

# European SARS-CoV-2 and Influenza Bioinformatics EQA 2024

Fields marked with \* are mandatory.



## European SARS-CoV-2 and Influenza virus Bioinformatics EQA 2024

### Participation Registration Form

**Please complete this online registration form before 12 February 2024 to register your laboratory's participation in the ESIB-EQA 2024.**

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#### Contact Information

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## Which SARS-CoV-2 EQA components will your laboratory participate in?

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\* SARS1 (Nanopore reads, 20 fastq files):

- ☒ Yes  
☐ No

\* SARS2 (Illumina reads, around 20 fastq files):

- ☐ Yes  
☒ No

\* SARS3 (clustering and classification, around 20 genomes):

- ☒ Yes  
☐ No

\* SARS4 (mutation detection, around 10 genomes):

- ☒ Yes

☐ No

## Which influenza virus EQA components will your laboratory participate in?

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\* INFL1/3/4 (Nanopore reads for 10 samples generated as part of wetlab EQA; clustering and classification; resistance prediction):

☒ Yes

☐ No

\* INFL2/3/4 (Illumina reads for 10 samples generated as part of wetlab EQA; clustering and classification; resistance prediction):

☐ Yes

☒ No

## Additional Comments

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In case you have particular points for us to keep in mind about your participation, please add them here.

**Thank you for completing this participation registration form!**

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## Contact

[Contact Form](#)

