

Specimen Tracking with Galaxy

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#usegalaxy @galaxyproject

Motivation

Sequencers are hiding behind every corner.

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Challenge

Possibilities

Approach

Solution

Outcome

Future

Resources

New genomic project @ Penn State w samples from 100s of families.

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Too small for enterprise LIMS but too complicated for low-end solutions.

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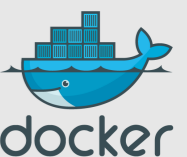
A need to track the hierarchy of derivates.

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A need to track the hierarchy of derivatives.

Wish: 'Just start sequencer and see the data in Galaxy'



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Heavyweight

Turn Galaxy into customer<->provider sequencing management platform.

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Galaxy tried talking to sequencers in the past.

(see Greg's `demo_sequencer` dir in Galaxy or <http://screencast.g2.bx.psu.edu/pacBio-demo/>)

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Limited to samples for sequencing = possibly missing context.

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Derived projects operational.

(Brad Chapman's work , TRON LIMS)

Lightweight

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Let people extend.

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Smartphones instead of barcode scanners.

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Smartphones instead of barcode scanners.

Scripts instead of integration.

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Less instead of more.

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Scripts instead of integration.

Less instead of more.

Start small and listen to users. Change known paths slowly. Don't try to solve too much.

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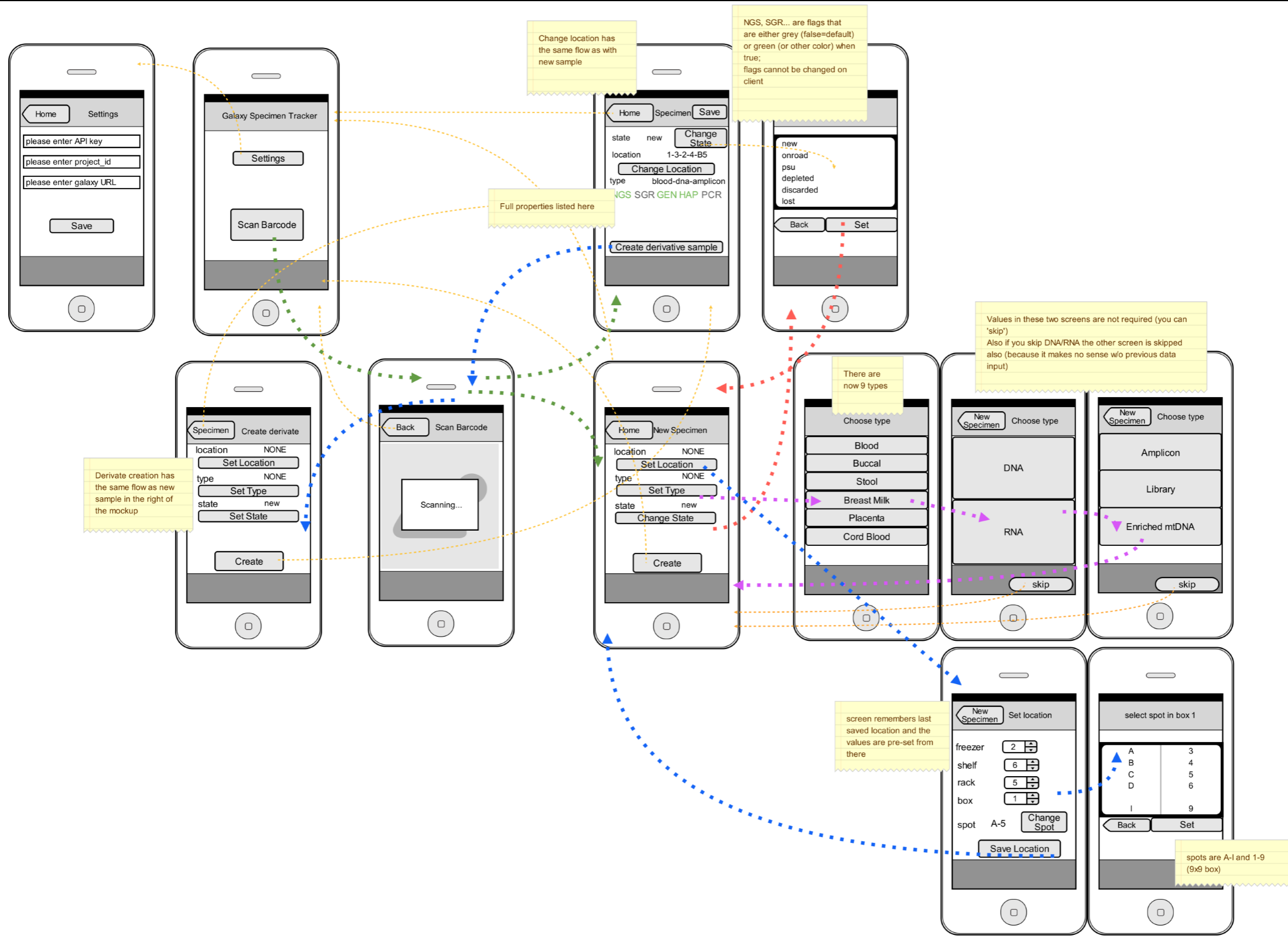
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99 00:58

Specimen DETAILS

Location box_1-shelf_5-rack_0-fridge_1-spot_F4
Change Location

Type blood
Change Type

State depleted
Change State

GEN HAP SGR NGS PCR

Update

Create derivative specimen

100 01:33

Choose type

blood	buccal	hair
breastmilk	stool	vaginal_swab
placenta	cord_blood	tissue

dna	rna
-----	-----

amplicon	library	enriched_mtdn a
----------	---------	--------------------

Set

99 00:58

Specimen CLOSE

Location Family 114

Derivative No

Relationship M

Type Participant dob 9/8/1977

Create time Apr 1, 2014 1:58:17 PM

State Update time Apr 1, 2014 1:58:17 PM

Date sent 11/8/2013

GEN Date of collection 2/11/2013

Note Imported from Sarah's XLS sheet on 3/17/2014.

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Hide Filter Clear Filter

123 barcode

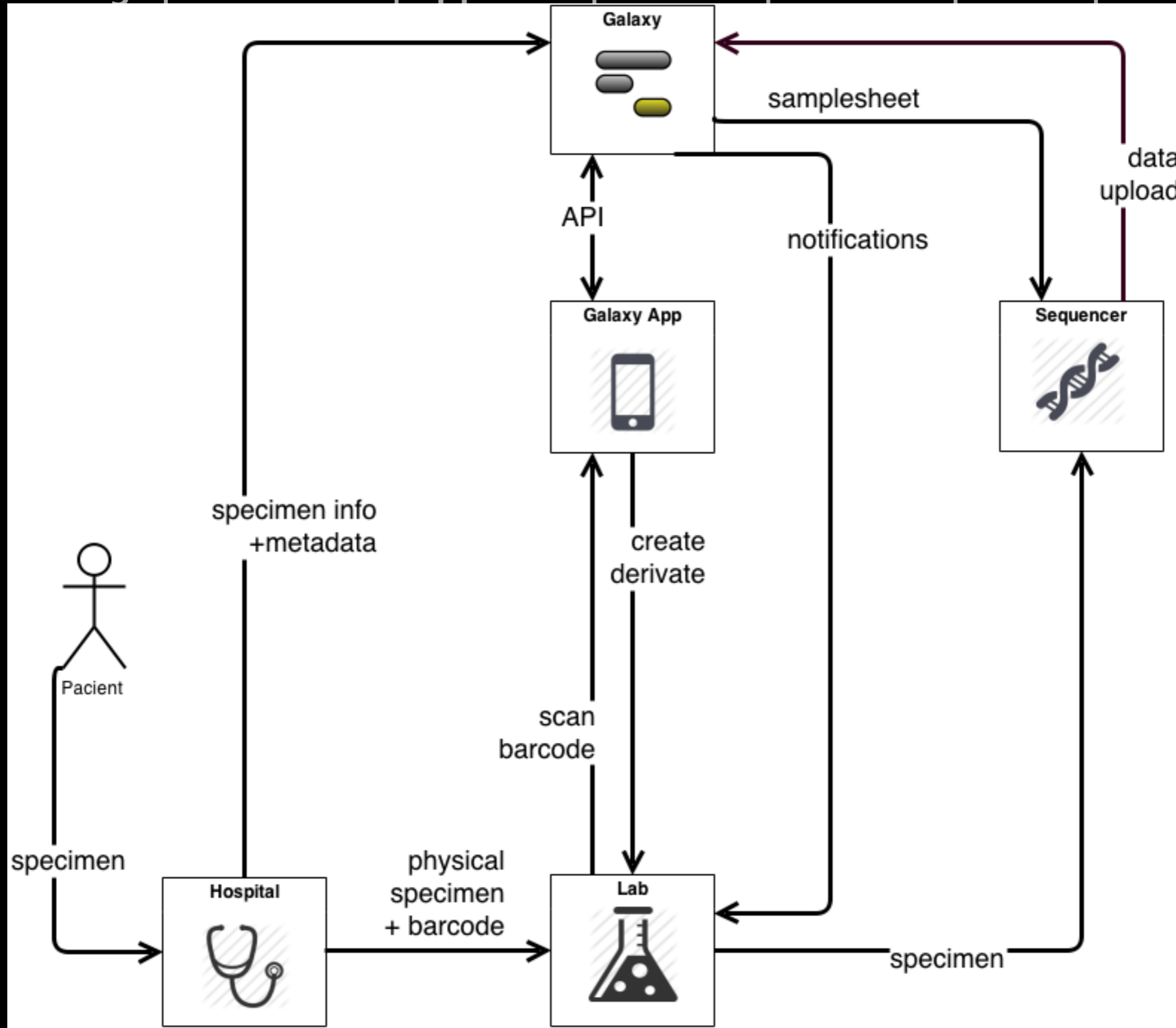
new onroad psu depleted

blood buccal hair breastmilk

dna rna

library amplicon enriched_mtdna

fam	rel	sex	state	type	location	derivate	note
123	C1	M	psu	blood	fridge_1-shelf_5-rack_0-box_2-spot_A2	no	Imported from Sarah's XLS sheet on 3/17/2014.
123	M		depleted	blood	fridge_1-shelf_5-rack_0-box_2-spot_B2	no	Imported from Sarah's XLS sheet on 3/17/2014.
123	D		psu	blood	fridge_1-shelf_5-rack_0-box_2-spot_C2	no	Imported from Sarah's XLS sheet on 3/17/2014.
123	C1	M	psu	blood-dna	fridge_2-shelf_2-rack_0-box_3-spot_B6	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	D		psu	blood-dna	fridge_2-shelf_2-rack_0-box_2-spot_H8	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	M		psu	blood-dna	fridge_2-shelf_2-rack_0-box_3-spot_B7	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	D		psu	blood-dna	fridge_2-shelf_2-rack_0-box_8-spot_H1	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	D		psu	blood-dna	fridge_2-shelf_2-rack_0-box_3-spot_B8	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	D		psu	blood-dna	fridge_2-shelf_2-rack_0-box_9-spot_B1	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	C1	M	psu	blood-dna	fridge_2-shelf_2-rack_0-box_9-spot_A9	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	C1	M	psu	blood-dna	fridge_2-shelf_2-rack_0-box_8-spot_G9	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	C1	M	psu	blood-dna	fridge_2-shelf_2-rack_0-box_2-spot_H6	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	C1	M	psu	stool		no	frozen Received at HMC: 3/24/2014 Imported from Sarah's XLS sheet on 6/9/2014.
123	C1	M	psu	buccal		no	Facility Buccal Barcode: KMCO0038 Imported from Sarah's XLS sheet on 6/9/2014.
123	M		psu	buccal		no	Facility Buccal Barcode: KMCO0039 Imported from Sarah's XLS sheet on 6/9/2014.
123	M		psu	blood-dna	fridge_2-shelf_2-rack_0-box_2-spot_H7	yes	Imported from Sarah's XLS sheet on 4/1/2014.
123	C1	M	psu	buccal-dna	fridge_2-shelf_5-rack_0-box_14-spot_D5	yes	Facility Buccal Barcode KMCO0038 Imported from Sarah's XLS sheet on 6/17/2014.
123	M		psu	buccal-dna	fridge_2-shelf_5-rack_0-box_14-spot_D6	yes	Facility Buccal Barcode KMCO0039 Imported from Sarah's XLS sheet on 6/17/2014.



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Control

**Network
neutrality**

Consistency?

Reliability

Convenience



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Continuous support and evolution of the pilot project.

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Connecting to and reviving the 'old' sample tracking?

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Dataset collections integration?

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Extending the smartphone apps with more general features for Galaxy public?

Trello card

bit.ly/specimen-tracking-trello

Sources of modified Galaxy, Android & iOS apps

<https://bitbucket.org/martenson>

Sample Tracking API documentation

docs.galaxysampletracking.apiary.io

Mobile app mockups

<https://moqups.com/martenson/ehuJNy2b/>

Thank you, Galaxy Team

@galaxyproject #usegalaxy



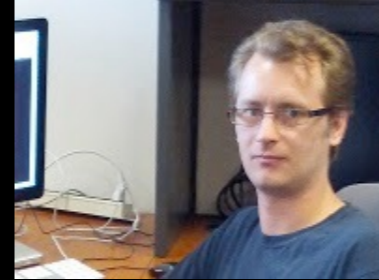
Enis Afgan



Dannon Baker



Dan Blankenberg



Dave Bouvier



Martin Čech



John Chilton



Dave Clements



Nate Coraor



Carl Eberhard



You?



Jeremy Goecks



Sam Guerler



Jen Jackson



Greg Von Kuster



Ross Lazarus



Nick Stoler



Anton Nekrutenko



James Taylor

<http://bit.ly/specimen-tracking-trello>