

## DNA-Dependent Protein Kinase Defects are Linked to Deficiencies in DNA Repair and V(D)J Recombination [and Discussion]

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# DNA-dependent protein kinase defects are linked to deficiencies in DNA repair and V(D)J recombination

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## SUMMARY

DNA-dependent protein kinase is a nuclear serine/threonine kinase whose catalytic properties are expressed only when the enzyme is bound to DNA ends or other discontinuities in the DNA. DNA-PK comprises two components: one mediates binding to DNA and corresponds to the heterodimeric human autoimmune antigen Ku; the other, DNA-PK catalytic subunit (DNA-PK<sub>cs</sub>), is a polypeptide of approximately 450 kDa. DNA-PK deficiencies are associated with certain mutant rodent cell lines that display defects in DNA double strand break repair and V(D)J recombination. Specifically, hamster *xrs-6* cells lack Ku function, whereas murine scid and hamster V3 cells lack functional DNA-PK<sub>cs</sub>. Furthermore, the phenotypes of *xrs-6* and V3 cells can be corrected by the expression of the genes encoding the 80 kDa component of Ku or DNA-PK<sub>cs</sub>, respectively. These results imply that DNA-PK is an important component of the DNA double strand break repair/recombination apparatus. Possible roles for DNA-PK in these processes are discussed.

## 1. INTRODUCTION

In recent years, several protein kinases have been shown to exist within the cell nucleus, suggesting that these enzymes may play important roles in modulating nuclear events. One such enzyme is DNA-dependent protein kinase (DNA-PK), a serine/threonine kinase that has been detected in organisms ranging from man to *Drosophila* (for a review see Finnie *et al.* 1995, also see references in Anderson & Lees-Miller 1992; Gottlieb & Jackson 1994). An important feature of DNA-PK is that it binds to DNA directly and needs to be DNA-bound to be active (Walker *et al.* 1985; Carter & Vancurova 1990; Jackson *et al.* 1990; Lees-Miller & Chen 1990). Biochemical fractionation studies have revealed that DNA-PK is a multiprotein complex comprising a large catalytic subunit (DNA-PK<sub>cs</sub>) and a DNA binding component called Ku (Dvir *et al.* 1993; Finnie *et al.* 1993; Gottlieb & Jackson 1993). Ku is an antigen recognized by antisera derived from patients with various autoimmune diseases and consists of two polypeptides of approximately 70 and 80 kDa (Ku70 and Ku80, respectively) in a tightly associated 1:1 complex.

Though physiological targets for DNA-PK have not been defined unequivocally, effective substrates for DNA-PK *in vitro* include the 34 kDa subunit of DNA replication/repair factor RPA (Brush *et al.* 1994; Pan *et al.* 1994; G. Smith & S. P. Jackson, unpublished

data), and transcription factors Sp1, c-Jun and p53 (Lees-Miller 1992; Bannister *et al.* 1993; Finnie *et al.* 1993; Jackson *et al.* 1993). DNA competition studies indicate that DNA-PK must bind *in cis* to the same DNA molecule as its protein substrate in order to function effectively (Gottlieb & Jackson 1993). The ability of DNA-PK to recognize several transcription factors suggests that it might play a role in transcriptional regulation. Consistent with this, DNA-PK has recently been shown to be a potent repressor of transcription by RNA polymerase I (Kuhn *et al.* 1995; Labhart 1995).

An intriguing feature of Ku is that it binds to DNA termini or other discontinuities in the DNA structure (for example, Paillard & Strauss 1991; Falzon *et al.* 1993). Consequently, DNA-PK is activated *in vitro* by linear but not by closed circular or supercoiled DNA (Carter & Vancurova 1990; Jackson *et al.* 1990, 1993; Gottlieb & Jackson 1993; Morozov *et al.* 1994). These properties suggest that a physiological function for DNA-PK is to recognize DNA strand interruptions in the genome such as occur in recombination intermediates and at the sites of DNA damage (Anderson 1993; Gottlieb & Jackson 1994). This has led ourselves and others to test whether DNA-PK/Ku defects exist in a set of radiation hypersensitive mutant rodent cell lines that share the common phenotype of being impaired specifically in DNA double-strand break (dsb) repair (Kemp *et al.* 1984; Giaccia *et al.* 1985; Whitmore *et al.* 1989; Fulop & Phillips 1990; Biedermann *et al.* 1991; Hendrickson *et al.* 1991). These

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cell lines fall into three distinct complementation groups. Ionizing radiation complementation group 4 (IR4) contains the XR-1 cell line, IR5 includes the Chinese hamster ovary (CHO)-derived cell line *xrs-6*, and IR7 contains the hamster cell line V3 and Scid cells that derived from the murine severe combined immune deficient (Scid) mouse. Scid mice lack a functional immune system due to an inability to perform V(D)J recombination, the site-specific genomic rearrangement process employed by developing B and T lymphocytes to generate the vast diversity of functional antigen receptor genes (for example, Schuler *et al.* 1986). Cell transfection approaches have established that cells of IR groups 4 and 5 are also defective in V(D)J recombination and that, as in Scid cells, this defect is in the effective ligation of recombination intermediates (Pergola *et al.* 1993; Taccioli *et al.* 1993). Taken together, these results imply that the ubiquitous DNA DSB repair apparatus is employed by the cell type-specific V(D)J recombination machinery and suggest that the gene products defined by the mutations in XR-1, *xrs-6* and Scid cells operate in the same biochemical pathway. Here, we review recent work demonstrating that cells of IR groups 5 and 7 are defective in Ku80 and DNA-PK<sub>cs</sub>, respectively, and assess the possible mechanisms by which DNA-PK functions in DNA repair and V(D)J recombination.

## 2. KU DNA BINDING ACTIVITY IS ABSENT FROM *XRS-6* CELLS

One approach to investigate the status of Ku in rodent cells has been to measure its non-specific DNA binding activity using the electrophoretic mobility shift assay. As shown in figure 1*a*, several DNA-protein complexes are formed when hamster cell extracts are incubated with a radiolabelled dsDNA oligonucleotide probe. Strikingly, one of these complexes (indicated by an arrow) is absent in *xrs-6* extracts but present in extracts from the parental cell line K1. From analysing the results of extract titration experiments, we estimate that the DNA binding activity is at least 50-fold more abundant in K1 cells than in *xrs-6* cells (see figure 1). The *xrs-6* defect appears to be linked to the mutation causing the DNA repair and recombination deficiencies of this cell line rather than some other mutation, because the DNA binding activity is also undetectable in extracts of the independently derived radiosensitive cell line XR-V15B, which falls into the same complementation group as *xrs-6* (see figure 3; compare XR-V15B and its parental line V79). Importantly, mixing *xrs-6* and K1 extracts does not result in the DNA binding activity of K1 cells becoming extinguished (data not shown), implying that the *xrs-6* defect is due to the lack of this factor rather than the presence of an inhibitory activity.

Several lines of evidence indicate that the DNA binding activity that is absent from *xrs-6* extracts is Ku. First, and consistent with the fact that Ku binds DNA ends, this complex is competed effectively by an oligonucleotide but not by supercoiled DNA (see figure 2*a*). Second, this complex is recognized specifically by anti-Ku monoclonal antibody N3H10 (see figure 2*b*).

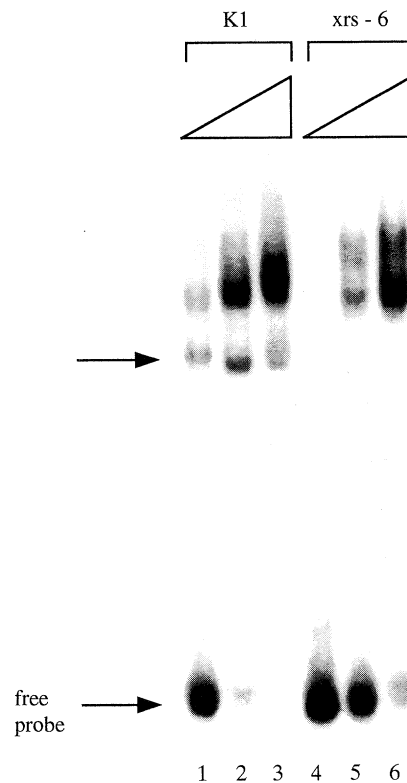


Figure 1. A DNA binding activity is defective in extracts of *xrs-6* cells. Electrophoretic mobility shift assays (EMSA) were performed as described previously using 10 fmol [ $\alpha$ - $^{32}$ P]-labelled ds oligonucleotide M1/M2 (Taccioli *et al.* 1994; Blunt *et al.* 1995). Reactions contained 0.5  $\mu$ g (lanes 1 and 4), 5  $\mu$ g (lanes 2 and 5), or 50  $\mu$ g (lanes 3 and 6) of whole cell extract (WCE) derived from K1 (lanes 1–3) or *xrs-6* (lanes 4–6) cells. The protein-DNA complex formed in K1 but not *xrs-6* extracts is indicated by an arrow.

Third, uv protein-DNA crosslinking reveals that an abundant DNA-binding protein of approximately 70 kDa is depleted in extracts of *xrs-6* cells (see figure 2*c*). This polypeptide comigrates with the major crosslinked protein in human HeLa cell extracts and with Ku70 present in a purified DNA-PK preparation. Fourth, this crosslinked polypeptide is restored in *xrs-6* derived hybrid cell lines '38' and 'H22' that have had their radiosensitivity and V(D)J recombination defects complemented by fragments of human chromosome 2 (see figure 2*c*; and for a description of cell hybrids see Finnie *et al.* 1995). Finally, *xrs-6* cells have been shown to be complemented by expression of the Ku80 cDNA (Taccioli *et al.* 1994; Smider *et al.* 1994; Boubnov *et al.* 1995). The human Ku80 gene has thus been designated as *XRCC5* (X-ray cross complementing gene for IR group 5). These results, taken together with those presented in other recent reports (Getts & Stamato 1994; Rathmell & Chu 1994; Taccioli *et al.* 1994), imply that Ku plays an important role in DNA DSB repair and V(D)J recombination.

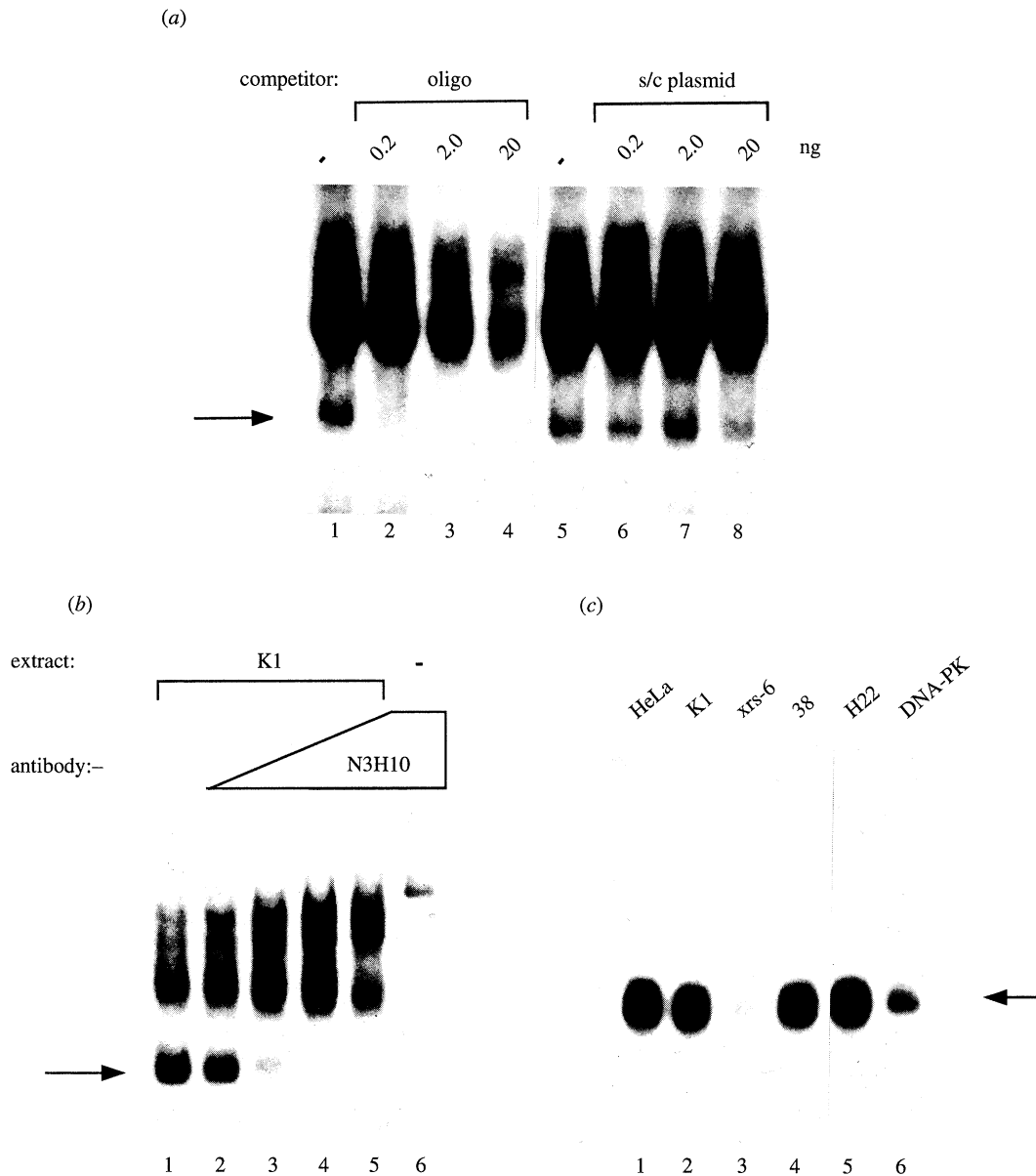


Figure 2. The DNA binding activity absent from *xrs-6* extracts corresponds to Ku. (a) The DNA binding activity binds to DNA ends. Whole cell extract (5 µg) derived from K1 cells was preincubated in the absence (–; lanes 1 and 5) or presence (lanes 2–4 and 6–8) of non radiolabelled competitor DNA at 30 °C for 10 min. [ $\alpha$ - $^{32}$ P]-radiolabelled ds oligonucleotide M1/M2 (0.2 ng) was then added and, after a further incubation for 10 min at 30 °C, DNA-protein complexes were detected by the EMSA. Reactions 2, 3, and 4, contained 0.2, 2, and 20 ng, respectively, of ds oligonucleotide M3/M4. Reactions 6, 7, and 8, contained 0.2, 2 and 20 ng, respectively, of closed circular plasmid DNA. The protein/DNA complex indicated is that which forms using K1 but not *xrs-6* extracts. (b) The protein/DNA complex that correlates with radiation resistance contains Ku70. K1 extract (10 µg) was preincubated at 4 °C for 10 min followed by room temperature (RT) for 10 min with no antibody (lane 1), or 0.02 µl, 0.1 µl, 0.5 µl, or 2.5 µl of anti-Ku70 monoclonal antibody N3H10 (lanes 2, 3, 4 and 5, respectively). Preincubation reaction 6 contained 2.5 µl of N3H10 alone. Next, 20 fmol radiolabelled M1/M2 was added and, after a further 10 min incubation at 30 °C, DNA-protein complexes were analysed by the EMSA. The protein/DNA complex indicated by the arrow is that which is absent in *xrs-6* extracts. (c) A 70 kDa DNA-binding polypeptide is deficient in *xrs-6* extracts. Purified DNA-PK (lane 6) or whole cell extract (lanes 1–5) was incubated at 30 °C for 10 min with 20 fmol [ $\alpha$ - $^{32}$ P]-end-labelled oligonucleotide M1/M2. Reactions were then subjected to uv-crosslinking for 10 min in a Stratilinker (Stratagene) and analysed by SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and autoradiography. Extracts employed were: HeLa (lane 1); K1 (lane 2); *xrs-6* (lane 3); hybrid *xrs-6* derived cell line 38 that contains a complementing fragment of human chromosome 2 (lane 4); or hybrid *xrs-6* derived cell line H22 that contains a complementing fragment of human chromosome 2 (lane 5). The major polypeptide labelled in HeLa and K1 extracts is deficient in extracts of non-complemented *xrs-6* extracts and comigrates with Ku70 in the DNA-PK preparation (indicated by an arrow).



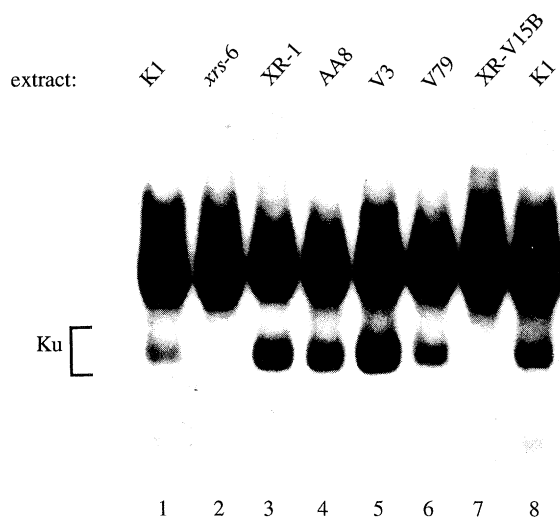


Figure 3. Cells of IR groups 4 and 7 possess Ku DNA end binding activity. EMSAs were conducted as described in the legend for figure 1 using 10 µg of whole cell extract (WCE) derived from the following cells: parental hamster K1 (lanes 1 and 8), hamster *xrs-6* mutant falling into IR5 (lane 2), hamster XR-1 mutant falling into IR4 (lane 3), parental hamster AA8 (lane 4), hamster V3 mutant falling into IR7 (lane 5), parental hamster cell line V79 (lane 6), and radiosensitive mutant cell line XR-V15B that is derived from V79 and whose mutation falls into IR group 5 (lane 7). The bracket indicates the position of the Ku/DNA complex.

### 3. CELLS OF IR GROUPS 4 AND 7 POSSESS KU DNA BINDING ACTIVITY

The results described above suggest that IR groups 4 and 7 might also lack Ku activity. To investigate this possibility, the Ku status of these cells was assessed by the electrophoretic mobility shift assay. In stark contrast to cells of IR group 5, Ku DNA binding activity is present at wild-type levels in extracts of all other mutant rodent cell lines so far tested (see figure 3, and also see Getts & Stamato 1994; Rathmell & Chu 1994). These include extracts from the hamster XR-1 cell line that falls into IR group 4, and extracts of hamster V3 and murine Scid cells that both fall into IR group 7.

### 4. CELLS OF IR GROUPS 4 AND 7 LACK DNA-PK ACTIVITY

The presence of Ku DNA binding activity in extracts of IR group 4 and 7 cells does not necessarily indicate that DNA-PK is functional: it is possible that these cells are defective in DNA-PK<sub>cs</sub>, the ability of Ku to activate DNA-PK<sub>cs</sub>, or in factors that regulate DNA-PK activity. As an approach to investigate these possibilities, we developed an assay system that is capable of detecting DNA-PK in extracts of rodent cells (Finnie *et al.* 1995). As shown in figure 4, although XR-1 extracts contain DNA-PK activity, *xrs-6* extracts essentially lack detectable DNA-PK. Furthermore, DNA-PK activity is absent from extracts of hamster V3 cells and murine Scid cells. Several lines of evidence

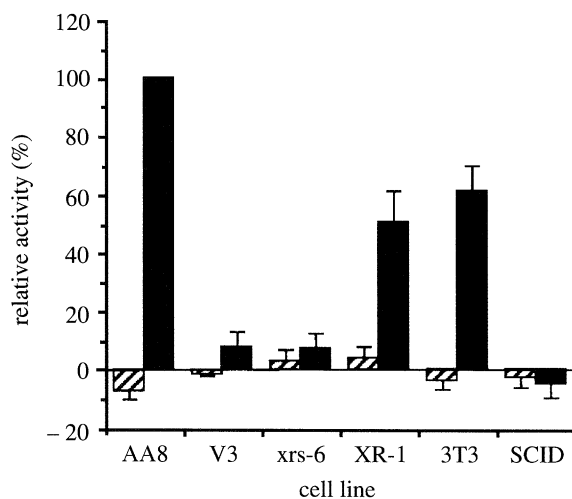


Figure 4. Cells falling into IR groups 5 and 7 lack DNA-PK activity. Whole cell extracts (100 µg) derived from the hamster or mouse cell lines indicated were tested for DNA-PK activity using assay conditions described previously (Finnie *et al.* 1995). Reactions were performed either with wild-type p53 peptide (solid columns) that is recognized effectively by DNA-PK, or with a mutated peptide (hatched columns) that is an ineffective DNA-PK substrate. Counts incorporated into peptide are plotted as a percentage of the value obtained with wild-type AA8 extract using the wt peptide.

indicate that this phenotype of V3 and scid cells is due to defects in DNA-PK<sub>cs</sub>. First, addition of purified DNA-PK<sub>cs</sub> but not Ku complements V3 and Scid extracts for DNA-PK activity (Blunt *et al.* 1995). Second, introduction of the DNA-PK<sub>cs</sub> gene into V3 cells results in complementation in terms of DNA-PK activity, DNA repair capacity, and ability to mediate V(D)J recombination (Blunt *et al.* 1995). Third, the DNA-PK<sub>cs</sub> gene maps to the region of human chromosome 8 that contains the gene *XRCC7* that complements the *scid* mutation (Kirchgessner *et al.* 1995; Siple *et al.* 1995). Finally, the gene encoding the mouse homologue of DNA-PK<sub>cs</sub> has been shown to co-segregate with the *scid* mutation in mouse genetic crosses (Miller *et al.* 1995).

### 5. HAMSTER V3 AND MURINE SCID CELLS CONTAIN DIMINISHED LEVELS OF DNA-PK<sub>cs</sub>

The above data suggest strongly that the mutations present in scid and V3 cells lie within the gene for DNA-PK<sub>cs</sub>. Potentially, these mutations could lead to decreased expression of DNA-PK<sub>cs</sub>, or could cause the production of a mutated polypeptide with reduced specific activity. To determine whether the V3 and Scid mutations affect DNA-PK<sub>cs</sub> expression, we employed anti-DNA-PK<sub>cs</sub> antibodies in Western blotting studies. As shown in figure 5, antisera raised against human DNA-PK<sub>cs</sub> recognize polypeptides in extracts of mouse 3T3 and hamster AA8 cells that comigrate on SDS-polyacrylamide gels with human DNA-PK<sub>cs</sub>. This polypeptide, however, is apparently present in much lower levels in rodent extracts compared to extracts of human cells (compare lanes 1 and 3 with lane 5),

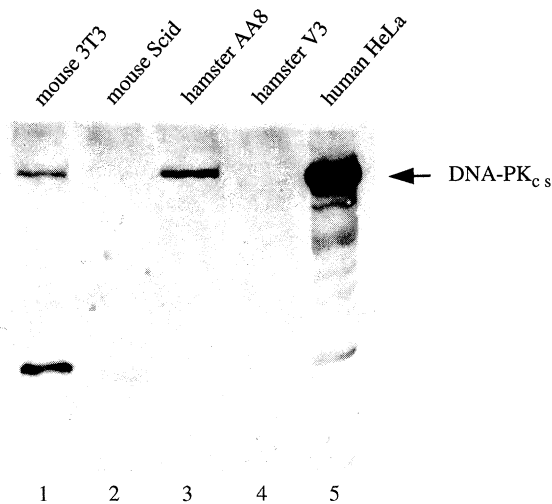


Figure 5. Hamster V3 and murine scid cells lack detectable DNA-PK<sub>cs</sub>. Whole cell extract derived from mouse 3T3 cell (lane 1), murine scid cells (lane 2), parental hamster cell line AA8 (lane 3), V3 mutant cells (lane 4), or human HeLa cells (lane 5) were electrophoresed on a 5% SDS-polyacrylamide gel together with a sample of human DNA-PK (lane 6). The samples were then subjected to Western immunoblot analysis using a rabbit polyclonal antiserum raised against human DNA-PK<sub>cs</sub>. The position of human DNA-PK<sub>cs</sub> and the crossreactive polypeptide present in wild-type rodent extracts is indicated by an arrow.

which is consistent with our previous finding that rodent cells possess 20- to 50-fold lower DNA-PK activity than human cells (Blunt *et al.* 1995; Finnie *et al.* 1995). Significantly, Western blot assays fail to detect a DNA-PK<sub>cs</sub> crossreactive protein in extracts of Scid or V3 cells (see figure 5). These results are consistent with the findings of Kirchgessner *et al.* (1995) and imply that the mutations in Scid and V3 cells either abolish DNA-PK<sub>cs</sub> synthesis, or result in the generation of an unstable protein. It will obviously be of great interest to analyse the DNA-PK<sub>cs</sub> genes in these two cell lines to define the precise nature of their mutations.

## 6. POSSIBLE MECHANISMS FOR DNA-PK ACTION *IN VIVO*

The data discussed above are most simply interpreted by concluding that the DNA-PK holoenzyme, comprising DNA-PK<sub>cs</sub> and Ku plays a central role in DNA DSB repair. DNA-PK activity also appears to be necessary for the DNA end joining steps of V(D)J recombination. There are a several ways in which DNA-PK may be envisaged to operate in these two processes. First and foremost, it is likely that the recognition of DNA DSBs by the Ku component of DNA-PK is central to its mode of action. For example, interactions between the DNA-PK holoenzyme and DNA termini might prevent exonucleases from destroying important genetic information before ligation events are able to be completed. Another way in which DNA-PK might function in DNA repair is by phosphorylating and activating other components of the DNA repair apparatus. Indeed, the restriction of

kinase activity in the vicinity of DNA ends could ensure that the repair machinery is only activated in the site of a DNA DSB. Alternatively, or in addition, DNA-PK could recruit other DNA repair components to the site of DNA damage by protein-protein interactions. Because DNA-PK<sub>cs</sub> is over 450 kDa in size (Hartley *et al.* 1995), the possibility exists that it fulfils a scaffolding role, mediating contacts with several other factors and orienting these appropriately with respect to one another to facilitate effective DNA repair. Polypeptides that might bind to DNA-PK<sub>cs</sub> include the products of the V(D)J recombination activating genes *RAG1* and *RAG2* (Oettinger *et al.* 1990), the product that is mutated in the XR-1 cell line, DNA ligases, and enzymes such as nucleases and terminal deoxynucleotidyl transferase that function in V(D)J recombination by mediating the controlled modification of coding ends prior to their joining.

A final possible role for DNA-PK is that of a DNA end sensor to alert the cell to the fact that it has sustained injury. Indeed, it is tempting to speculate that DNA-PK activation by DNA damage might trigger a protein phosphorylation cascade that results in pleiotropic effects on cellular metabolism. These effects potentially include the transcriptional or post-transcriptional induction of components that directly or indirectly mediate DNA repair, and the activation of cell cycle checkpoint controls. Although there is at present no evidence to suggest that checkpoint control mechanisms are defective in cells that lack DNA-PK function, it is possible that multiple, partially redundant pathways for detecting DNA DSBs exist in higher eukaryotic cells and that DNA-PK constitutes just one of these. Clearly, the availability of clones for DNA-PK components and DNA-PK defective cell lines will prove invaluable in addressing the various possibilities discussed above and in defining the precise physiological roles of DNA-PK.

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### Discussion

Y. A. HANNUN (*Department of Medicine & Cell Biology, Duke University Medical Center, North Carolina, U.S.A.*). Does DNA-PK phosphorylate nucleic acid?

S. P. JACKSON. We have never done that experiment.

M. KARIN (*Department of Pharmacology, University of California, San Diego, U.S.A.*). Have you ever tried to link Ku's kinase recognition sequence onto different DNA-binding sequences, to see whether you can recruit DNA-PK<sub>cs</sub> to regions other than double-stranded breaks.

S. P. JACKSON. We do not yet know the answer to that, but there is evidence that DNA-PK<sub>cs</sub> may itself need to contact DNA: if it is targeted to DNA-bound Ku, it can itself get uv cross-linked to the DNA. DNA-PK may even have itself to contact DNA ends. We are now trying to do yeast two-hybrid studies to try and narrow down the interaction domain between the kinase and Ku.

M. KARIN. How does DNA-PK influence PolI transcription?

S. P. JACKSON. Our work on this was published earlier this year (Kuhn *et al.* 1995). DNA-PK inhibits transcription, as a result of phosphorylating some elements of the PolI transcriptional apparatus. It certainly inhibits elongation, but it is not clear whether it also inhibits initiation. It also inhibits transcription by PolIII.

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*Question.* Is DNA-PK ubiquitous, or is its abundance regulated in relation to cell proliferation or tumorigenesis?

S. P. JACKSON. There is some evidence that the Ku concentration increases in rapidly proliferating cells. However, this effect is not large, and DNA-PK and Ku are abundant proteins. Human cells have about half a million copies of Ku and not many less of the DNA-PK<sub>cs</sub>. For some reason, rodents have about 50-fold less DNA-PK than humans. My current hypothesis is that this may be related to longevity, in that mice may not need to repair DNA as effectively as humans. There is no evidence that DNA-PK levels change in response to radiation or in cell-lines. However, there are derived cell-lines that have reduced DNA-PK levels, and these should prove very useful in analysing its function. Carl Anderson, with whom we have been collaborating on the cloning of DNA-PK, has now got to the promoter region, which looks like the promoter of a housekeeping gene.



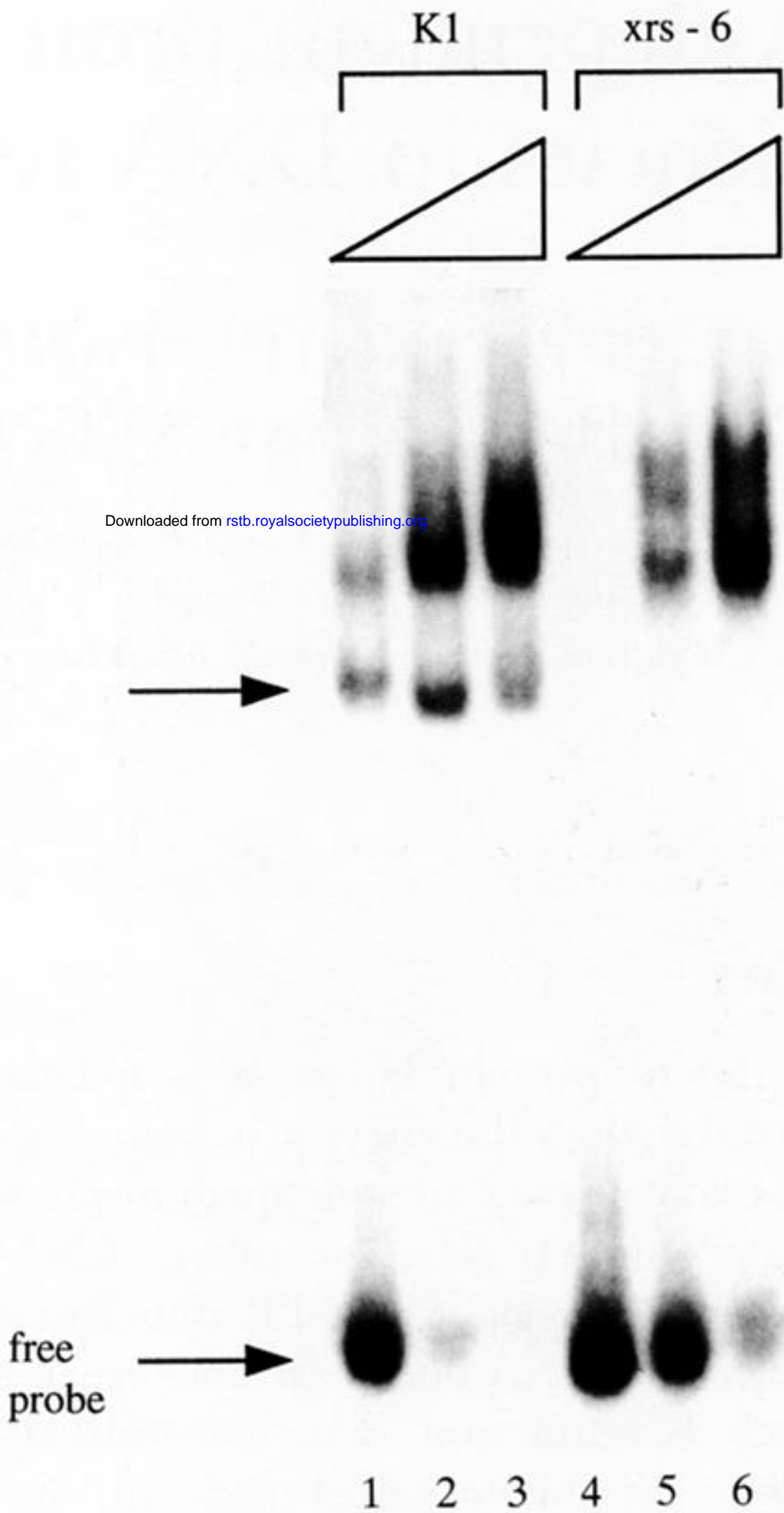
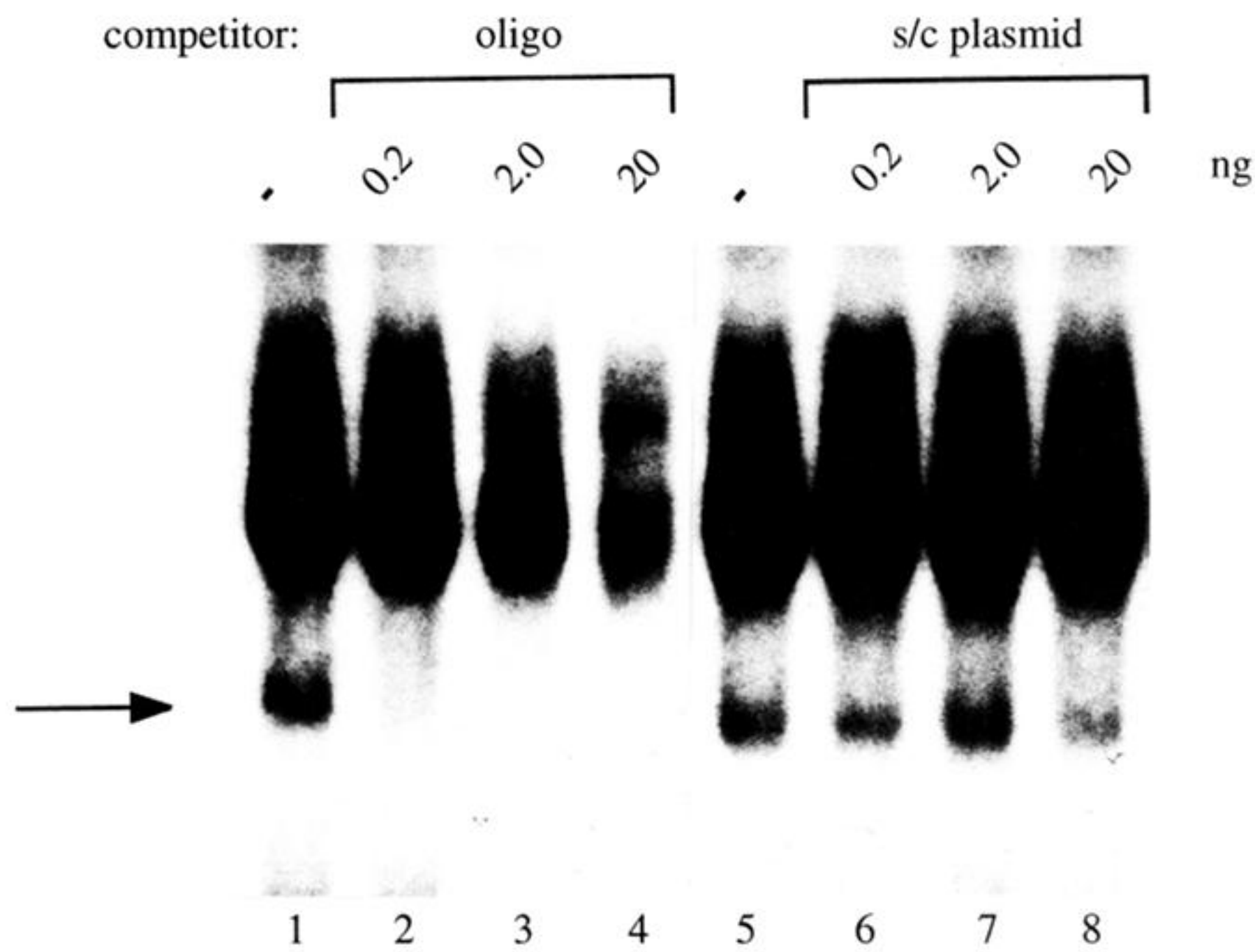


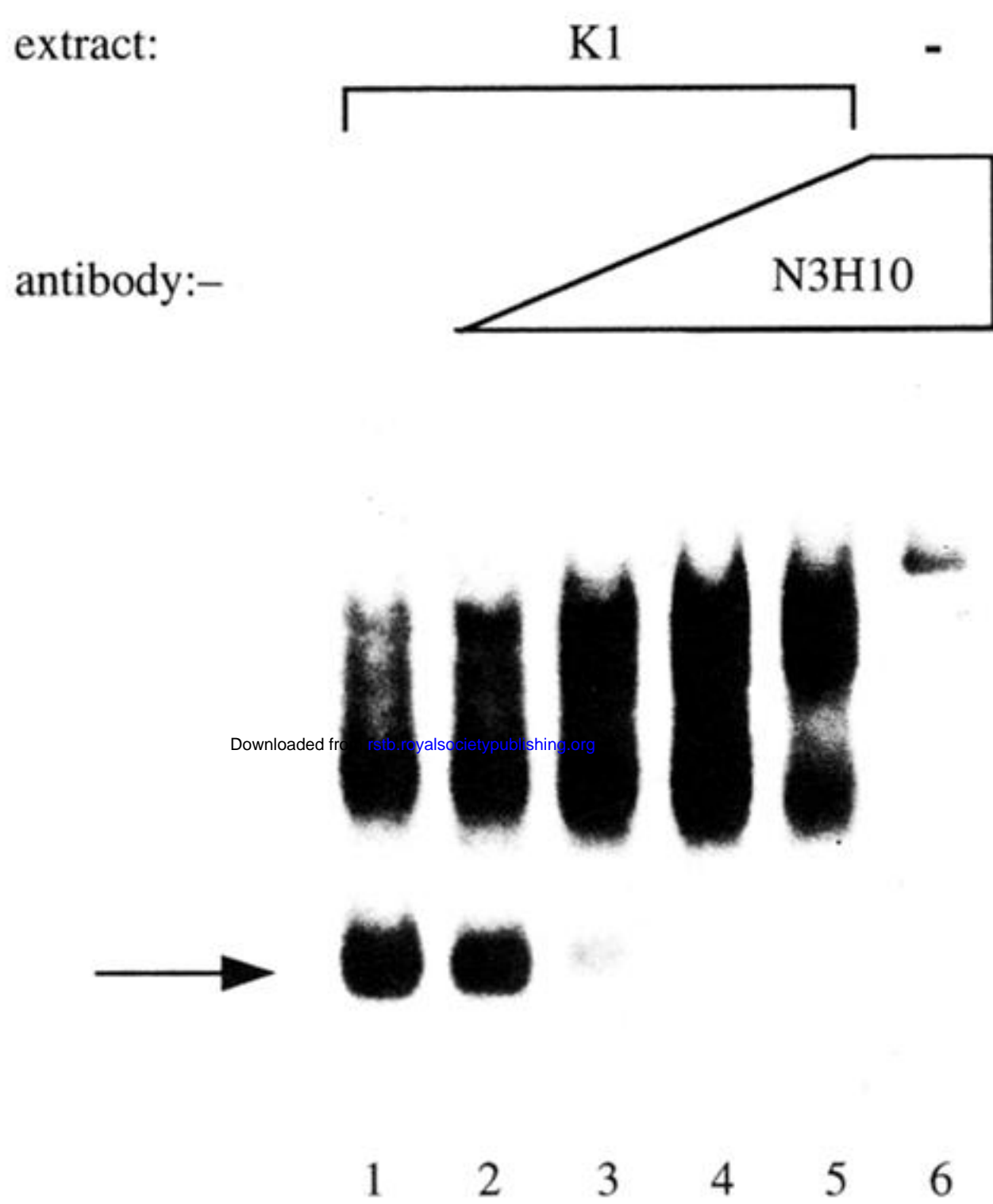
Figure 1. A DNA binding activity is defective in extracts of *xrs-6* cells. Electrophoretic mobility shift assays (EMSAs) were performed as described previously using 10 fmol [ $\alpha$ - $^{32}$ P]-labelled ds oligonucleotide M1/M2 (Taccioli *et al.* 1994; Hunt *et al.* 1995). Reactions contained 0.5  $\mu$ g (lanes 1 and 4), 5  $\mu$ g (lanes 2 and 5), or 50  $\mu$ g (lanes 3 and 6) of whole cell extract (WCE) derived from K1 (lanes 1–3) or *xrs-6* (lanes 4–6) cells. The protein-DNA complex formed in K1 but not in *xrs-6* extracts is indicated by an arrow.



(a)



(b)



(c)

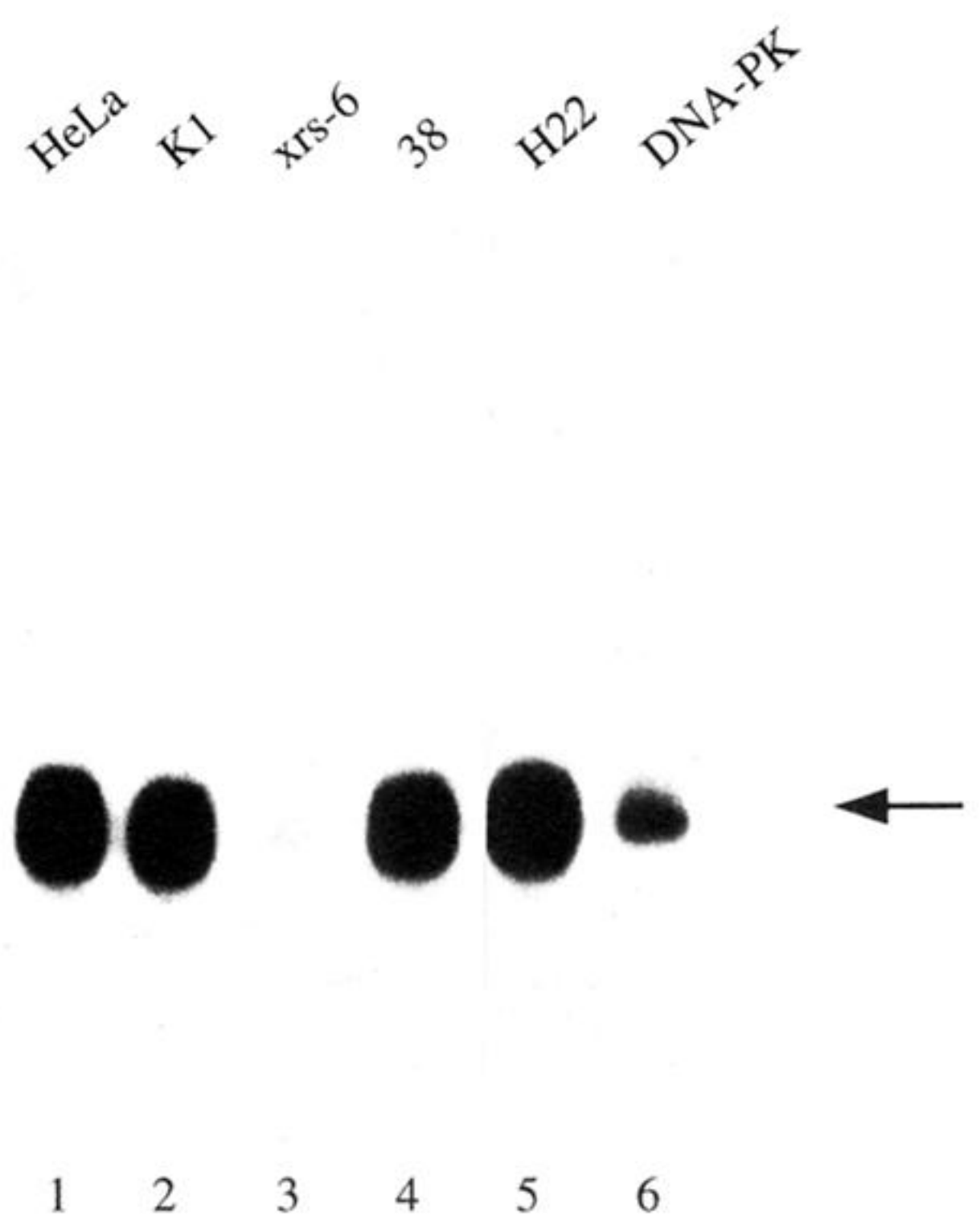


Figure 2. The DNA binding activity absent from *xrs-6* extracts corresponds to Ku. (a) The DNA binding activity corresponds to DNA ends. Whole cell extract (5  $\mu$ g) derived from K1 cells was preincubated in the absence (–; lanes 1 and 5) or presence (lanes 2–4 and 6–8) of non radiolabelled competitor DNA at 30 °C for 10 min. [ $\alpha$ - $^{32}$ P]-radiolabelled oligonucleotide M1/M2 (0.2 ng) was then added and, after a further incubation for 10 min at 30 °C, DNA-protein complexes were detected by the EMSA. Reactions 2, 3, and 4, contained 0.2, 2, and 20 ng, respectively, of ds oligonucleotide M3/M4. Reactions 6, 7, and 8, contained 0.2, 2 and 20 ng, respectively, of closed circular plasmid DNA. The protein/DNA complex indicated is that which forms using K1 but not *xrs-6* extracts. (b) The protein/DNA complex that correlates with radiation resistance contains Ku70. K1 extract (10  $\mu$ g) was preincubated at 4 °C for 10 min followed by room temperature (RT) for 10 min with no antibody (lane 1), or 0.02  $\mu$ l, 0.1  $\mu$ l, 0.5  $\mu$ l, or 2.5  $\mu$ l anti-Ku70 monoclonal antibody N3H10 (lanes 2, 3, 4 and 5, respectively). Preincubation reaction 6 contained 0.5  $\mu$ l of N3H10 alone. Next, 20 fmol radiolabelled M1/M2 was added and, after a further 10 min incubation at 30 °C, DNA-protein complexes were analysed by the EMSA. The protein/DNA complex indicated by the arrow is that which is absent in *xrs-6* extracts. (c) A 70 kDa DNA-binding polypeptide is deficient in *xrs-6* extracts. Purified DNA-PK (lane 6) or whole cell extract (lanes 1–5) was incubated at 30 °C for 10 min with 20 fmol [ $\alpha$ - $^{32}$ P]-end-labelled oligonucleotide M1/M2. Reactions were then subjected to UV-crosslinking for 10 min in a Stratelinker (Stratagene) and analysed by SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and autoradiography. Extracts employed were: HeLa (lane 1); K1 (lane 2); *xrs-6* (lane 3); hybrid *xrs-6* derived cell line 38 that contains a complementing fragment of human chromosome 2 (lane 4); or hybrid *xrs-6* derived cell line H22 that contains a complementing fragment of human chromosome 2 (lane 5). The major polypeptide labelled in HeLa and K1 extracts is deficient in extracts of non-complemented *xrs-6* extracts and comigrates with Ku70 in the DNA-PK preparation (indicated by an arrow).



Extract:

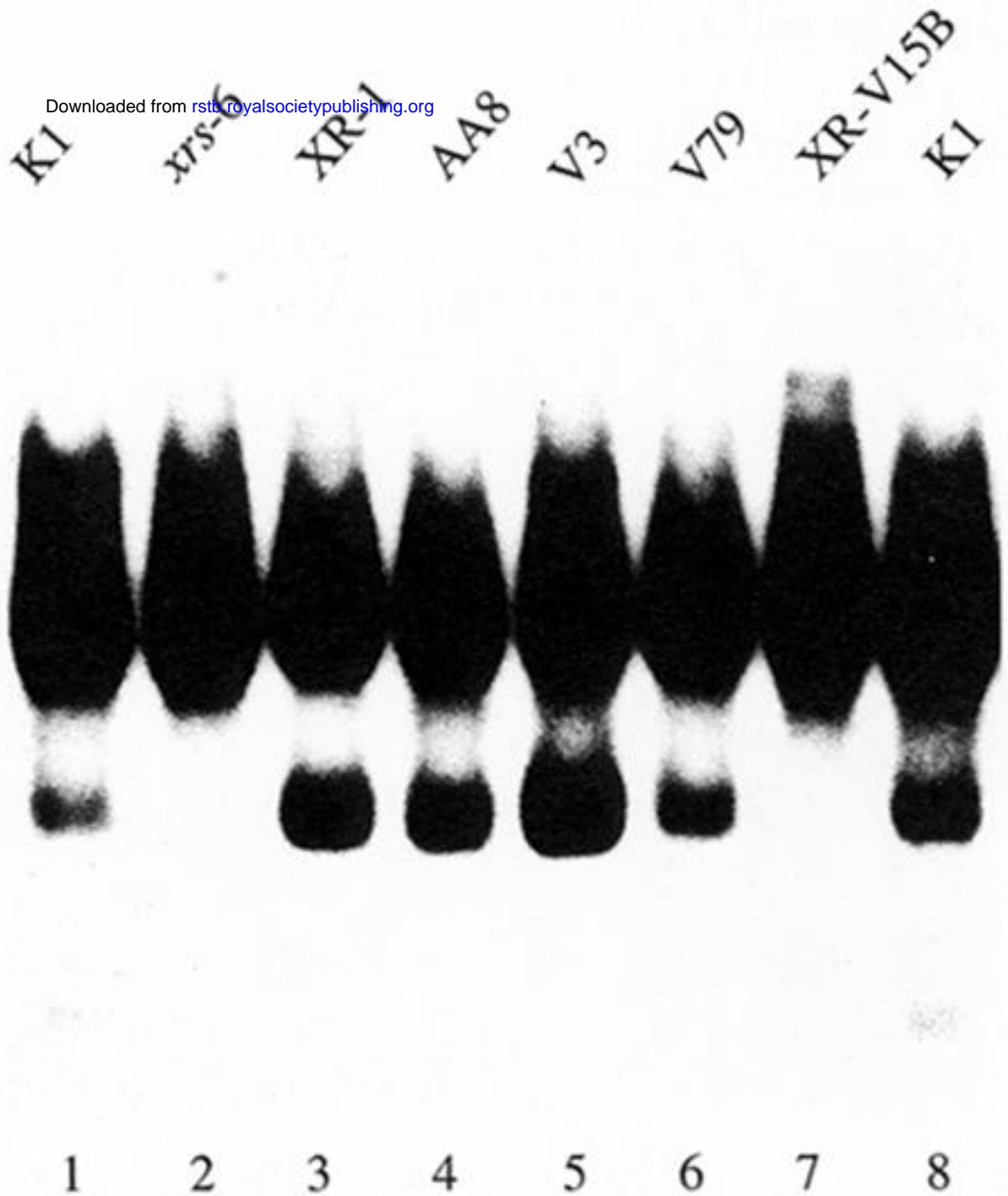


Figure 3. Cells of IR groups 4 and 7 possess Ku DNA end binding activity. EMSAs were conducted as described in the legend for figure 1 using 10  $\mu\text{g}$  of whole cell extract (wce) derived from the following cells: parental hamster K1 (lanes 1 and 8), hamster *xrs-6* mutant falling into IR5 (lane 2), hamster XR-1 mutant falling into IR4 (lane 3), parental hamster AA8 (lane 4), hamster V3 mutant falling into IR7 (lane 5), parental hamster cell line V79 (lane 6), and radiosensitive mutant cell line XR-V15B that is derived from V79 and whose mutation falls into IR group 5 (lane 7). The bracket indicates the position of the Ku/DNA complex.

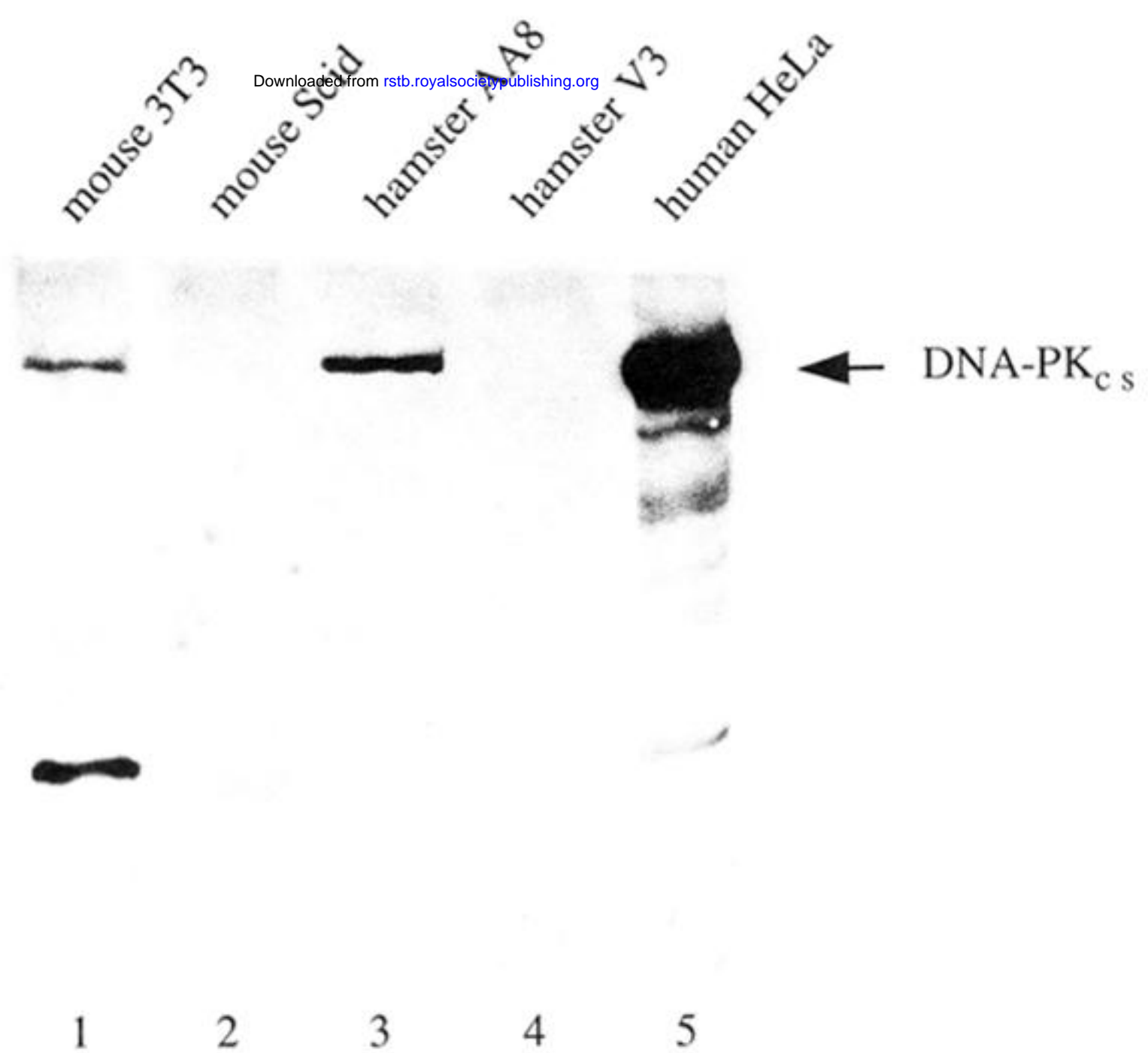


Figure 5. Hamster V3 and murine scid cells lack detectable DNA-PK<sub>cs</sub>. Whole cell extract derived from mouse 3T3 cell (lane 1), murine scid cells (lane 2), parental hamster cell line AA8 (lane 3), V3 mutant cells (lane 4), or human HeLa cells (lane 5) were electrophoresed on a 5% SDS-polyacrylamide gel together with a sample of human DNA-PK (lane 6). The samples were then subjected to Western immunoblot analysis using a rabbit polyclonal antiserum raised against human DNA-PK<sub>cs</sub>. The position of human DNA-PK<sub>cs</sub> and the crossreactive polypeptide present in wild-type rodent extracts is indicated by an arrow.