CDKs in late G1 defines a "point of no return" after which Cdc6 synthesis cannot promote DNA replication in yeast. *Genes Dev.* **10**, 1516–1531 (1996).

11. Desdouets, C. *et al.* Evidence for a Cdc6p-independent mitotic resetting event involving DNA polymerase α . *EMBO J.* **17**, 4139–4146 (1998).
12. Perkins, G., Drury, L. S. & Diffley, J. F. X. Separate SCF^{CDC4} recognition elements target Cdc6 for proteolysis in S phase and mitosis. *EMBO J.* **20**, 4836–4845 (2001).
13. Nguyen, V. Q., Co, C. & Li, J. J. Cyclin-dependent kinases prevent DNA re-replication through multiple mechanisms. *Nature* **411**, 1068–1073 (2001).
14. Wilmes, G. M. *et al.* Interaction of the S-phase cyclin Clb5 with an 'RXL' docking sequence in the initiator protein Orc6 provides an origin-localized replication control switch. *Genes Dev.* **18**, 981–991 (2004).
15. Adams, P. D. *et al.* Retinoblastoma protein contains a C-terminal motif that targets it for phosphorylation by cyclin-cdk complexes. *Mol. Cell. Biol.* **19**, 1068–1080 (1999).
16. Chen, J., Saha, P., Kornbluth, S., Dynlacht, B. D. & Dutta, A. Cyclin-binding motifs are essential for the function of p21CIP1. *Mol. Cell. Biol.* **16**, 4673–4682 (1996).
17. Drury, L. S., Perkins, G. & Diffley, J. F. X. The cyclin dependent kinase Cdc28p regulates distinct modes of Cdc6p proteolysis during the budding yeast cell cycle. *Curr. Biol.* **10**, 231–240 (2000).
18. Pawson, T. Specificity in signal transduction: from phosphotyrosine-SH2 domain interactions to complex cellular systems. *Cell* **116**, 191–203 (2004).
19. Elia, A. E. *et al.* The molecular basis for phosphodependent substrate targeting and regulation of Plks by the Polo-box domain. *Cell* **115**, 83–95 (2003).
20. Cheng, K. Y., Lowe, E. D., Sinclair, J., Nigg, E. A. & Johnson, L. N. The crystal structure of the human polo-like kinase-1 polo box domain and its phosphopeptide complex. *EMBO J.* **22**, 5757–5768 (2003).
21. Durocher, D. *et al.* The molecular basis of FHA domain:phosphopeptide binding specificity and implications for phospho-dependent signaling mechanisms. *Mol. Cell* **6**, 1169–1182 (2000).
22. Halford, K. A. *Biochemical Analysis of Yeast Pre-Replicative Complex Assembly*. Thesis, Univ. College London (2003).
23. Li, J. J. & Kelly, T. J. Simian virus 40 DNA replication *in vitro*. *Proc. Natl Acad. Sci. USA* **81**, 6973–6977 (1984).

Supplementary Information accompanies the paper on www.nature.com/nature.

Acknowledgements We are grateful to members of our laboratory for discussions and for critical reading of the manuscript. We are also grateful to J. Gannon and T. Hunt for discussions as well as human CDK antibodies and p13^{suc1} beads. We thank L. Drury, K. Labib, J. Li, G. Perkins and S. Reed for yeast strains. We also thank N. O'Reilly and the Peptide Synthesis Facility at the London Research Institute. This work was supported by Cancer Research UK and the Human Frontier Science Program Organization. S.M. is supported by JSPS postdoctoral Fellowships for Research Abroad.

Competing interests statement The authors declare that they have no competing financial interests.

Correspondence and requests for materials should be addressed to J.E.X.D. (john.diffley@cancer.org.uk).

corrigendum

Cloning of adiponectin receptors that mediate antidiabetic metabolic effects

Toshimasa Yamauchi, Junji Kamon, Yusuke Ito, Atsushi Tsuchida, Takehiko Yokomizo, Shunbun Kita, Takuya Sugiyama, Makoto Miyagishi, Kazuo Hara, Masaki Tsunoda, Koji Murakami, Toshiaki Ohteki, Shoko Uchida, Sato Takekawa, Hironori Waki, Nelson H. Tsuno, Yoichi Shibata, Yasuo Terauchi, Philippe Froguel, Kazuyuki Tobe, Shigeo Koyasu, Kazunari Taira, Toshio Kitamura, Takao Shimizu, Ryozi Nagai & Takashi Kadowaki

Nature **423**, 762–769 (2003).

In this Letter, Fig. 1 is an illustration of the sorting procedure, rather than an original data set, which we did not explicitly describe. Because the *x*-axis was used for FACS analysis of both FITC- and PE-labelled cells, only the gated population was shown; these data were extracted from the analyses and inserted into a FACS profile. The same file of a single gated population (from Fig. 1c) was mistakenly misused for a part of the plot in Fig. 1b by inserting it into both Fig. 1b and Fig. 1c in the original figure. In order to clear up the confusion surrounding this figure, the original primary data are shown here in the Supplementary Information. The Supplementary Information also includes further details of our cell-sorting procedure, which were not provided in the published protocol. Although these corrections do not affect the conclusions of our paper, we apologize to readers who have been misled by these mistakes. □

Supplementary Information accompanies the corrigendum on www.nature.com/nature.