

**To:** [redacted] [redacted] [redacted] [redacted]@ecdc.europa.eu]; [redacted] [redacted] [redacted] [redacted]@rivm.nl]  
**From:** [redacted] [redacted] [redacted]  
**Sent:** Wed 1/13/2021 12:41:58 PM  
**Subject:** RE: Assay type info from molecular EQA  
**Received:** Wed 1/13/2021 12:42:00 PM  
[number of labs mol. EQA SARS-CoV-2 different assays.xlsx](#)

Hi [redacted]

Sorry, I misunderstood. I had previously extracted this information from the raw data. I quickly added it to the table (attached).

Best wishes,

[redacted]

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**From:** [redacted] [redacted] [redacted] [redacted]@ecdc.europa.eu>  
**Sent:** woensdag 13 januari 2021 13:31  
**To:** [redacted] [redacted] [redacted] [redacted]@rivm.nl>; [redacted] [redacted] [redacted] [redacted]@rivm.nl>  
**Subject:** RE: Assay type info from molecular EQA

Hi again,

Unfortunately it doesn't contain the number of labs that used each assay, but rather the list of assays used. Do you maybe have this info in an excel? or even raw data will be fine.

Many thanks,

[redacted]

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**From:** [redacted] [redacted] [redacted] [redacted]@rivm.nl>  
**Sent:** 13 January 2021 13:24  
**To:** [redacted] [redacted] [redacted] [redacted]@ecdc.europa.eu>; [redacted] [redacted] [redacted] [redacted]@rivm.nl>  
**Subject:** RE: Assay type info from molecular EQA

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Dear [redacted]

The data are presented in our manuscript. I attached the current 'early' version of the published pdf. Please check Supplementary Table 2. Hopefully it has the information that you are looking for.

Best wishes,

[redacted]

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**From:** [redacted] [redacted] [redacted] [redacted]@ecdc.europa.eu>  
**Sent:** woensdag 13 januari 2021 11:52  
**To:** [redacted] [redacted] [redacted] [redacted]@rivm.nl>; [redacted] [redacted] [redacted] [redacted]@rivm.nl>  
**Subject:** Assay type info from molecular EQA

Dear [redacted] and [redacted]

May I kindly ask you to share any data collected during our first molecular EQA round on the types of assays used? Apologies if you have shared this with me already, but I cannot retrieve it.

We were wondering if this could be useful to identify labs that are using S-gene based protocols.

Many thanks and best regards,

[redacted]



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Disease Prevention and Control (ECDC)**

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