

Feedback

FragmentStore: Genome, Annotations



1

Please give us the first

4 characters of your ID: _____

2

How would you rate your SeqAn skills concerning

the topic addressed in this tutorial before having done it.

☐

No Experience

☒

Basic Skills,
Few Experience

☐

Ordinary Experience

☐

Much Experience

☐

Extraordinary
Experience

3

What was your motivation to do this tutorial? How helpful was the tutorial for you?

learn about organization of & access to
annotation

4

Could you have finished this tutorial without help by the SeqAn team?

☐

No

☐

Yes, but only with a
significant amount of
more time

☒

Yes, with slightly to
no more additional
time

5

Do you think that the documentation web pages

(www.seqan.de and trac.seqan.de) were helpful to solve the exercises of the tutorial?

☐

Hardly

☐

Average

☒

Above Average

☐

Extraordinary

6

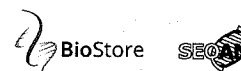
What did you particularly like about this tutorial?

7

What did you particularly dislike about this tutorial? What posed the biggest problems?

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• get to know the concept

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What did you particularly like about this tutorial?

• good complexity level of assignments,
good presentation

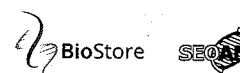
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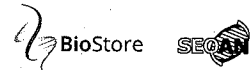
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Interested in understanding these techniques.

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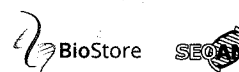
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What did you particularly dislike about this tutorial? What posed the biggest problems?

there are 2 kinds of `goNext()` and the documentation is different (one for serial iterators and another for tree iterators) a bit confusing but OK.

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Annotation are quite important and so,
I was interested in this topic.

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What did you particularly like about this tutorial?

There was enough time for the
sub-tutorials (well explained)

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None, just curious.

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