

Minutes of Pre-Bid meeting for the tender whole genome sequencing (WGS) services

Bid reference: WGS – Tender ID: 2020_DARE_572668_1

Date and time of pre-bid meeting: 27-07-2020 at 11:15AM

Venue of pre-bid meeting: online, hosted at ICAR-NBPGR, New Delhi

The following bidder's representatives attended the pre-bid meeting:

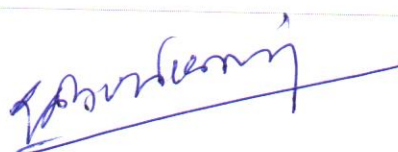
S. No	Name of prospective bidder / firm	Name of the representative
1	AgriGenome Labs Pvt Ltd, Cochin	Dr. Ram Nageena Singh
2	Nucleome Informatics Private Limited, Hyderabad	Mr. Dushyant Singh Baghel Mr. Bipin Mr. Abhishek

The following members from NBPGR and ICRISAT were present at the meeting:

1. Dr. Parimalan R, ICAR-NBPGR,
2. Dr. S. Rajkumar, ICAR-NBPGR,
3. Dr. Rakesh Singh, ICAR-NBPGR,
4. Dr. S. Sivasubramani, ICRISAT,

Minutes of the meeting were given herewith.

1. To set the tone for the meeting, the indenter, Dr. Parimalan R, clearly explained the technical requirements, expectations, and terms and conditions for the WGS tender process, as given in the tender document and invited the representatives, for their queries, if any.
2. Thereafter, the prospective bidders had their queries related to the technical details and terms and conditions given in the tender document. They are given in **Annexure A**, of this document
3. The queries raised, and the response made were categorized into technical, and terms and conditions type, and enclosed in **Annexure-A**, following this page.



Annexure – A

Clarifications regarding queries / suggestions received from the representatives of the firms who had participated in the pre-bid meeting for the WGS tender, were provided below in two sections, namely, **technical (pg. 2)**, and **terms and conditions (pg. 2)**.

TECHNICAL

S. No	Para / clause under reference as per the tender document	Content of para / clause under reference as per the tender document	Query / suggestions	Response
WGS – Tender ID: 2020 DARE 572668 1				
1	Pg. 14, point a), second bullet point.	Single molecule real time (SMRT) long-read 10kb+ sequencing (20-30Gb)	It is better to have 30Gb as minimal level of data.	Tender document provided the range from 20-30Gb data which can cover the suggested minimal level of data.
2	Pg. 14, point a), sixth bullet point.	Iso-seq sequencing for generating reference transcriptome... for each subspecies	What is the data requirement for this?	The data generated from SMRT cell for each tissue type will be used (irrespective of amount of data) for gene annotation purpose only, and a minimum of <u>10Gb</u> per SMRT cell is required.
3	Pg. 16, point 9.	The firm to provide the final data outputs in any standard external hard drive... other bioinformatics analysis.	Please clarify the details of other bioinformatics analysis.	The other bioinformatics analysis include, gene coordinates identification and sequence extraction for the three reference-based assemblies.

TERMS AND CONDITIONS

S. No	Para / clause under reference as per the tender document	Content of para / clause under reference as per the tender document	Query / suggestions	response
WGS – Tender ID: 2020 DARE 572668 1				
1	Pg. 16, point 6	Payment (100%) will be made in 30-45 days' time upon satisfactory receipt of the raw data and output results.	Will it be possible to give part payment?	Not possible.

[Handwritten Signature]