

PAUL SCHERRER INSTITUT



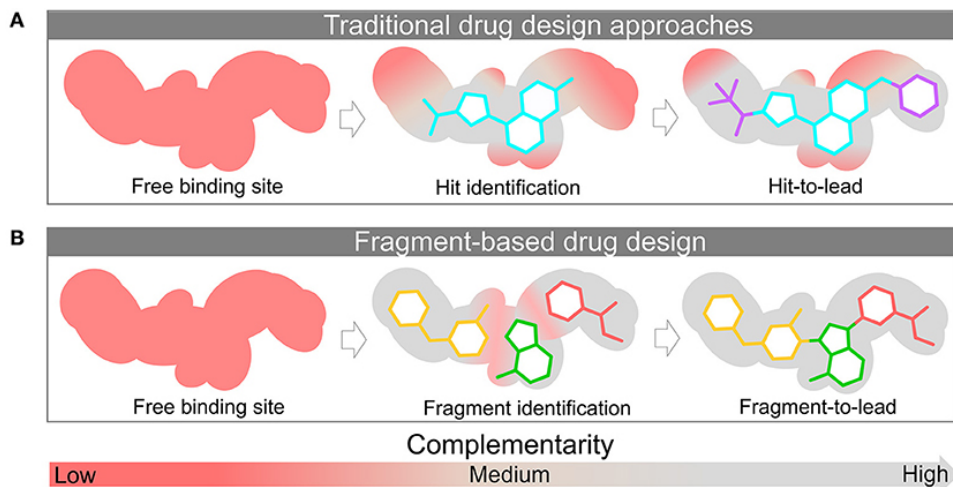
May Sharpe :: Group Leader MX Samples :: Paul Scherrer Institut

HTP Crystallographic Screening for Fragment Based Drug Discovery at the SLS

SLS MX User Meeting 2022, 1st March 2022, PSI

Fragment Based Drug Discovery (FBDD)

- FBDD is a productive alternative to traditional screening (HTS)
- Screen much smaller collections of much simpler compounds
- Allows efficient exploration of chemical space
- Identify fragments hits, then grow or link to increase potency
- Pharma partners already use FBDD
- Increasing use in academia
- Usually screen with biophysical techniques to identify hits



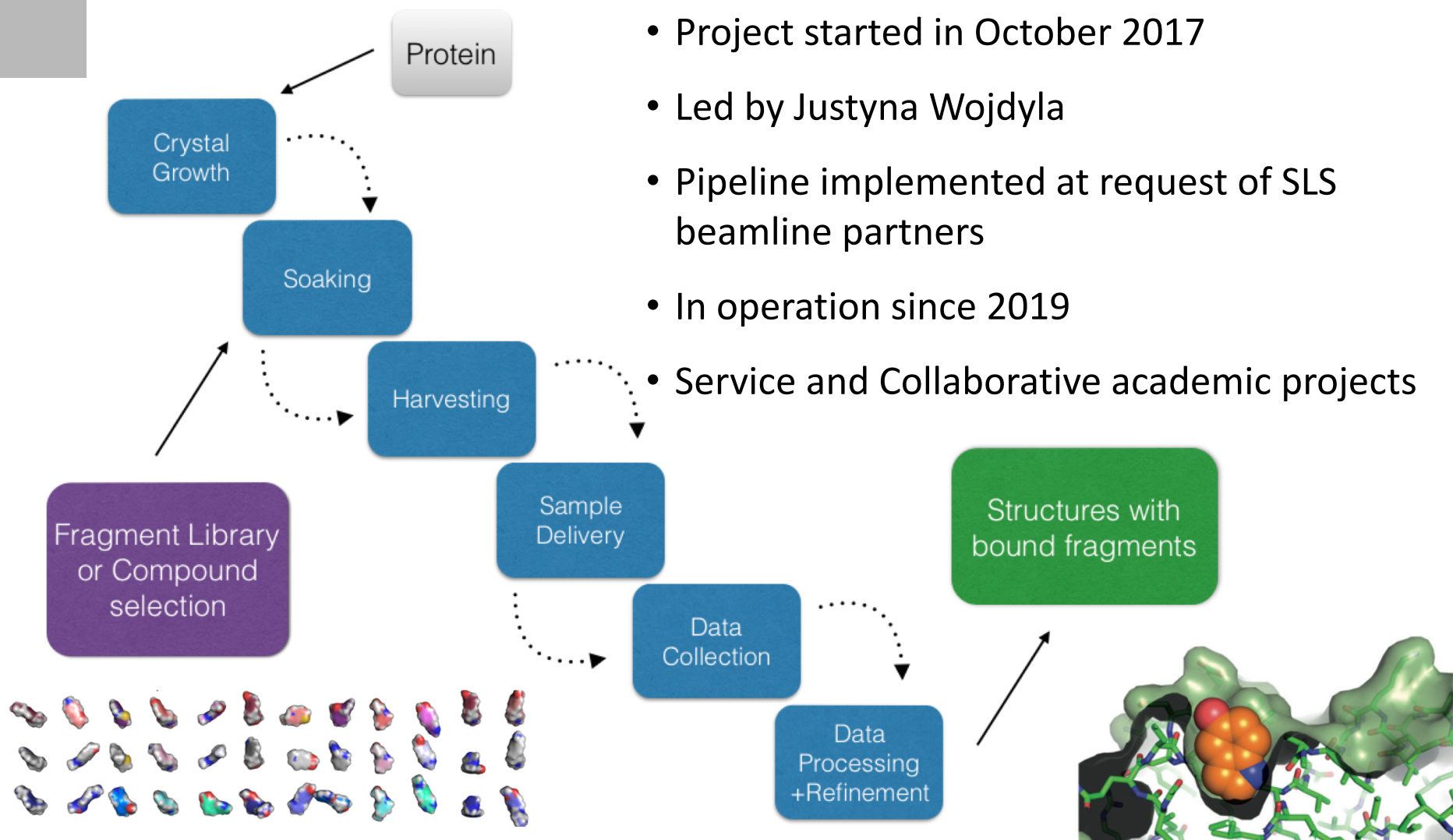
De Souza Neto *et al.*, (2020).
Frontiers in Chemistry 8

Crystallography for FBDD

- Crystallography works extremely well as a primary screen
- Crystallography essential for follow-up of hits from other techniques
- Conditions favour occupancy for weak binders
- High information content (hit, poise, feasibility of elaboration)
- Other synchrotrons have also set up facilities for this i.e Diamond (Xchem), ESRF (CrystalDirect), Max IV (FragMax) and HZB Fragment screening facility
- Technology enabled by increasing automation and speed of data collection
- **Crystallographic FBDD Pipeline implemented at the SLS**

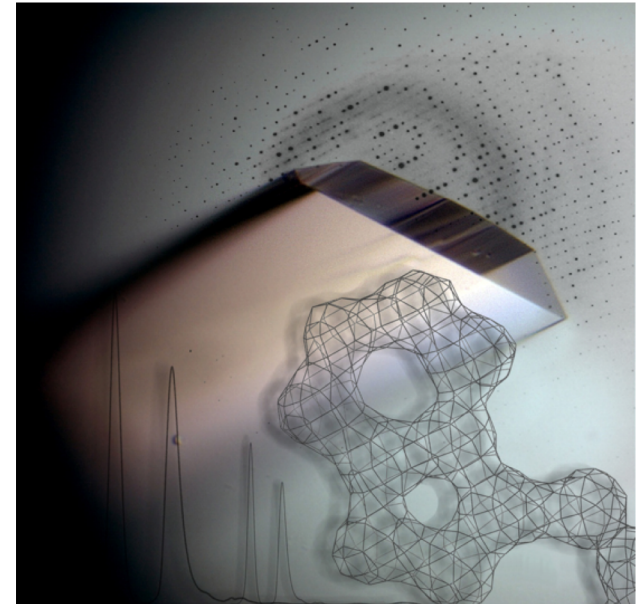


Fast Fragment and Compound Screening (FFCS) Pipeline at the SLS



- Project started in October 2017
- Led by Justyna Wojdyla
- Pipeline implemented at request of SLS beamline partners
- In operation since 2019
- Service and Collaborative academic projects

- Key component of pipeline
- Link compounds to crystals to datasets
- Automated, detailed and robust
- In-house design: FFCS Database + GUI
- Relational database as fragment campaign hub
- Generates input files and collects output files for all devices
- Organises crystal plates, cryo protection, soaking and fishing



Crystal
Growth

Soaking

Harvesting

Sample
Delivery

Data
Collection

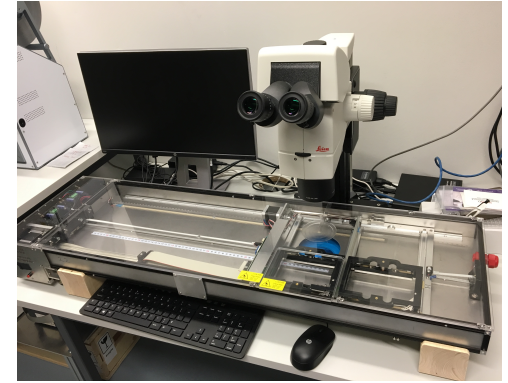
Data
Processing
+Refinement

Structures!

Fragment Logistics



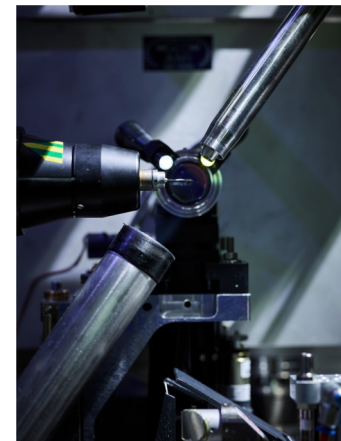
Rockimager



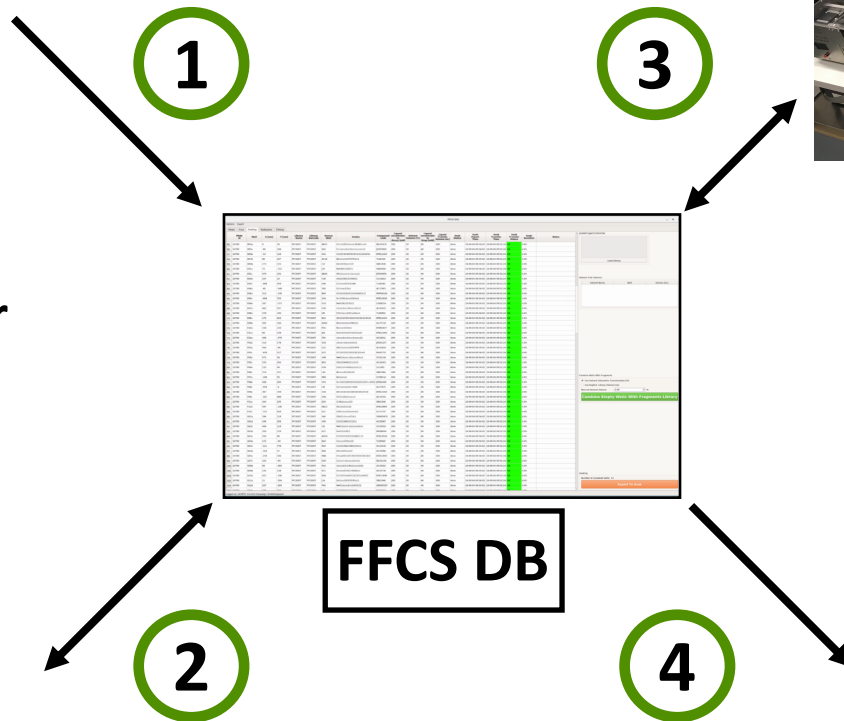
Shifter



Echo



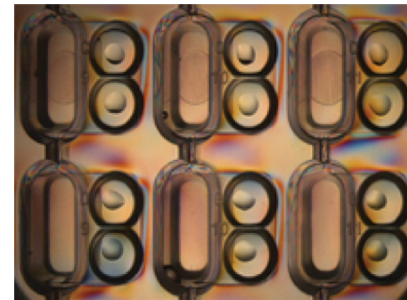
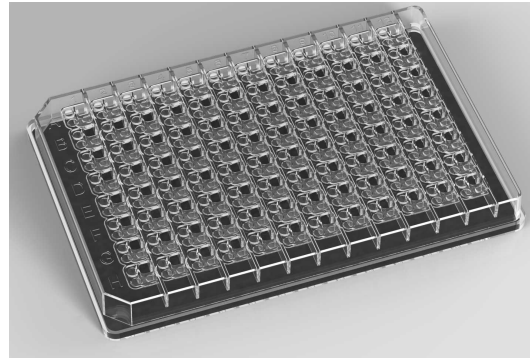
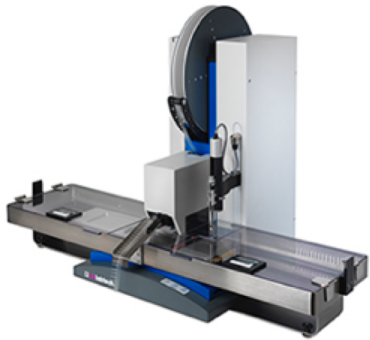
Beamline



FFCS DB

Growing and Locating Optimal Crystals

- Need large numbers of high-quality protein crystals
- Need to reproduced in 96 well plates
- Typical drop volumes 200- 400nl
- Need to identify which drops contain suitable crystals and location in drop



**Crystal
Growth**

Soaking

Harvesting

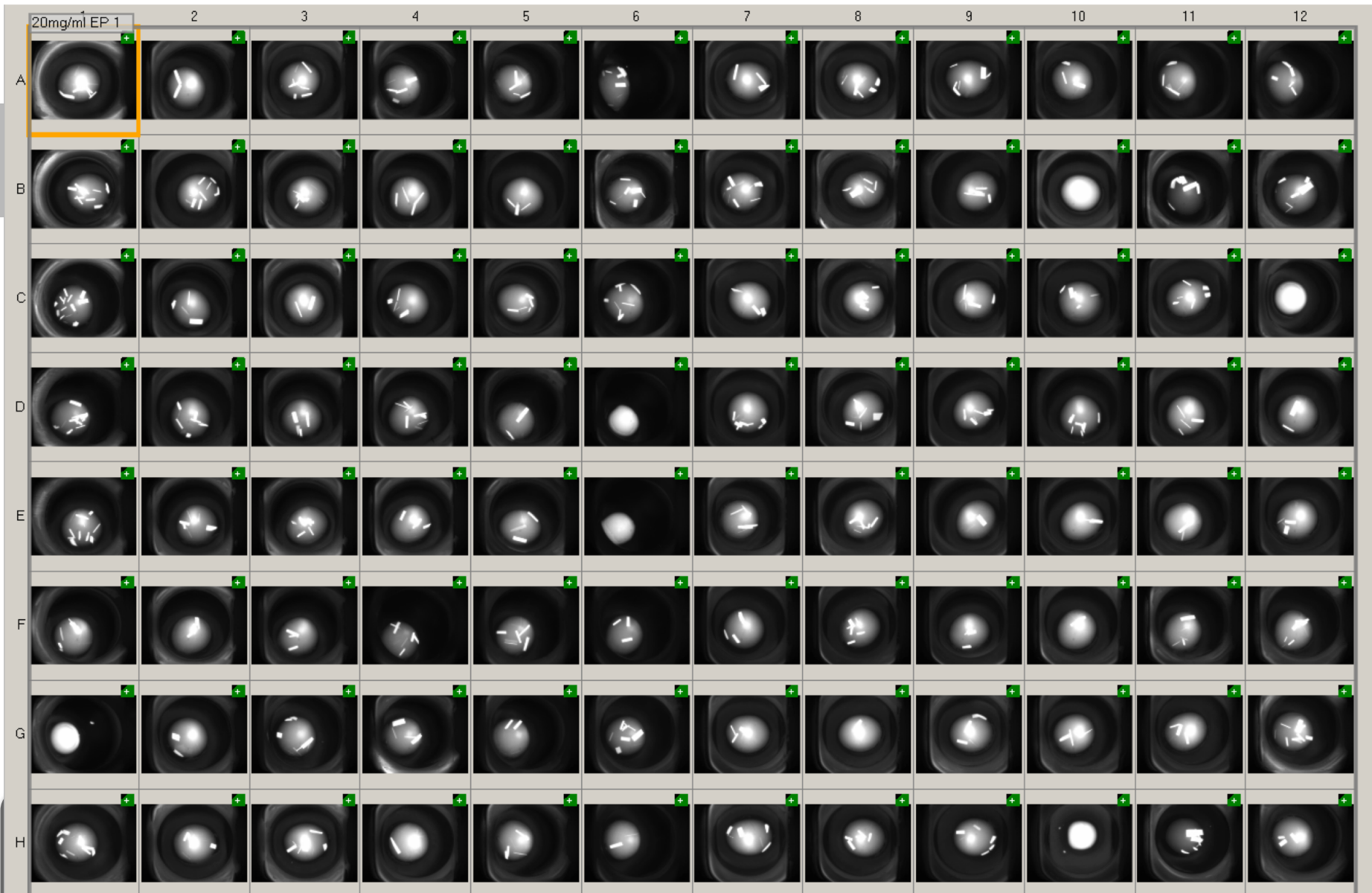
Sample
Delivery

Data
Collection

Data
Processing
+Refinement

Structures!

Crystal Identification



Projects

Endothiapepsin

Load campaign

0%

Plates

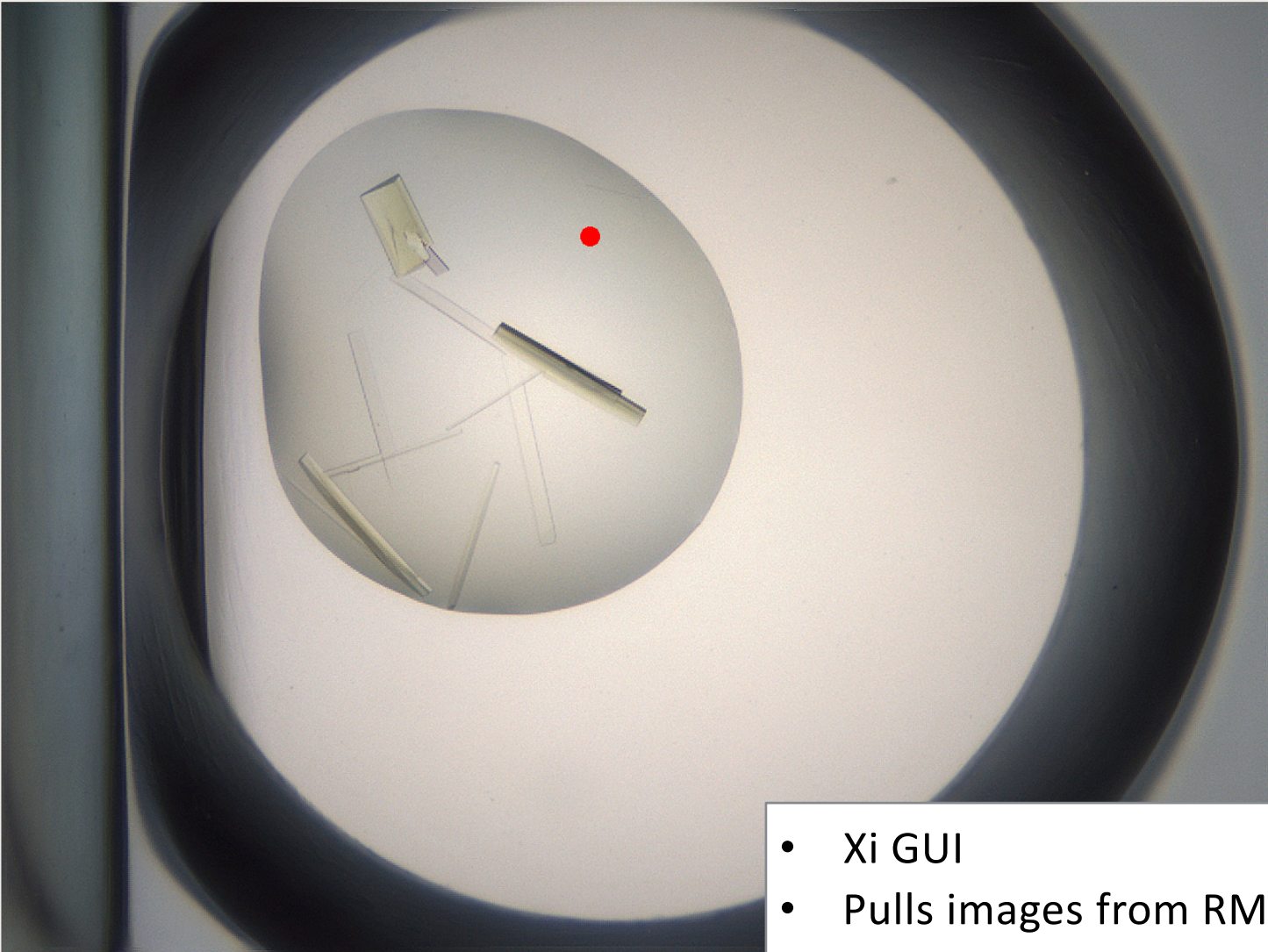
10807

Load plate

Current Plate

Campaign Id	Endothiapepsin
Plate Id	10807
Plate type	MRC3
Last imaged	2018-09-03 17:55:10
Well name	A04c
Nr of wells	288
Targeted wells	7

Create Echo Input File



• Xi GUI
• Pulls images from RM PC
• Select drops to target
• Send information to DB

Options		Expert												
Plates		Cryo	Soaking	Redissolve	Fishing									
Plate Id	Well	X [um]	Y [um]	Library Name	Library Barcode	Source Well	Smiles	Compound Code	Ligand concentration In Library [mM]	Solvent Volume [%]	Ligand concentration In Drop [mM]	Ligand Transfer Volume [nL]		
55	10780	D05a	9	26	FFCS057	FFCS057	AB33	<chem>CCc1c(O)c2ccccc2[nH]c1=O</chem>	RJC01476	200	20	40	100	
56	10780	D05c	-60	166	FFCS057	FFCS057	S43	<chem>Cl.c1cnc2c(c1)ccc1ccnc12</chem>	JFD03909	200	20	40	100	
57	10780	D06a	-14	126	FFCS057	FFCS057	K44	<chem>C1C2C3C4OC5C3C1C1C2C4C51</chem>	BTB12415	200	20	40	100	
58	10780	D07a	86	267	FFCS057	FFCS057	AC18	<chem>NCC1ccc(C(F)(F)F)cc1</chem>	TL00150	200	20	40	100	
59	10780	D09a	273	215	FFCS057	FFCS057	I14	<chem>Oc1c(Cl)cccc1Cl</chem>	S801838	200	20	40	100	
60	10780	E01a	75	-152	FFCS057	FFCS057	J35	<chem>N#CN1CCOCC1</chem>	S800580	200	20	40	100	
61	10780	E01c	479	232	FFCS057	FFCS057	AB26	<chem>OCc1ccccc1-clccccc1</chem>	JFD03959	200	20	40	100	
62	10780	E03a	247	23	FFCS057	FFCS057	H35	<chem>CN1CCN(C)C(CN)C1</chem>	CC13813	200	20	40	100	
63	10780	E03c	-408	459	FFCS057	FFCS057	H46	<chem>Cc1ccccc1Ck1C#N</chem>	TL00285	200	20	40	100	
64	10780	E04a	-40	-548	FFCS057	FFCS057	Y08	<chem>Clc1nc(Cl)n1</chem>	AC37803	200	20	40	100	

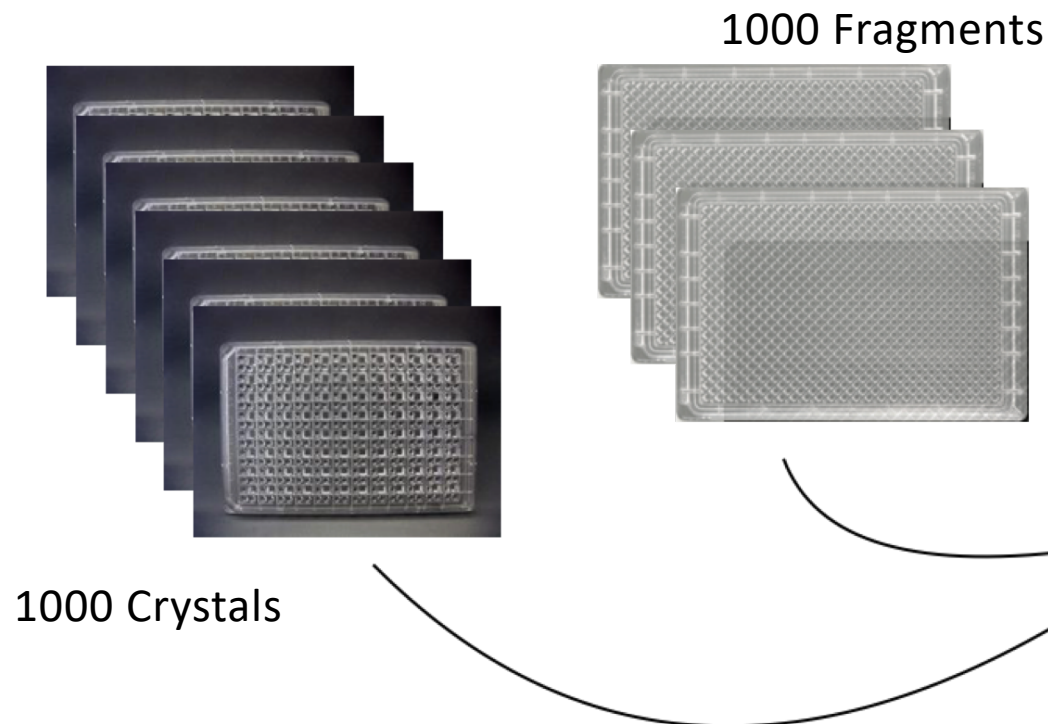
- Soaking coordinates transferred to DB
- Fragment Library loaded and assigned to drops
- Input file for Echo created and transferred to Echo PC
- Echo run results transferred to DB

Collins *et al.*, (2017). *Acta Cryst.* D73, 246-255



Crystal Soaking

- Fragments transferred to the crystal containing drops using a Labcyte Echo 550 acoustic liquid handler
- Coordinates transferred from FFCS Database



Crystal
Growth

Soaking

Harvesting

Sample
Delivery

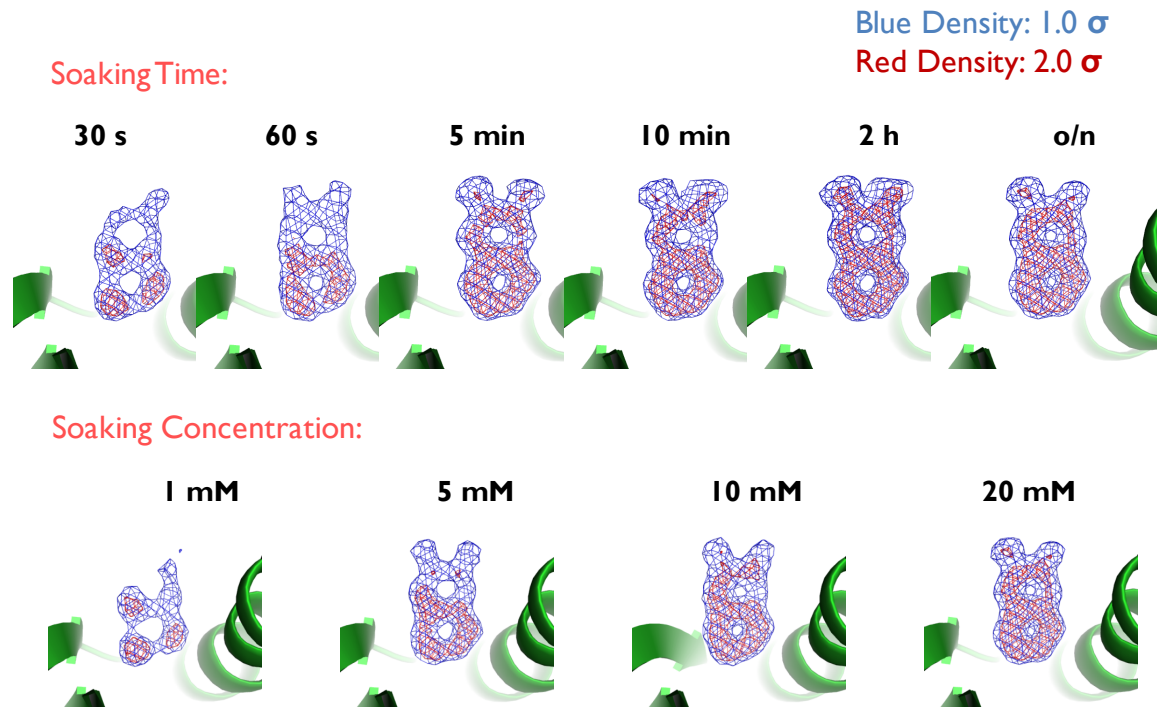
Data
Collection

Data
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+Refinement

Structures!

Crystal Soaking

- Soaking parameters first need to be optimized
- Maximum soaking time (1– 24 hours) and maximum DMSO soaking concentration (5 -30%) determined



Crystal
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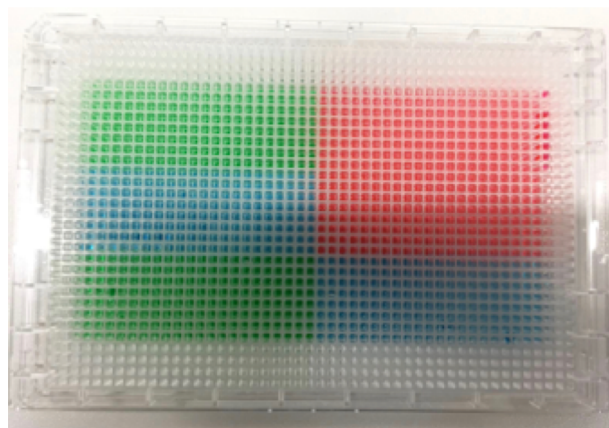
Data
Collection

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Structures!

Fragment and Compound Library Storage

- Stored in hermetically sealed dry environment (N₂ purged)
- Heat sealed plates
- Limit number of times opened
- Extensive advice from Idorsia Compound Library and Screening
- Idorsia 1056 library and Maybridge 2500 library
- Utilize any library stored in Echo compatible plates



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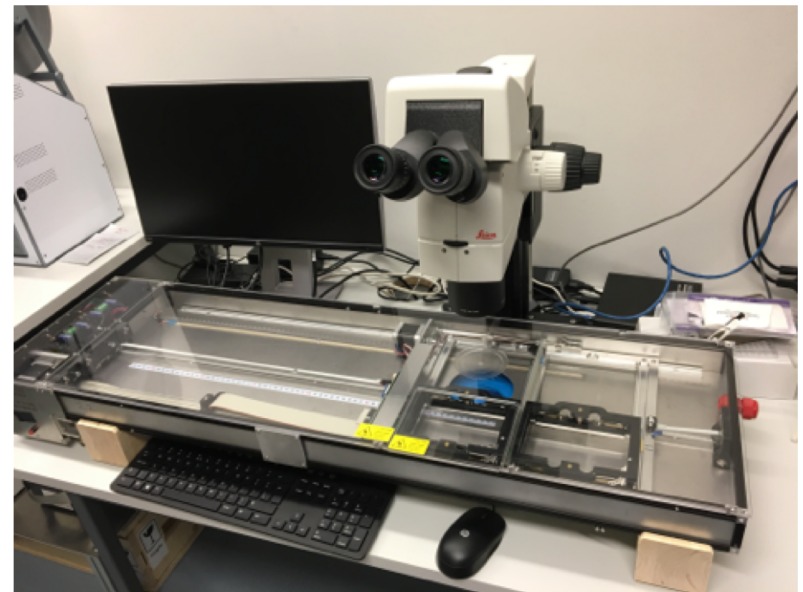
Data
Collection

Data
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Structures!

- Utilize Shifter robot (Oxford Lab Technology)
- Manually assisted harvesting
- Shifter input file generated by FFCS DB
- Automates all repetitive steps
- Can freeze 100-150 crystals per hour
- Barcoded pucks

Developed at Diamond Light Source



Wright *et al.*, (2021). Acta Cryst. D77, 62-74

Crystal
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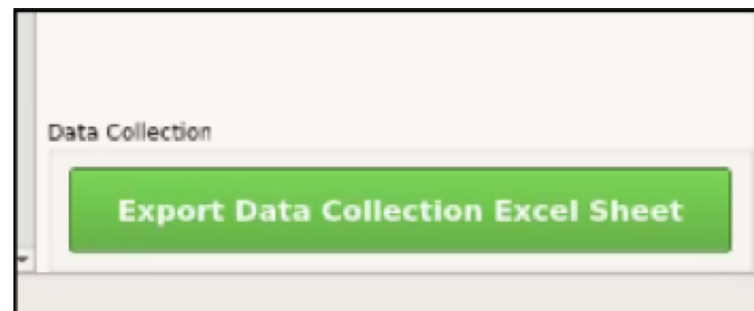
Data
Collection

Data
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Structures!

Options Expert

Plates	Cryo	Soaking	Redissolve	Fishing					
Plate Id	Well	Fishing Result	Fishing Time	Fishing Duration	Puck Barcode	Puck Position			
49	10780	C12a	OK: Mounted Clear	18-08-30 15:45:59	00:01:20	CFSL5010	3		-C
50	10780	D01a	OK: Mounted Clear	18-08-30 15:47:27	00:00:32	CFSL5010	4		-C
51	10780	D01c	OK: Mounted Clear	18-08-30 13:39:15	00:00:47	CFSL5009	5		-C
52	10780	D02a	OK: Mounted Clear	18-08-30 13:40:04	00:00:27	CFSL5009	6		-C
53	10780	D02c	OK: Mounted Clear	18-08-30 15:48:02	00:00:40	CFSL5010	5		-C
54	10780	D03c	OK: Mounted Clear	18-08-30 15:48:44	00:01:00	CFSL5010	6		-C
55	10780	D05a	OK: Mounted Clear	18-09-04 13:01:24	00:00:37	CFSL5003	1		-C
56	10780	D05c	OK: Mounted Clear	18-09-04 13:02:04	00:00:41	CFSL5003	2		-C
57	10780	D06a	OK: Mounted Clear	18-09-04 13:02:47	00:00:31	CFSL5003	3		-C
58	10780	D07a	OK: Mounted Clear	18-09-04 13:03:19	00:00:19	CFSL5003	4		-C
59	10780	D09a	OK: Mounted Clear	18-09-04 13:03:41	00:00:19	CFSL5003	5		-C
60	10780	E01a	OK: Mounted Clear	18-09-04 13:04:02	00:01:21	CFSL5003	6		-C
61	10780	E01c	OK: Mounted Clear	18-09-04 13:05:24	00:00:29	CFSL5003	7		-C



Crystal Growth

Soaking

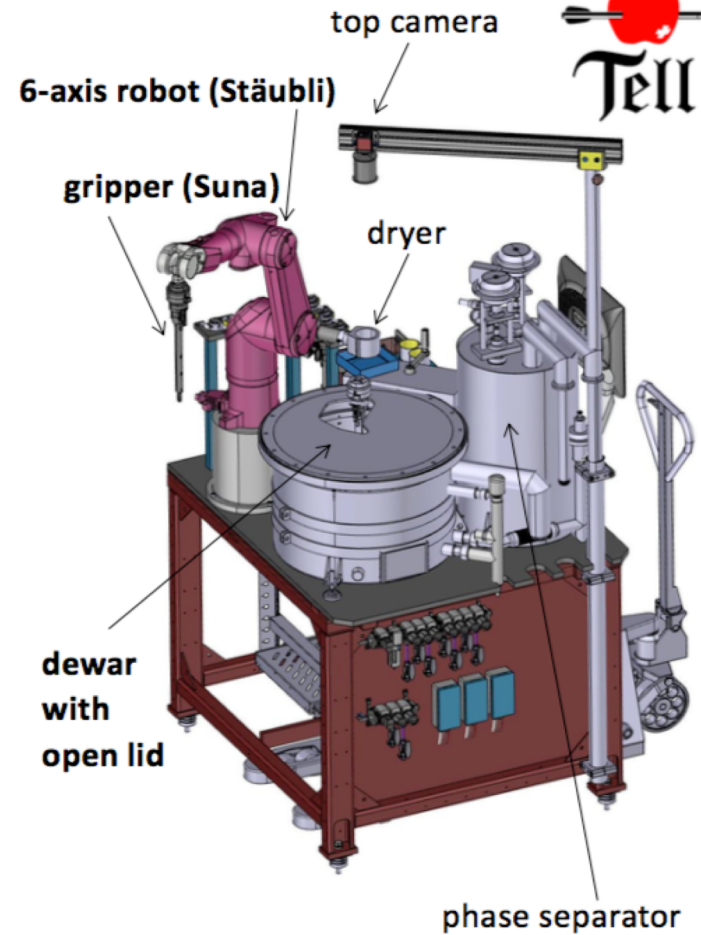
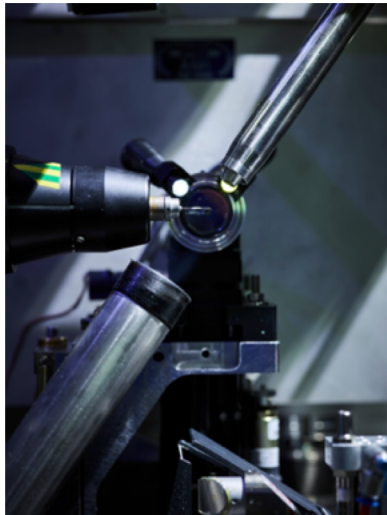
Harvesting

Sample Delivery**Data Collection**Data Processing
+Refinement

Structures!



- Crystals delivered to beam using in house designed sample changer robot
- 20-30 samples / hour
- Unattended data collection



Martiel *et al.*, (2020). *J. Sync. Rad.* 27, 860-863

Crystal
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Collection

Data
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+Refinement

Structures!

Options		Expert	
Plates	Cryo	Soaking	Redissolve Fishing
Plate Id		Well	
49	10780	C12a	
50	10780	D01a	

- DMSO can block binding sites and damage crystals
- The pipeline is based around DMSO stocks of compounds
- Can we dispense from DMSO stocks and then get rid of the DMSO?
- Dry fragment onto crystallisation plate, re-suspend in crystal soaking solution
- Transfer crystal to plate → soak → harvest
- Echo used to transfer both fragments and soaking solution
- Shifter expedites transfer of crystals to soaking plate

Bedi *et al.*, (2020). ACS Chem Biol. 15, 618-625

ACS
chemical
biology

pubs.acs.org/acschemicalbiology

Letters

Selectively Disrupting m⁶A-Dependent Protein–RNA Interactions with Fragments

Rajiv Kumar Bedi, Danzhi Huang, Lars Wiedmer, Yaozong Li, Aymeric Dolbois, Justyna Aleksandra Wojdyla, May Elizabeth Sharpe, Amedeo Caffisch,* and Pawel Sledz*

 Cite This: ACS Chem. Biol. 2020, 15, 618–625

 Read Online

Crystal
Growth

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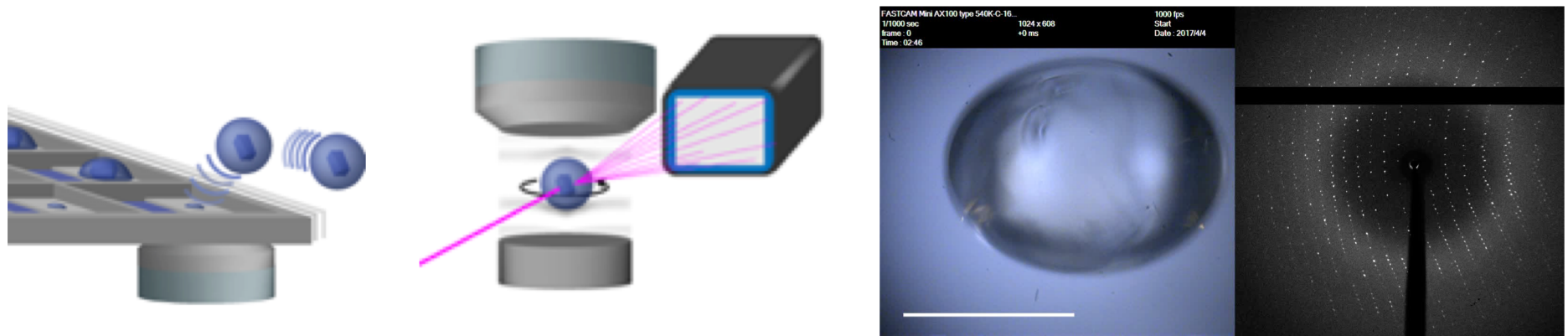
Data
Collection

Data
Processing
+Refinement

Structures!

Acoustic Droplet Levitation for Sample Delivery

- Crystal handling and freezing takes a lot of time and manual effort.
- Medicinal chemistry would benefit from RT data (e.g. RT water structure).
- Crystal delivery by acoustic droplet levitation would transform crystal handling.
- Would enable rapid, automated collection of RT data for fragment screening.



S. Tsujino & T. Tomizaki, Scientific Reports 6, 25558 (2016).

Crystal
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Harvesting

Sample
Delivery

Data
Collection

Data
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+Refinement

Structures!

- PSI
 - Justyna Wojdyla
 - Kuba Kaminski
 - Laura Vera
 - Chia-Ying Huang
 - Deniz Eris
 - Meitian Wang
 - Nathalie Meier
 - Dominik Buntschu
 - Roman Schneider
 - Sonia Reber
- Idorsia
 - Oliver Peter
 - Aengus Mac Sweeney
 - Geoffroy Bourquin

Sharpe *et al*, Nippon Kessho Gakkai-Shi (2021) **63**(3) 232
Kaminski *et al.*, (2022). Acta Cryst. **D78**, 328-336

