

SEEDSUPPLY for Drug Discovery

Supply hit compounds and target proteins

September, 2017

SEEDSUPPLY Inc.

CEO Naoki Tarui

Company Overview

Established: May 15, 2017 (Spin-out from Takeda Pharmaceutical Company)

Location: Japan (Fujisawa, Kanagawa)

Business: · Support for drug discovery

· in-house drug discovery (Joint projects with pharma/academia)

Customers: From big pharma to small biotech, worldwide

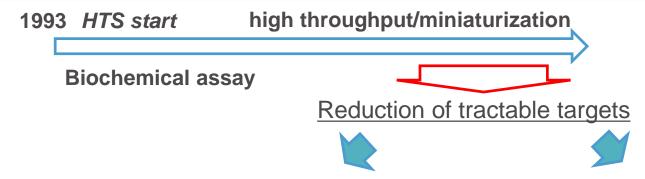
Key expertise: · Finding of binding compounds to a target

· Identification of binding proteins to a compound (by using a unique binder selection technology)

Ownership: privately owned, TAKEDA<20%

Employees: 5

New solution for drug discovery



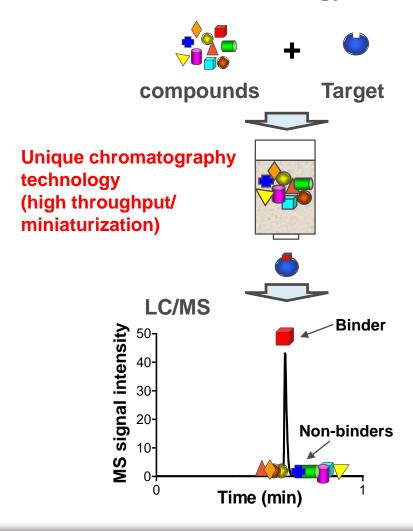
Increase of targets that cannot be handled by conventional HTS

	Conventional HTS	Our HTS
Target	Target applicable to HTS	Every target
Assay method	Each method for each target	One method
Detection	Various detectors	LCMS

Target deconvolution (Identification of proteins that bind to compounds from our protein library without labeling compounds/proteins)

Binder selection technology

Binder selection technology



Hit binder finding

Target

Soluble protein **Membrane fraction Microsomes Organelle**

DNA/RNA

X

- **Enzyme**
- **GPCR**
- Ion channel
- **Transporter**
- **Transcription** factor

Takeda library compounds

(400K)

- High capacity
- Low cost

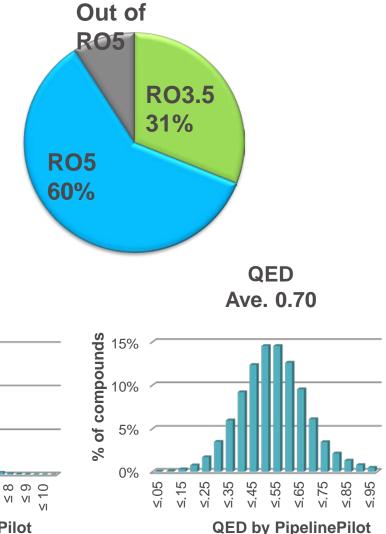
Hit binder

Inhibitor Activator Agonist Antagonist Modulator Stabilizer

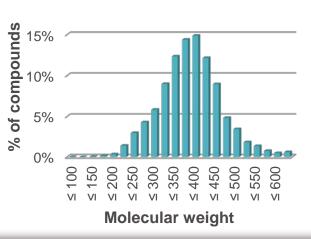
Profile of library

Number of compounds: 400K

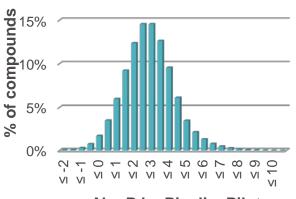
- ✓ Drug-like/lead-like compounds
- ✓ LC/MS-parameters confirmed
- √ Soluble at 30 uM in buffer
- ✓ Cell-growth inhibition tested
- √ No reactive compounds



Molecular Weight Ave. 374.7



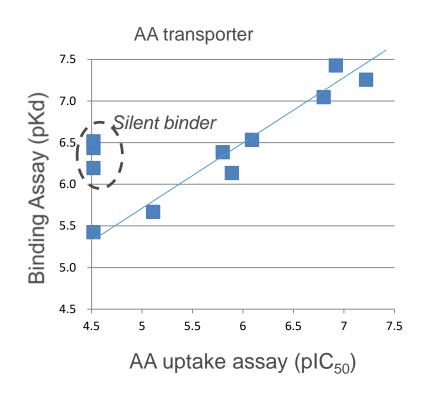
logP Ave. 2.60



AlogP by PipelinePilot

Hit binder finding

Biochemical vs. binding



Fkinase inhibitor

Hit binder	Binding Kd (nM)	IC ₅₀
SS-409	241	4.6E-06
SS-731	18	3.1E-09
SS-869	25	1.7E-09
SS-968	51	1.3E-08
SS-438	18	1.9E-09
SS-013	21	2.7E-09
SS-713	28	4.6E-09

NR agonist

Hit binder	Binding Kd (nM)	EC ₅₀
SS-931	33	7.1E-09
SS-994	34	2.2E-08
SS-798	99	1.0E-08
SS-703	453	6.9E-08
SS-285	88	4.6E-09
SS-134	20	2.9E-08
SS-539	37	3.4E-07

Target deconvolution (Binding protein finding)

