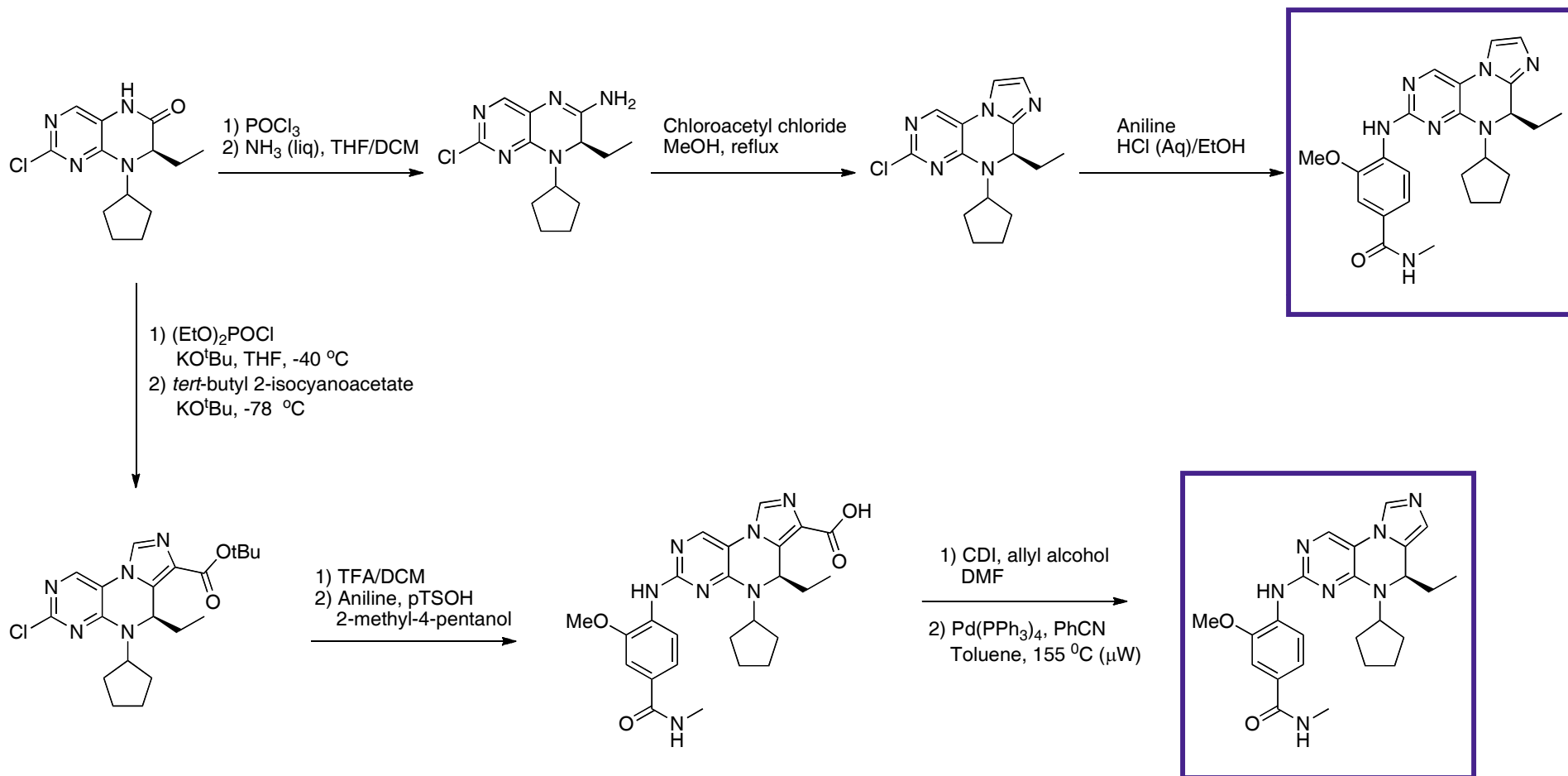
HCT-116 IC_{50} : 7 nM

Solubility at $pH_{7.4}$: > 200 μM

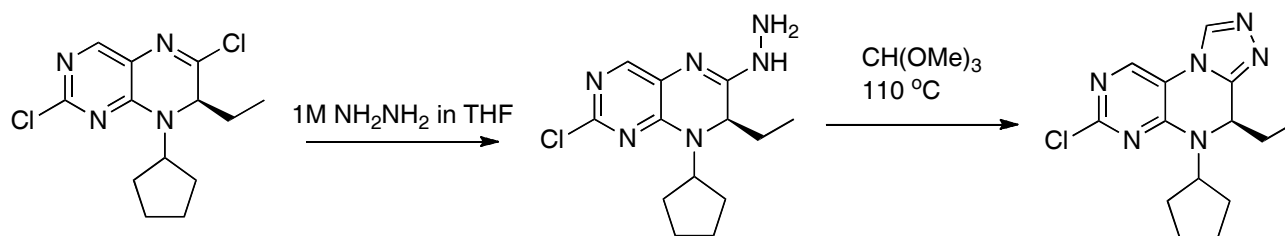
Mw: 448.5 $gmol^{-1}$
cLogP: 1.54
PSA: 110 \AA^2

- Fused ring systems provides potent Plk inhibitors
- Good solubility with no traditional solubilising group
 - Triazolo variant preferred for its optimal physical properties

Synthesis of Imidazolo 6:6:5 scaffold variants

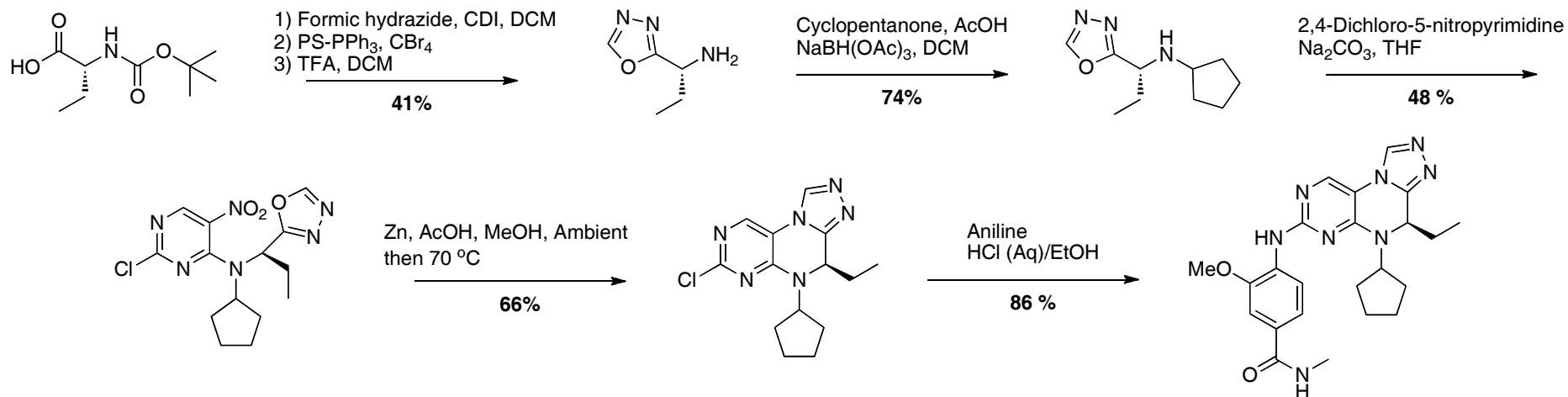


Triazolo 6:6:5 scaffold - pro's and con's



- ✓ 4,5-Dihydro-[1,2,4]triazolo[4,3-f]pteridine core scaffold allows very potent druglike inhibitors to be developed
- ✗ Route requires use of non-aqueous hydrazine
 - Introduces safety concerns particularly on large scale
- In order to safely produce large quantities of the core scaffold an alternative route was required

A safe synthesis of triazolo 6:6:5 *via* key reductive cyclisation



- Use of 1,3,4-oxadiazole allows incorporation of hydrazine motif in a safe manner
- Key step: One-pot reduction of nitro group with concomitant cyclisation leads to formation of triazole ring in good yield

For similar cyclisation reaction see WO2005121152



Conclusions

- 5 novel scaffolds have been identified for potent kinase inhibition
- Inherent three dimensionality confers good solubility to all these scaffolds without need to append traditional solubilising groups
- Safe route developed to triazolo 6:6:5 scaffold
 - Key reductive cyclisation step avoids the use of non-aqueous hydrazine



Acknowledgements

Chemistry

- Jean-Damien Charrier
- Guy Brenchley
- David Kay
- Shazia Keily
- Chau Mak
- Michael O'Donnell
- Françoise Pierard
- Joanne Pinder
- Sharn Ramaya
- John Studley
- Heather Twin
- Anisa Virani

Project Management

- Peter Weber
- Julian Golec
- Stephen Young

Biology

- Matthew Griffiths
- Catherine Hudson

Crystallography and Modelling

- Kieron Brown
- Ronald Knegtel



Vertex: A Global Health Sciences Company

- Founded in 1989; public since 1991
- 6 major locations worldwide including 5 R&D sites with standalone discovery capabilities and numerous field locations
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 - San Diego, CA
 - Coralville, IA
 - Milton Park, UK
 - Laval, Canada
 - Washington, D.C.
- 1,700+ employees worldwide, 70% in R&D

