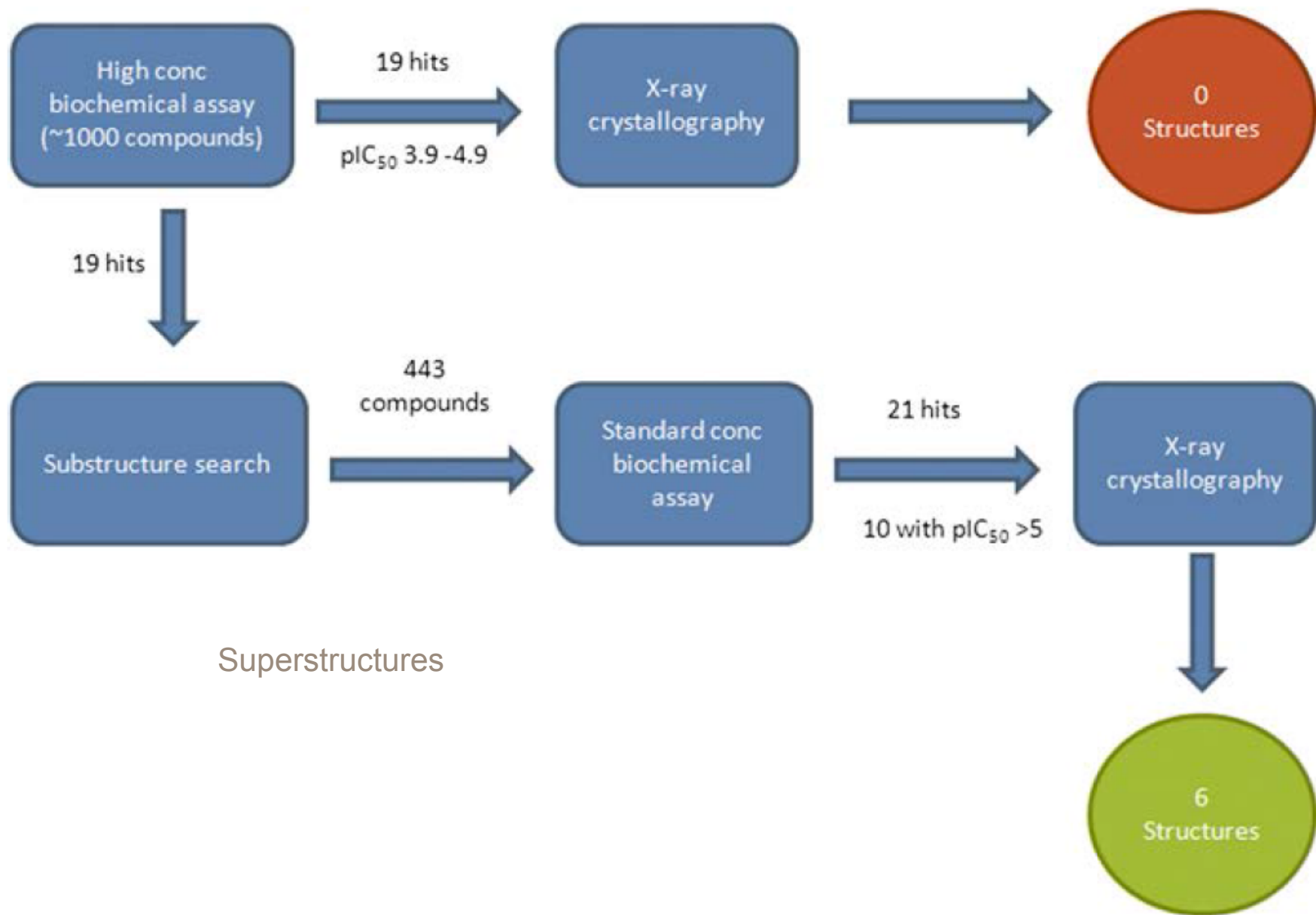
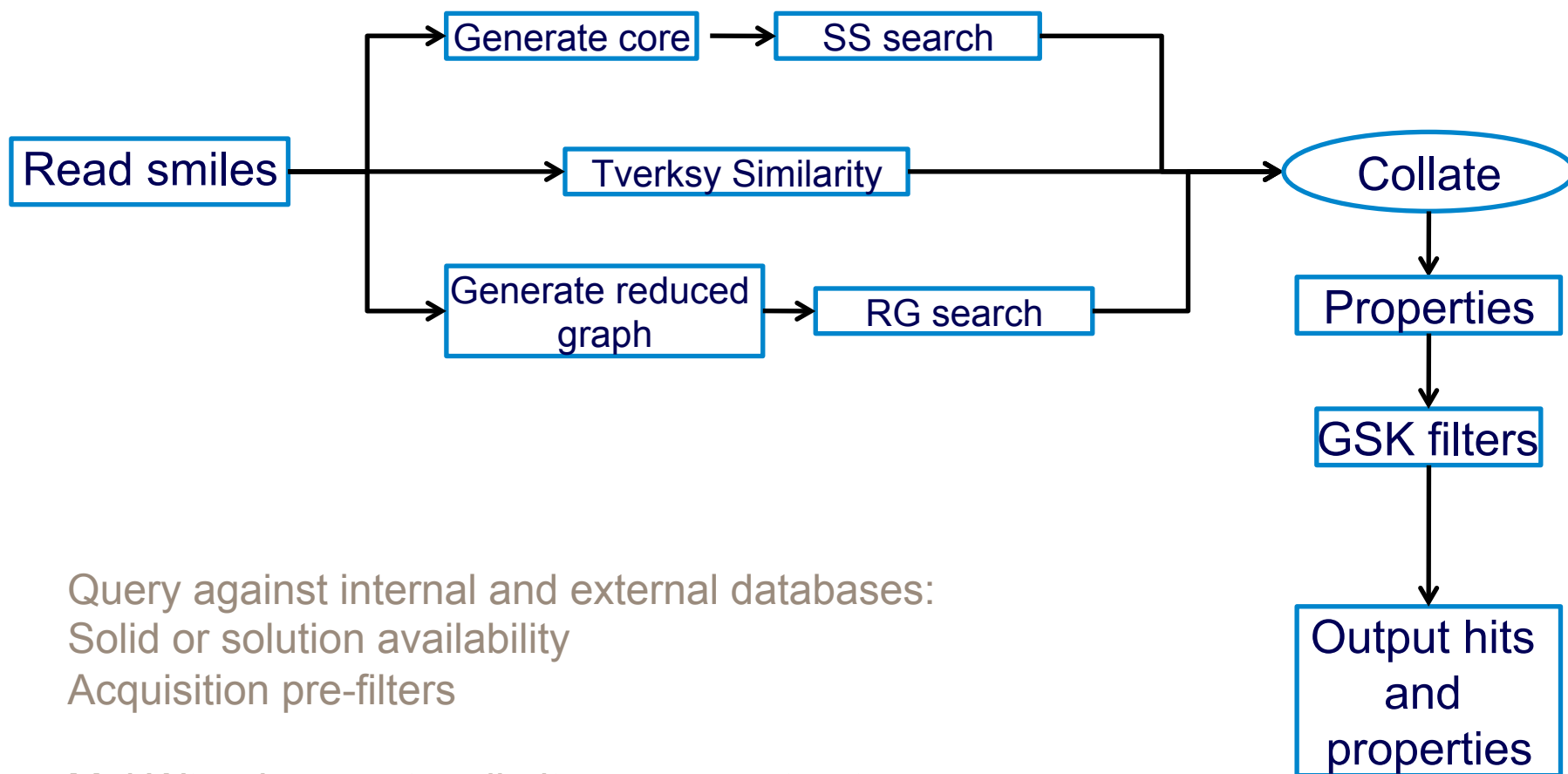


Fragment hit prioritisation

Ian Wall

Analogue Searching increases X-ray Success

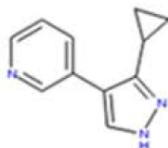




Query against internal and external databases:
Solid or solution availability
Acquisition pre-filters

Mol Wt or heavy atom limit per query structure
Total hits per sub-query (most similar or lowest mol wt)

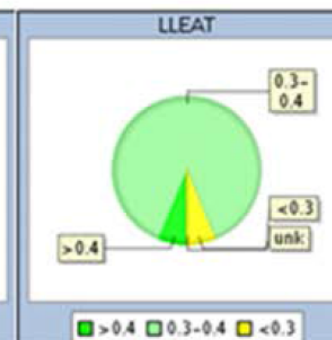
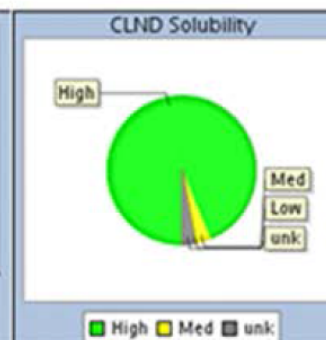
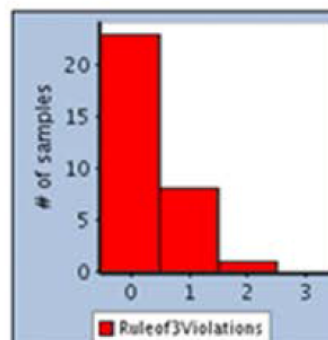
Multi parameter decision making



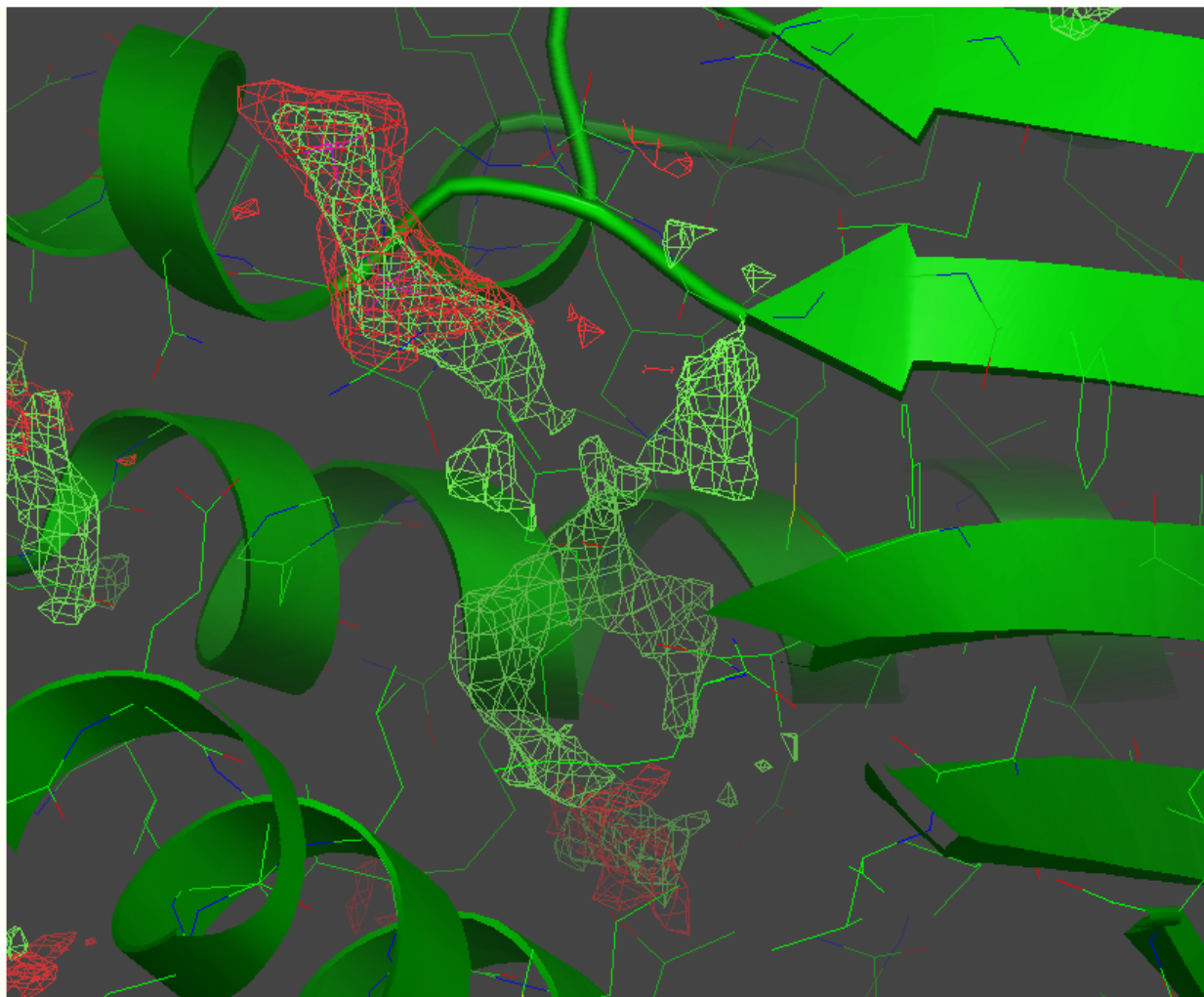
Calculated Properties	
Regno	
mw	185.23
clogP	1.2
tPSA	41.57
NumHeavy	14
Acceptors	2
Donors	1
Rotbonds	2

Measured Properties	
Regno	
HPGDS pIC50	4.8
LigEff	0.47
LigEffAT	0.46
CLND_sol	>=391 uM
CHROM_LOGD_PH74	1.6
HSA_PCT_BIND	84.2
AGP_PCT_BIND	53.0

Summary Comments					
Properties of Validated Hit Series	Current data pIC50	LE	LLEAT	Does the series hit the criteria	Comments
LE between 0.3 & 4.3					
Best in cluster					
Clear X-ray binding mode					
Astex rule of 3					
Synthetic tractability					
Expectation that elaborated molecules will be patentable					
Off target effects					
EXP					
Stability data					



Active site mapping (eg GRID)



Viewing	GRID	HotSpots	Structures
<input checked="" type="checkbox"/>	Methyl (-10.0)		-3.0
<input type="checkbox"/>	Hydrophobic (-10.0)		-1.0
<input type="checkbox"/>	amide NH (-10.0)		-6.0
<input type="checkbox"/>	sp3 amine cation (-15.0)		-8.0
<input type="checkbox"/>	Water (-16.0)		-8.0
<input checked="" type="checkbox"/>	carbonyl O (-10.0)		-5.0
<input type="checkbox"/>	carboxy O, e.g. carboxylic acid (-10.0)		-2.0

[AstexViewer2 documentation](#)

- Hit validation? (active in orthogonal assays)
- Judge the series not the hit
- Generate as much data as possible to inform decisions
- Potency a consideration, but not the primary factor
- Consider potential for growth
 - Growth vectors
 - Known structures/SAR
 - Hot spotting tools
- Synthetic tractability
- Chemical stability
- Not worried about selectivity at this point

- Collect all screening data before committing to work on a series
 - Screen available analogues to generate SAR before starting chemistry
 - Core optimisation (most LE starting point)
 - Grow slowly
 - Monitor properties (use of LE, LLE, etc.)
 - Use of structure based design to drive chemistry plans
 - Growing and hybridisation preferred to linking fragments
 - Avoid flat highly aromatic compounds
-

Was my change “worthwhile”?

- Required potency change to maintain LLE_{AT} for common aromatic substituents

