

analyze workflow

adage-server v0.8.1b

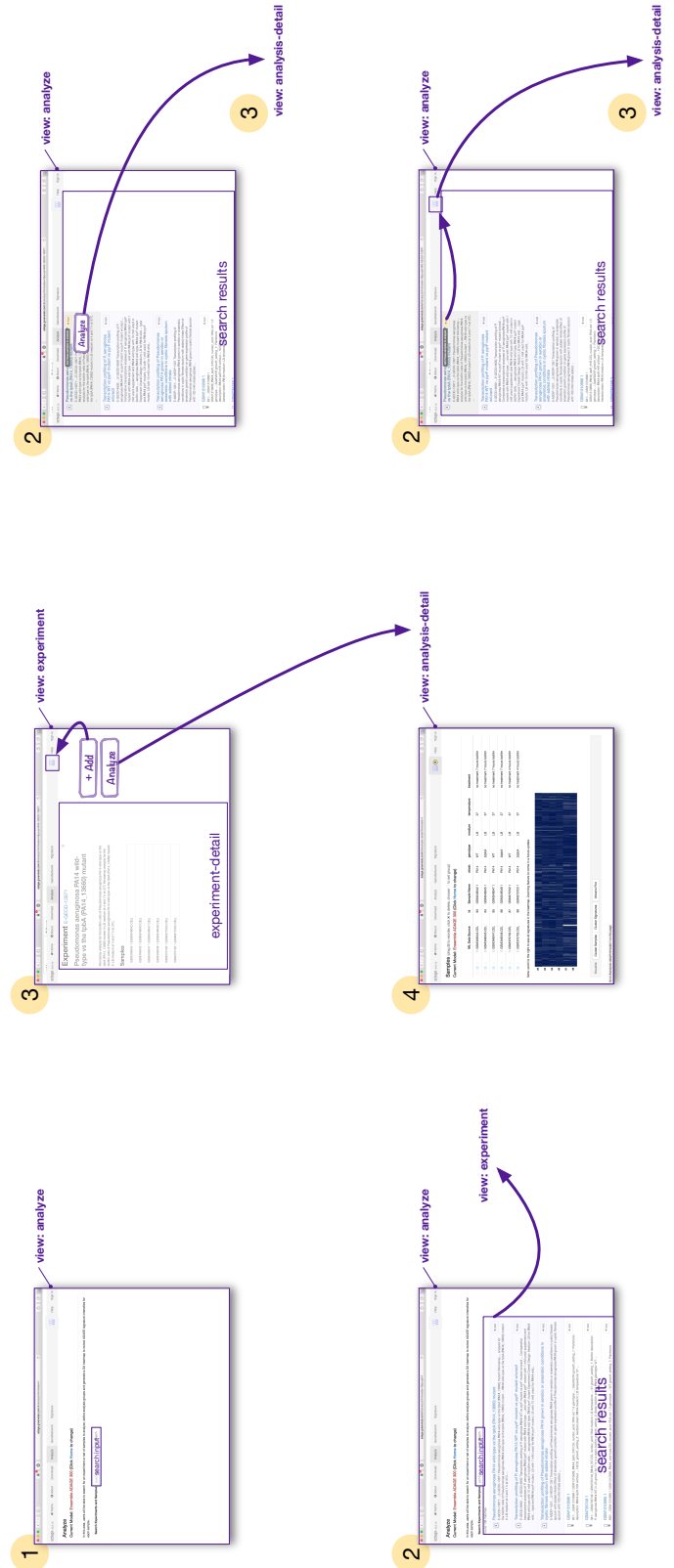
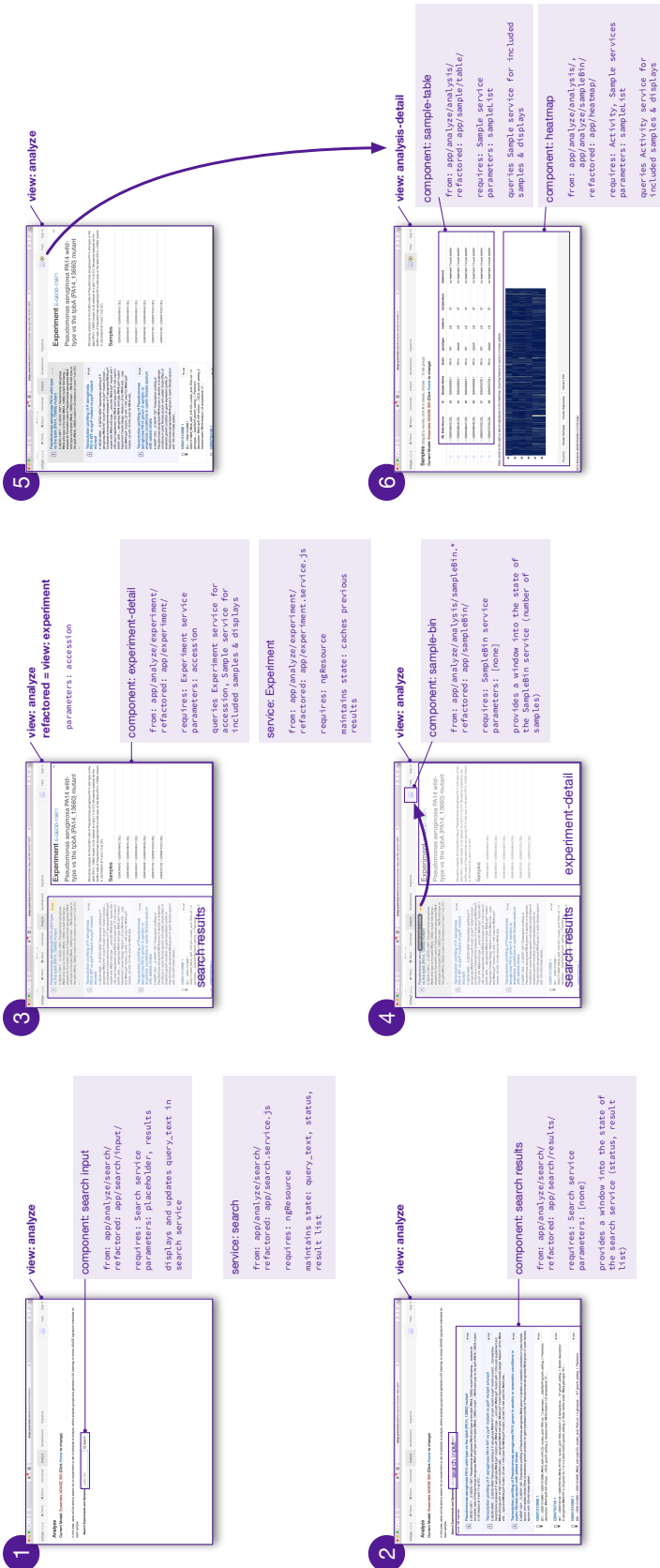
refactoring design 26-Sep-2017

existing workflow:

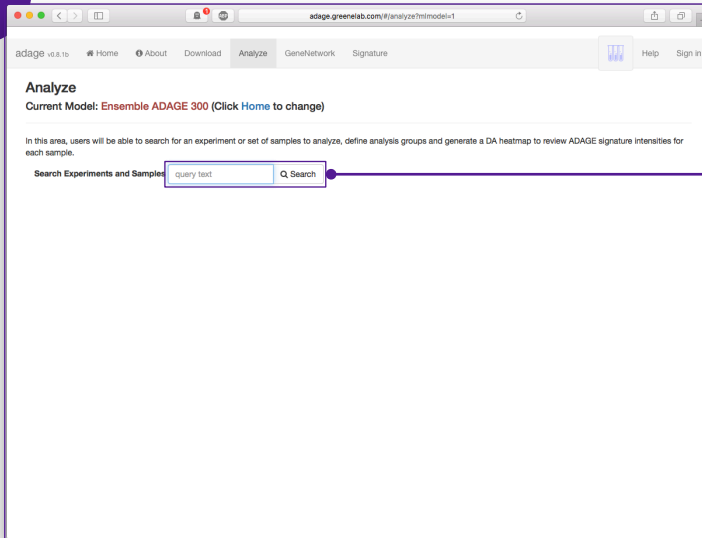
1 2 3 4 5 6

revised workflow:

1 2 3 4



1



view: analyze

component: search input

from: app/analyze/search/
refactored: app/search/input/

requires: Search service
parameters: placeholder, results

displays and updates query_text in
search service

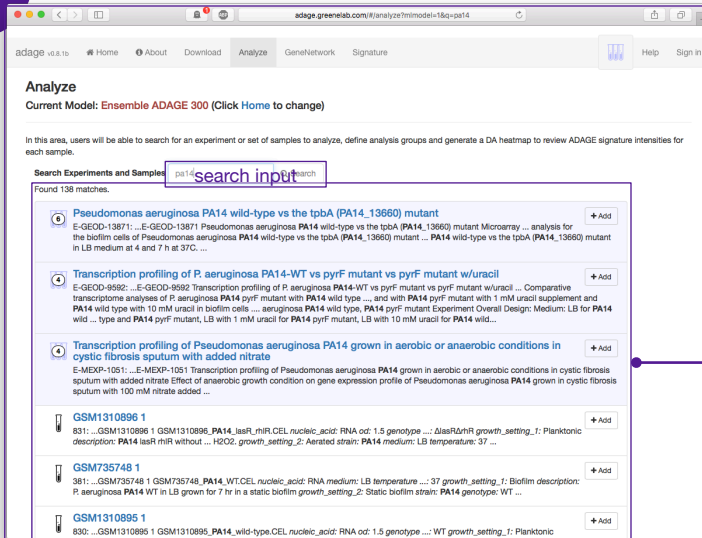
service: search

from: app/analyze/search/
refactored: app/search.service.js

requires: ngResource

maintains state: query_text, status,
result list

2



view: analyze

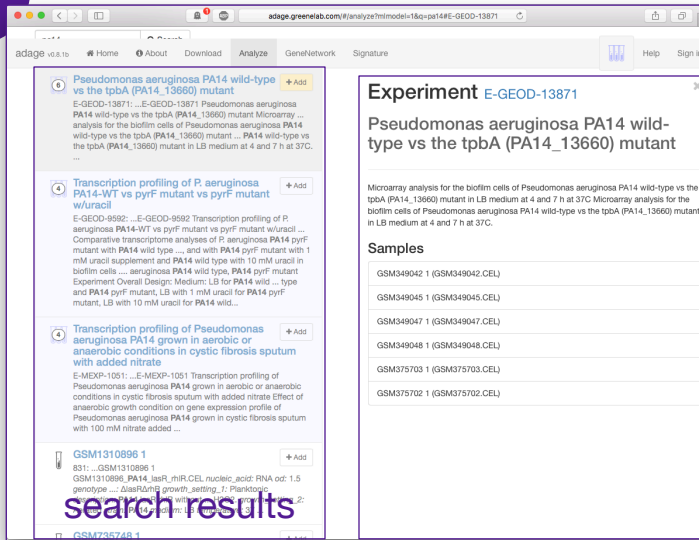
component: search results

from: app/analyze/search/
refactored: app/search/results/

requires: Search service
parameters: [none]

provides a window into the state of
the search service (status, result
list)

3



view: analyze
refactored = view: experiment

parameters: accession

component: experiment-detail

from: app/analyze/experiment/
refactored: app/experiment/

requires: Experiment service
parameters: accession

queries Experiment service for
accession, Sample service for
included samples & displays

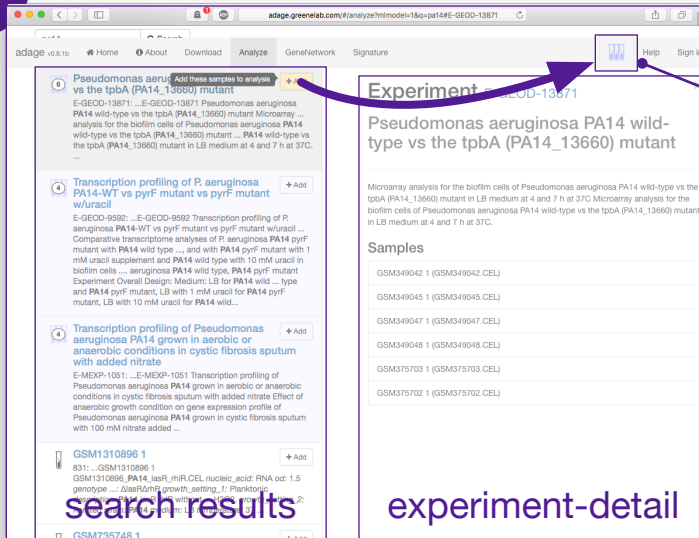
service: Experiment

from: app/analyze/experiment/
refactored: app/experiment.service.js

requires: ngResource

maintains state: caches previous
results

4



view: analyze

component: sample-bin

from: app/analyze/analysis/sampleBin.*
refactored: app/sampleBin/

requires: SampleBin service
parameters: [none]

provides a window into the state of
the SampleBin service (number of
samples)

5

Experiment E-GEOD-13871

Pseudomonas aeruginosa PA14 wild-type vs the tpbA (PA14_13660) mutant

Microarray analysis for the biofilm cells of Pseudomonas aeruginosa PA14 wild-type vs the tpbA (PA14_13660) mutant in LB medium at 4 and 7 h at 37C. Microarray analysis for the biofilm cells of Pseudomonas aeruginosa PA14 wild-type vs the tpbA (PA14_13660) mutant in LB medium at 4 and 7 h at 37C.

Samples

GSM349042	1	(GSM349042.CEL)
GSM349045	1	(GSM349045.CEL)
GSM349047	1	(GSM349047.CEL)
GSM349048	1	(GSM349048.CEL)
GSM375703	1	(GSM375703.CEL)
GSM375702	1	(GSM375702.CEL)

view: analyze

6

Samples (drag to reorder, click X to delete, choose to set group)

Current Model: Ensemble ADAGE 300 (Click Home to change)

ML Data Source	Id	Sample Name	strain	genotype	medium	temperature	treatment
GSM349042.CEL	83	GSM349042.1	PA14	WT	LB	37	no treatment 7 hours biofilm
GSM349045.CEL	84	GSM349045.1	PA14	ΔtpbA	LB	37	no treatment 7 hours biofilm
GSM349047.CEL	85	GSM349047.1	PA14	WT	LB	37	no treatment 7 hours biofilm
GSM349048.CEL	86	GSM349048.1	PA14	ΔtpbA	LB	37	no treatment 7 hours biofilm
GSM375702.CEL	87	GSM375702.1	PA14	WT	LB	37	no treatment 4 hours biofilm
GSM375703.CEL	88	GSM375703.1	PA14	ΔtpbA	LB	37	no treatment 4 hours biofilm

Note: scroll to the right to see all signatures in the heatmap. Zooming feature to come in a future update.

Visualize Cluster Samples Cluster Signatures Volcano Plot

view: analysis-detail

component: sample-table

from: app/analyze/analysis/
refactored: app/sample/table/

requires: Sample service
parameters: sampleList

queries Sample service for included
samples & displays

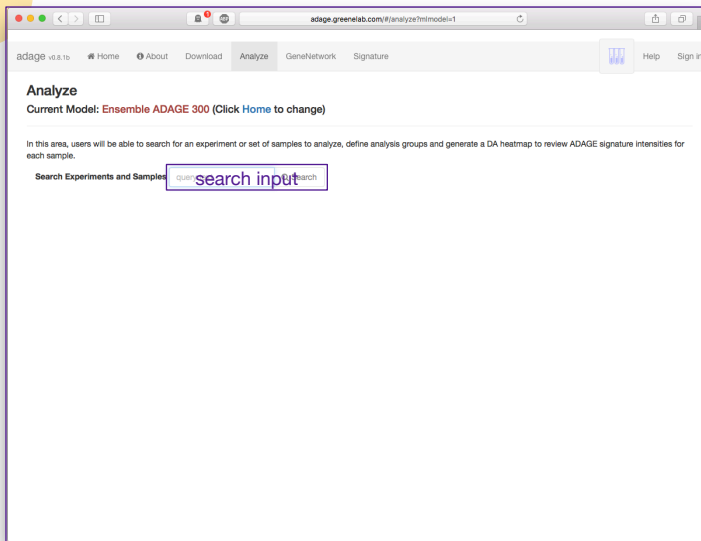
component: heatmap

from: app/analyze/analysis/
app/analyze/sampleBin/
refactored: app/heatmap/

requires: Activity, Sample services
parameters: sampleList

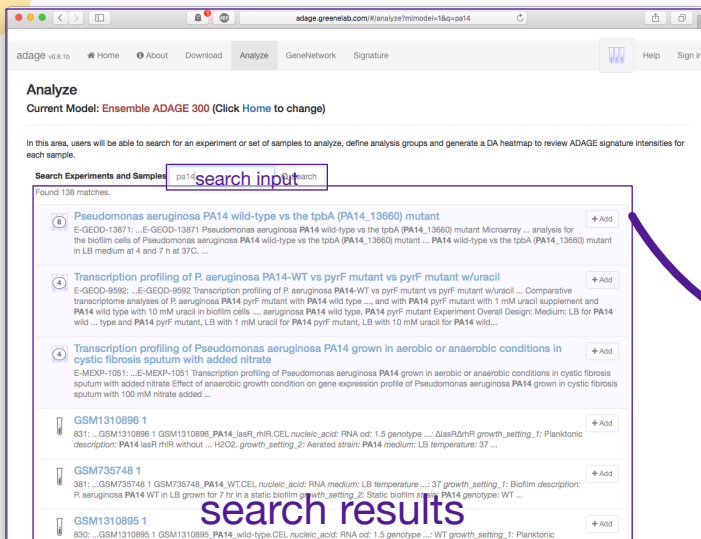
queries Activity service for
included samples & displays

1



view: analyze

2



view: analyze

view: experiment

3

Experiment E-GEOD-13871

Pseudomonas aeruginosa PA14 wild-type vs the tpbA (PA14_13660) mutant

Microarray analysis for the biofilm cells of Pseudomonas aeruginosa PA14 wild-type vs the tpbA (PA14_13660) mutant in LB medium at 4 and 7 h at 37C. Microarray analysis for the biofilm cells of Pseudomonas aeruginosa PA14 wild-type vs the tpbA (PA14_13660) mutant in LB medium at 4 and 7 h at 37C.

Samples

GSM349042.1 (GSM349042.CEL)
GSM349045.1 (GSM349045.CEL)
GSM349047.1 (GSM349047.CEL)
GSM349048.1 (GSM349048.CEL)
GSM375703.1 (GSM375703.CEL)
GSM375702.1 (GSM375702.CEL)

experiment-detail

+ Add

Analyze

view: experiment

4

Samples (drag to reorder, click X to delete, choose to set group)

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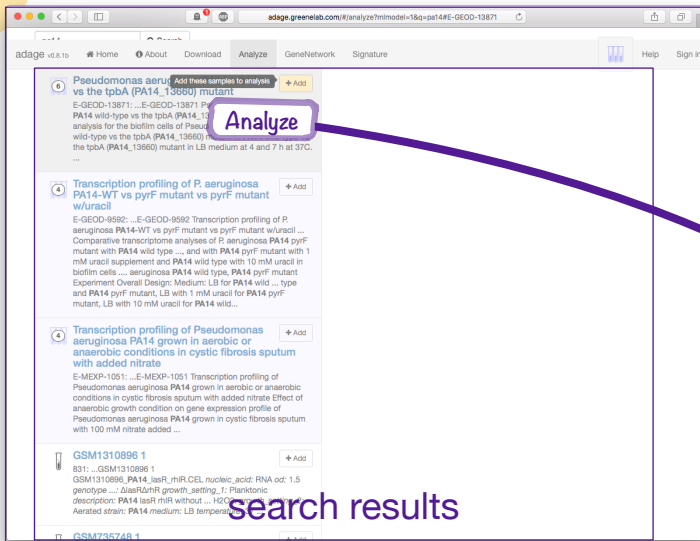
Note: scroll to the right to see all signatures in the heatmap. Zooming feature to come in a future update.

Visualize Cluster Samples Cluster Signatures Volcano Plot

Go to #analysis-detail?model=1 on this page

view: analysis-detail

2

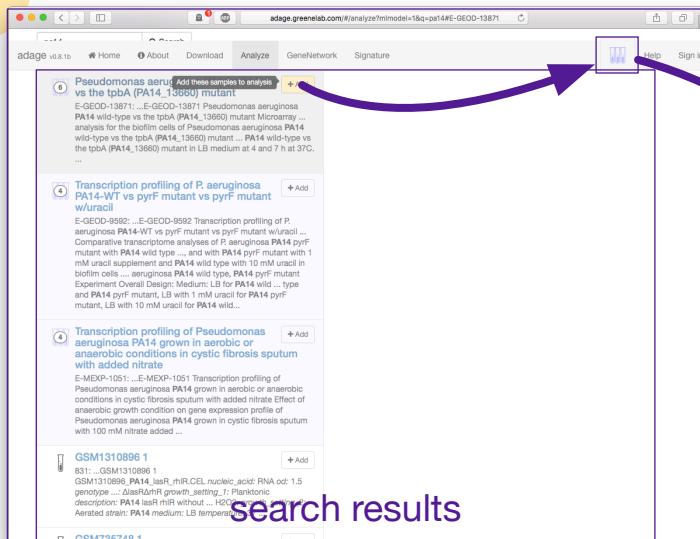


view: analyze

3

view: analysis-detail

2



view: analyze

3

view: analysis-detail