

2019 Cresset User Group Meeting Cambridge

A Tale of Hide and Seek: Hit to Drug Candidate

June 20, 2019 Inhee Choi, RPh, PhD



We are Pasteurians fighting disease in Korea for all mankind

Institut Pasteur International Network





Institut Pasteur Korea Mission

Institut Pasteur Korea **Discovery Approach**

S. Sold States Driving convergence of BT, IT and Chemistry in the pursuit of Innovative drug discovery

Discovery Biology

Institut Pasteur Korea has been fighting infectious and neglected diseases.

- · Tuberculosis Research Lab - Hepatitis Research Lab Dengue Research Lab Cancer Biology Research Lab
- · Antibacterial Resistance Research Lab
- * Institut Pasteur Korea is studying MERS-CoV, Ebola, and Malaria as well.

- Respiratory Virus Research Lab - Leishmania Research Lab

Discovery Chemistry

Once Institut Pasteur Korea has identified promising molecules, our chemists turn them into more effective drug candidates. Lead Optimization - Early Safety Profiling Testing in Animal Models

Screening Science & Novel Assay Technologies

Institut Pasteur Korea has been building the best High Content Screening platform in both BSL-2 and BSL-3 laboratories to enable discovery biology to understand disease, and develop novel therapies.

· Technology Development Platform · Assay Development & Screening

Computational Biology

· Lab Support Operation

Sample Management & Automation

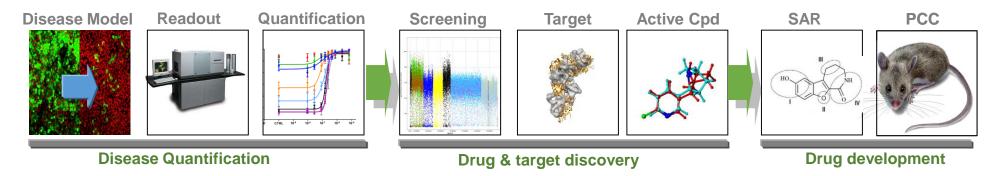
Industrial Partnerships

Institut Pasteur Korea works closel	y with industry and academia to incre	ase	
collaboration and partnerships. Wit	h these connections, our technologie	s will be	
developed into future medication a	nd therapies.		
Invention Disclosure	- Patent	 Consulting 	
- R&D Collaboration	- Intellectual Property	- Licensing	
- Knowledge and Practice	- Material Transfer Agreement		
* Please visit our website to see m	ore on our institute's inventions and te	chnologies.	

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Institut Pasteur Korea RND Infrastructure

General workflow





Outline

- I. Strategic procurement of HTS chemical library
- II. Success story of development of antitubercular drug
- **III. DFRS virtual libraries**



Outline

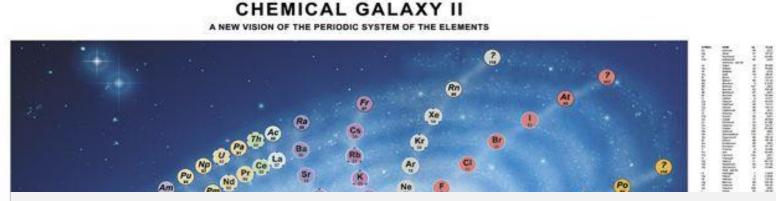
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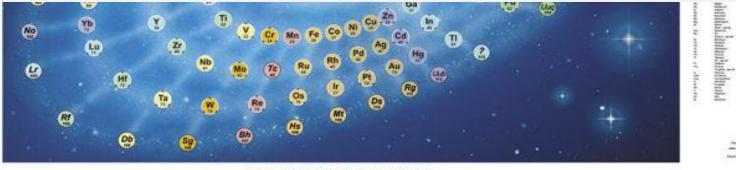


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What is "Chemical" Space?



Chemical space is the space spanned by **all possible** (i.e. energetically stable) **molecules and chemical compounds** – that is, all stoichiometric combinations of electrons and atomic nuclei, in all possible topology isomers. – Wikipedia –



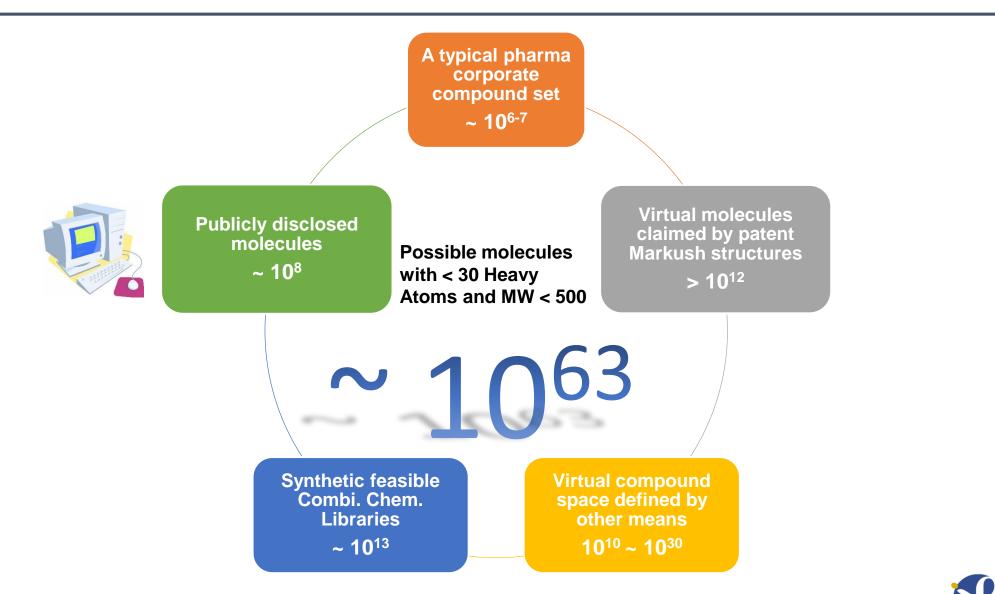


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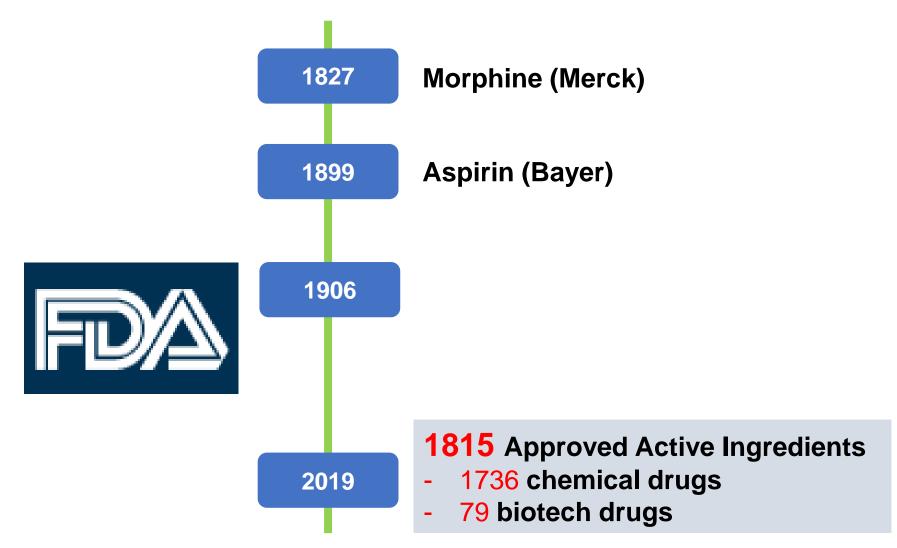
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What is the Size of Chemical Space?



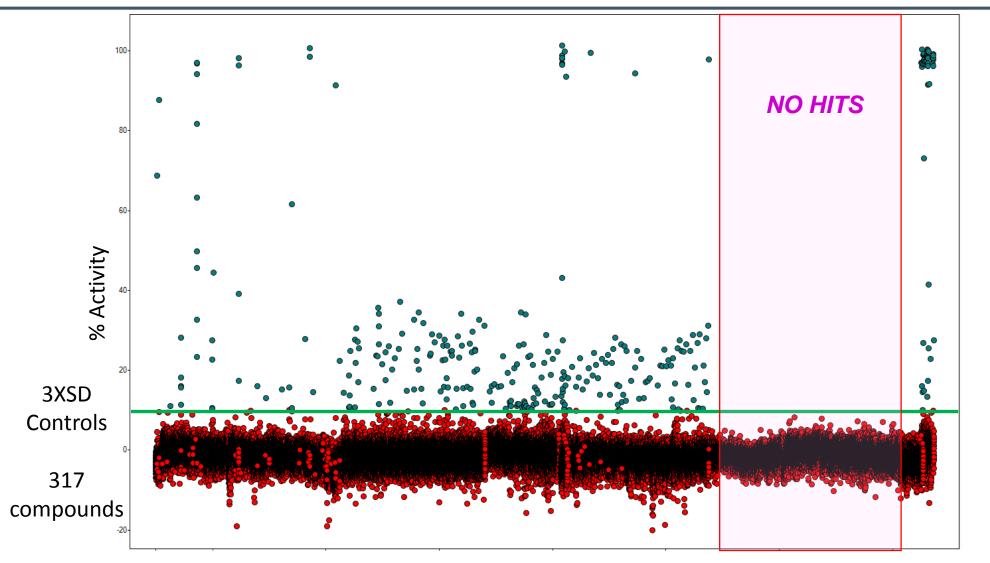
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How many FDA approved drugs today?



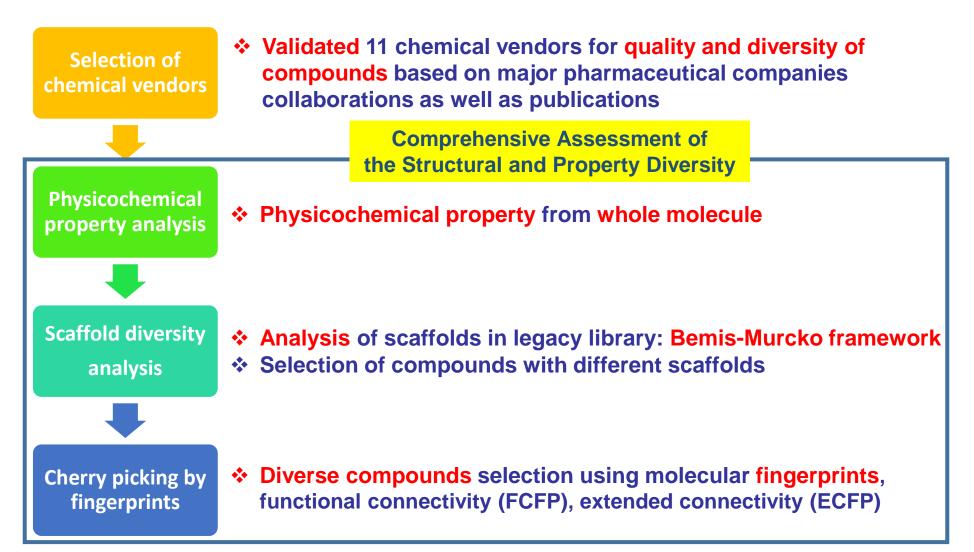


Case study: Hits from HTS





Compound Procurement Workflow



Chemical Vendors

Validated chemical vendors for quality and diversity of compounds based on major pharmaceutical companies collaborations as well as publications



ChemBridge









Maybridge







Specs





Vitas-M







Filtering Criteria 1: Drug-like Rules

In order to discriminate between drug-like compounds and non-drug compounds, a combination of Lipinski's, Veber's and Oprea's rules were applied

Lipinski Rule of 5

MW 250-500 LogP 0-5 H-bond donor ≤ 5 H-bond acceptor ≤ 10

Vebers' Rule

Polar Surface Area ≤ 140 Å No. Rotatable Bonds 0-10

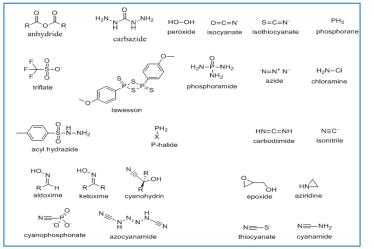
Oprea's Rules (MDDR-like)

<u>Drug-like</u> No. Rings ≥ 3 No. Rotatable bonds ≥ 6 <u>Nondrug-like</u> No. Rings ≤ 2 No. Rotatable bonds ≤ 5

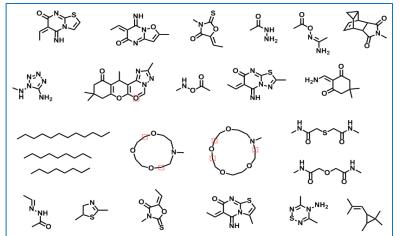


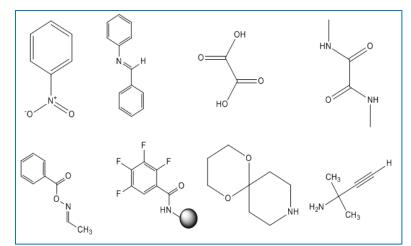
Filtering Criteria 2: Fragments

Oprea's Filter: Non-drug like fragments & toxicophore filter



Chemically not interesting fragments







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Filtering Criteria 3: Scaffold Diversity Analysis

Bemis-Murcko scaffolds (BMS) were generated and used as filter to achieve

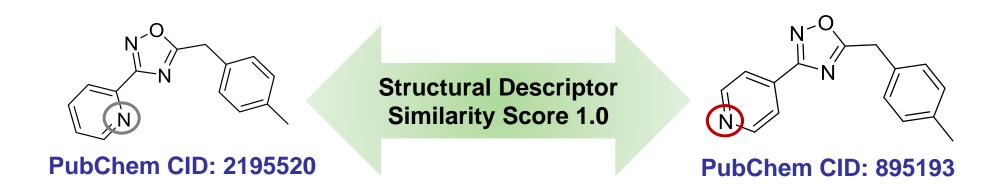
lead-likeness of a compound library Scaffold Representations 60000 NVY-AUY922 Number of Substructures 50000 HO 40000 56983 30000 HÓ HÓ ΗŃ HN 20000 Side Chains **Ring systems** Linkers 10000 12911 HO 0 300-400 400-500 HÓ MW of Substructures **Bemis Murcko** HN Framework Framework

- ✓ Compounds similar to "Lead-like" Bemis-Murcko scaffolds were selected
 - Excluding compounds with MW ≥ 400 Bemis-Murcko scaffolds



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Deception of Similarity Prediction

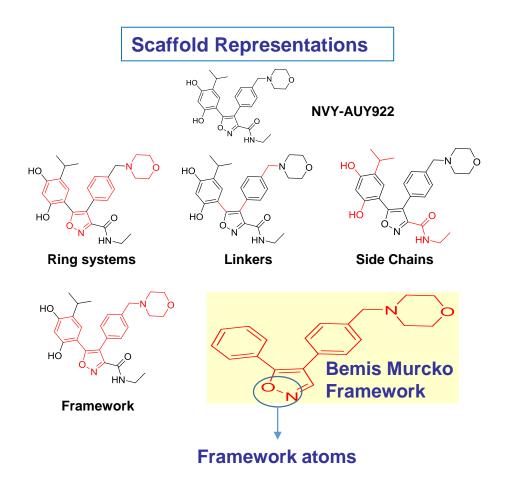


- BioAssay AID: 1321
- BioAssay Name: Primary Cell-based High Throughput Screening Assay for Inhibitors of Wee1 Degradation
- Target Protein: WEE1



Scaffold Diversity Analysis

***Bemis-Murcko scaffolds (BMS)** were generated and used as filter to achieve scaffold diversity of a compound library



- Ring systems and linkers are identified for potential use in a combinatorial-type approach to compound library generation.
- Consideration of atom properties (atom type, hybridization, and bond order) for framework atoms.
- Compound libraries may be evaluated for their relationship to the shapes of compounds. In other words, we can evaluate how well the diversity space of a library is.



Diversity: BMS Chemical Space

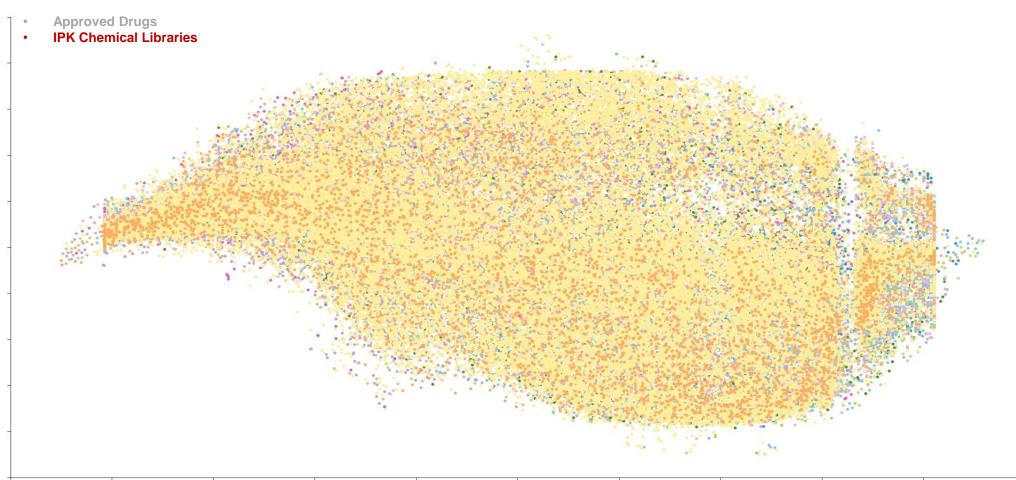
Bemis-Murcko scaffolds of approved drugs





Diversity: BMS Chemical Space

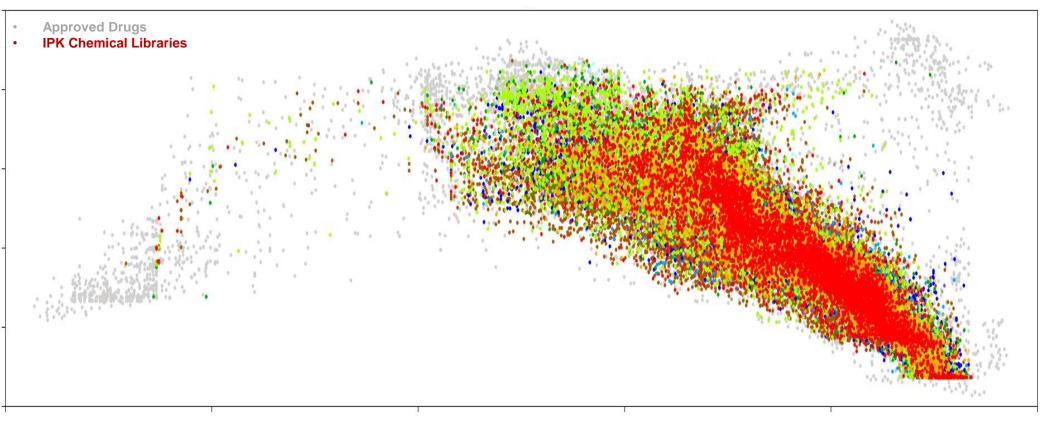
Bemis-Murcko scaffolds in IPK library are diversely distributed and covering the realm of FDA approved drugs.





Diversity: Whole Compounds' Chemical Space

Compounds in IPK library are diversely distributed and covering the realm of FDA approved drugs.





Analysis of IPK Chemical Libraries

Number of compounds in IPK collection

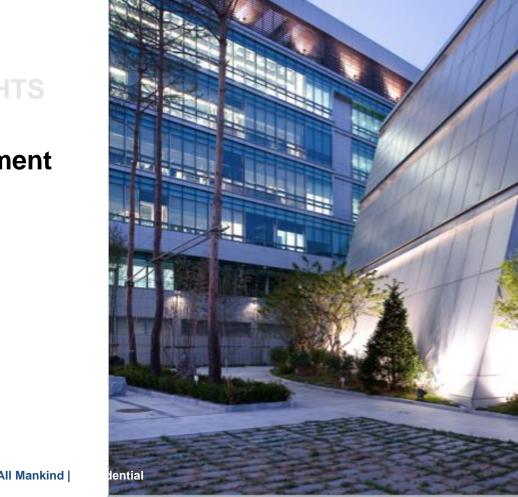
Four major chemical libraries from IPK collection

IPK Collection	Total number of compounds*	Number of Bemis-Murcko Scaffolds
IPK Legacy	~122,300	~68,600
IPK Bioactives	~7,500	~1,800
IPK Chemistry	~8,000	~3,400
IPK 2015	~100,200	~35,700
" Duplicates and salt forms have been removed	238K	110K

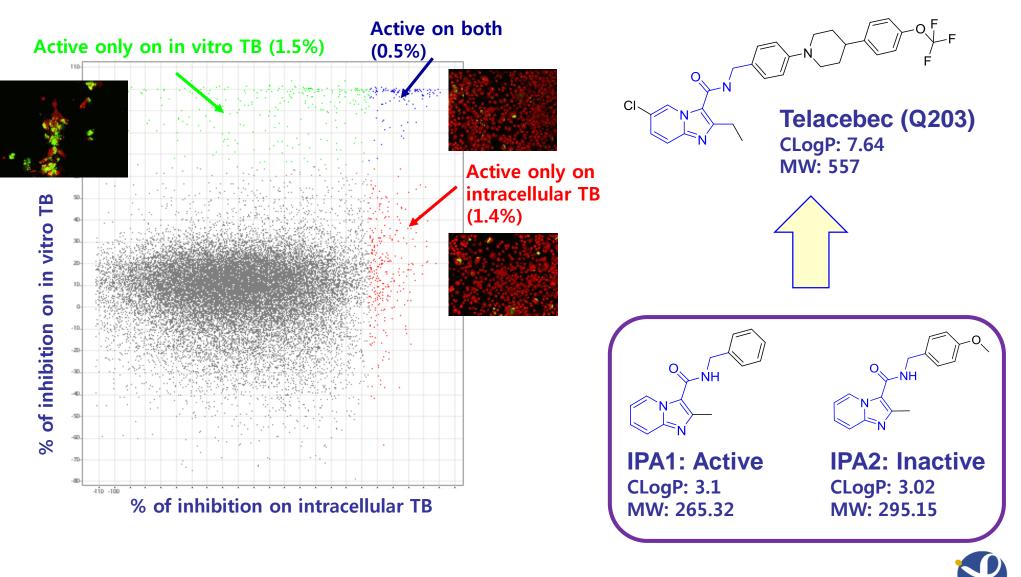


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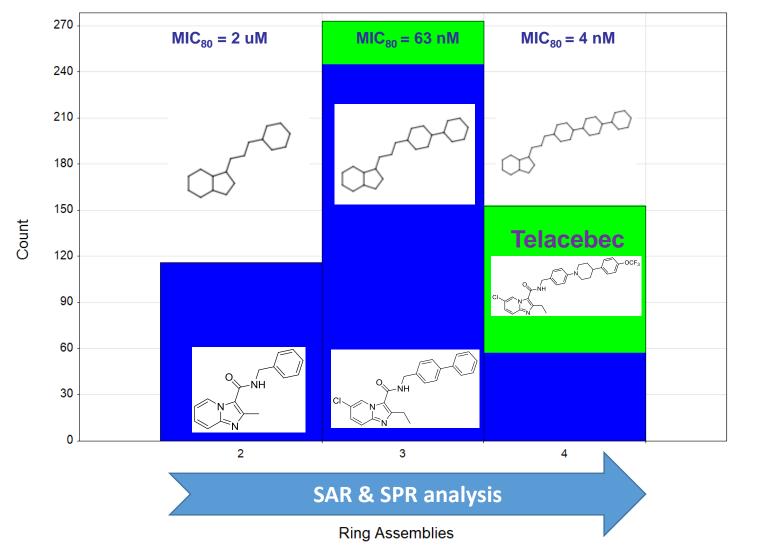
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Success @ IPK: Hit to Lead



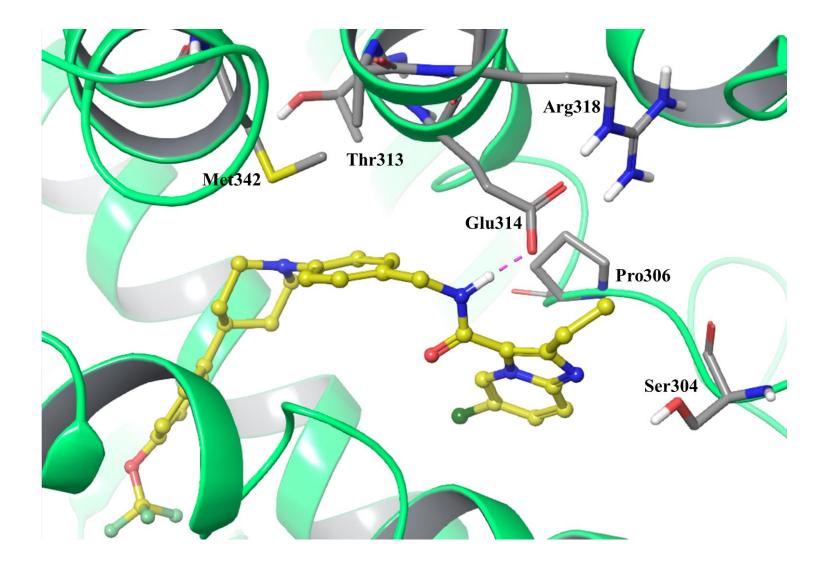
IPA series: Bemis Murcko Scaffolds



Blue: Lipinski pass Green: Lipinski fail



Target of Telacebec: QcrB

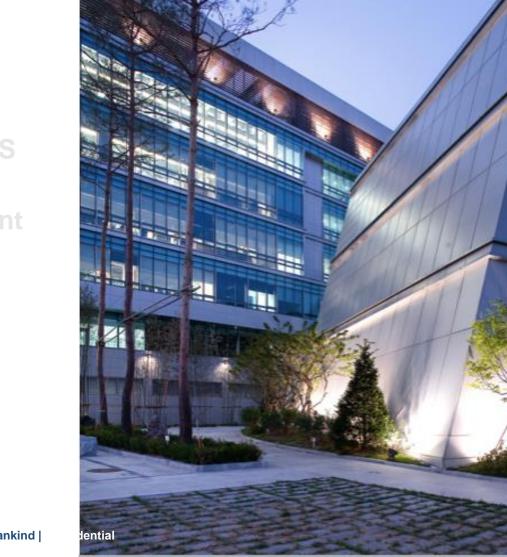




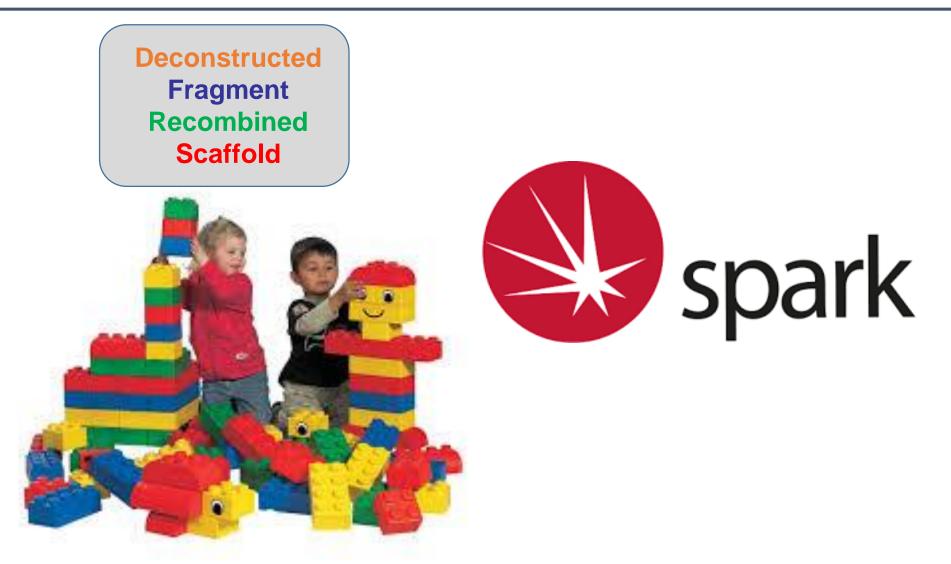
Institut Pasteur Korea - Over 10 Years in Korea Fighting Disease for All Mankind | Confidential Choi I et al., Bull. Korean. Chem. Soc., 2016

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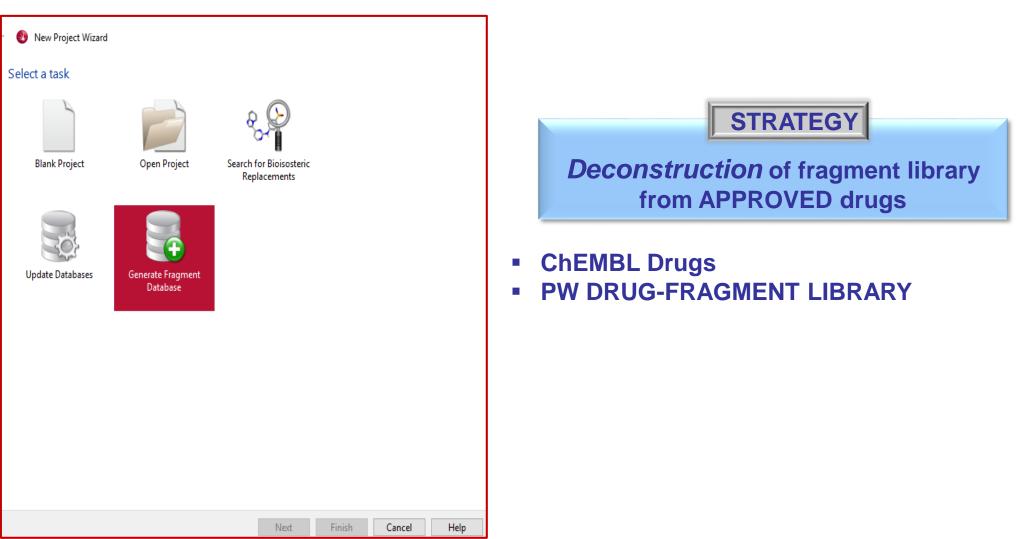


Combinatorial Virtual Chemical Libraries





1. Generate Fragment Databases





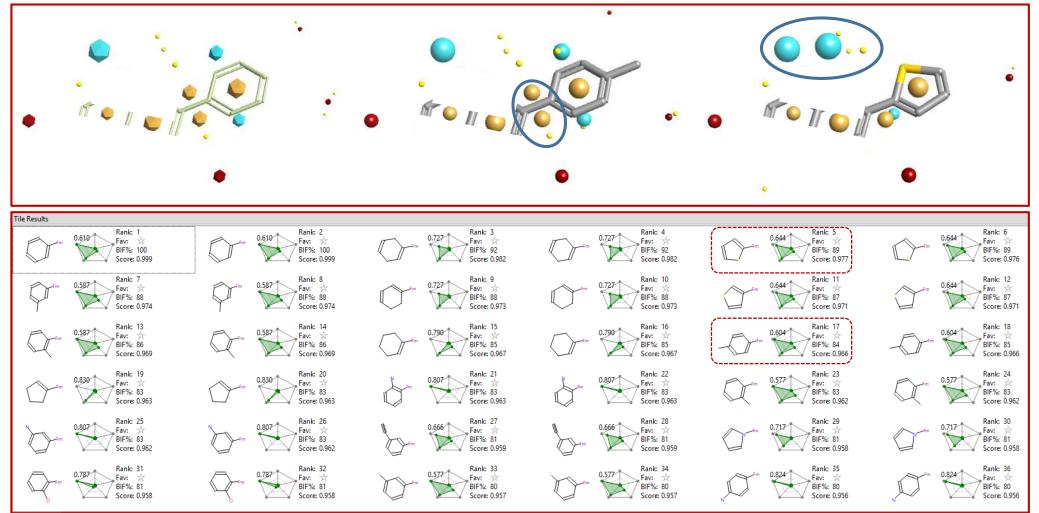
2. Selection and Edit Attachment Points

Edit allowed atom types for attachment point 1	#Atoms: 17 MV: 2222 logP: 3.295 TPSA: 28.3 Rof5: 0	Carbon All None ☑ Csp3 ☑ Csp2 ☑ Car ☑ Csp Nitrogen All ▲ Nsp3 ☑ Nsp2 Other ▲ II None ☑ Osp3 ☑ S ☑ P ☑ Hal Select All Select None Ring constraints: No Ring Constraints
	Next Finish	Cancel Help



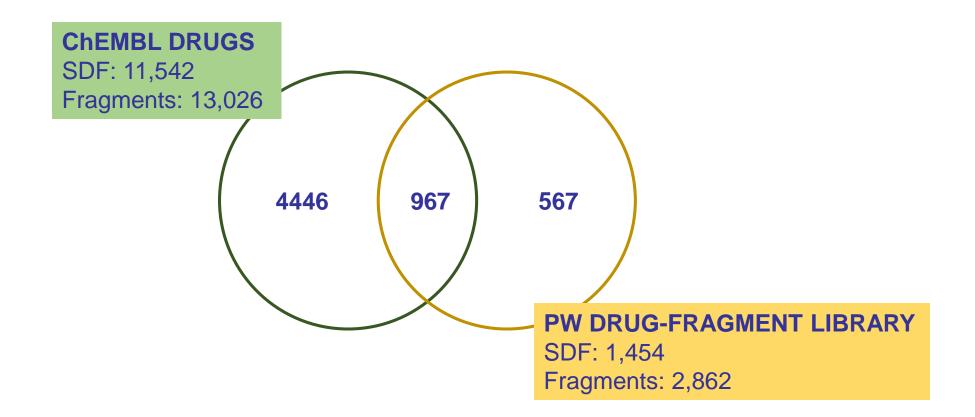
3. Search Fragments

 Results Interpretation: Field points, Bio-Isostere Factor (BIF%), Score (Field score + Shape score)



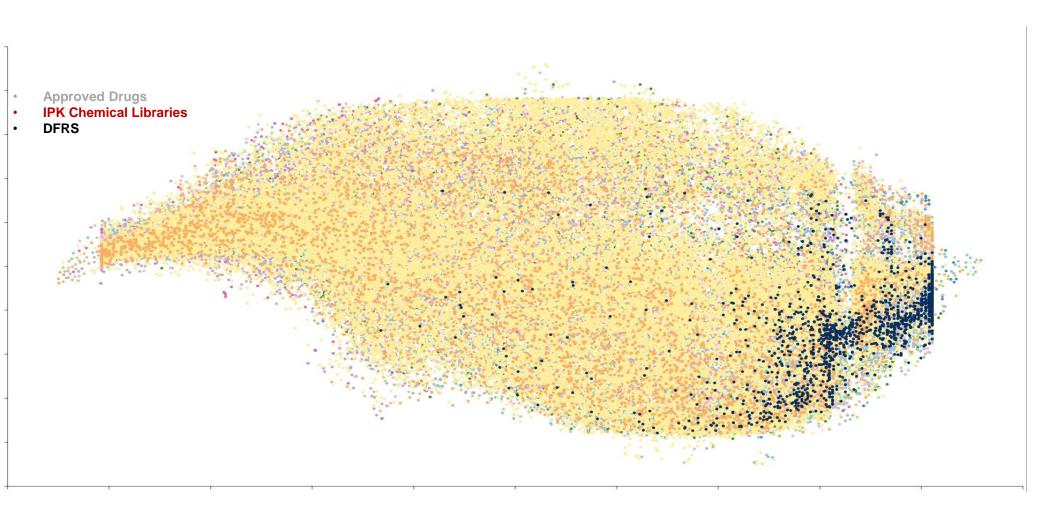
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Generated DFRS Virtual Chemical Libraries





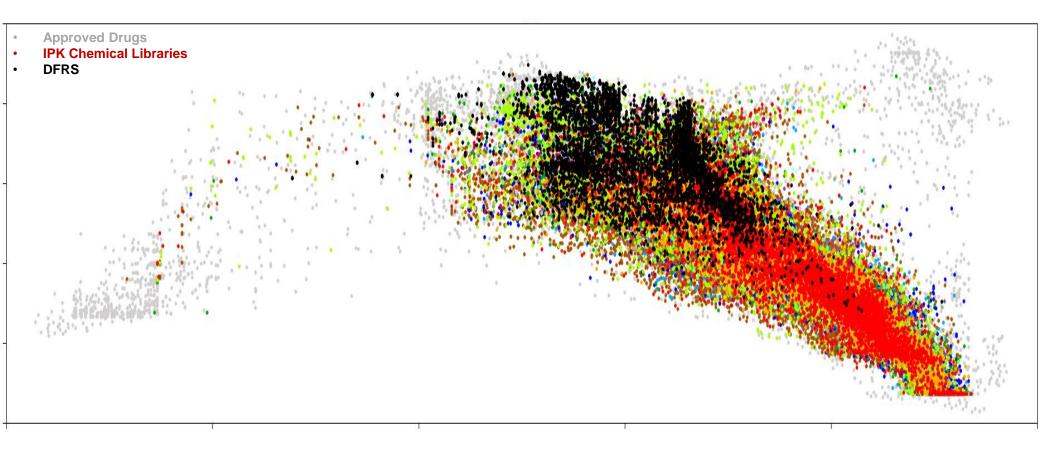
DFRS Virtual Chemical Libraries: BMS Chemical Space





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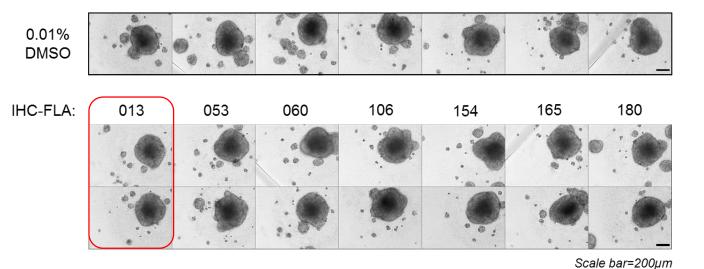
DFRS Virtual Chemical Libraries Whole Compounds' Chemical Space



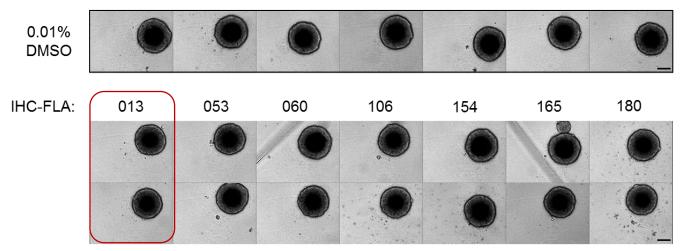


Biological activities of SPARK designed compounds: Liver cancer

Huh7.5 Spheroids



Hep3B spheroids



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Scale bar=200µm



Acknowledgement

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