

Information for contractors 2

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: Do you have the expected set / path for the preparation of RNA libraries-
seq? eg are libraries to be prepared from mRNA or from total RNA?

Answer: total RNA

Question: are stranded libraries?

Answer: YES

Question: Are depleted libraries?

Answer: No, polyA enrichment

Question: Do you accept libraries to be made with the "Illumina Stranded Total
RNA Prep with Ribo-Zero Plus"? We use it routinely with RNAseq libraries with
amazing effects.

Answer: YES we accept

Question: RNA should meet the necessary batch requirements - concentrations (optimally,
at least less 20-25 ng / μ l), volume (at least 20-30 μ l), RINu > 8. We will assess how-bone of
the obtained material before entering the library preparation protocol. One-does the Em-
ployer ensure that the above-mentioned parameters of the samples are met? What inin case
the samples do not meet the batch criteria required in the protocol?

Answer: The samples will meet the batch requirements.

Question: Does the Employer accept the sequencing of the received bi-
libraries also in the mode of paired readings 2x75 bp?

Answer: YES

Question: Will the 24 samples be delivered at once or in tranches? If
divided, how many samples per tranche and at what time interval will the tranches be sent?

Answer: Samples will be provided at one time

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

Question: Will the Ordering Party accept the use of the Illumina NovaSeq6000 technology of
the length? 2 x 250 bp read?

Answer: We do not accept this solution.

Due to the large number of questions, we are extending the deadline for submitting offers to 26.07.2022 to 3 p.m.

Vice-Rector for Development

prof. dr hab. inż. Adam Szewczuk