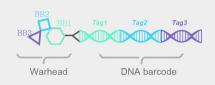


A New Method of Hit Discovery

In contrast to high throughput screening, DEL requires a very small amount of target protein and minimum assay development. More importantly, it disrupts the concept of "cost-per-well" and allows testing billions of compounds in one tube.







Sequencing & decoding the binders

Allowing access to more chemical space with lower cost

Novel Business Model

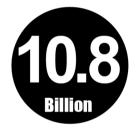
Three Types of Services Designed For Different Needs



Delopen

Free access to DEL data sharing

Target confidentiality For academic users
Open source



DElight

Unprecedented data release

Target confidentiality For all drug developers
Easy access with reduced risk



DELPIO

One- stop solution customizable

Target exclusivity For all drug developers
Access to unique scaffolds

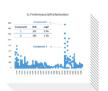
Extended Research Field

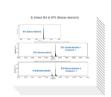
More Than Conventional DEL Screening

Irreversible Covalent Screening



Optimized selection method to enhance the **signal-to-noise** ratio

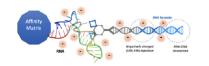




Pre-incubation Time (h)	Cmpd 1 IC ₅₀ (nM)	Cmpd 2 IC ₅₀ (nM)	Ibrutinib IC ₅₀ (nM)
0	14	2,600	49
3	0.1	50	11

RNA Target

Optimized selection procedure to counter charge, RNA/DNA interaction and structural integrity of RNA during DEL screen



Compound	Library	SPR (K _D)
RNA Hit 5 & 6	3 cycle library	Strong binding
RNA Hit 7	2 cycle library	1.7 uM
RNA Hit 8	2 cycle library	1.3 <u>uM</u>
Reference	Positive Control	144 nM

PROTAC and Molecular Glue

Optimized selection method to specifically enrich **dual-binding** molecules

On-DNA Tool compound 1 binds to POI 1 & 2 On-DNA Tool compound 2 only binds to POI 2



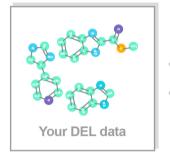


DEL-enhanced Virtual Screening









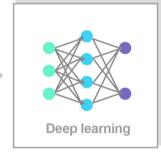




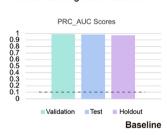
32,167



242.817

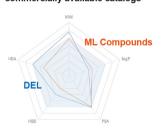


Model training and evaluation





Drug-like test compounds from commercially available catalogs







Al/MLmodel access by web desktop app

DEL Structure Based Virtual Screen Compound library DNA-encoded compound library Virtual + Commercial library Selection method Experimental Computational Computational Protein structure Compound acquisition Resynthesis Off-the-shelf or Resynthesis Screen time 4 weeks 3-4 weeks 2-3 weeks Hit rate 5-50%* 1-25%** 5-32%					
Selection method Experimental Computational Computational Protein structure Compound Resynthesis Off-the-shelf or Resynthesis Screen time 4 weeks 3-4 weeks 2-3 weeks			DEL		DEL + ML
method Experimental Computational Computational Protein structure Compound acquisition Resynthesis Off-the-shelf or Resynthesis Screen time 4 weeks 3-4 weeks Computational Computational Computational Not needed Not needed Not needed 2-3 weeks				Til taal Tooliiilooda	Til taal - commissional
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acquisition Resynthesis Resynthesis Resynthesis Screen time 4 weeks 3-4 weeks 2-3 weeks			Not needed	Required	Not needed
Colocal time - A state - Colocal time - Colocal tim			Resynthesis		
Hit rate 5-50%* 1-25%** 5-32%		Screen time	4 weeks	3-4 weeks	2-3 weeks
		Hit rate	5-50%*	1-25%**	5-32%

* In house statistics ** J. Med. Chem. 2013, 56, 17, 6560-6572

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