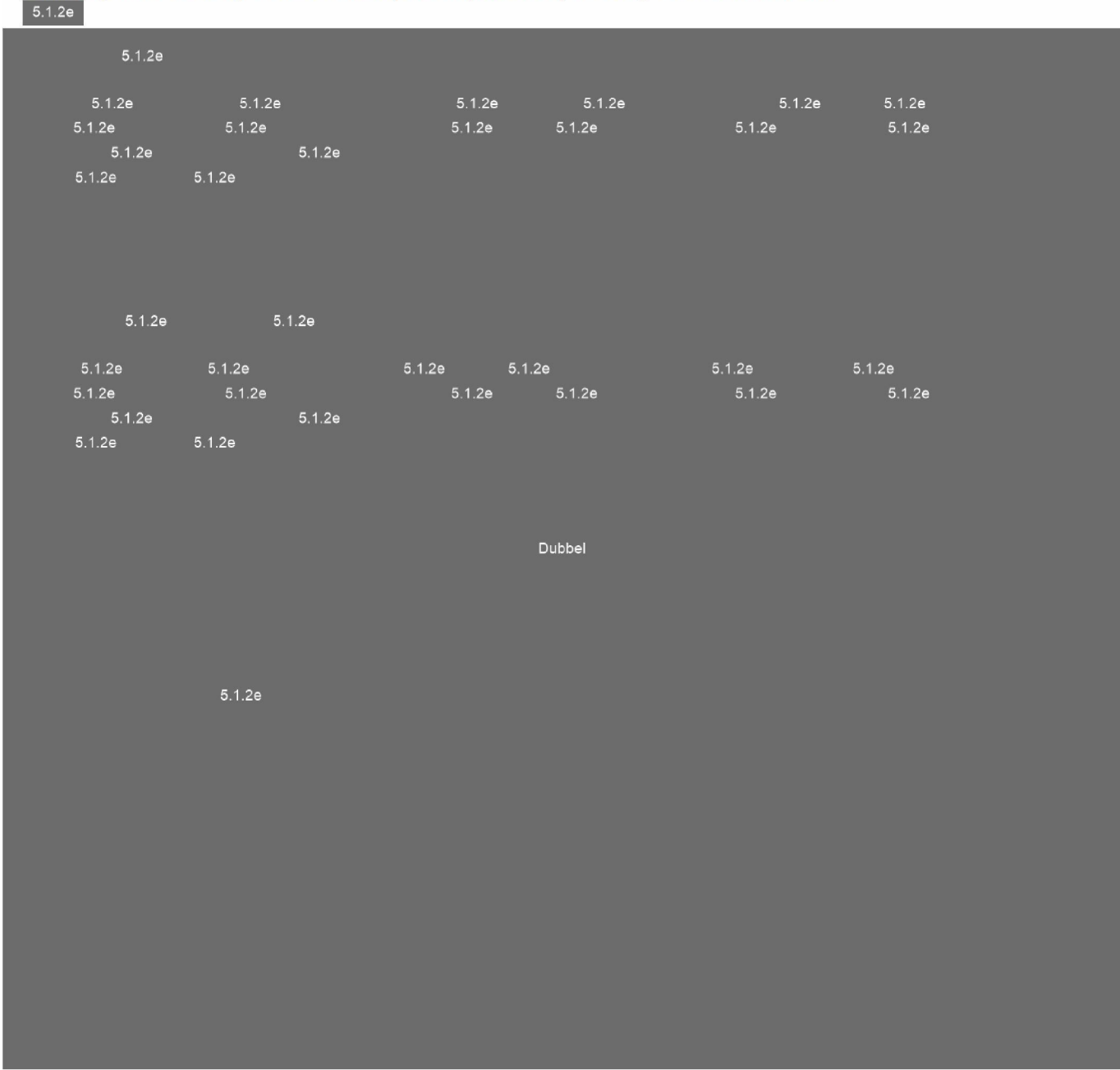


To: 5.1.2e | 5.1.2e | @rivm.nl]; 5.1.2e | 5.1.2e | @rivm.nl]; 5.1.2e | 5.1.2e | @rivm.nl]; 5.1.2e
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From: 5.1.2e
Sent: Fri 9/4/2020 9:16:47 AM
Subject: RE: Presentation on reported re-infections at COVID network call on Friday?
Received: Fri 9/4/2020 9:16:49 AM
[snps3pairs.pptx](#)

Ik heb de figuur up-to-date gemaakt met afname data
Analyses die hier aan vooraf gaan zijn met veel meer sequenties gedaan maar die allemaal tonen voegt niets toe aan de boodschap, het maakt het alleen maar onoverzichtelijker. Voor de figuur zijn uit de achtergrond dataset sequenties geselecteerd die illustreren dat er bij 5 van de 6 sequenties nauwverwante sequenties aanwezig zijn in dezelfde periode in Nederland. Alleen bij het laatst afgenomen monster van Saltro is geen nauwverwante sequentie aanwezig in de dataset, tot dat moment, die sequenties zijn we nog aan het verzamelen



2 - 2

Dubbel

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 5.1.2e < 5.1.2e @gmail.com>; 5.1.2e < 5.1.2e @erasmusmc.nl>

Subject: RE: Presentation on reported re-infections at COVID network call on Friday?

Dear 5.1.2e

We have several cases (n=4) with sequence evidence in the Netherlands in different patient backgrounds. For three cases comparative seq info was determined at RIVM and for one at Earsmus MC. Serology profiling is being done this week for some of those cases to provide more clarity also vs lingering co-infection vs re-infection. Also the phylogenetic trees might support the differentiation between co-infection and re-infection in some cases. Also we have a register with approx. 19 suspected cases that need to be more thoroughly looked at.

From RIVM we prefer to give the complete picture when we have all data collected. I'll get back to you if that will be possible this Friday, but probably not.

Best 5.1.2e

From: 5.1.2e < 5.1.2e @ecdc.europa.eu>

Sent: woensdag 26 augustus 2020 12:11

To: 5.1.2e @kuleuven.be; 5.1.2e @sciensano.be; 5.1.2e @sciensano.be; 5.1.2e @uzleuven.be;
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 < 5.1.2e @erasmusmc.nl>

Subject: Presentation on reported re-infections at COVID network call on Friday?

Importance: High

Dear colleagues,

We hope you are doing well. We saw the recent media reports of the first cases in Belgium and the Netherlands of confirmed SARS-CoV-2 re-infections (<https://nos.nl/artikel/2345309-ook-herbesmettingen-in-nederland-en-belgie.html>). We think it would be very interesting for colleagues in other countries to hear more about these cases and the investigations. We are therefore considering organising a brief COVID-19 network call on Friday around this issue and the wider implication of re-infections. Would you or colleagues be available to present your findings to the COVID network at the call on Friday 28 August at 14:00?

We would appreciate if you could let us know by the end of the day so we can send out invitations.

Apologies for the short notice,

Kind regards

5.1.2e

On behalf of the COVID-19 surveillance and microbiology groups



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