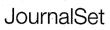




	Please give us the first 4 characters of your ID: $0 w 4 q \cdots \cdots$
2	How would you rate your SeqAn skills concerning the topic addressed in this tutorial before having done it.
	No Experience Basic Skills, Few Experience Ordinary Experience Much Experience Extraordinary Experience
3	What was your motivation to do this tutorial? How helpful was the tutorial for you? Very mce and helpful futorial
4	Could you have finished this tutorial without help by the SeqAn team? 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?
	O O O O 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() on the extension of 2) Add interface to check if Sequence in Journal Set 15, joined.
	sequence in Sournal Set 15, joined

Feedback

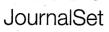






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