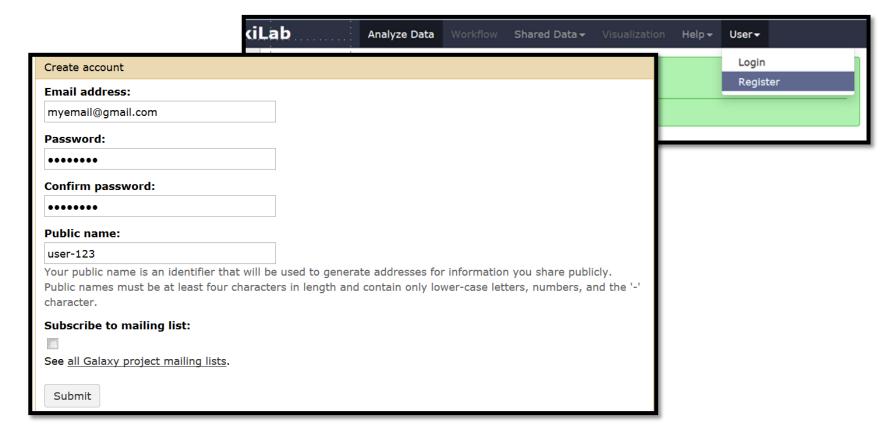
ExomeAl

- ✓ WES data
- ✓ Only tumor samples are required
- ✓ BAF or VCF file
- ✓ detecting AI (including LOH) segments shared among all or some of the tumor samples
- ✓ Removing false positives using a large internal control database

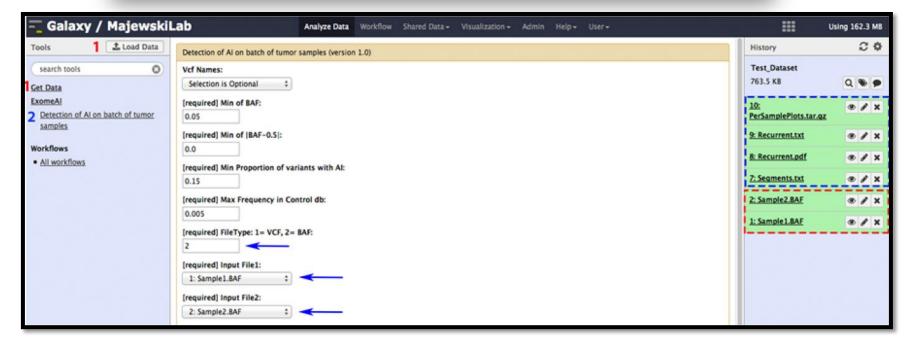
Registering at ExomeAl

- only email address is required
- use all features: managing your data, space, results, histories, ...
- come back to previous sessions after logout
- ...



Using the simulated dataset

=_ Galaxy / Ma	ijewskiLa	ь	Analyze Data	Workflow	Shared Data 🕶	Visualization	Help▼	User▼
Published Histories					Data Libraries Data Libraries Beta			
search name, annotation, Advanced Search	owner, and tags				Published Hist			
Name	Anno	tation	<u>Owner</u>		Published Workflows Published Visualizations Published Pages			Community Tags
Test Dataset AI 1			javad					



Some results for the simulated data

ExomeAl detects a recurrent segment, the blue rectangle, shared between the two samples at about 15-20 Mb (length ~ 5Mb).

