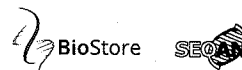


Feedback

RNA-Seq, FragmentStore



1

Please give us the first 4 characters of your ID:

Repj5

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?

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?

I was entirely unfamiliar with the commonly used data (types) used in quantitative seq workflows, learned a lot!

7

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

Too less time. It is required to copy + paste the hints, the time is too short to figure out your own approach

Feedback

RNA-Seq, FragmentStore



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?

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?

useful example

7

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

- too little time
- difficult tutorial maybe better in the morning than late afternoon

Feedback

RNA-Seq, FragmentStore



1

Please give us the first

4 characters of your ID: j3dl

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?

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?

Feedback

RNA-Seq, FragmentStore



1

Please give us the first
4 characters of your ID:

phro

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?

Practically very relevant

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?

Combines many different parts of SeqAn

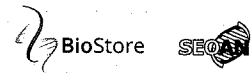
7

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

*- OPEN MP makes the solutions harder to understand
↳ should be a version w.o.
- Complexity of the tutorial was too high
↳ RPKM computation should be additional*

Feedback

RNA-Seq, FragmentStore



1

Please give us the first

4 characters of your ID: ghuu

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?

*Just wanted to do the 'Advanced' lecture.
Maybe helpful, I learned about IntervalTrees. Also
got an idea of the generell problem.*

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

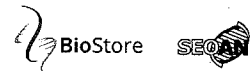
What did you particularly like about this tutorial?

7

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

Feedback

RNA-Seq, FragmentStore



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.

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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 RNA seq algy's

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?

nice to have a working program!

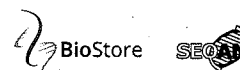
7

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

bit to rushed.

Feedback

RNA-Seq, FragmentStore



1

Please give us the first 4 characters of your ID:

jwzr

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?

Working on RNA-seq data.

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?

practical application
real-life

7

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

Feedback

RNA-Seq, FragmentStore



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?

Understand approach and workflow
for RNA-seq applications

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?

present a simplified view & starting point on
a real world example

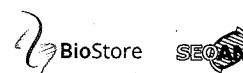
7

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

Mostly about chasing the doc,
a bit more background on library
structure would have been useful.
e.g. more specifics about the store

Feedback

RNA-Seq, FragmentStore



1

Please give us the first 4 characters of your ID:

0 Wh g

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 how to use SeqAn in complex analysis of NGS Data

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?

Very good practical example

7

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

~~Not clear~~ From documentation
not easy to understand how to
use data structures