

Harnessing Robotic Microscopes, Al, And Machine Learning To Increase The Speed Of Drug Discovery

August Allen | Automation Scientist

Ben Miller | Director High Throughput Screening

September 26, 2017 | Washington, DC

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Overview

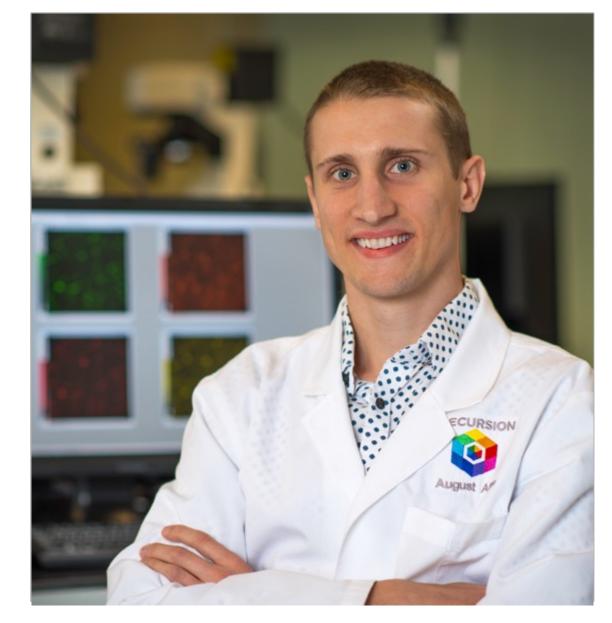
What we'll be talking about

- About Recursion
 - Mission: Accelerate Drug Discovery
 - Process: Robotics, Deep Learning, AI
- Challenges of rapid growth
 - Scaling and improving existing processes
 - Implementing new procedures and automation
- ► How Splunk helps
 - ETL
 - Manufacturing Metrics Tracking



August Allen

- Work Experience
 - Joined Recursion in June 2016
 - Johnson and Johnson Co-op 2015
 - Bristol Myers-Squibb Co-op 2013
- Splunk
 - Dec 2016-Present
- Education
 - Rochester Institute of Technology
 - B.S. in Biomedical Engineering
- Recreation
 - Soccer, snowboarding, biking
 - Co-host podcast on space exploration
- https://www.linkedin.com/in/august-allen-a8239841





Ben Miller

- ► Work:
 - Recursion: Director of High Throughput Science
 - Myriad Genetics
 - Velocity11
 - GlycoFi
- ► Splunk:
 - Dec 2014-Present
 - 2016 Innovation award
- School:
 - Dartmouth College: MS Genetic Engineering
 - BA Engineering
- Fun:
 - Biking, Skiing, Hiking, Climbing
- in linkedin.com/in/bentonjmiller





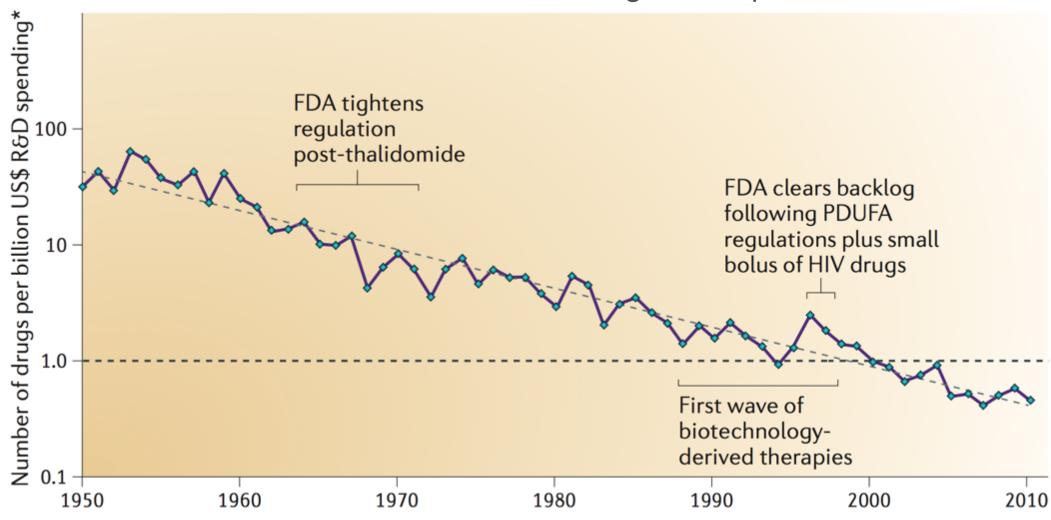
Recursion Pharmaceuticals

Our Mission and Process



Our Mission

Reverse Eroom's Law of Drug Development



Scannell, et al. Diagnosing the decline in pharmaceutical R&D efficiency. Nature Reviews Drug Discovery, 2012



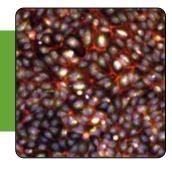
Recursion Technology

Turning cell biology into a facial recognition problem



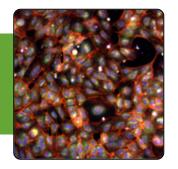
Healthy child

Healthy cells





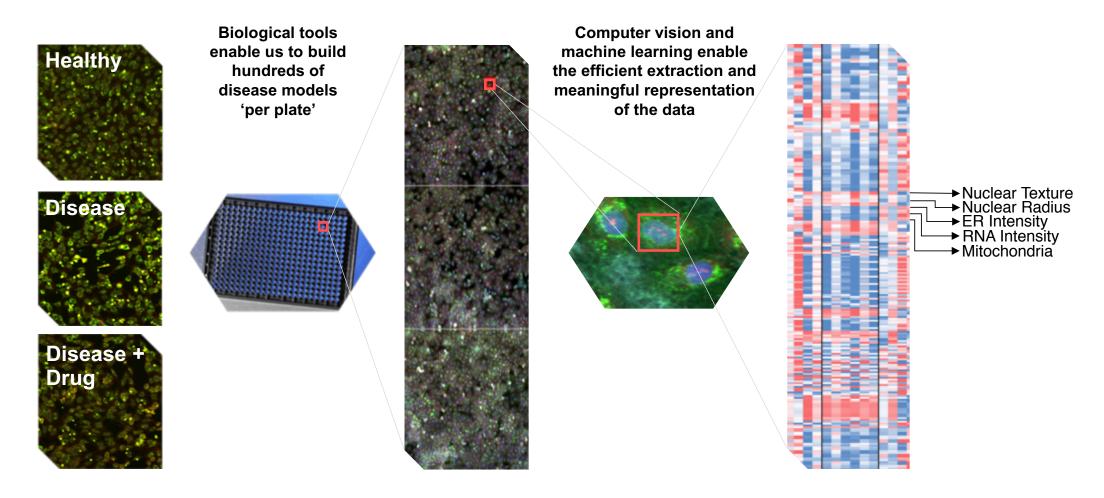
Child with rare genetic disease (Cornelia de Lange Syndrome) Genetic disease model cells (Cornelia de Lange Syndrome)





Artificial Intelligence In Drug Discovery

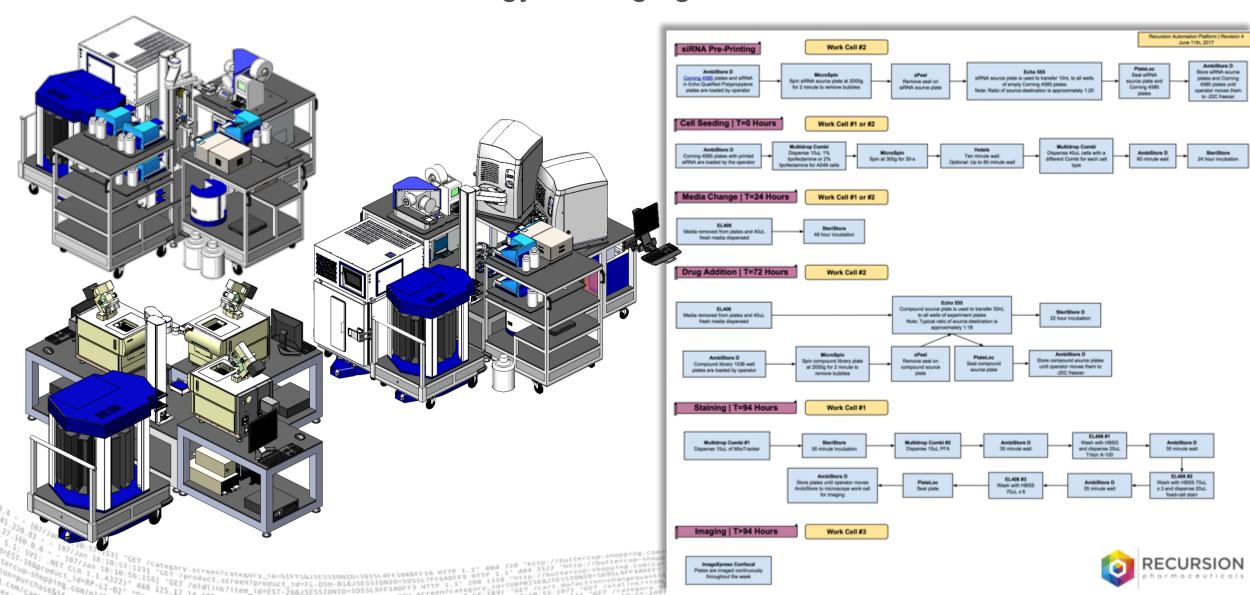
Automated high-dimensional feature extraction

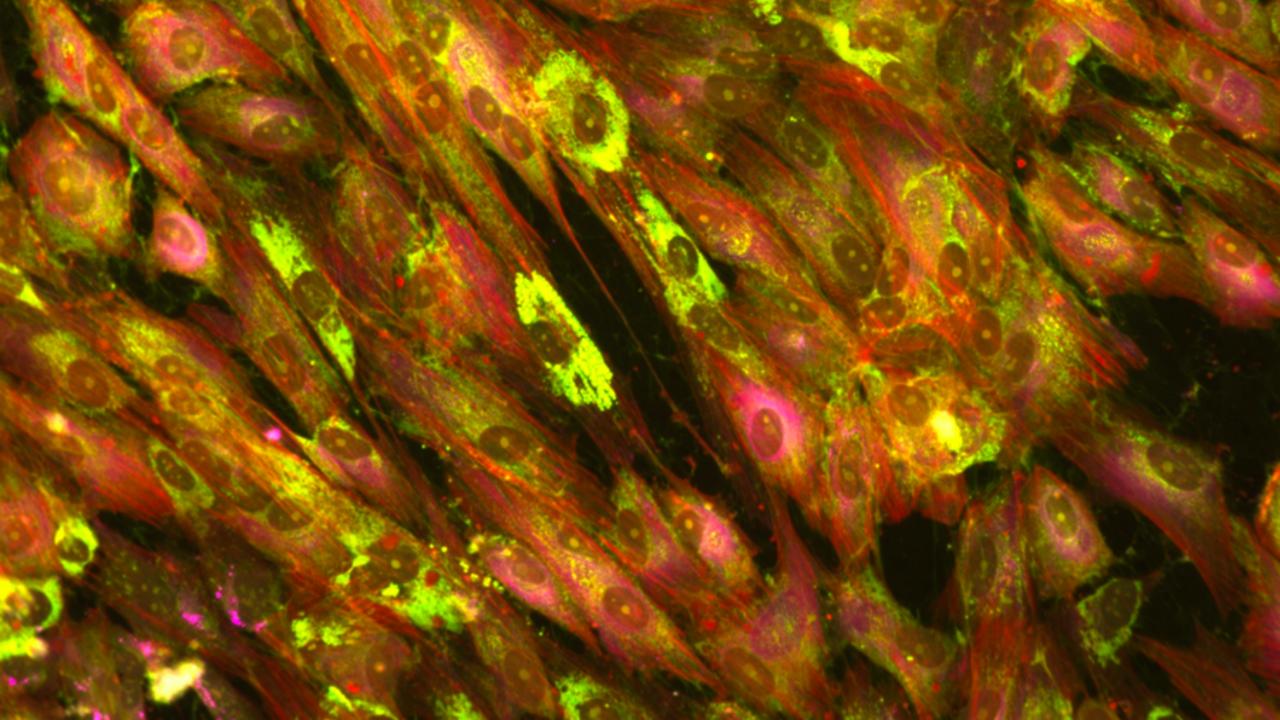


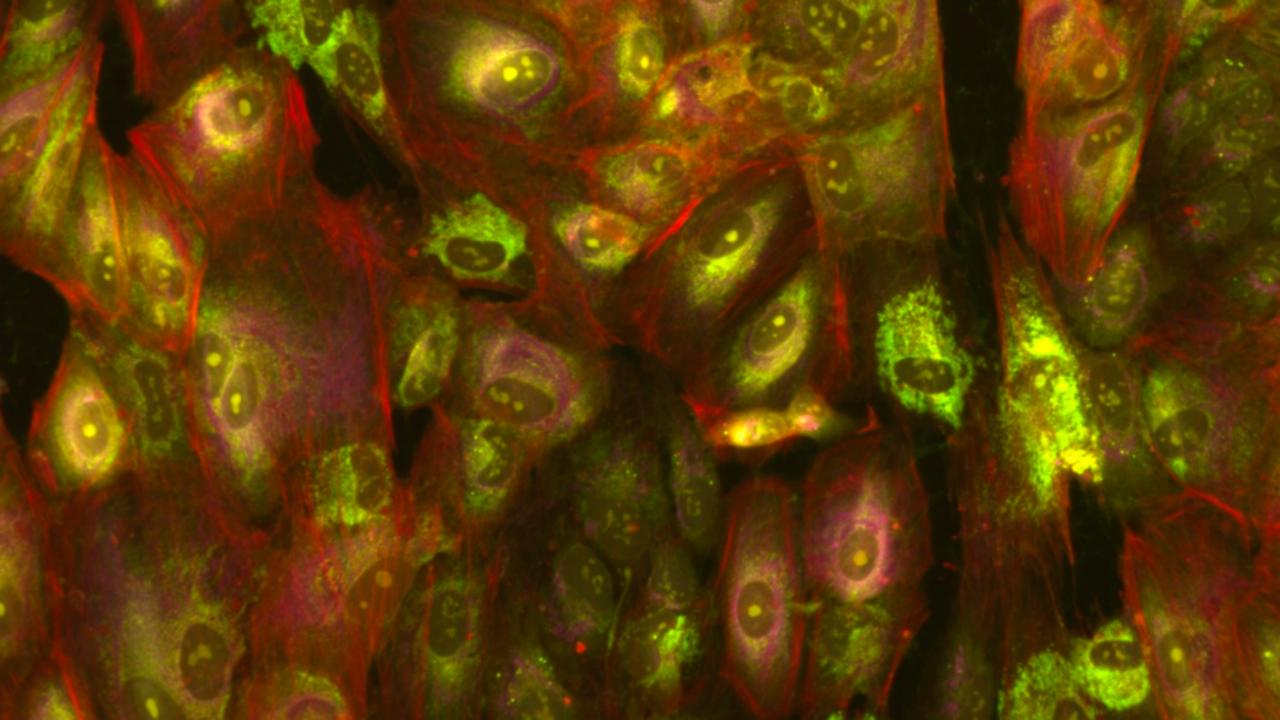


Generating Massive Data Sets

Automated biology leveraging advanced robotics







Our Vision



RARE GENETIC DISEASE

SHORT TERM

100+ genetic disease treatments in 10 years



EXPAND FOCUS

Disrupt drug discovery across new disease areas and new applications



SYSTEMS BIOLOGY

Leverage technology to build a map of human biology



Recursion Pharmaceuticals

Challenges of Rapid Growth and the Evolution of a Splunk implementation





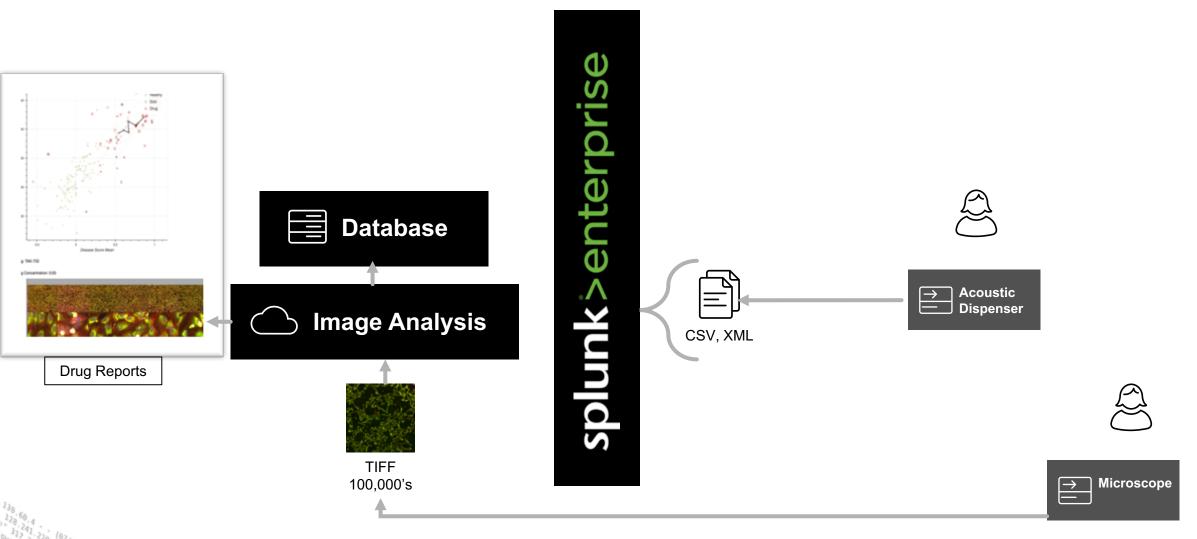
Oct 2016: Platform 1.0

18:18:183] "GET /Product.screen?category_id=GLFTS&LSESSIONID=SSISLAFFIONSFIO HTTP 1.2
222) "GET /Product.screen?product_id=FL-DSH-01&JSESSIONID=SDSSLFFT6ADFF9 HTTP 1.7 200 1318
-82 "468 125.17 /Oldlink?item_id=EST-26&JSESSIONID=SDSSL9FF1ADFF3 HTTP 1.7 200 1318

- ▶ 34 Plates per week max
- ► Limited instrumentation
- Manual labor

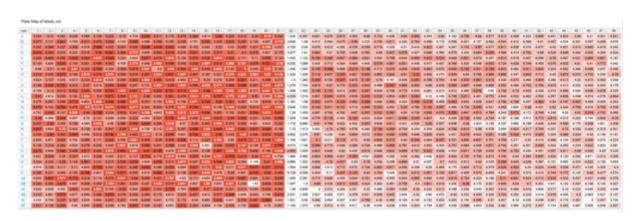


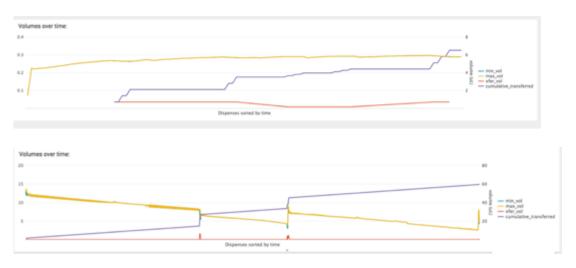
Laboratory Information Management





Compound Management

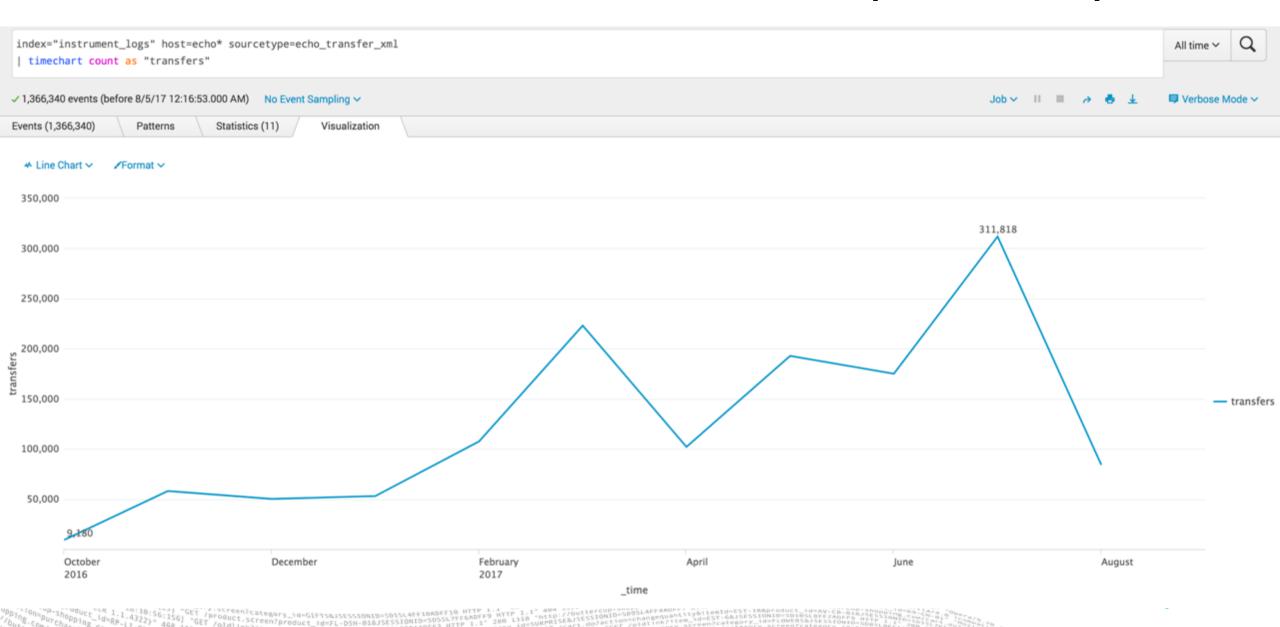


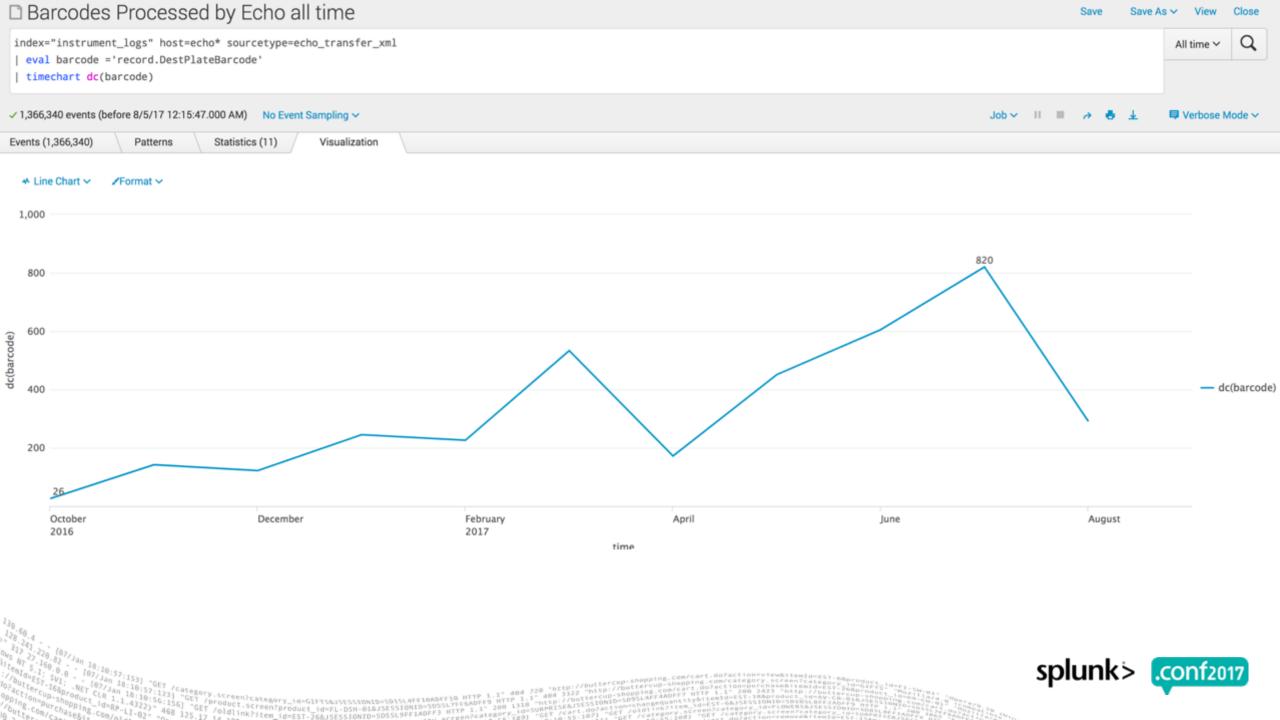


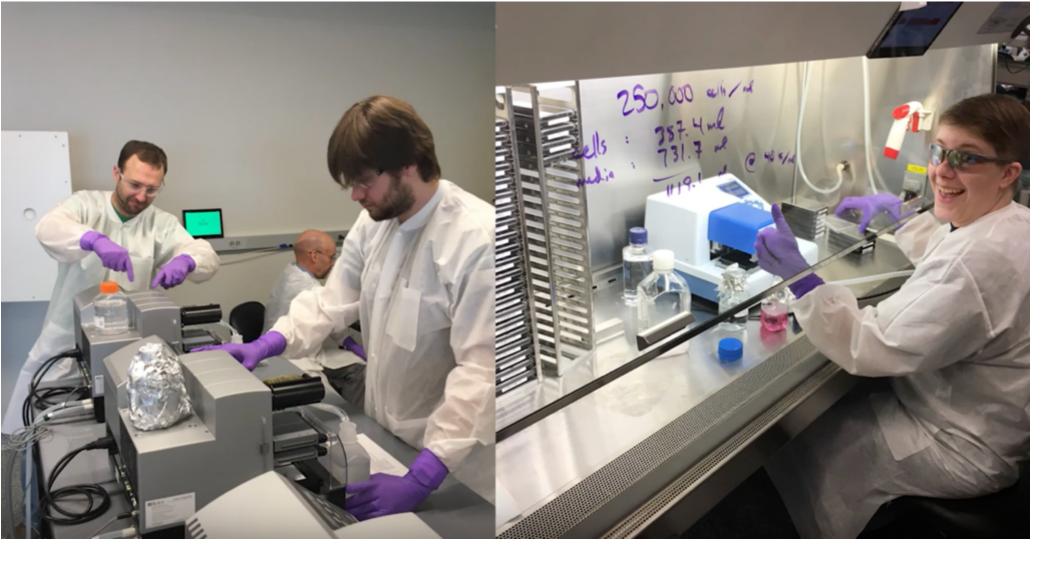




Acoustic Transfers Over Time (1,366,340)





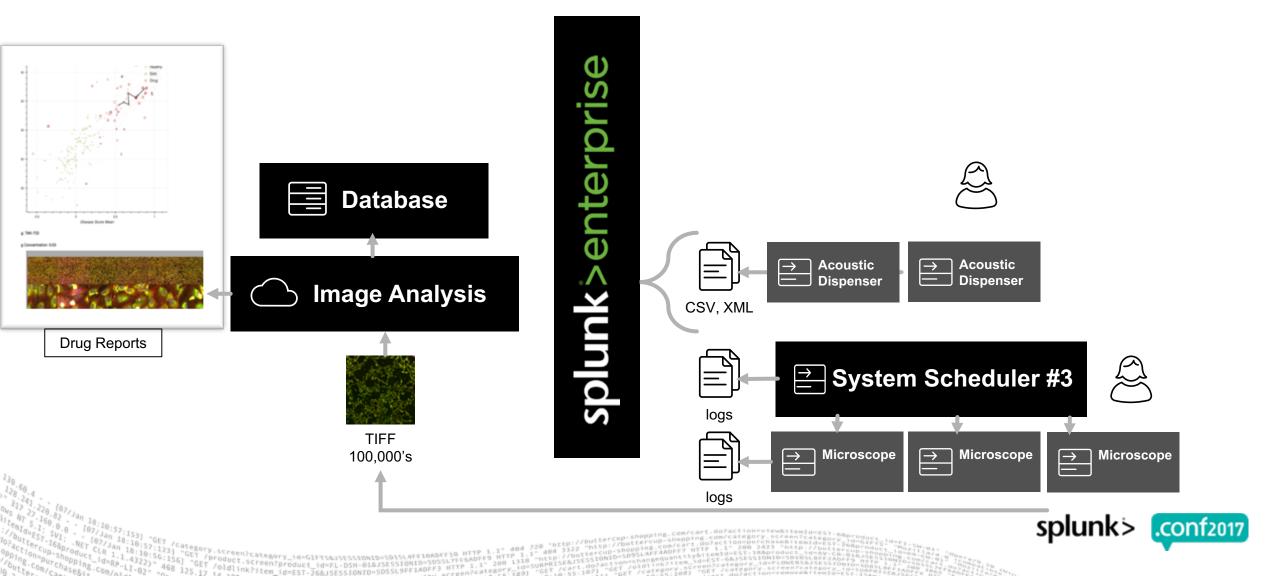


April 2017: Platform 1.5

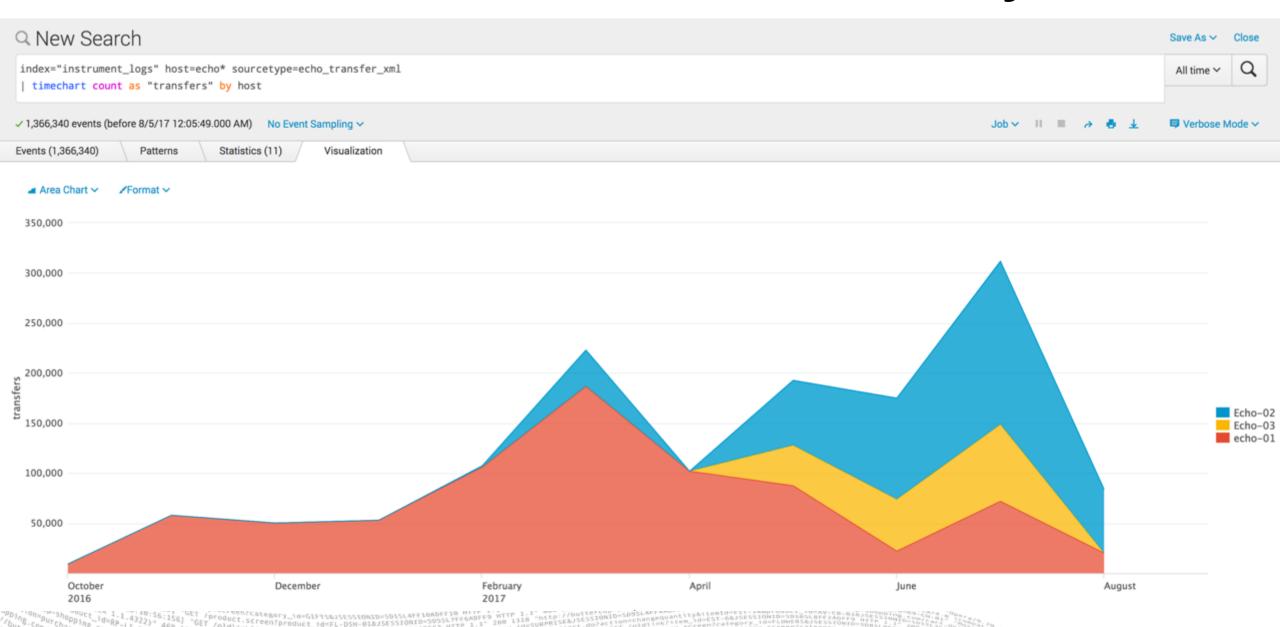
- ▶ 100 Plates per week max
- Multiple instruments allow parallel processing
- Manual processes streamlined



Laboratory Information Management



Acoustic Transfers Performed Over Time By Machine



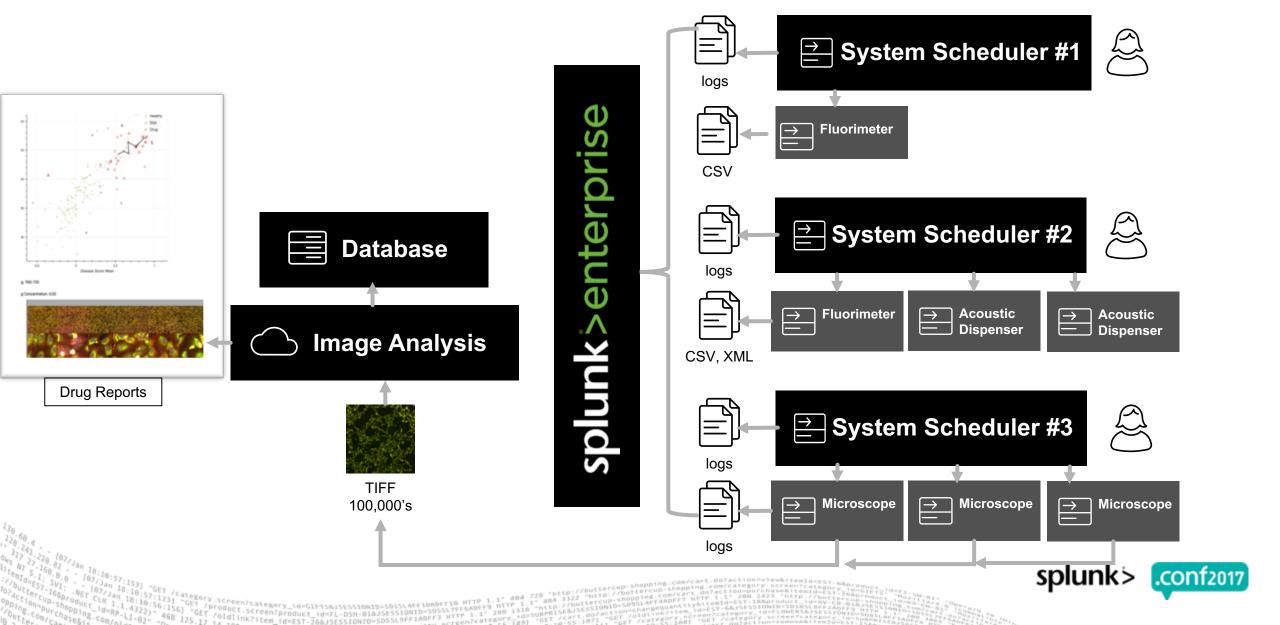


July 2017: Platform 2.0

- ▶ 165 Plates per week Max
- Instrumentation re-allocated
- Processes automated

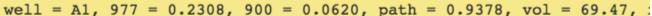


Laboratory Information Management



Dispenser QC



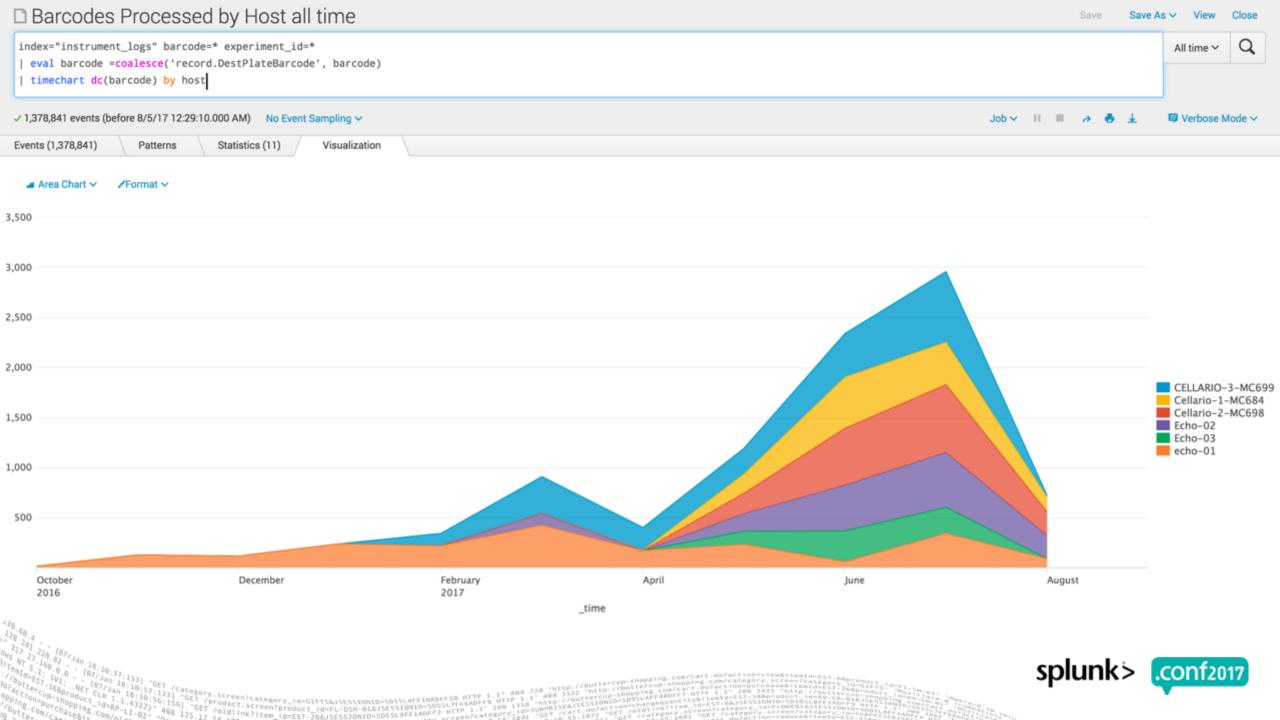


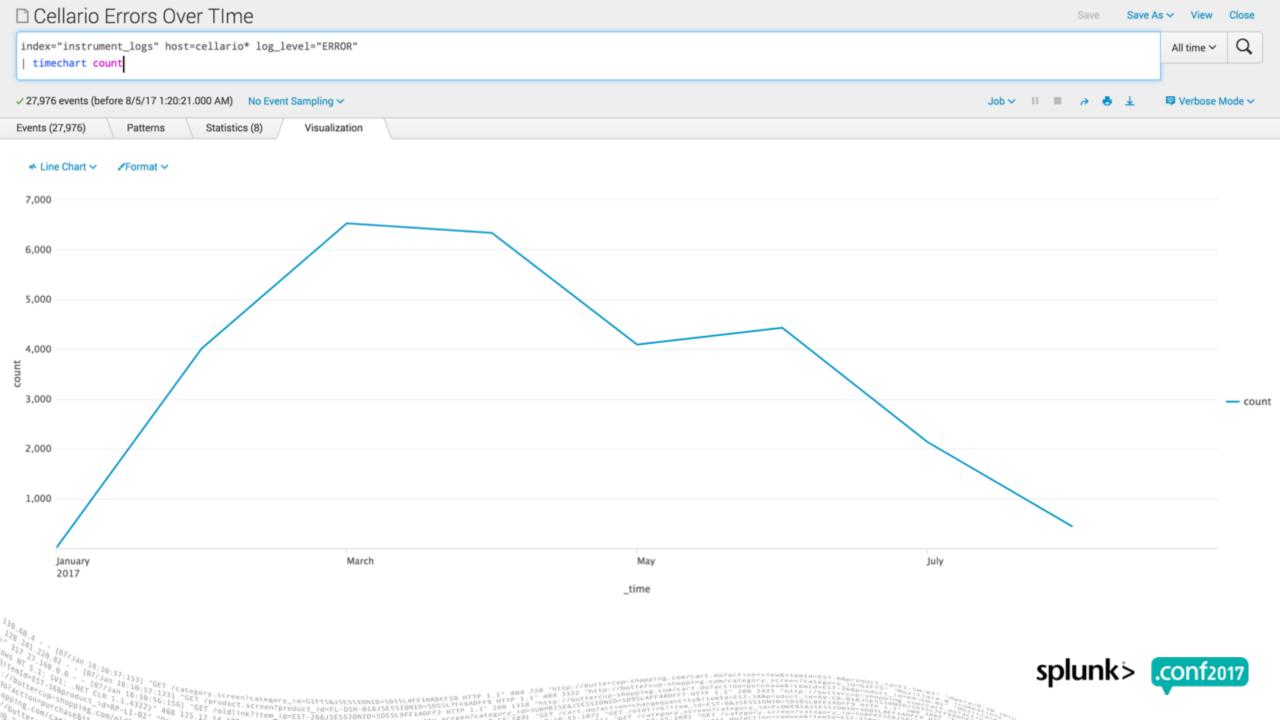
"Y.Screen?category_id=GIFTS&1StSSIONID=S01SL4Ffl0AGFF10 HTTP 1.2 /Product.screen?product id=EL-DSH-018JSESSIONID=SDSSL7F6 1.1 20

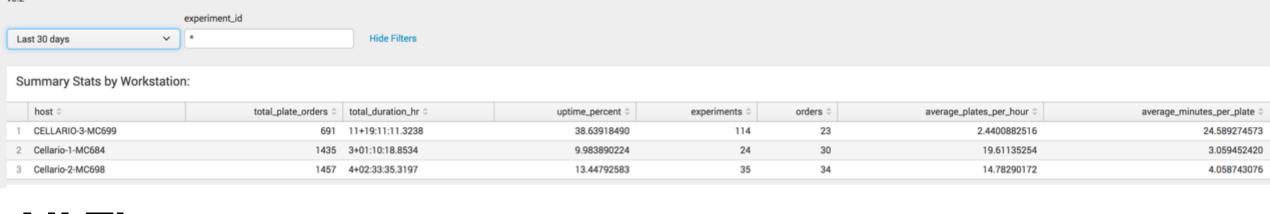
2017-08-04 08:38:56.9249 TRACE Diagnostic - well = B1, 977 = 0.2350, 900 = 0.0620, path = 0.9810, vol = 71.19, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9249 INFO Diagnostic - well = C1, 977 = 0.2302, 900 = 0.0612, path = 0.9389, vol = 69.55, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.909 TRACE Diagnostic - well = C1, 977 = 0.2302, 900 = 0.0612, path = 0.9389, vol = 69.55, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.909 INFO Diagnostic - well = D1, 977 = 0.2348, 900 = 0.0620, path = 0.9600, vol = 71.11, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9579 INFO Diagnostic - well = D1, 977 = 0.2333, 900 = 0.0630, path = 0.9610, vol = 71.011, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9579 INFO Diagnostic - well = E1, 977 = 0.2333, 900 = 0.0630, path = 0.9610, vol = 70.08, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9740 INFO Diagnostic - well = E1, 977 = 0.2331, 900 = 0.0604, path = 0.9517, vol = 70.49, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9910 INFO Diagnostic - well = F1, 977 = 0.2317, 900 = 0.0604, path = 0.9517, vol = 70.49, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9910 INFO Diagnostic - well = C1, 977 = 0.2317, 900 = 0.0604, path = 0.9517, vol = 70.49, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9910 INFO Diagnostic - well = C1, 977 = 0.2317, 900 = 0.0604, path = 0.9517, vol = 70.49, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:56.9910 INFO Diagnostic - well = C1, 977 = 0.2317, 900 = 0.0604, path = 0.9517, vol = 68.27, resource = Combi_02, dispenser = 836-39448M, cassette = 832057609
2017-08-04 08:38:57.0071 INFO Diagnostic - well = C1, 977 = 0.2317, 900 = 0.0604, path = 0.9517, vol = 68.27,









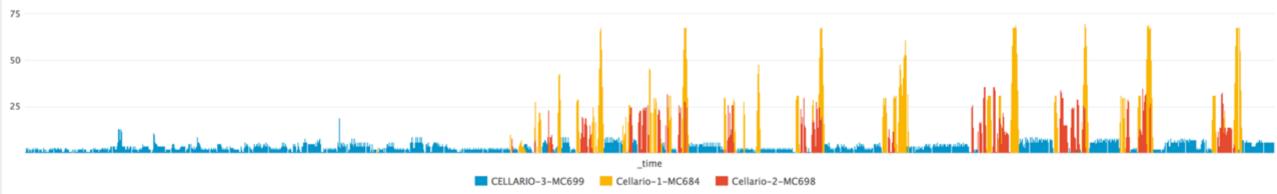


Export ~

All Time:

Workstation Usage

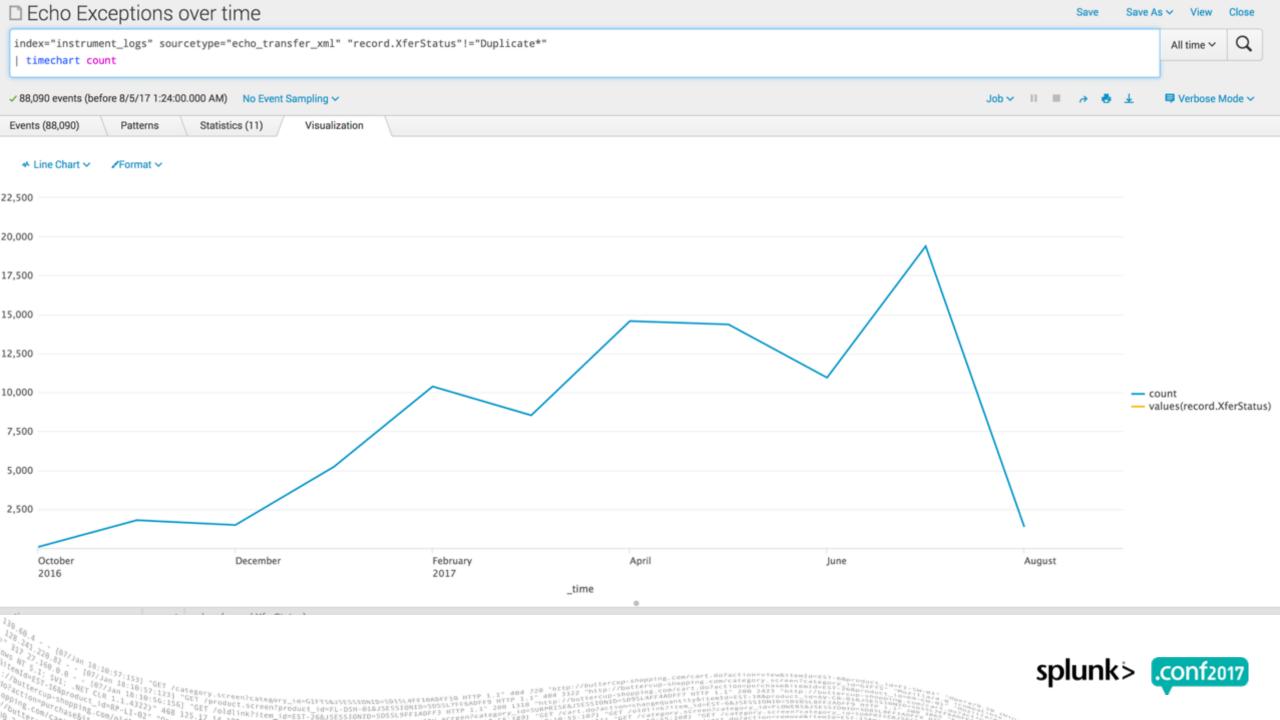
Distinct count of barcodes by host per hour:



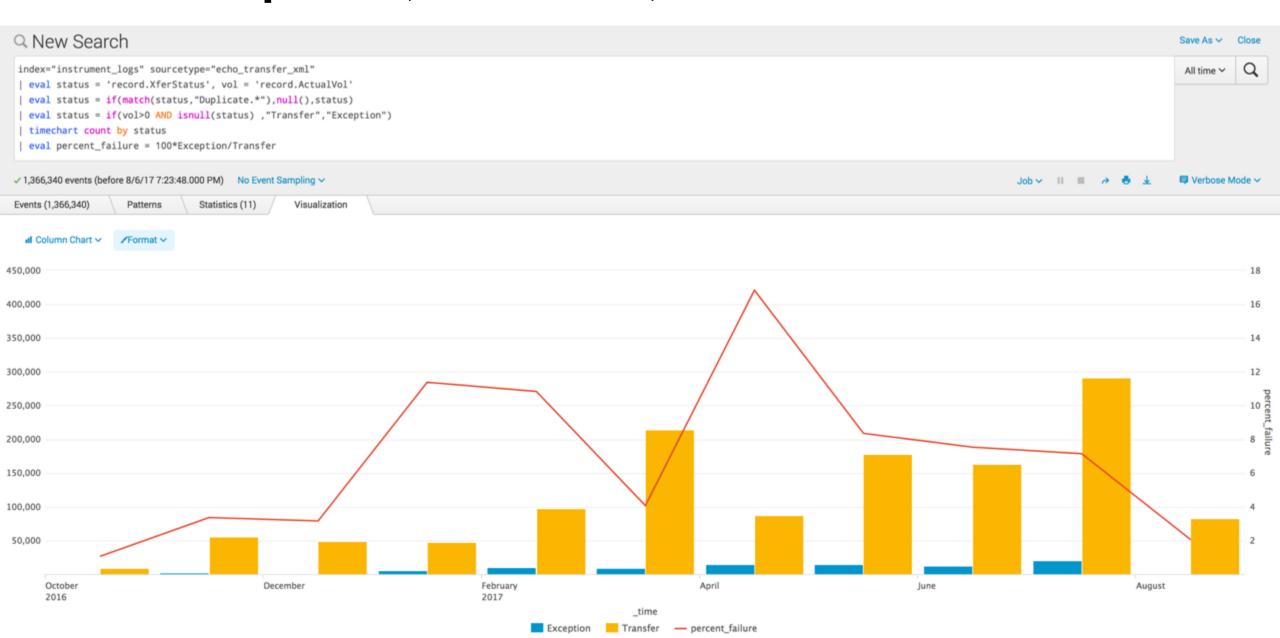
Timeline view of Cellario orders by workstation:

468 125.17 tallink?item_id=E3T-26AJ\$E\$5510NID=SD\$SLTFF6ADF# 1.1" 288 1318

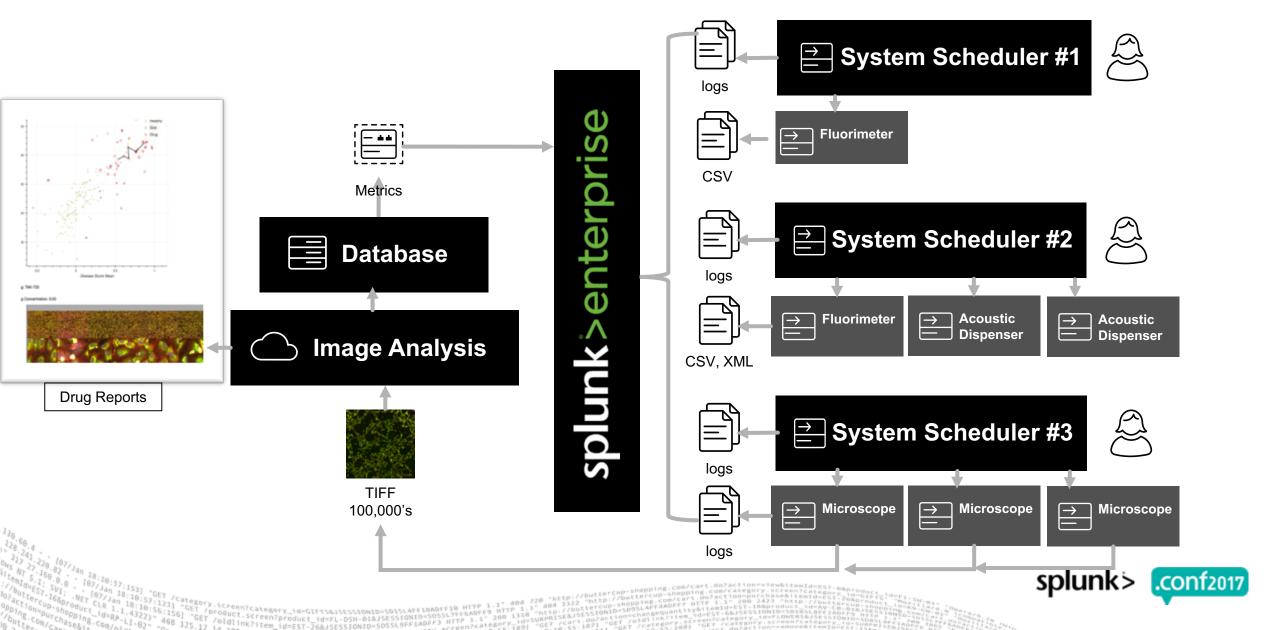




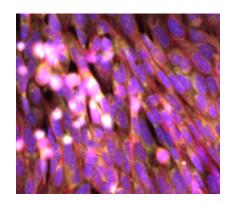
Echo Exceptions, Transfers, And Error Rate Over Time

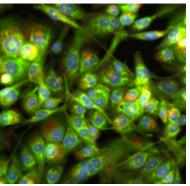


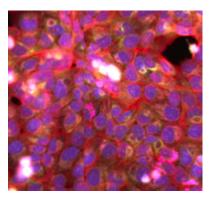
Laboratory Information Management

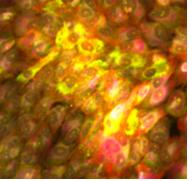


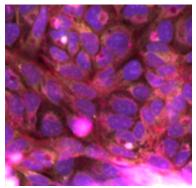
Stain Intensity

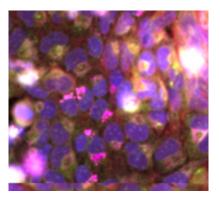










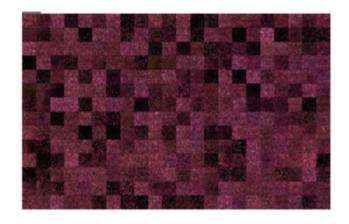


Raw	Data
naw	Data

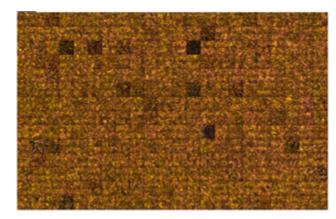
avg_channel1_intensity •	avg_channel2_intensity ‡	avg_channel3_intensity 0	avg_channel4_intensity 0	avg_channel5_intensity 0	avg_channel6_intensity 0	barcode 0	cell_type 🗘	disease_reagent 0
25808.1469727	12194.8449707	19789.0744629	16661.4907227	28730.1542969	24949.9677734	mSBXhh	U20S	s27484
21852.8208008	10148.588623	18187.9821777	13255.1413574	32997.8261719	23067.0234375	XK4smS	U20S	s29290
20721.7006836	13160.0712891	18416.5869141	14442.5844727	25518.8139648	22767.4267578	cGELoq	U2OS	s195324
19697.0734863	12475.0661621	16913.873291	13995.5244141	20908.534668	22823.34375	LPSf2G	U2OS	s18292
19629.8933105	11648.0418701	19421.2907715	11859.1444092	18833.6164551	20804.0861816	vkNcjq	U2OS	s533397



Stain Intensity Principle Component







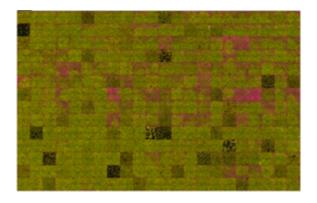
```
| foreach *_intensity [ eval norm_<<MATCHSTR>> = <<FIELD>> / n_cells]
| fit StandardScaler norm_*
| fit PCA SS_* k=1
| stats avg(metric) as average max(_time) as _time by barcode
```

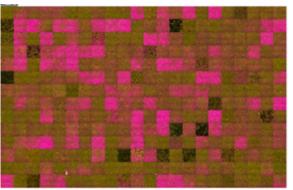


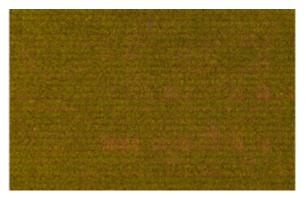


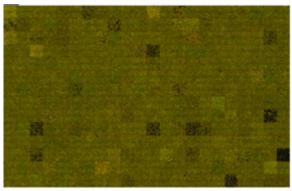


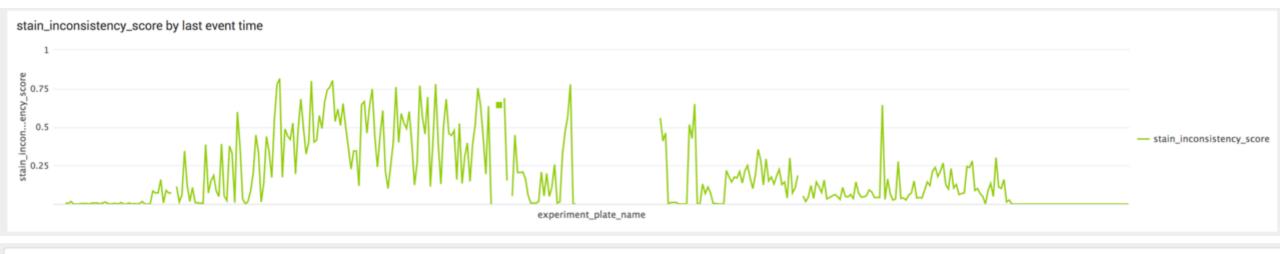
Stain Score











| dbxquery query="select wm.*, e.execution_date from metrics.well_metrics AS wm INNER JOIN design.experiments AS e ON wm.experiment_id = e.experiment_id where wm.barcode IS NOT null and e.execution_date > now () - interval '60 day'; connection="Metrics" maxrows=500000



Plate Flatness

table _time well fine_position site axotrace_barcode acquisition_name host
streamstats last(acquisition_name) as acquisition_name last(well) as well last(site) as site by host global=false



```
Iteration, 0, Fine Position, 0, 0.00,
Iteration, 0, Fine Position, 1, 10270.64,
Site Elapsed Time: 1"
```

```
Iteration, 0, Fine Position, 0, 0.00, Time: 1° 17:1 17:1 17:1 Site Elapsed Time: 2"

17:1 17:1 0, Iteration, 0, Fine Position, 1, 10269.92, 17:1 17:1 0, Iteration, 0, Fine Position, 0, Time: 1° 0, Fine Position, 0, Iteration, 0, Fine Position, 1, 10269.98, 0, Site Elapsed Time: 3"

0, Iteration, 0, Fine Position, 1, 10268.80, 0, Site Elapsed Time: 4"

Well Elapsed Time: B07"
```





What's Next:

- More functionality
 - Alerts with Slack integration
 - Scheduled Reports
- ► More Metrics:
 - Define control limits
 - By Experiment
 - By Well
- More connections
 - Incubation timing
 - Environmental monitoring

- More Data
 - Inventory control
 - Protocol changes
 - Application performance
- More automation
 - Integrating reports to schedule workloads
 - Using alerts for automation control





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