### Feedback - Module 3 Quiz

Help Center

Thank you. Your submission for this guiz was received.

You submitted this quiz on **Wed 12 Aug 2015 9:58 PM PDT**. You got a score of **25.00** out of **28.00**. You can attempt again, if you'd like.

### **Question 1**

Where are basic analyses you can perform found in the Galaxy interface?

Your Answer		Score	Explanation
O Data Libraries			
The Workflow Tab			
The Tool Panel	~	1.00	
The History			
Total		1.00 / 1.00	

### **Question 2**

Where are the datafiles and output of analysis you have performed located in the Galaxy interface?

Your Answer	Score	Explanation
The Tool Panel		
The Workflow Tab		
O Data Libraries		

The History	<b>~</b>	1.00
Total		1.00 / 1.00

### Question 3 Where can I find tools that download or upload data to Galaxy? Your Answer Score Explanation Shared Data tab in Galaxy's top navigation bar The Workflow Tab In the Galaxy History • Get Data in the tool panel ✓ 1.00

1.00 / 1.00

## Question 4 What file format was used for both the exon and repeat data in the exercise? Your Answer Score Explanation TXT file format GFF file format FASTA file format ■ BED file format ✓ 1.00 Total 1.00 / 1.00

Total

### **Question 5**

What information is not contained within a BED file?

Your Answer		Score	Explanation
O Position on chromosome for each feature			
A name for each feature	×	0.00	
The genome the feature belongs to			
The strand on which the feature is encoded			
Total		0.00 / 1.00	

### **Question 6**

What does the "Operate on Genomic Intervals: Join" tool do?

Your Answer		Score	Explanation
Combine overlapping regions within a single dataset			
Match rows in one or more datasets that share the same key			
Ocombine regions that are located close to each other along the chromosome			
Join overlapping region from two or more datasets	~	1.00	
Total		1.00 /	
		1.00	

### **Question 7**

What question are we trying to answer using the exon and repeat data?

Your Answer		Score	Explanation
Which exons overlap a repetitive element?			
How many repeat regions overlap each exon region?	~	1.00	
O How many exons overlap each repeat region?			
O How many exons in each gene?			
Total		1.00 / 1.00	)

### **Question 8**

Which of the following can be done with the group tool?

	Score	Explanation
<b>~</b>	1.00	
	1.00 /	
	1.00	
	~	✓ 1.00

### **Question 9**

How does Galaxy assist in the reproducibility of analysis?

Your Answer		Score	Explanation
Allows repeating the entire analysis			
All of these options	~	1.00	

Automatically maintains records of tools and parameters	
Archives data from external databases that may change	
Total	1.00 / 1.00

### **Question 10**

The "pencil" icon in the Galaxy history allows you to do all but

Your Answer		Score	Explanation
Edit a dataset's contents	~	1.00	
Modify the name of a datasets			
Provide an annotation for a datasets			
Modify metadata about a datasets			
Total		1.00 / 1.00	

### **Question 11**

The file Quiz3\_Genes.bed contains locations of how many transcripts?

Your Answer		Score	Explanation
O 1810			
<ul><li>1720</li></ul>	~	1.00	
O 1620			
O 1510			
Total		1.00 / 1.00	

### **Question 12**

The file Quiz3\_Genes.bed contains locations of how many genes?

Your Answer		Score	Explanation
O 1810			
O 1320			
<u> </u>			
<ul><li>1510</li></ul>	~	1.00	
Total		1.00 / 1.00	

### **Question 13**

What is a workflow?

Your Answer		Score	Explanation
A summary of the provenance of a given piece of data			
An abstract representation of a multi-step analysis	~	1.00	
A description of how to carry out an analysis on a specific dataset			
A tutorial for analyzing a type of data			
Total		1.00 /	
		1.00	

### **Question 14**

How can a workflow be added to a Galaxy instance?

Your Answer		Score	Explanation
Import a workflow from a file			
All of these options	~	1.00	
Create a new workflow from scratch with the workflow editor			
Create a workflow from an existing history			
Total		1.00 /	
		1.00	

### **Question 15**

Which of the following statements is true?

	Score	Explanation
~	1.00	
	1.00 / 1.00	)
	•	✓ 1.00

### **Question 16**

Which are advanced options available in workflows?

Your Answer	Score	Explanation
O Hide datasets that represent intermediate steps		
Neither of these options		

Both of these options	<b>✓</b> 1.00
Rename output datasets	
Total	1.00 / 1.00

# Question 17 Which of the following is true of tags? Your Answer Score Explanation ● An item can have more than one of them ✓ 1.00 They can contain arbitrary free form text They can only be attached to datasets They can only be created by a Galaxy administrator Total 1.00 / 1.00

### 

### **Question 19**

Which of the following is true of tags and annotations?

Your Answer		Score	Explanation
They can be used for search			
All of these options	~	1.00	
They can be attached to Datasets, Histories, and Workflows			
They allow users to provide additional information about entities in Galaxy			
Total		1.00 /	
		1.00	

### **Question 20**

What does it mean to share a galaxy history?

Your Answer		Score	Explanation
Only certain users will be able to access it	~	1.00	
Automatically send an email/tweet/facebook with a link to it			
Make is possible to download an entire history			
Make the history automatically available on any Galaxy server			
Total		1.00 /	
		1.00	

### **Question 21**

What does it mean to publish a galaxy history?

Your Answer
Score Explanation

Email the history

Post it on social media (twitter/facebook)

The history will be listed in a central, public place within Galaxy and all users can access it

Post it on a blog

Total

1.00 /
1.00

### What do you need to share a history with just one user? Your Answer Access to a server where the data can be uploaded to It cannot be done; a history can only be shared with all

users	
Galaxy administrator access	
The email associated with the user's Galaxy account	<b>✓</b> 1.00
Total	1.00 / 1.00

### **Question 23**

Who can see a history that's been made 'accessible'?

Your Answer Score Explanation

**Explanation** 

Only the person that owns (i.e., has created) this history	
Only people I have given explicit permission	
It is visible on the 'Published Histories' page so anyone that clicks on the history name	
Anyone who knows the URL (the browser address)	<b>✓</b> 1.00
Total	1.00 /
	1.00

### **Question 24**

Where are published histories found on Galaxy?

Your Answer		Score	Explanation
◯ In a Data Library			
On the Galaxy's wiki			
All of these options			
Shared Data -> Published Histories	~	1.00	
Total		1.00 / 1.00	

### **Question 25**

What can you do if you no longer want a history to be accessible?

Your Answer		Score	Explanation
It must be deleted			
<ul> <li>From the same interface used to make it accessible: use the 'Share or Publish' link in the history options</li> </ul>	~	1.00	
Need to send an email to helpdesk and ask to make it private			

Rename the history	
Total	1.00 /
	1.00

### **Question 26**

What do Galaxy pages allow you to do?

Your Answer		Score	Explanation
All of these options	~	1.00	
Add and format text			
None of these options			
<ul><li>Embed histories</li></ul>			
Total		1.00 / 1.00	

### **Question 27**

Why can annotations be important in a Galaxy page?

Your Answer		Score	Explanation
They are needed to attach histories and workflow to a page			
They are required to share the page with others			
They provide machine readable provenance information	×	0.00	
They can be used to describe why an analysis was performed in a particular way			
Total		0.00 /	

	Score	Explanation
×	0.00	
,		
	0.00 / 1.00	
	•••	<b>×</b> 0.00