

# Feedback

## JournalSet



1

Please give us the first 4 characters of your ID: ow4g . . . . .

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?

Very nice and helpful tutorial

4

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



No

Yes, but only with a  
significant amount of  
more timeYes, with slightly to  
no more additional  
time

5

Do you think that the documentation web pages ([www.seqan.de](http://www.seqan.de) and [trac.seqan.de](http://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?

The time estimation was almost perfect.

7

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

Suggestions: 1) Name `join()` is misleading maybe better would be `attach()` or ~~and~~ or  
2) Add interface to check if  
sequence in `JournalSet` is "joined"

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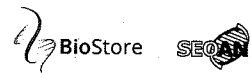
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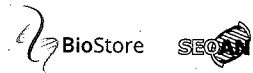
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What did you particularly like about this tutorial?

Überblick  
Habe den Aufbau eines Journalstrings gelernt.

7

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

toHostPosition und toViewPosition fehlen in Doku

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Good introduction  
might have been nice to have more descr.  
of algorithm

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not enough time!

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