

Physics research as a hobby

I am amused to read the proposal put forward by P. Chaddah (*Curr. Sci.*, 2001, **81**, 868–869) to encourage physics research as an avocation or a hobby. He pleads that bright young persons may opt to do physics research in their spare time for its creative pleasure. I think he has the case of C. V. Raman at the back of his mind and he intends to create more Ramans by his utopian proposal.

In the same issue, T. V. Ramakrishnan (*Curr. Sci.*, 2001, **81**, 977–982) has summed up the Indian physics scenario with the remark that there is a catastrophic decline in the number of students

opting for physics at the undergraduate and the postgraduate levels, thus creating a vacuum in the research laboratories. It is most unfortunate that when India has created world-class facilities and infrastructure for physics research by setting up Inter-University Centres, viz. Nuclear Science Centre at Delhi, IUCCA at Pune and IUC-DAEF at Indore, the trend has changed suddenly in favour of information technology, management and engineering disciplines.

I do not reject Chaddah's idealism for promotion of avocational research *in toto*. My apprehensions are based on per-

sonal experience of thirty-eight years in experimental physics. However, it may be possible in theoretical physics to do something worthwhile making use of the internet, computers and library services, while experimental research is a full-time job and cannot be pursued as a part-time hobby.

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Rhizobia of the β -subclass of proteobacteria: A tale of losing the race

Rhizobia are the traditional soil bacteria capable of forming root or stem nodules on various leguminous plants, where they undertake symbiotic fixation of atmospheric nitrogen. Currently, they are divided into six genera with approximately 30 species, including *Allorhizobium*, *Azorhizobium*, *Bradyrhizobium*, *Mesorhizobium*, *Rhizobium* and *Sinorhizobium*. Phylogenetically, all these bacteria belong to the α -subclass of proteobacteria¹. Till recently, none of the bacteria belonging to the β -subclass of proteobacteria were known to form root or stem nodules on legumes. Moulin *et al.*² reported the ability of *Burkholderia* species, a member of the β -subclass of proteobacteria, to nodulate the African legumes *Aspalathus* and *Machaerium*. Later, Chen *et al.*³ described *Ralstonia taiwanensis* as the first member of the β -subclass of proteobacteria capable of nodulating two species of *Mimosa*, i.e. *M. pudica* and *M. diplo-tricha*. Careful reading of the two papers indicates that authors of these two papers were in touch with each other and yet, each group claims to be the first to report the α -subclass of proteobacteria as being involved in nodulation in legumes.

During the UNESCO/AICOPTAX sponsored practical training course on 'Molecular techniques in diversity, phylogeny and taxonomy of plant-associated bac-

teria' organized by us at the School of Biotechnology, Banaras Hindu University between 5 and 18 November 2000, one of the groups of trainees was assigned the task of characterizing the root nodule bacteria of *M. pudica*. The group consisted of Subhash Verma, Abha Mishra (BHU, Varanasi), Parag Vaishampayan (MACS, Pune) and Deepak Sharma (UDSC, New Delhi). The bacterial community present in the *Mimosa* nodules was found to be of only one type as characterized by PCR fingerprinting technique. Some of these isolates were further subjected to identification on the basis of their carbon source utilization ability, using the BIOLOG system which identified all of them as *Ralstonia eutropha*. The finding that some species of *Ralstonia*, rather than *Rhizobium*, could be present in the legume nodule was surprising and unexpected. Since we did not have an easy access to DNA sequencing facility at that time to identify these bacteria on the basis of their 16S rDNA sequence (one of the definitive methods for identifying bacteria), we could not follow up this interesting finding further. Keeping in line with the existing dogma¹ that no bacteria other than the members of Rhizobiaceae can nodulate legumes, we dismissed the intriguing observation possibly resulting from contamination.

Nevertheless, in view of the observation being unexpected and novel, we kept wondering if it could be true. After a few weeks when we thought of reinvestigating the bacteria from the *Mimosa* nodules, it was realized that the proper time for getting *M. pudica* plants with nodules was September–November. Therefore, we had to wait till September 2001 to isolate bacteria from the surface-sterilized nodules of *M. pudica* plants. The isolates were handled with utmost microbiological care and subjected again to BIOLOG way of identification. This, once again confirmed these bacteria to be *R. eutropha*. The 16S rDNA from these isolates was PCR-amplified and the RFLP of amplicons showed that all of them were genetically identical. The 16S rDNA amplicon was then sequenced and compared with the sequences in the databases. To our surprise we found 99% sequence identity with a very recently described *Ralstonia taiwanensis*, showing that bacteria isolated by us from *M. pudica* nodules was indeed closely related to *R. taiwanensis*. With respect to the 16S rDNA sequence, the next most similar species to our isolates was *R. eutropha*. When we started searching the description of this species, we failed to find any publication describing this species in the BLAST information database. While searching for publica-