

# THE CFL: MAXIMIZING FRAGMENT-TARGET INTERACTION SPACE WITH A MINIMAL SCREENING SET

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## FBLD AT BIOBLOCKS

Many fragment library compounds are unique but not designed to enable follow-up. BioBlocks's Comprehensive Fragment Library (CFL) is a next-generation set of small, rigid, medicinally interesting fragments designed to address this common problem in fragment SAR. Extensive clustering analysis allows broad coverage of chemistry space and ensures that an immediate follow-up strategy is available from every screening hit.

## DESIGNING THE CFL

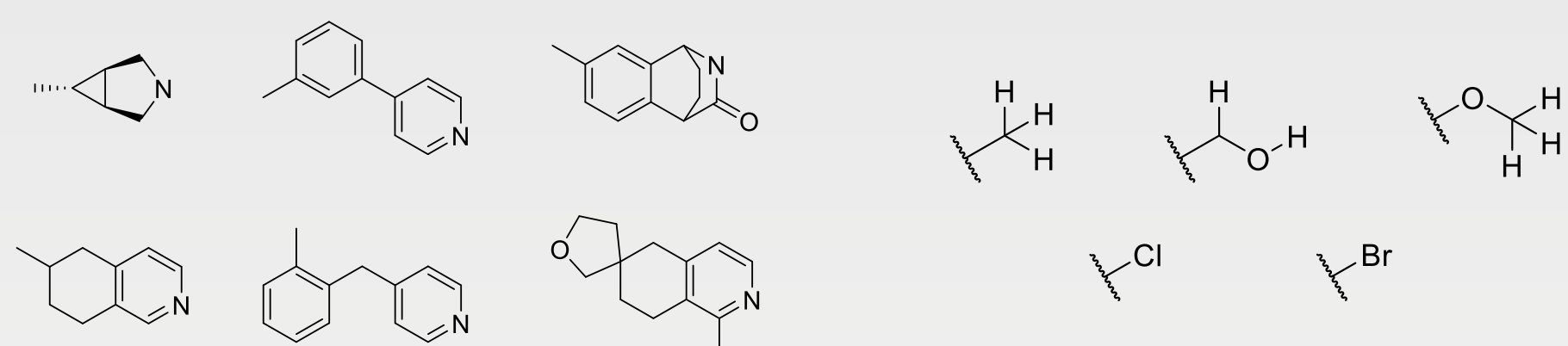
The CFL originates from a set of >3 million potentially synthesizable virtual fragments designed from first principles and 3D enabled to maximize exploration of target interactions. An exhaustive virtual set of ring structures containing ≤18 heavy atoms and 1 handle atom was generated.

Key inclusion criteria:

- ≥ 1 ring, including bridged, spiro and fused ring connections
- ≤ 3 unique rings
- ≤ 2 rotatable bonds
- C,N,S,O atoms only
- ≤ 2 ring assemblies
- 1 atom has all rotatable bonds
- ≤ 1 S

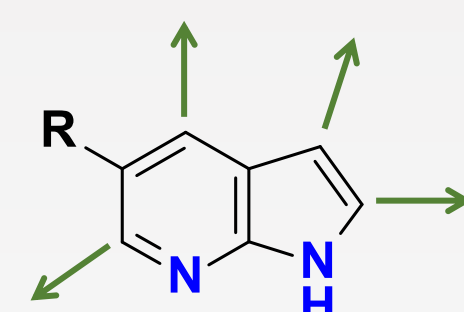
### Example CFL Fragments

### Example CFL Handles



## FRAGMENT ANALOGS FROM A CFL HIT

| Analog Category      | CFL000025 | 10 Fragment Hits | Plate 1 |
|----------------------|-----------|------------------|---------|
| Handle Analogs       | 10        | 100              | 550     |
| Handle Vectors       | 4         | 47               | 257     |
| Same 2D Scaffold     | 37        | 343              | 3308    |
| Same HBond Pattern   | 274       | 12857            | 42667   |
| Same 3D Cluster      | 47        | 573              | 6371    |
| Total CFL Compounds  | 873       | 13676            | 50669   |
| % Unique to Compound | 100%      | 74%              | 26%     |
| CFL Analogs          | >30,000   | >1 M             | >25 M   |



Five categories give access to orthogonal analog sets from a hit

- Handle – variable synthesis options
- HBond Pattern – core options
- Vector – handle position options
- 3D Cluster – similarity options
- Scaffold – heavy atom options

L2L also uses an analog generator to explore sp3 ring variations

## CURRENT FRAGMENT SET

Available screening collection:

120 Fragments for initial screen

Soluble at 200 mM in DMSO

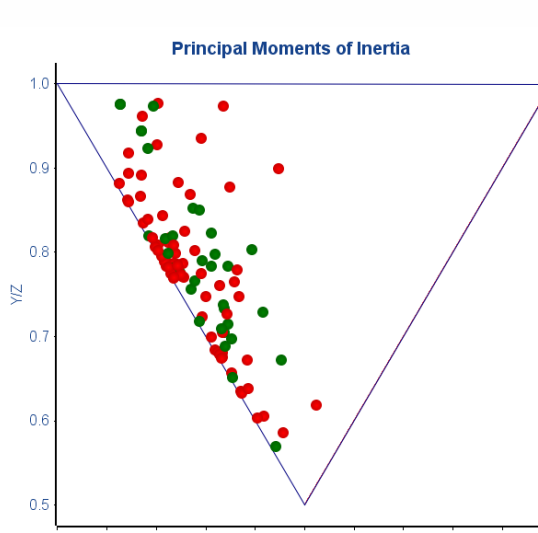
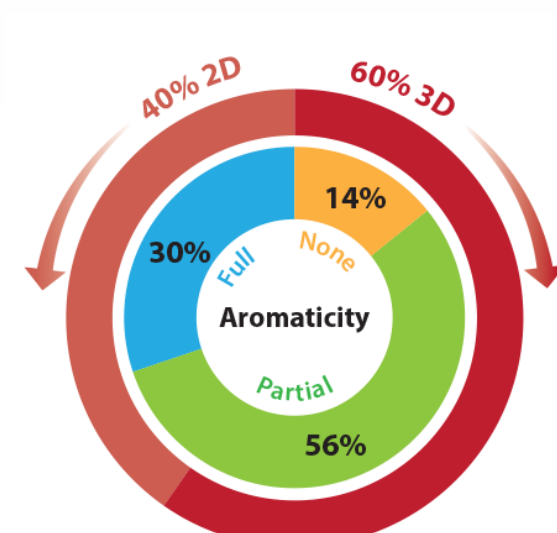
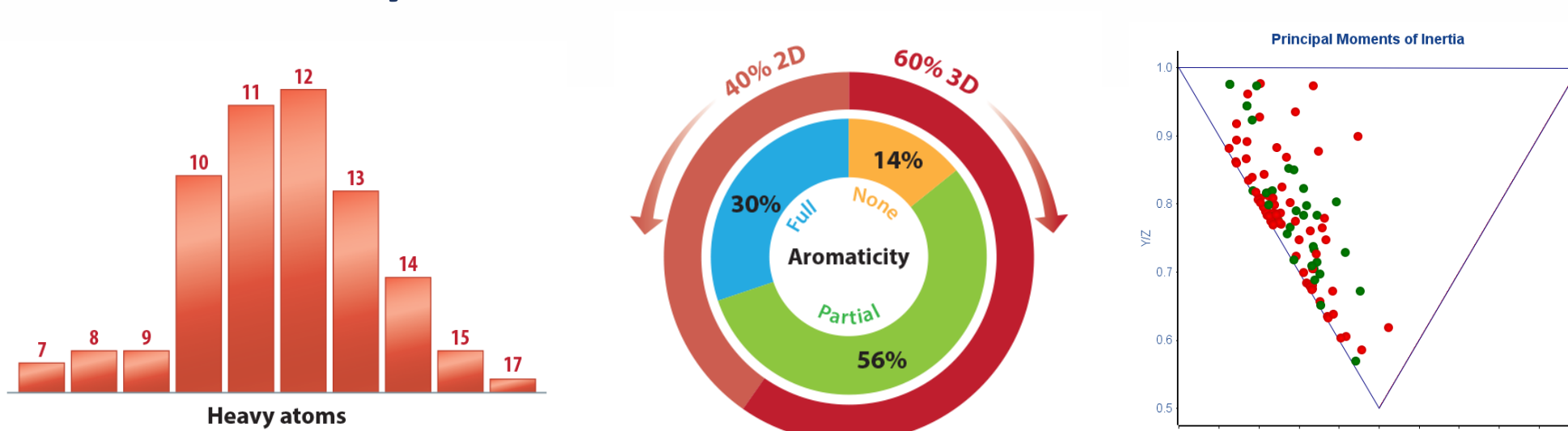
Well behaved at 2 mM in multiple assay formats

>100 Analogs for follow up

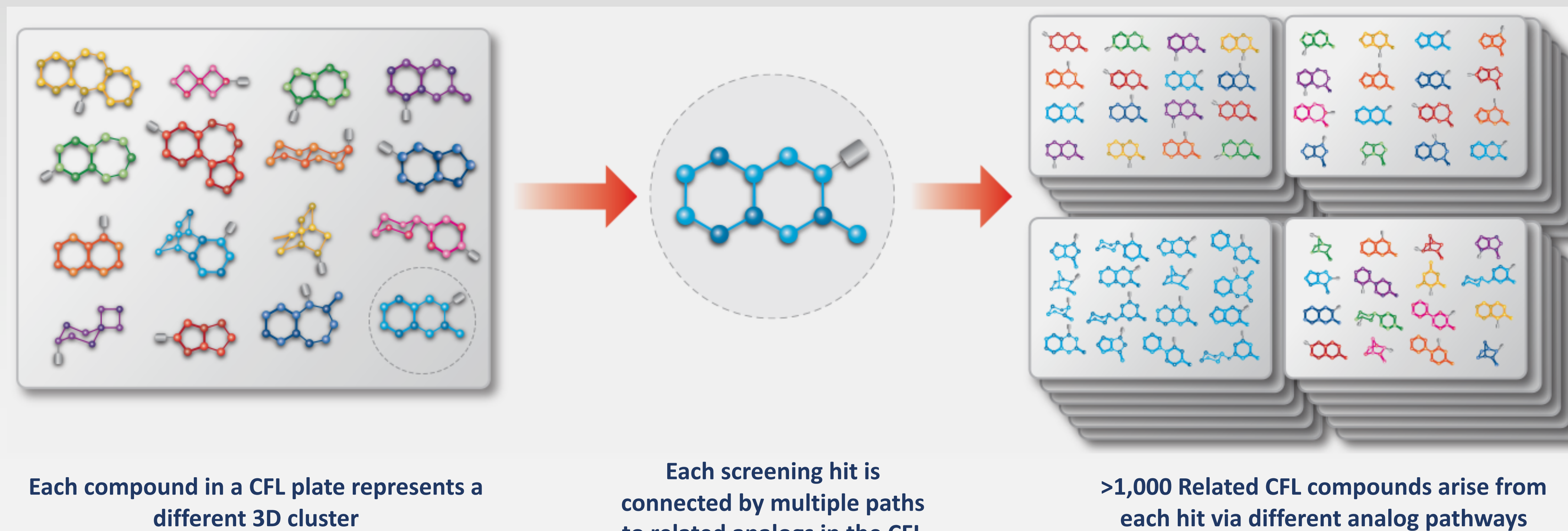
Soluble at ≥100 mM in DMSO

Alternative handles for multiple expansion chemistries

Plate Summary:



## THE LEAP-TO-LEAD™ PLATFORM REVEALS IMPROVED PATHS FORWARD



Each compound in a CFL plate represents a different 3D cluster

Each screening hit is connected by multiple paths to related analogs in the CFL

>1,000 Related CFL compounds arise from each hit via different analog pathways

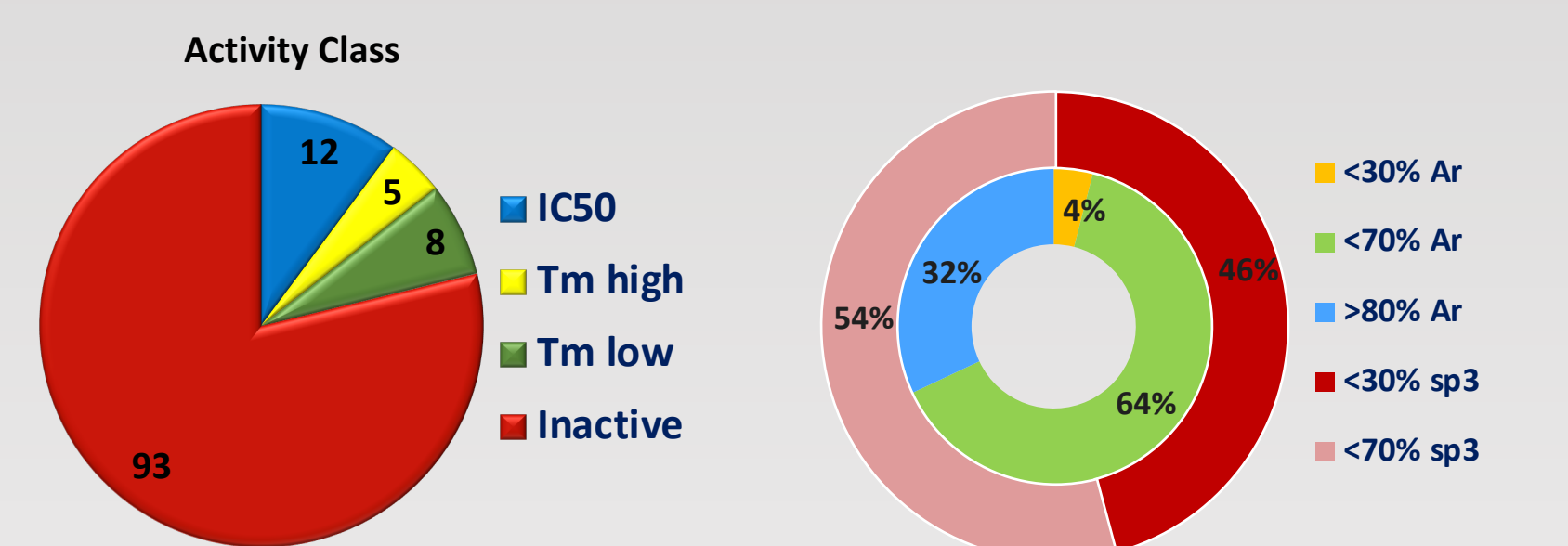


## TARGET CASE STUDY 1



Prostaglandin D synthase (HPGDS) mediates inflammatory signaling in asthma. The CFL library set was screened against HPGDS in a thermal shift assay (TSA). Followed by a competitive FP assay with a known active site inhibitor.

Screening Summary:



Hit fragments are more flat and more aromatic than the overall plates. Published inhibitors for this target have high % aromatic character. The majority of the CFL fragment hits are only partially aromatic.

## ANALOG GENERATION

4/6 Analog Categories Assayed

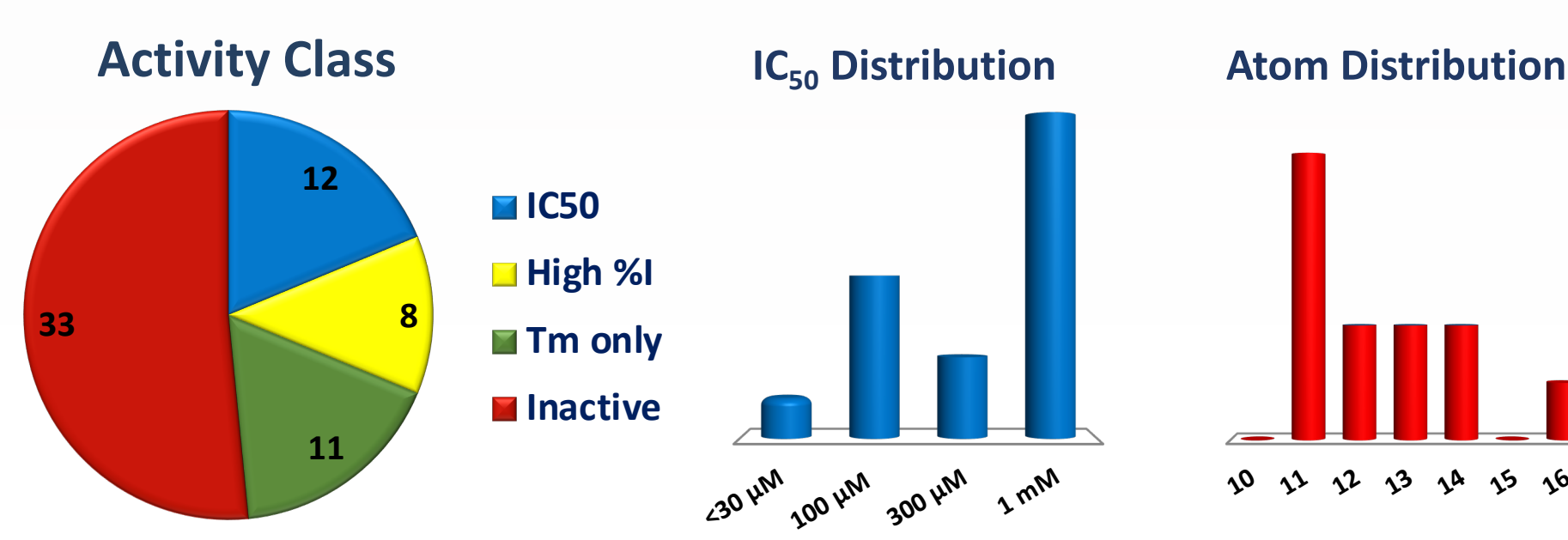
Handle Scaffold  
Vector L2L analoger  
3D reserved for next round

17/25 Fragments had readily accessible analogs

<1% of possible analogs were commercially available  
Supplemented with 1 step synthesis

64 Analogs tested in follow up activity assays

## ANALOG ACTIVITY

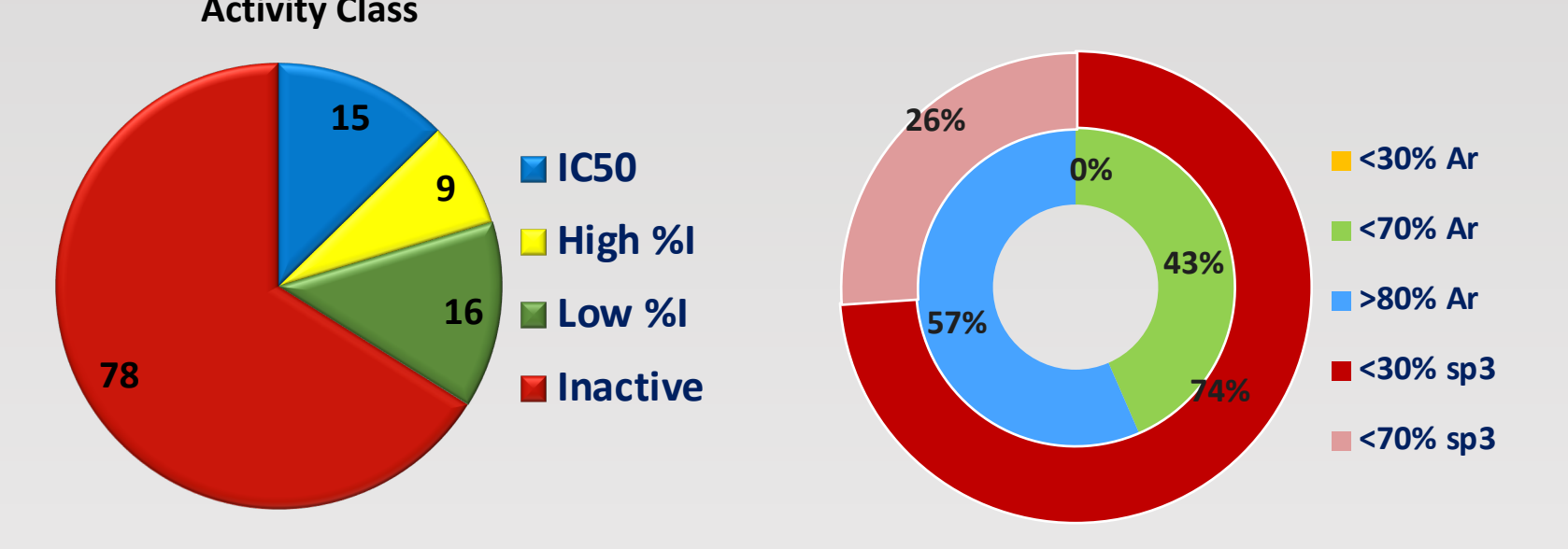


## TARGET CASE STUDY 2



SGK1 is an emerging kinase target in oncology and inflammation. Amenable to high concentration biochemical assays. The CFL library set was screened for %inhibition at 2 mM against active SGK1.

Screening Summary:



Hit fragments with IC<sub>50</sub> values are predominantly aromatic. Published inhibitors for this target are also predominantly aromatic. Some hits identified from CFL with higher Fsp<sup>3</sup>. Most hits biased towards typical hinge-binding HBond pattern DA-2.

## ANALOG GENERATION

4/6 Analog Categories Assayed

Handle Scaffold  
Vector  
HBond pattern – varied core with DA-2

12/15 Fragments had readily accessible analogs

<1% of possible analogs were available commercially  
>95% of available analogs were predominantly aromatic

95 Analogs tested in follow up activity assays

## ANALOG ACTIVITY

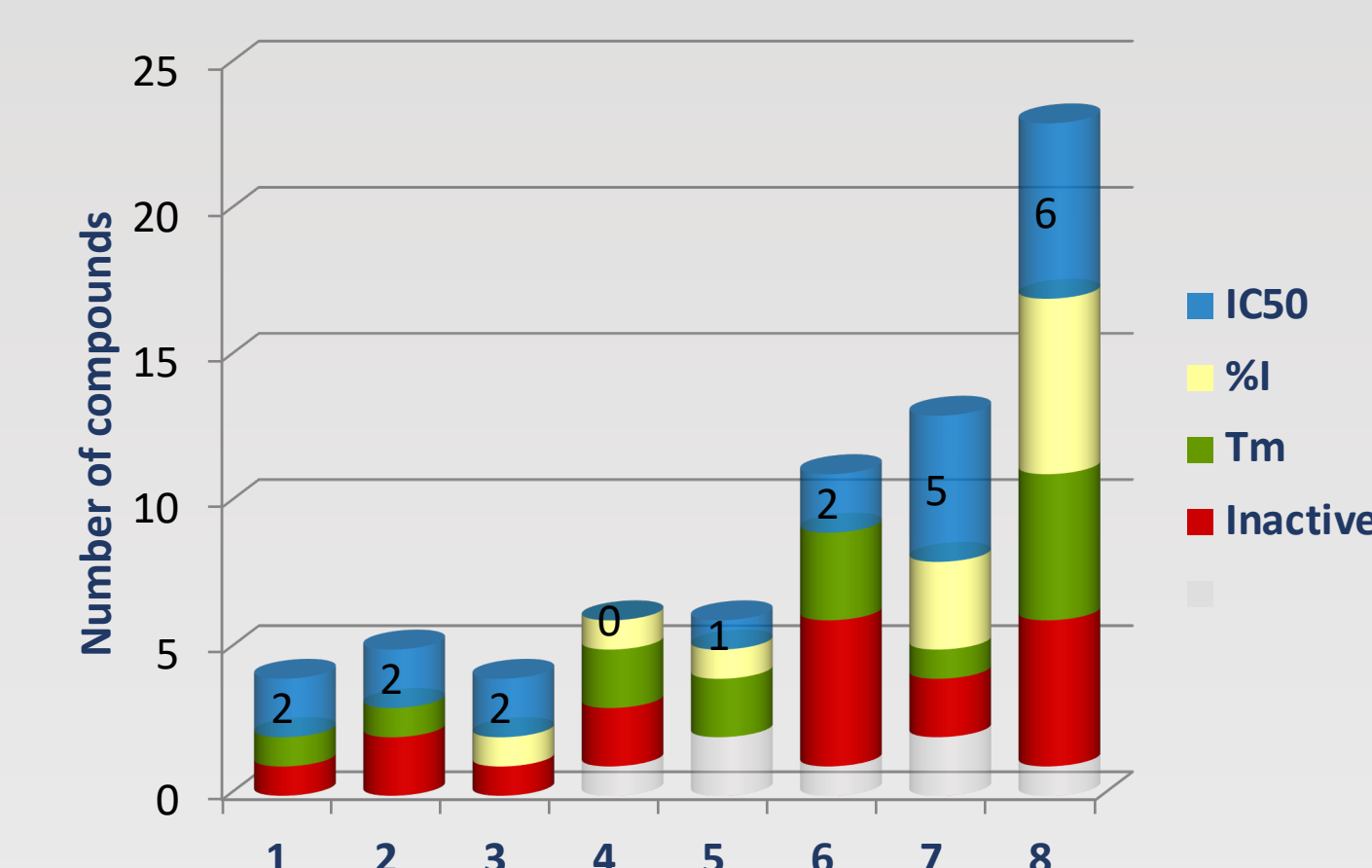


## ANALOG FAMILIES FOR LEAD FINDING

Analysis pathways designed to distribute analogs into families. Focus on ligand efficient families to start with desirable properties. Tactical med chem allows improved potency while maintaining properties. L2L strategy provides superior entry points to lead optimization.

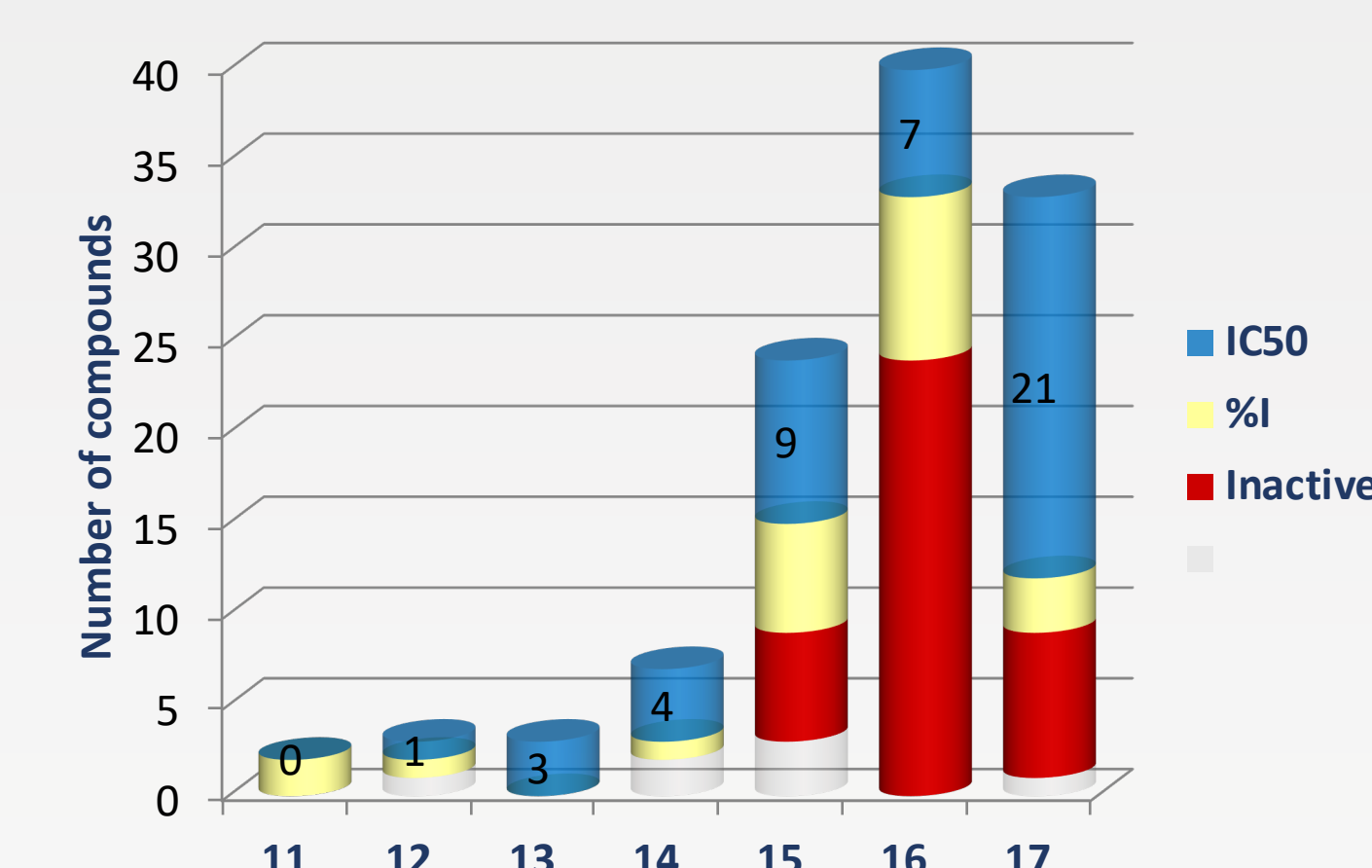
HPGDS analogs distribute into 8 structural families

Remaining singletons had no available analogs  
Partially aromatic families had limited availability  
Next phase requires synthesis of related analogs



SGK1 analogs distribute into 7 structural families

Remaining singletons had no available analogs  
Predominantly aromatic families had high availability  
Next phase requires expansion of nonaromatic regions



% hit overlap between targets in initial screens

Only minor activity overlap between targets (0-3 compounds/family)

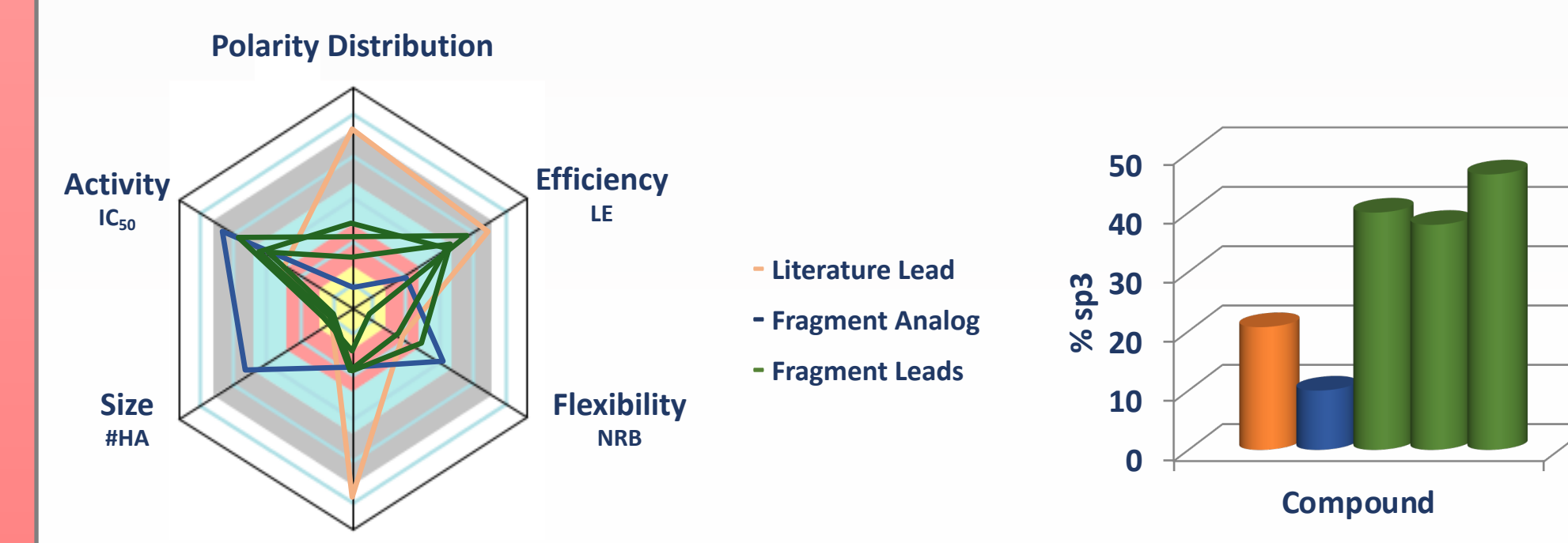
Each target had 3-4 families for next phase of fragment evolution

## NOVEL LEAD SERIES IDENTIFIED

One SGK1 analog family was progressed to medicinal chemistry. Superior drug-like properties compared to literature leads. Novel series is now in early lead optimization for oncology indications.

## ProperType Optimization Plot

A good lead hits the bull's eye



## CONTACT



BioBlocks is developing Leap-to-Lead™ for use in Lead Discovery Collaborations. While the CFL is not available for independent purchase, we welcome new collaborators. For additional information or to discuss using the CFL and the Leap-to-Lead™ platform in a drug discovery effort, please contact [wwade@bioblocks.com](mailto:wwade@bioblocks.com) or visit our website: [www.bioblocks.com](http://www.bioblocks.com)