

increased up to a maximum of 18. However, it is felt that wherever the deviation values are higher, it is necessary to subject more tooth samples to that particular system so as to obtain mean strength values with acceptable deviation ranges.

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Errata

Mechanism of ATP synthesis by protonmotive force

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The numbering of amino acid residues of the ϵ subunit of ATP synthase corresponds to *Escherichia coli* (and not to bovine heart mitochondria, as inadvertently implied). Thus, lines 36-37 on p. 718 should read, 'Further Ser-108 of the rotating ϵ subunit (*Escherichia coli* numbering) interacts covalently^{14,17} with Glu-381 (*Escherichia coli* numbering, corresponding to Glu-395 in bovine heart mitochondria) of β_E '. Similarly, in figure 1, the numbering of the important amino acid residues is for *Escherichia coli*, while the labelling in the figure is for mitochondria. Therefore, in Figure 1, the label, 'Inner membrane' should be substituted by 'Inner membrane/periplasm', while the label, 'Matrix' should be replaced by 'Matrix/cytoplasm'. The second line in the legend to Figure 1 should read, 'The important amino acid residues are shown'. These corrections do not in any way alter the results or conclusions of the communication.

Atomic Energy in India – 50 years

reviewed by

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The opening sentence of the concluding paragraph should read as, 'Underperformance of the PHWRs and the cost and time overruns in installing them may have been the underlying reasons for the decision to purchase two Russian reactors of 1000 MWe each'.