

CORRIGENDUM

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A positive-feedback-based bistable 'memory module' that governs a cell fate decision

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Nature 426, 460–465 (2003)

In Box 1, equation (1) should read

$$\frac{d[A^*]}{dt} = \{\text{stimulus} \times ([A_{\text{tot}}] - [A^*])\} + f \frac{[A^*]^n}{K^n + [A^*]^n} ([A_{\text{tot}}] - [A^*]) - k_{\text{inact}}[A^*]$$

$$\text{Setting } \frac{d[A^*]}{dt} = 0, \text{ it follows that stimulus} = \frac{f \times [A^*]^n [A_{\text{tot}}] - k_{\text{inact}} K^n [A^*] - (f + k_{\text{inact}}) [A^*]^{n+1}}{([A^*] - [A_{\text{tot}}]) ([A^*]^n + K^n)}$$

which implicitly defines all of the possible steady state values of $[A^*]$ for any given value of the stimulus. The plots in Box 1 show only the stable steady states (the sections of the curves with positive slopes).

ERRATUM

doi:10.1038/nature06114

RNA-templated DNA repairFrancesca Storici, Katarzyna Bebenek, Thomas A. Kunkel,
Dmitry A. Gordenin & Michael A. Resnick*Nature* 447, 338–341 (2007)

In Figure 1, the column header indicating the repair frequencies should read “Repair frequency (Leu^+) $\times 10^{-7}$ ” rather than “Repair frequency (Leu^+) $\times 10^{-3}$ (per 10^7 viable cells)”.

CORRIGENDUM

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Structure of the *E. coli* signal recognition particle bound to a translating ribosomeChristiane Schaffitzel, Miro Oswald, Imre Berger, Takashi Ishikawa,
Jan Pieter Abrahams, Henk K. Koerten, Roman I. Koning & Nenad Ban*Nature* 444, 503–506 (2006); doi:10.1038/nature05182 (published online 29 October 2006)

During the preparation of the manuscript, we inadvertently mislabelled ribosomal protein L32 as ribosomal protein L18 when interpreting the density based on the 50S coordinates (PDB accession number 2AW4). Therefore, whenever L18 is mentioned in the text and in Figs 3 and 4, it should be considered to refer to ribosomal protein L32. Our results and conclusions are not affected.