


Journal club

A HISTONE CODE FOR DNA REPAIR

In 1998, Rogakou, Bonner and colleagues reported that irradiation of mammalian cells causes a subtle but immediate alteration in the mobility of a small fraction of histones, as revealed by the analysis of samples using two-dimensional (2D) gel electrophoresis. The novel mobility histone species were attributable to radiation-induced phosphorylation of the histone H2A variant, H2AX, on Ser139 of its carboxy-terminal tail. The same authors later showed, in 1999, that antibodies specific for γ H2AX (H2AX phosphorylated on Ser139) decorate the chromatin flanking mammalian DNA double-strand breaks. They calculated that megabase tracts of

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chromatin might be modified at Ser139 of H2AX in the vicinity of a chromosome break — a new idea that immediately caught the attention of the field. Although relationships between chromatin structure and DNA repair had been suggested by previous studies, the discovery of a single, simple, locally break-responsive histone mark seemed to provide the ‘ocular proof’ of this concept.

Today we have an increasingly sophisticated understanding of the biochemical events of the γ H2AX response and its importance in preventing genome instability. We appreciate that γ H2AX, together with the adaptor protein mediator of DNA damage checkpoint protein 1 (MDC1), forms part of a ‘histone code’ of DNA repair, which recruits DNA damage response proteins to regions of damaged chromatin.

It is instructive to recall that there was a time when the chromatin response to chromosome breaks was considered of marginal interest. I take pleasure in the thought that one of the cornerstones of this emergent field was a tiny spot on a 2D gel (shown in Figure 1 of the 1998 paper by Rogakou *et al.*), which Rogakou, Bonner and colleagues had the confidence not to ignore.

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ORIGINAL RESEARCH PAPERS Rogakou, E. P. *et al.* DNA double-stranded breaks induce histone H2AX phosphorylation on serine 139. *J. Biol. Chem.* **273**, 5858–5868 (1998) | Rogakou, E. P. *et al.* Megabase chromatin domains involved in DNA double-strand breaks *in vivo*. *J. Cell Biol.* **146**, 905–916 (1999)