

1/2022/APM Wrocław 20.07.2022

Information for contractors

Concerning: proceedings conducted in the framework of request for quotation, the subject of which is:

Part 1. Sequencing service for RNA isolated from canine lymphoma and leukemia cell lines. The order includes sequencing of 24 samples.

Question: Moreover do you need bioinformatic analysis either for RNAseq or/and whole ge-

nome?

Answer: No, additional bioinformatics analyzes are not required.

Question: Will the 24 samples be delivered at once or in tranches? If split, how many samples

per tranche and at what time interval will the tranches be sent?

Answer: Samples will be delivered at one time.

Question: Could you kindly confirm that you would like to submit total RNA as starting ma-

terial?

Answer: Yes the Ordering Party will submit total RNA

Question: Do you require library preparation with polyA selection (includes polyadenylated

messenger RNAs) or rRNA depletion (may also include long non-coding RNAs)?

Answer: the Ordering Party require polyA selection

Question: Could you kindly confirm that you request a total turnaround time of 21 work-

ing/business days?

Answer: Yes the Ordering Party confirms.

Question: Will the Ordering Party agree to change the order completion date in part 1 to: up

to 2 months from the passing of the quantitative and quantitative control of samples?

Answer: The Ordering Party authority does not agree.

Question: Part 2 concerns the preparation of the library for sequencing using the RNA-Seq technique (enrichment in poly (A)) from the RNA isolated from canine lymphoma and leukemia cell lines. RNASeq sequencing will not yield whole genomic sequencing results. In view of the above, will the Ordering Party consent to the deletion of the word "whole genomes" in the AGREEMENT model Part 1, par. 3 sec. 4: "Whole genome sequencing results will be made available to the Contractor in the cloud."

Answer: The Ordering Party agrees to amend to the Agreement model Part 1.

Question: Do you allow the possibility of using a different sequencing technology than the one offered by the Illumina platform? Do you allow the use of nanopore sequencing?

Answer: We do not allow any technology other than that specified in the specification of the inquiry.

Question: Do you need bioinformatics analysis?

Answer: No, we do not need bioinformatic analyses other than already specified.



Question: What is the sample type? Total RNA?

Answer: YES, it is total RNA.

Question: What Sequencing service you are looking for? Now we can provide: Eukaryotic Strand-specific Transcriptome Sequencing/mRNA sequencing /lncRNA sequenc-

ing/smallRNA sequencing

Answer: Eukaryotic Strand-specific mRNA sequencing

Question: You can tell us what the purpose of your research is and we can recommend a solution for you.

Answer: Differential gene expression analysis

Question: The submission deadline is relatively close we would like to inquire whether an extension of the deadline would be possible?

Answer: Yes, we are extending the deadline for submitting offers until 07/25/2022 until 15.00

Question: Would it be possible to negotiate the contract in the case of an awarded bid or does the non-acceptance of the contract lead to an exclusion?

Answer: The draft contract may be subject to minor changes only, the selected offer will be an attachment to the contract, so the changes may not apply to essential elements of the price, delivery date, warranty period, scope of the contract.

Question: Does the ordering party indicate the number of paired reads (20M paired reads) / unpaired per sample in the inquiry?

Answer: Yes, the ordering party indicates the number of paired readings (20M paired reads) / unpaired readings per sample

Part 2. De novo sequencing of staphylococcal genomes. Number of samples: 100.

Question: Regarding De novo sequencing do you need also DNA extraction from pure cultures?

Answer: No, purified, bacterial gDNA will be provided by us to the sequencing service.

Question: Moreover do you need bioinformatic analysis either for RNAseq or/and whole genome?

Answer: No, additional bioinformatic analyses are not requested.

Question: Does the Ordering Party plan to deliver all samples at once or will they be divided into tranches? If split, how many samples per tranche and at what time interval will the tranches be sent?

Answer: Samples will be delivered once

Question: Could you kindly confirm that you would like to submit purified genomic DNA? **Answer:** Yes, we will submit purified gDNA.

Question: Will all 100 samples be submitted in a single batch? If not, what is the expected batch number and size?

Answer: Yes, all 100 samples will be submitted in a single bath.



Question: Could you kindly confirm that you request a total turnaround time of 45 working/business days?

Answer: Yes, we request a total turnaround time of 45 working/business days.

Question: Do you require a separate de novo genome assembly for all 100 samples? Is Illumina 2x150bp sequencing sufficient for your project?

Answer: Yes, we request de novo genome assembly for all 100 samples

Question: If, no please let us know if you require 2x250bp, or 2x300bp sequencing. What insert length is required?

Answer: We request a 2x300bp sequencing.

Question: Will the Ordering Party agree to clarify the order completion date in part 2 for: up to 45 working days from the moment of passing the quantitative control of the samples? **Answer:** the Ordering Party does not agree to the changes.

Question: Do you allow the possibility of using a different sequencing technology than the one offered by the Illumina platform? Do you allow the use of nanopore sequencing? **Answer:** We do not allow any technology other than that specified in the specification of the inquiry.

Question: Do you need bioinformatics analysis?

Answer: No, we do not need bioinformatic analyses other than already specified.

Question: 1 What is the sample type? gDNA?

Answer: The sample will be bacterial genomic DNA

Question: The submission deadline is relatively close we would like to inquire whether an extension of the deadline would be possible?

Answer: Yes, we are extending the deadline for submitting offers until 07/25/2022 until 15.00

Question: Would it be possible to negotiate the contract in the case of an awarded bid or does the non-acceptance of the contract lead to an exclusion?

Answer: The draft contract may be subject to minor changes only, the selected offer will be an attachment to the contract, so the changes may not apply to essential elements of the price, delivery date, warranty period, scope of the contract.

Vice-Rector for Development

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