

# Reactive oxygen-mediated damage to a human DNA replication and repair protein

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Ultraviolet A (UVA) makes up more than 90% of incident terrestrial ultraviolet radiation. Unlike shorter wavelength UVB, which damages DNA directly, UVA is absorbed poorly by DNA and is therefore considered to be less hazardous. Organ transplant patients treated with the immunosuppressant azathioprine frequently develop skin cancer. Their DNA contains 6-thioguanine—a base analogue that generates DNA-damaging singlet oxygen (<sup>1</sup>O<sub>2</sub>) when exposed to UVA. Here, we show that this <sup>1</sup>O<sub>2</sub> damages proliferating cell nuclear antigen (PCNA), the homotrimeric DNA polymerase sliding clamp. It causes covalent oxidative crosslinking between the PCNA subunits through a histidine residue in the intersubunit domain. Crosslinking also occurs after treatment with higher—although still moderate—doses of UVA alone or with chemical oxidants. Chronic accumulation of oxidized proteins is linked to neurodegenerative disorders and ageing. Our findings identify oxidative damage to an important DNA replication and repair protein as a previously unrecognized hazard of acute oxidative stress.

Keywords: azathioprine; PCNA; reactive oxygen; skin cancer; thioguanine

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## INTRODUCTION

DNA lesions such as cyclobutane pyrimidine dimers and 6,4 pyrimidine photoproducts caused by the absorption of UVB (wavelength 290–320 nm) from sunlight, are important contributors

to skin cancer (Friedberg *et al*, 2006). UVA (320–400 nm)—the predominant energy source in incident sunlight—is absorbed poorly by DNA and is therefore considered to be less harmful. The danger from UVA is mainly indirect; it produces thymine:thymine cyclobutane photoproducts in cellular DNA (Young *et al*, 1998; Courdavault *et al*, 2005), probably through photosensitized reactions involving non-DNA chromophores. UVA also generates reactive oxygen species (ROS)—principally singlet oxygen (<sup>1</sup>O<sub>2</sub>) (Cadet *et al*, 2005)—which damage proteins, lipids and nucleic acids. Although essential for normal cellular processes, excessive ROS are hazardous and cause oxidative stress. Persistent oxidative DNA damage is linked to mutation and to human cancer (Al-Tassan *et al*, 2002); the accumulation of oxidized proteins is associated with neurodegenerative disorders (Floyd & Hensley, 2002).

Unlike canonical DNA bases, some base analogues such as 6-thioguanine (6-TG), are strong UVA chromophores. 6-TG, 6-mercaptopurine and azathioprine (Aza) are anticancer and immunosuppressant thiopurines (Aarbakke *et al*, 1997). Aza, which is used extensively as an immunosuppressant following organ transplantation and is increasingly prescribed for inflammatory disorders (Lichtenstein *et al*, 2006), causes 6-TG to accumulate in patients' DNA (Warren *et al*, 1995; Cuffari *et al*, 1996; O'Donovan *et al*, 2005). Organ transplant patients have a high risk of skin cancer for which sunlight exposure is a cofactor. DNA 6-TG interacts with UVA to generate ROS, which oxidize the 6-TG to guanine-6-sulphonate (G<sup>SO3</sup>; O'Donovan *et al*, 2005), a powerful block to DNA polymerases *in vitro* (O'Donovan *et al*, 2005; Zhang *et al*, 2006). Here, we report that combined 6-TG–UVA also causes a new oxidative modification of the DNA replication and repair protein, proliferating cell nuclear antigen (PCNA).

## RESULTS

The main DNA 6-TG photoproduct G<sup>SO3</sup>, blocks primer extension by Klenow fragment DNA polymerase. *In vivo*, doses of UVA as low as 10 kJ/m<sup>2</sup> cause a rapid inhibition of DNA synthesis in human cells containing DNA 6-TG (Zhang *et al*, 2006; data not shown). As Y family DNA polymerases bypass a replication-blocking template G<sup>SO3</sup> *in vitro* (O'Donovan *et al*, 2005), we examined whether 6-TG–UVA treatment induced the cellular

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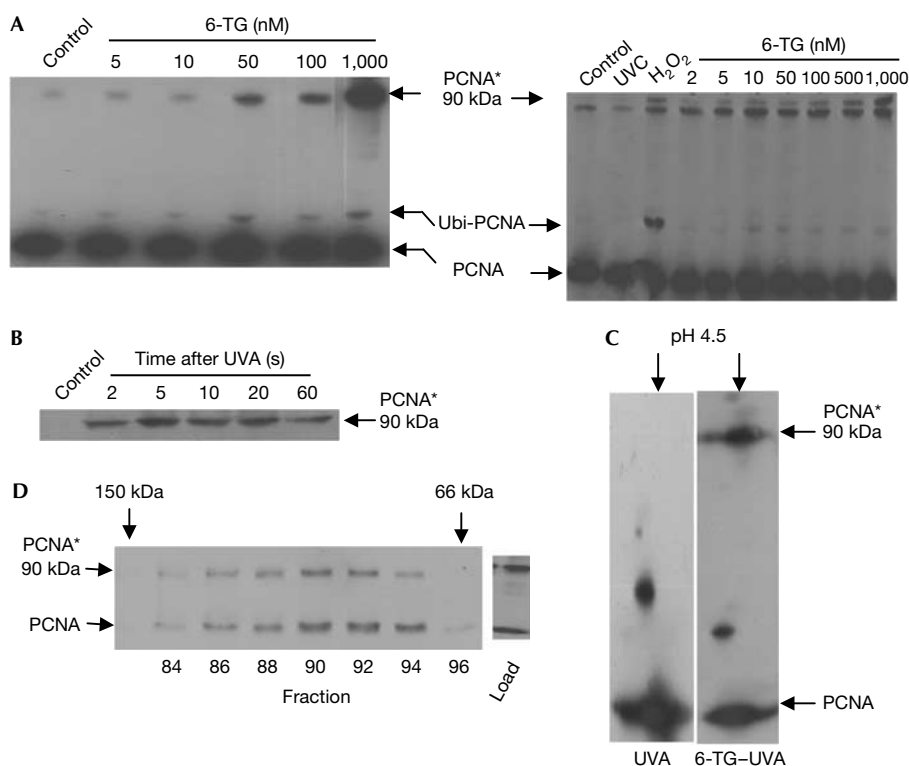
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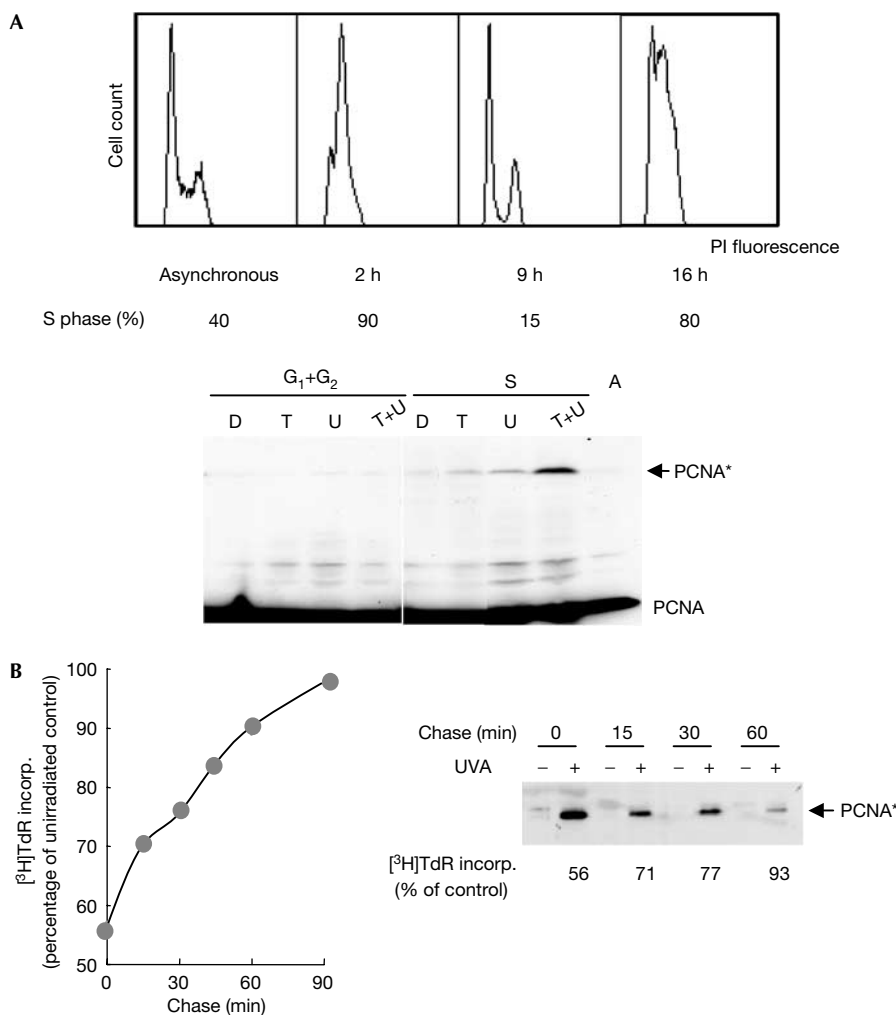


**Fig 1** | Formation and characterization of a new high-molecular-weight PCNA species. (A) CCF-CEM (left panel) or HCT116 (right panel) cells were grown for 48 h in medium containing 6-TG at the concentrations indicated and irradiated with 3 kJ/m<sup>2</sup> UVA. Both cell lines are mismatch repair defective and accumulate significant levels of DNA 6-TG ( $\geq 0.1\%$  of DNA guanine) without toxic effect; control cells were unirradiated. HCT116 cells grown in the absence of 6-TG were also treated with H<sub>2</sub>O<sub>2</sub> (0.03%, 2 min) or 10 J/m<sup>2</sup> UVC. Chromatin-associated proteins extracted 6 h after irradiation, or immediately after H<sub>2</sub>O<sub>2</sub> treatment, were analysed by western blotting with PC10 PCNA antibodies. The position of monoubiquitinated PCNA (Ubi-PCNA) is indicated. Immunoprecipitation experiments followed by western blotting with ubiquitin antibodies confirmed the presence of ubiquitin. The 90 kDa PCNA complex (PCNA\*) is also shown. This complex migrated slightly more slowly than a nonspecific band that is more prominent in HCT116 than in CCF-CEM extracts. (B) Kinetics of PCNA\* formation. HCT116 cells grown for 48 h in 1  $\mu$ M 6-TG were irradiated with 1 kJ/m<sup>2</sup> UVA at a high dose rate (0.5 kJ/m<sup>2</sup>/s). Extracts prepared at the time points indicated were analysed by western blotting with PC10. (C) PCNA\* *pI*. Chromatin-associated proteins from CCF-CEM cells grown in the presence or absence of 6-TG (1  $\mu$ M, 48 h), as indicated, and UVA irradiated (10 kJ/m<sup>2</sup>) were analysed by two-dimensional IEF/PAGE analysis followed by western blotting with PC10. (D) Size-exclusion chromatography. Chromatin-associated proteins extracted from CCF-CEM cells treated with 6-TG (1  $\mu$ M, 48 h) and UVA (50 kJ/m<sup>2</sup>) were separated by chromatography on AcA44 precalibrated with alcohol dehydrogenase (150 kDa), BSA (66 kDa) and carbonic anhydrase (30 kDa). Fractions containing PCNA were identified by dot blot and analysed further by western blotting with PC10. 6-TG, 6-thioguanine; Control, unirradiated; IEF/PAGE, isoelectric focusing-polyacrylamide gel electrophoresis; PCNA, proliferating cell nuclear antigen; UVA, ultraviolet A.

PCNA monoubiquitination associated with polymerase switching from replication to lesion bypass (Kannouche *et al*, 2004). Western blots showed an altered PCNA form that was consistent with monoubiquitination (Ubi-PCNA; Fig 1A). Strikingly, the same blots revealed a new high-molecular-weight PCNA species (PCNA\*) of approximately 90 kDa (Fig 1A), which was particularly abundant in partly tetraploid CCF-CEM cells. It was also formed in HCT116 (Fig 1A), A2780 and Raji cells (supplementary Fig S1a online). Immunoprecipitated PCNA\* was unreactive towards a ubiquitin antibody, eliminating polyubiquitination (data not shown). It was consistently observed as a discrete band rather than the smear that would be expected from DNA-protein crosslinking. PCNA\* formed rapidly (Fig 1B) and persisted for several hours. It was not detectable after 3 kJ/m<sup>2</sup> UVA or 6-TG treatment alone (Fig 1A and see below). The *pI* of PCNA\* was approximately 4.5

and indistinguishable from that of a PCNA monomer (Fig 1C). On size-exclusion chromatography under nondenaturing conditions, PCNA\* coeluted with native trimeric PCNA with a *V<sub>e</sub>* corresponding to a globular protein of approximately 100 kDa (Fig 1D). We conclude that PCNA\* is a covalent PCNA complex with a native mass and charge similar to that of an unmodified PCNA trimer. Thus, in addition to introducing potentially replication-blocking DNA lesions, 6-TG-UVA causes a rapid covalent modification of an important DNA replication and repair protein.

To examine whether PCNA\* formed at active replication forks, where it is likely to be more hazardous, we compared the effect of 6-TG-UVA in S-phase and non-S-phase cells. Synchronized HCT116 cells released into S phase were allowed to incorporate 6-TG for 60 min. Fluorescence-activated cell sorting (FACS) analysis confirmed that more than 90% of the cells were in S phase



**Fig 2** | Formation of a new high-molecular-weight PCNA species at active replication forks. (A) The 90 kDa PCNA complex (PCNA\*) in non-S-compared with S-phase cells. Synchronized HCT116 cells were 6-TG labelled (60 min, 10 μM) 2 h after release from a double thymidine block. At this time ≥90% of cells were in S phase, as determined by FACS of propidium iodide (PI)-stained cells (upper panel). Cell-cycle distributions were assigned using Watson Pragmatic. Aliquots of 6-TG-treated cells were UVA irradiated (10 kJ/m<sup>2</sup>) either 9 h (when 85% of cells were in G<sub>1</sub> + G<sub>2</sub> phase) or at 16 h (>80% S phase) after 6-TG treatment. Cell extracts were analysed by western blots probed with PC10 (lower panel); the positions of PCNA and PCNA\* are indicated. Left: G<sub>1</sub> + G<sub>2</sub> cells; right: S-phase cells. (B) 6-TG pulse-chase. Exponentially growing CCF-CEM cells were pulse labelled with 6-TG (10 μM, 15 min), followed by a return to normal growth medium without 6-TG. At the time points indicated, aliquots of cells were irradiated with UVA (10 kJ/m<sup>2</sup>). Immediately after irradiation, one sample of each aliquot was used to monitor DNA replication by [<sup>3</sup>H]thymidine incorporation, which is expressed as a percentage of that in unirradiated control. The remaining cells were extracted and proteins analysed by western blotting with PC10. 6-TG, 6-thioguanine; A, asynchronous untreated cells; D, double thymidine block only; FACS, fluorescence-activated cell sorting; PCNA, proliferating cell nuclear antigen; T, 6-TG only; T + U, 6-TG + UVA; U, UVA only; UVA, ultraviolet A.

during 6-TG treatment and that incorporated 6-TG did not affect subsequent cell-cycle progression. The first S phase was complete (≤15% S-phase cells) 9 h after 6-TG labelling. By 16 h, approximately 80% of 6-TG-treated and untreated cells had entered a second S phase (Fig 2A). Extracts were prepared from cells exposed to 10 kJ/m<sup>2</sup> UVA at 9 h (G<sub>1</sub> + G<sub>2</sub>) or 16 h (S) after 6-TG treatment. Western blots indicated that despite similar levels of PCNA, irradiated G<sub>1</sub> + G<sub>2</sub>-phase cells contained little PCNA\*, whereas the complex was readily detectable in cells irradiated in S phase (Fig 2A). These observations are consistent with the formation of PCNA\* from PCNA engaged in replication; pulse-chase

experiments confirmed this possibility. UVA irradiation of asynchronous cells, immediately after a 6-TG pulse, caused significant PCNA crosslinking (Fig 2B; supplementary Fig S1b online) and inhibited DNA synthesis. If irradiation was delayed to allow replication forks (with PCNA) to migrate away from the 6-TG-containing DNA, the amount of PCNA\* declined together with the inhibition of DNA synthesis. DNA itself and active replisome proteins are the most likely targets of the highly unstable ROS generated from DNA 6-TG. A short stretch of DNA 6-TG close to the replisome is sufficient for a low dose of UVA to generate replication-blocking DNA lesions and PCNA\*.

The UVA–6-TG interaction generates  $^1\text{O}_2$  (Zhang *et al*, 2006), a form of ROS implicated in protein crosslinking (Shen *et al*, 1996), and PCNA\* was formed when chromatin proteins were treated with rose bengal and visible light—an acknowledged source of  $^1\text{O}_2$  (supplementary Fig S2 online). UVA also generates  $^1\text{O}_2$  in cells (Cadet *et al*, 2003) through absorption by non-DNA chromophores. Although the UVA doses ( $<10\text{ kJ/m}^2$ ) that we used did not generate PCNA\* in the absence of DNA 6-TG, the complex was readily detected after higher—although still modest—doses of UVA alone (Fig 3A; supplementary Fig S1 online). We examined whether the trimeric structure of PCNA juxtaposes amino acids susceptible to crosslinking by  $^1\text{O}_2$ . Inspection of the intersubunit region shows several candidate residues (Fig 3B). The proximity of His 153 and Lys 77 is particularly intriguing, as histidine–lysine crosslinking by  $^1\text{O}_2$  is highly favoured (Au & Madison, 2000). To investigate the possible involvement of His 153, we took advantage of the replacement of this otherwise highly conserved residue by glutamine in insects and *Xenopus*. Although PCNA\* is detectable after  $30\text{ kJ/m}^2$  UVA in human cells, it was not detected in extracts of SF9 (*Spodoptera frugiperda*) insect cells irradiated with up to  $300\text{ kJ/m}^2$  UVA (Fig 3C); it was not possible to examine PCNA\* formation by 6-TG–UVA in SF9 cells, which do not incorporate 6-TG. PCNA\* was not formed following irradiation of replicating or nonreplicating *Xenopus* egg extracts (Fig 3C). This indicates that His 153 is required for photochemical PCNA crosslinking. PCNA can be crosslinked by aldehydes through the conserved Lys 110 residue (Wenz *et al*, 1998; Balajee & Geard, 2001; Naryzhny *et al*, 2005). Formaldehyde treatment of SF9 cells and *Xenopus* egg extracts generated high-molecular-weight PCNA complexes (Fig 3C), confirming the presence of trimeric PCNA. Formaldehyde and UVA (or UVA–6-TG) produced different PCNA complexes. This might reflect differences in the crosslinked amino acids (supplementary Fig S2b online). These findings implicate PCNA His 153 in the photochemical formation of intersubunit covalent bonds, probably through Lys 77.

It has been suggested that aldehydes generate a covalent circular PCNA trimer of approximately 86 kDa (Wenz *et al*, 1998; Balajee & Geard, 2001; Naryzhny *et al*, 2005); the size and *pI* of PCNA\* are consistent with this possibility. In preliminary analyses, immunoprecipitated PCNA\* co-purified with polypeptides of approximately 83 kDa on SDS–polyacrylamide gel electrophoresis (supplementary Table 1 online). No co-purifying protein of 54 kDa was identified, indicating that PCNA\* does not comprise a 29 kDa PCNA monomer and a non-PCNA protein. The observation is consistent with covalent crosslinking between the three PCNA subunits; however, non-circular PCNA multimers are not excluded by our findings.

## DISCUSSION

Ultraviolet in sunlight, of which more than 90% is UVA, is a significant risk factor for skin cancer in organ transplant patients (Euvrard *et al*, 2003). Aza immunosuppression produces DNA 6-TG—a source of  $^1\text{O}_2$  on UVA exposure. Photochemical PCNA crosslinking at replication forks is a previously unsuspected sunlight-related hazard for these patients. Higher, but nevertheless still moderate, doses of UVA that generate ROS through cellular photosensitizers also form PCNA\*. PCNA might be particularly susceptible to oxidative stress. In agreement with this possibility, the chemical oxidants  $\text{H}_2\text{O}_2$  and  $\text{KBrO}_3$  also crosslinked PCNA (supplementary Fig 2c online; Fig 1A).

PCNA improves the speed and processivity of DNA polymerases. It provides a scaffold for DNA–protein interactions in cellular signalling and DNA replication. It also participates in DNA mismatch repair (reviewed by Jiricny, 2006), and in nucleotide and base excision repair (reviewed by Friedberg *et al*, 2006). The extent to which the formation of PCNA\* compromises these important functions remains to be determined. It might have a bearing on curious and unexplained differences in the efficiency of thymine:thymine cyclobutane dimer removal. These DNA photo-products are reportedly excised less efficiently by nucleotide excision repair if they are induced by UVA compared with UVB (Courdavault *et al*, 2005). It is tempting to speculate that oxidative damage to PCNA or other repair proteins might contribute to this inefficiency. PCNA\* is an oxidation-related modification that depends on a reactive histidine in the intersubunit domain. A highly favoured reaction might be compatible with a rapid joining of all PCNA subunits to form the proposed covalent circular trimer (Wenz *et al*, 1998; Balajee & Geard, 2001; Naryzhny *et al*, 2005). PCNA\* is easily detected by its significantly changed electrophoretic mobility. It seems unlikely that this is the only form of protein damage caused by  $^1\text{O}_2$  generated in DNA, and PCNA\* might be a marker for general, perhaps more subtle, oxidative modifications of replication proteins. Chronic protein oxidation is implicated in ageing and in some neurological disorders (Floyd & Hensley, 2002). The relationships between oxidative DNA damage—the formation, repair and biological impact of oxidized DNA bases—and cancer have been extensively studied. Our findings indicate that examination of the impact of oxidative stress on the efficiency of DNA replication and repair proteins might also be warranted.

## METHODS

**Cells and cell culture.** A2780 SC5 ovarian carcinoma cells (Zhang *et al*, 2006) and HCT116 colorectal carcinoma cells were grown in DMEM, Raji Burkitt's lymphoma and CCF-CEM leukaemia cells in RPMI, and SF9 cells in Grace's medium, all supplemented with 10% FCS. HCT116 cells were synchronized by double thymidine block, and Raji cells by 2% dimethyl sulphoxide (Fiore *et al*, 2002). Replication was measured by [ $^3\text{H}$ ]TdR labelling ( $1\text{ }\mu\text{Ci/ml}$ , 15 min,  $1 \times 10^6$  cells), followed by the determination of trichloroacetic acid-insoluble radioactivity.

*Xenopus* egg extracts (Kubota & Takisawa, 1993) were induced to replicate by the addition of sperm nuclei.

**Chemicals and reagents.** Chemicals were obtained from Sigma-Aldrich (Poole, UK). 6-TG in 0.1 N NaOH was diluted in culture medium before cell treatment. PCNA (PC10) and ubiquitin (P4D1) antibodies were obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

**Irradiation.** UVA from a UVH 253 lamp (UVLIGHT Technology, Birmingham, UK), fitted with a black glass filter to remove wavelengths less than 320 nm, was routinely delivered at a dose rate of  $0.1\text{ kJ/m}^2/\text{s}$ . UVC (254 nm) was delivered from a germicidal lamp with a dose rate of  $1\text{ J/m}^2/\text{s}$ .

**Chemical treatment.** Cells were treated with formaldehyde in PBS. For  $\text{H}_2\text{O}_2$  and  $\text{KBrO}_3$  treatments, cells were in growth medium.

**Cell extracts and analysis.** A total of  $1 \times 10^6$  cells were lysed in  $100\text{ }\mu\text{l}$  cold 1% NP-40, 10 mM NaF, 1 mM  $\text{Na}_3\text{VO}_4$ , 1 mM  $\text{Na}_4\text{P}_2\text{O}_7$  plus protease inhibitor cocktail (Roche Diagnostics, Lewes, UK). After 1 h incubation on ice, insoluble material was removed by centrifugation (30 min, 11,000g) and discarded.





For chromatin-enriched extracts (Balajee & Geard, 2001), cells were suspended in cold 10 mM Tris-HCl (pH 7.5), 2.5 mM MgCl<sub>2</sub>, 0.5% NP-40 plus protease inhibitors. After 8 min incubation on ice, extracts were centrifuged (750g, 5 min) and the pellet was resuspended in chromatin extraction buffer (25 mM Na<sub>4</sub>P<sub>2</sub>O<sub>7</sub> pH 7.4, 0.5 M NaCl, 1 mM EDTA, 0.5% Triton X-100, 10% glycerol, 5 mM MgCl<sub>2</sub> and protease inhibitors). After a further 20 min incubation on ice, the extracts were centrifuged (5 min, 12,000g) and the supernatant was retained.

Proteins were resolved on 8% SDS gels for western blotting. Complexes were detected by using enhanced chemiluminescent substrate and visualized on Hyperfilm (Amersham Pharmacia, Little Chalfont, UK). Protein size estimates are based on Prestained Low Molecular Weight Range Markers (Bio-Rad, Hemel Hempstead, UK).

**Supplementary information** is available at *EMBO reports* online (<http://www.emboreports.org>).

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