



भा.कृ.अनु.प. – राष्ट्रीय कृषि उपयोगी सूक्ष्मजीव ब्यूरो
ICAR - NATIONAL BUREAU OF AGRICULTURALLY IMPORTANT MICROORGANISMS
 कुसमौर, मऊ नाथ भंजन, मऊ. उत्तर प्रदेश-275103
 Kusmaur, Mau Nath Bhanjan, Uttar Pradesh- 275103
 (दूरभाष): (0547) 2530080, (फैक्स): (0547) 2530381,
 e-mail: nbaimicar@gmail.com, (web): www.nbaim.org.in / www.mgportal.org.in



F. No. 4-13/Rectt. of RA/SRF/NBAIM/2016-17/ **76**

Date 12/04/2016

WALK-IN-INTERVIEW

Selection will be held for following positions of Research Associate (01) and Senior Research Fellow (05) in different ICAR funded projects at **ICAR-National Bureau of Agriculturally Important Microorganisms, Maunath Bhanjan, Mau. U.P.** on **06.05.2016** at 10.00 AM. Interested candidates may appear before the selection committee along with their bio-data, photograph and self-attested copies of mark sheet and certificate {**Format for bio-data: Name of the Candidate; Father's Name; Date of Birth; Sex; Permanent Address; Corresponding Address; Email/Phone no.; Educational Qualification [Matriculation onwards with details of subjects taken, Year of Passing, Board/University and Marks obtained with percentage; Research experience (details, if any); List of publications and a write up of research work done earlier (200 words)]**}. The original documents in support should also be presented before the committee. The application can also be send through email ID. **nbaimicar@gmail.com**

S.No	Name of the projects	No. of Position	Essential/Desirable Qualification Required
1.	Reference and de Novo genome sequencing	One (RA) Ph.D. holder: 40000/-+HRA M.Sc. holder: 38000/-+HRA One (SRF) 25,000/- P.M. + H.R.A.	Essential Qualification: RA: Ph.D. in the relevant subject OR Master degree in the relevant subject (M.Sc. /M. Tech. in Biotechnology /Bioinformatics, M.Sc. in Ag. Microbiology/Life Sciences subjects/Biochemistry) with 4 year/ 5 years of Bachelor's degree having 1 st division or 60% marks or equivalent overall grade point average with atleast two years of research experience as evidenced from Fellowship/ Associateship/ Training / other engagements as circulated vide F.No. 2-9/ 2012-HRD dated 25.04.2014. SRF: Master degree in the relevant subject (M.Sc. /M Tech. in Biotechnology /Bioinformatics, M.Sc. in Ag. Microbiology/Life Sciences subjects/Biochemistry) with 4 year/ 5 years of Bachelor's degree or Ph.D degree in relevant subject. Candidates having post graduate degree in Basic Sciences with 3 years Bachelor's degree and 2 years Master degree should have NET qualification as mentioned in the OM SR/S9/Z-09/2012 of DST dated 21.10.2014. Desirable Qualification (for both positions): Working research experience in molecular biology/DNA or RNA sequencing/ experience in the assembly/ annotation/ analysis of sequencing data/ comparative genomic analysis etc. using different bioinformatics tools.
2.	Microbial Genomic Resource Repository (MGRR)	One (SRF) 25,000/- P.M. + H.R.A.	Essential Qualification SRF: Master degree in the relevant subject (M.Sc. /M Tech. in Biotechnology /Bioinformatics, M.Sc. in Ag. Microbiology/Life Sciences Subjets/Biochemistry) with 4 year/ 5 years of Bachelor's degree or Ph.D degree in relevant subject. Candidates having post graduate degree in Basic Sciences with 3 years Bachelor's degree and 2 years Master degree should have NET qualification as mentioned in the OM SR/S9/Z-09/2012 of DST dated 21.10.2014. Desirable Qualification : Working research experience in molecular biology/DNA or RNA sequencing/ experience in the assembly/ annotation/ analysis of sequencing data/ comparative genomic analysis etc. using different bioinformatics tools.
3.	ICAR-CRP Genomics	One (SRF) 25,000/- P.M. + H.R.A.	Essential Qualification SRF: Master degree in the relevant subject (M.Sc. /M. Tech. in Biotechnology /Bioinformatics, M.Sc. in Ag. Microbiology/Life Sciences subjects/Biochemistry) with 4 year/ 5 years of Bachelor's degree or Ph.D degree in relevant subject. Candidates having post graduate degree in Basic Sciences with 3 years Bachelor's degree and 2 years Master degree should have NET qualification as mentioned in the OM SR/S9/Z-09/2012 of DST dated 21.10.2014. Desirable Qualification: Working research experience in molecular biology/DNA or RNA sequencing/ experience in the assembly/ annotation/ analysis of sequencing data/ comparative genomic analysis etc. using different bioinformatics tools.

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