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To
The Editor,
Sir,

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I request that the following message may kindly be published in your esteemed daily:

**Hands-on Training Programme on Next Generation Sequencing Data Analysis
Inaugurated at TNAU**

The Department of Plant Molecular Biology and Bioinformatics, Centre for Plant Molecular Biology and Biotechnology (CPMB&B), Tamil Nadu Agricultural University (TNAU), Coimbatore, is organizing a four-day Hands-on Training Programme on Next Generation Sequencing (NGS) Data Analysis from 22 to 25 September 2026.

The programme brings together participants from various academic and research institutions across India and abroad, including Tamil Nadu Agricultural University, Elizabeth Creak Horticultural Technology Centre, University of Warwick, UK, ARM Laboratory, ICAR-National Research Centre for Banana, Kumaraguru College of Technology, University of Delhi South Campus, Shri Sakthikailassh Women's College, and CEGS Foundation – Centre for Eastern Ghats Studies. The training programme commenced with an Inaugural Session. Dr L.Arul, Professor and Head, Department of Plant Molecular Biology and Bioinformatics, delivered the Welcome Address and highlighted the scope of the training programme and the growing importance of NGS technologies in modern biological research. This was followed by an introduction of the participants.

The Inaugural Address was delivered by Dr N. Senthil, Director, CPMB&B, who emphasised the rapid growth of genomics data and the importance of acquiring skills in advanced sequencing technologies and data annotation. Dr V.Balasubramani, Controller of Examinations, TNAU, felicitated the gathering and highlighted the significance and wide-ranging applications of NGS technologies in contemporary scientific research, with relevant real-world examples. The inaugural session concluded with the Vote of Thanks proposed by Dr. M. Jayakanthan, Assistant Professor, Department of Plant Molecular Biology and Bioinformatics. The four-day training programme focuses on providing participants with practical exposure to NGS data analysis and computational approaches, enabling them to develop essential skills for handling and interpreting high-throughput sequencing data.

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