

NOTES

Physiological Responses of the Hyperthermophilic Archaeon “*Pyrococcus abyssi*” to DNA Damage Caused by Ionizing Radiation

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The mechanisms by which hyperthermophilic *Archaea*, such as “*Pyrococcus abyssi*” and *Pyrococcus furiosus*, survive high doses of ionizing gamma irradiation are not thoroughly elucidated. Following gamma-ray irradiation at 2,500 Gy, the restoration of “*P. abyssi*” chromosomes took place within chromosome fragmentation. DNA synthesis in irradiated “*P. abyssi*” cells during the DNA repair phase was inhibited in comparison to nonirradiated control cultures, suggesting that DNA damage causes a replication block in this organism. We also found evidence for transient export of damaged DNA out of irradiated “*P. abyssi*” cells prior to a restart of chromosomal DNA synthesis. Our cell fractionation assays further suggest that “*P. abyssi*” contains a highly efficient DNA repair system which is continuously ready to repair the DNA damage caused by high temperature and/or ionizing radiation.

The hyperthermophilic *Archaea* thrive in geothermal environments with optimal growth temperatures approaching the boiling point of water (25). Under these conditions, DNA is expected to be unstable due to the accumulation of many different types of damage (e.g., hydrolytic depurination, deamination of cytosine and adenine, and single- or double-strand breakage [DSB]; note that ionizing radiation and/or genotoxic chemicals can also induce similar damage [13, 18]). During the last few years, it has been shown that some hyperthermophilic archaea, such as *Desulfurococcus amylolyticus* and *Thermococcus stetteri*, are resistant to ionizing radiation (17). More recently, DiRuggiero and collaborators have reported that a hyperthermophilic archaeon, *Pyrococcus furiosus*, did not show any loss of viability at a dose of 2,000 Gy (gamma radiation), while the viability of *Escherichia coli* was already drastically decreased at a dose of 100 Gy (9). We also showed earlier that no specific DNA protection mechanisms exist in “*Pyrococcus abyssi*” (12), suggesting that an efficient mechanism(s) for repair of radiation-induced DNA damage must exist in *Pyrococcus* species. Presumably, these yet to be elucidated mechanisms, and their possible coordination with DNA replication, are critical for the accurate maintenance of genetic information in the hyperthermophilic *Archaea*.

We were previously unable to find any evidence for radiation-induced gene expression in *Pyrococcus* species by use of two-dimensional sodium dodecyl sulfate-polyacrylamide gel electrophoresis (12), and two *recA/rad51*-like genes, *radA* and *radB*, of *Pyrococcus* sp. are constitutively expressed in *P. furiosus* (15). In this work we have further investigated physiological responses of “*P. abyssi*” to ionizing radiation by studying the kinetics of DSB repair and by investigating other possible responses to massive DNA damage in “*P. abyssi*” cells. Altogether, our results indicate that, despite the fact that the bacterium *Deinococcus radiodurans* and the archaeon “*P. abyssi*” grow under very different environments and contain nonhomologous repair and replication proteins, these two organisms respond in a similar fashion to DNA damage caused by ionizing radiation (1, 2).

Radiation-induced growth delay and DNA damage. Previous experiments indicated that “*P. abyssi*” cells were able to fully recover growth within 2 h after gamma-ray irradiation at 2,500 Gy under optimal growth conditions in YPS medium (11) (data not shown), suggesting that irradiated cells efficiently repaired massive DNA damage within this repair period (12). To investigate the fate of damaged chromosomes during this period, DSB analysis was performed by the pulsed-field gel electrophoresis technique as described earlier (12) to visualize the DNA restoration of fragmented chromosomes immediately after irradiation of “*P. abyssi*” cells.

Our results showed that the chromosomal DNA of “*P. abyssi*” was fragmented following irradiation (Fig. 1). Interestingly, chromosomal DNA degradation appeared reproducibly

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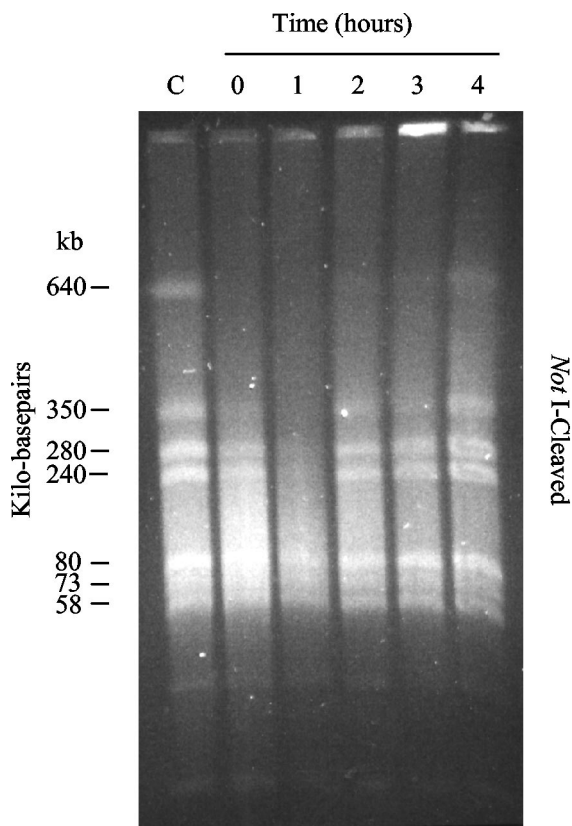


FIG. 1. Pulsed-field gel electrophoresis analysis of “*P. abyssi*” chromosomal DNA following gamma irradiation of 2,500 Gy. Irradiation was performed on ice with a ^{137}Cs gamma-ray source at a rate of 60 Gy/min (Institut Curie, Orsay, France). Each lane corresponds to DNA extracted from 10^7 cells/ml that was visualized by ethidium bromide staining (5 $\mu\text{g/ml}$). Lane C, control culture (without gamma irradiation); lanes 0 to 4, cells were subjected to 0, 1, 2, 3, and 4 h, respectively, of incubation at 96°C after irradiation. Chromosomal DNA was digested with *NotI*, and the sizes of the different *NotI* macrorestriction fragments are indicated in kilobase pairs (left side).

more extensive at 1 h postirradiation under optimal growth conditions (Fig. 1; 1 h) than directly after irradiation (Fig. 1; 0 h). This observation could indicate the presence of active nuclease activities in “*P. abyssi*” cells that function in DNA repair (4, 14, 23, 24). Alternatively, this postirradiation DNA degradation could result from cleavage by heat of the topologically relaxed fragmented chromosomal DNA. Our analyses also showed that the fragmented chromosomal DNA was fully restored within 2 h after gamma-ray irradiation at 2,500 Gy under optimal growth conditions, which is clearly visible by the reappearance of the largest genomic band of 640 kbp. Note also that the pattern of *NotI*-digested chromosomal DNA was identical before and after radiation and was in good agreement with earlier data (9, 10). Thus, “*P. abyssi*” is able to efficiently repair DSBs, similar to what has been observed for *P. furiosus* (9) and *Deinococcus radiodurans* (2, 5, 6, 7, 20).

DNA repair and DNA synthesis are uncoupled during the repair period. In *Deinococcus radiodurans*, DNA replication is known to cease following irradiation (8, 21), thus prompting us to investigate whether DNA repair and replication could also be coupled in our model organism. “*P. abyssi*” cells efficiently incorporate [^3H]uracil (1 $\mu\text{Ci/ml}$ [Amersham]; specific activity,

16.2 Ci/mmol) into DNA, a property which has been used to monitor DNA replication in this organism (22). To investigate whether ionizing irradiation inhibits DNA synthesis in “*P. abyssi*,” cells were irradiated at 2,500 Gy during stationary phase and were allowed to recover growth under optimal conditions. DNA synthesis in “*P. abyssi*” cells was monitored at different time points by measuring the incorporation of ^3H -labeled uracil into alkaline-resistant trichloroacetic acid (TCA) precipitates (5% ice-cold TCA solution and 20 mM sodium pyrophosphate mixture) in both irradiated and nonirradiated cells (22). As shown in Fig. 2A, irradiated cultures showed a delay in DNA synthesis of approximately 60 to 90 min when cells were irradiated with 2,500 Gy. The nonirradiated culture did not show a similar lag phase, indicating that the observed delay is somehow related to DNA repair. The notion of repair-coupled DNA synthesis arrest is also supported by the observation that the measured end point of the replication delay corresponds well with the time of growth recovery and chromosome restoration. In the stationary-phase control culture (Fig. 2A) and exponential-phase cells (22), the incorporation into DNA of ^3H -labeled uracil starts after 15 to 20 min (a similar delay is also observed for exponential-phase cells [Hannu Myllykallio, unpublished observations]) when cells are incubated at 96°C. These results suggest that DNA replication proteins, which are required for uracil incorporation into DNA, exist as functional forms in “*P. abyssi*” cells in exponential- and stationary-phase cultures.

Export from the cells of damaged and undamaged DNA. To investigate the fate of the damaged DNA in irradiated “*P. abyssi*” cells, cultures in the stationary growth phase were labeled with [^3H]uracil as described above, washed two times with artificial seawater, and then suspended in fresh growth medium. Irradiation was performed at a sublethal dose of 2,000 Gy to avoid detection of the DNA in the supernatant fraction due to lysis by cell death. Following irradiation, cultures were allowed to recover under optimal growth conditions. One-milliliter aliquots of cultures were removed at intervals corresponding to each incubation time point. The amount of radioactivity in alkaline-resistant form in the culture supernatants was measured using TCA precipitation and scintillation counting.

In the case of irradiated cultures, the radioactivity released in the supernatant increased after the first 30 min postirradiation and reached its maximal level after 90 min of incubation (Fig. 2B). The amount of radioactivity in TCA precipitates started to decrease 2 h after irradiation, and it became close to the control values at 2.5 and 3 h postirradiation. “*P. abyssi*” cells thus actively expelled damaged DNA (note that free nucleotides are not precipitated under these experimental conditions) into the growth medium just before a restart of DNA synthesis. As this material can be precipitated using TCA, the expelled DNA corresponds to DNA fragments longer than 20 nucleotides, and our preliminary estimation reveals that 0.3 to 0.4% of labeled genomic DNA is released from the irradiated “*P. abyssi*” cells. The increase in supernatant radioactivity appears higher than has been observed for radiation-resistant “*Chroococcidiopsis*” cells (4). The mechanisms generating and transporting these DNA fragments out of the cell are currently unclear (3, 26). They could possibly correspond to DNA molecules generated during the repair processes during enzymatic

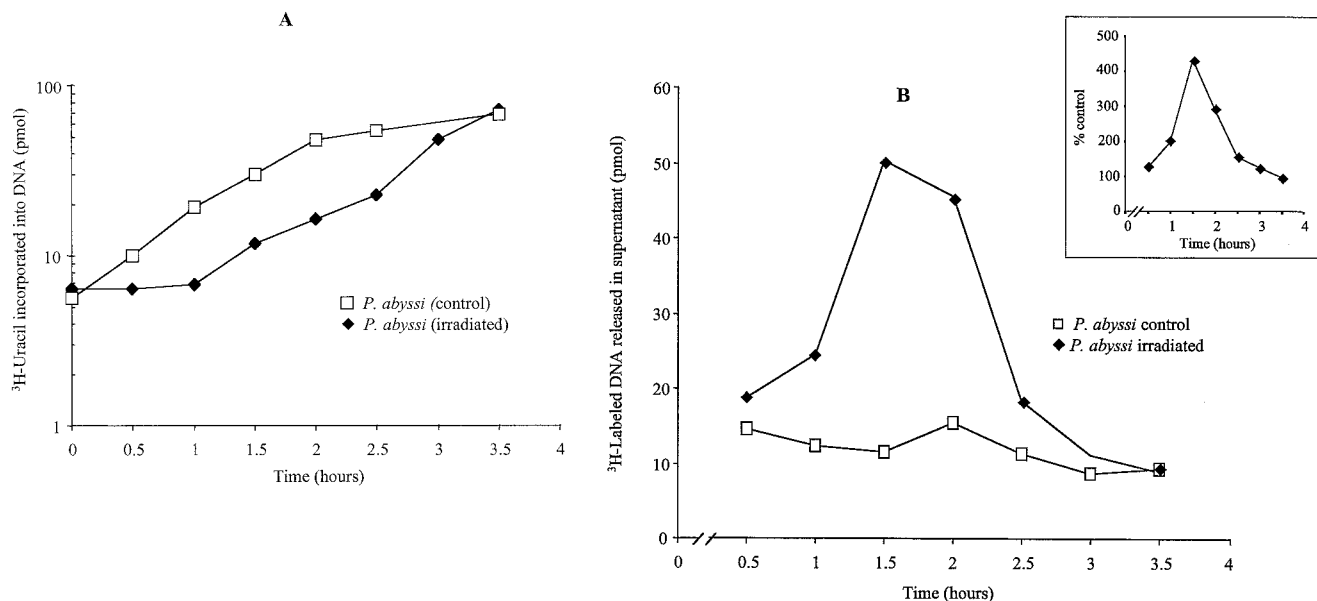


FIG. 2. Effect of gamma irradiation on DNA synthesis and export. (A) DNA synthesis restarts after gamma irradiation. The curves represent the restart of DNA replication in irradiated and nonirradiated cultures of “*P. abyssi*.” Cells were irradiated during stationary phase with 2,500 Gy of radiation. Fresh medium containing ³H-labeled uracil was inoculated and incubated at 95°C for 210 min. DNA replication was monitored by using scintillation counting to measure the incorporation of labeled uracil into DNA after TCA precipitation of whole cells. The curves show DNA replication in irradiated cells after irradiation at 2,500 Gy (◆) and DNA replication in unirradiated cultures (□). These results are the means of two independent experiments. (B) Export of damaged and undamaged DNA from the “*P. abyssi*” cells following gamma irradiation. This experiment shows the release of radioactivity in the supernatant from “*P. abyssi*” cells in early stationary phase after gamma irradiation at 2,000 Gy. The “*P. abyssi*” culture was previously labeled overnight with [³H]uracil at a final concentration of 1 μCi per ml before irradiation. After irradiation, the measurement of radioactivity in the supernatant was performed every 30 min until 210 min at 96°C. The inset shows the percentages of labeled DNA material from the irradiated cells versus the nonirradiated cells. The radioactivity released in the supernatant of irradiated cells (◆) was monitored by using scintillation counting to measure the labeled uracil in DNA after TCA precipitation of whole cells. Percentages were determined from control cultures treated under the same conditions but without gamma irradiation. These values are averages of two independent experiments.

processing of DSBs. A priori, the transport of damaged DNA should prevent the accumulation of genetic mistakes in irradiated *Pyrococcus* cells, similar to what was found earlier for *Deinococcus radiodurans* (1, 2) and “*Chroococciopsis*” cells (3).

A subset of “*P. abyssi*” DNA replication and repair proteins is chromatin bound before and after irradiation at 2,500 Gy. A possibility for repair of DSBs in *Pyrococcus* sp. is homologous recombination (10, 15) by chromatin-bound DNA repair complexes. As constitutively expressed *P. furiosus* RadA presumably functions in homologous recombination and physically interacts with several replication proteins (e.g., replication protein A [RPA] and replication factor C [RFC]) (16), we have tested whether these proteins (RadA, RPA, and RFC) dissociate from the chromatin in stationary-phase and irradiated cells, thus explaining the observed block in DNA synthesis (Fig. 2A).

We examined by simple cell fractionation and immunodetection assays the level of chromatin association of “*P. abyssi*” RadA, RPA, and RFC in irradiated cells (early stationary phase). After irradiation, “*P. abyssi*” cells from irradiated and nonirradiated control cultures were incubated for 120 min. Culture aliquots were removed at the indicated intervals. Soluble proteins (supernatant) and the chromosomal DNA-enriched insoluble fraction (pellet) were separated by centrifugation through a sorbitol cushion, as described previously (19).

The insoluble pellet fraction, containing DNA and chromatin-associated proteins, was washed once and dissolved in an equal volume of extraction buffer by brief sonication. This treatment yielded DNA fragments of 500 to 1,000 bp. Immunodetection of chromatin-associated proteins was performed as described previously (19).

In accordance with earlier results (15), our data indicated that a steady-state expression level of RadA protein (expected molecular mass of 38.4 kDa) was not massively induced after gamma irradiation (Fig. 3A). Moreover, our results indicate that the amount of RadA protein in soluble and chromatin-associated forms remained similar for 120 min postirradiation. Similar results were obtained for RPA (data not shown) and RFC (Fig. 3B), which are implicated in DNA replication. Therefore, the delay of DNA synthesis (replication) observed after irradiation does not result from dissociation of RadA, RPA, and RFC proteins from the chromatin.

Conclusion. In this study we have shown that “*P. abyssi*” cultures in stationary phase are able to completely repair their fully fragmented chromosomes following gamma irradiation within 2 h. On the basis of our experimental data, “*P. abyssi*” utilizes an efficient strategy for responding to DNA damage that includes (i) an uncoupling of DNA repair and DNA synthesis and (ii) a prevention of accumulation of genetic mistakes by exporting damaged DNA. In the light of our finding that RadA, RPA, and RFC stay chromatin bound before and after

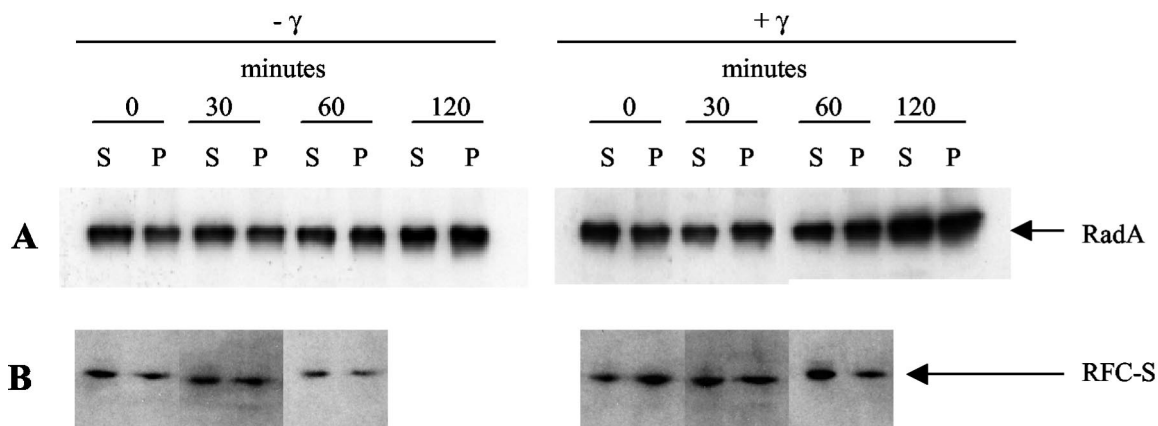


FIG. 3. Western blotting analysis of RadA and a subunit of the RFC complex. Experiments were performed with early-stationary-phase "*P. abyssi*" cells after the cells were irradiated at 2,500 Gy (+ γ) or not irradiated ($-\gamma$). Culture samples were removed every 30 min (up to 2 h at 96°C) and kept on ice. The soluble proteins (supernatant fraction [S]) and the DNA-bound proteins (pellet fraction [P]) were extracted from the cells as described earlier (19). Proteins were subjected to sodium dodecyl sulfate–10% polyacrylamide gel electrophoresis and were analyzed by Western blotting. The RadA (38.4 kDa) and RFC-S (37.4 kDa) proteins were immunodetected by specific rabbit antibodies by using the ECL Western blotting kit (Amersham Pharmacia Biotech). The protein band recognized by specific antibodies was visualized by exposure of the membranes to film.

irradiation, it is tempting to speculate that active chromatin-bound repair and replication complexes are continuously ready to counteract DNA damage in "*P. abyssi*." Identification of possible novel functions in adaptive responses of hyperthermophilic *Archaea* to radiation-induced DNA damage will now require efforts in functional genomics and/or genetics. In particular, it will be of interest to investigate whether the radiation-induced DNA synthesis block could result either from a physical blockage of replication forks or, alternatively, from the presence of a regulated mechanism (a check point) that would uncouple DNA replication and repair processes.

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