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Feb 22, 2022

Protocol in dev - Seq Analysis pipeline V.1

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¹protocols.io



dx.doi.org/10.17504/protocols.io.byhmpt46

Emmas new workspace



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test

[Example Doc.docx](#)

DOI

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Emma Ganley 2022. Protocol in dev - Seq Analysis pipeline. **protocols.io**
<https://dx.doi.org/10.17504/protocols.io.byhmpt46>



protocol ,

Sep 24, 2021

Feb 22, 2022

53517

Section 1 10m

1

A	B	C	D
10	2	7	
5			
15			19

A	B	C	D

 0 µL  0 µL

=sum(a1,b1,c1)

2



10m

Add  50 mL A

Add  100 mL B

Leave for  00:10:00

Add  50 µL

 50 µL

Important note info here

Step 2 includes a Step case.

Step case 1

Step case 2

Section 2

10m

step case

Step case 1

3

Cell number	<input type="text"/>
Dead cells	<input type="text"/>

4



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Linux

[source](#) by Wellcome Trust Sanger Institute

5

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6

7

Vöhringer HS, Sanderson T, Sinnott M, De Maio N, Nguyen T, Goater R, Schwach F, Harrison I, Hellewell J, Ariani CV, Gonçalves S, Jackson DK, Johnston I, Jung AW, Saint C, Sillitoe J, Suci M, Goldman N, Panovska-Griffiths J, Wellcome Sanger Institute Covid-19 Surveillance Team., COVID-19 Genomics UK (COG-UK) Consortium., Birney E, Volz E, Funk S, Kwiatkowski D, Chand M, Martincorena I, Barrett JC, Gerstung M (2021). Genomic reconstruction of the SARS-CoV-2 epidemic in England.. Nature.
<https://doi.org/10.1038/s41586-021-04069-y>

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<https://doi.org/10.1038/d41586-021-02401-0>

Step 7 includes a Step case.

Case 1

Case 2