Corrigenda

Mutations in yeast Rad51 that partially bypass the requirement for Rad55 and Rad57 in DNA repair by increasing the stability of Rad51–DNA complexes

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Correction to: *The EMBO Journal* (2002) **21**, 3160–3170. doi:10.1093/emboj/cdf293

In the above paper, the authors reported the identification and characterization of six *rad51* alleles that suppress the ionizing radiation sensitivity of *rad55* and *rad57* mutants. Each of these alleles was reported to contain a single nucleotide change responsible for the suppression phenotype.

In recent work to further characterize one of these alleles, *rad51-L119P*, they discovered a second mutation at position 983 changing valine 328 to alanine. The V328A substitution alone is responsible for the suppression phenotype. Thus, all of the suppressing mutations map to a 32-residue region of Rad51 that is proposed to be involved in DNA binding.

This error does not alter the major conclusion of the paper, that Rad51 mutant proteins with increased affinity for DNA can partially bypass the requirement for Rad55 and Rad57 in DNA repair. Because the six *rad51* alleles recovered change only four residues, it seems likely that the screen is close to saturating and the suppression of *rad55* and *rad57* can only be achieved by alterations in this region of Rad51.

The authors apologize for this error and hope that it has not misled other researchers in the field.

A conserved catalytic residue in the ubiquitin-conjugating enzyme family

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Due to a typesetting error, an author name in the above article was published incorrectly. The correct author listing for this article should be as follows:

PY Wu, M Hanlon, M Eddins, C Tsui, RS Rogersm, JP Jensen, MJ Matunis, AM Weisman, C Wolberger and CM Pickart