

# Caspases mediate nucleoporin cleavage, but not early redistribution of nuclear transport factors and modulation of nuclear permeability in apoptosis

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Val-Ala-aspartyl-fluoromethylketone; DEVD-CHO, N-acetyl-Asp-Glu-Val-Asp-aldehyde

## Introduction

The most evident morphological feature of apoptosis is the disassembly of the nucleus, which involves the condensation of chromatin and its segregation into membrane-enclosed particles.<sup>1</sup> Biochemical hallmarks of apoptotic nuclear execution are DNA cleavage in large and small (oligonucleosomal-sized) fragments, as well as the specific proteolysis of several nuclear substrates. Major effectors of apoptotic nuclear changes are members of the cysteine protease family of caspases. Nuclear substrates for caspases include nucleoskeletal elements like lamins,<sup>2,3</sup> and proteins involved in the organisation and replication of DNA, like SAF-A, MCM3 and RCF140.<sup>4-6</sup>

Cleavage of nuclear proteins may have important implications for the apoptotic process. For example, PARP cleavage shuts down its ADP-ribosylating activity, preventing a futile recruitment of the DNA-repair machinery and sparing ATP necessary for the apoptotic execution.<sup>7</sup> Similarly, cleavage of MCM proteins and replication factor 140 may inhibit unnecessary DNA replication cycles. Recent findings also show that a nuclear pore protein Nup153 is cleaved in apoptosis.<sup>8</sup> This protein, which belongs to the XFXFG-family of nucleoporins and resides at the nucleoplasmic side of the pore as a constituent of the nuclear basket<sup>9</sup> (see also<sup>10,11</sup>), has been implicated in various types of nuclear protein import as well as mRNA export from the nucleus.<sup>12,13</sup> Nup153 has been shown to bind soluble components of the nuclear transport machinery, such as members of the importin  $\beta$  family and the small GTPase Ran.<sup>14,15</sup> Therefore cleavage of Nup153 by caspases may affect nucleocytoplasmic transport in cells undergoing apoptosis.

A potential role of nucleocytoplasmic transport processes in apoptotic execution has been first inferred by Yasuhara *et al.*,<sup>16</sup> who showed that microinjection of different inhibitors of nuclear import including WGA (wheat germ agglutinin), a lectin binding to several O-glycosylated nucleoporins, protected from Fas-induced apoptotic nuclear changes. WGA was also reported to prevent nuclear condensation elicited by the apoptosis-inducing factor AIF<sup>17</sup> or by Acinus (apoptotic chromatin condensation inducer in the nucleus).<sup>18</sup> Even though WGA may bind other factors than nucleoporins,<sup>19</sup> these experiments suggest that active nuclear translocation of apoptosis executioners may be required for nuclear degradation. In line with this assumption is the finding that procaspase-1 and -2, as well as the caspase-activated DNase CAD, carry nuclear localisation sequences (NLSs).<sup>20-22</sup>

## Abstract

**In eukaryotic cells, both soluble transport factors and components of the nuclear pore complex mediate protein and RNA trafficking between the nucleus and the cytoplasm. Here, we investigated whether caspases, the major execution system in apoptosis, target the nuclear pore or components of the nuclear transport machinery. Four nucleoporins, Nup153, RanBP2, Nup214 and Tpr are cleaved by caspases during apoptosis. In contrast, the nuclear transport factors, Ran, importin  $\alpha$  and importin  $\beta$  are not proteolytically processed, but redistribute across the nuclear envelope independently and prior to caspase activation. Also, mRNA accumulates into the nucleus before caspases become active. Microinjection experiments further revealed that early in apoptosis, the nucleus becomes permeable to dextran molecules of 70 kD molecular weight. Redistribution of import factors and mRNA, as well as nuclear permeabilisation, occur prior to caspase-mediated nucleoporin cleavage. Our findings suggest that the apoptotic programme includes modifications in the machinery responsible for nucleocytoplasmic transport, which are independent from caspase-mediated degradation of nuclear proteins.**

**Keywords:** caspase; nuclear pore complex; nuclear import; mRNA export; microinjection

**Abbreviations:** aa, amino acids; NLS, nuclear localisation sequence; WGA, wheat germ agglutinin; STS, staurosporine; ActD, actinomycin D; DEVD-afc, N-acetyl-Asp-Glu-Val-aspartyl-aminofluoro-methylcoumarine; z-VAD-fmk, N-benzyloxycarbonyl-

Since the integrity of the nuclear pore may be essential for nucleocytoplasmic transport, we initially studied the fate of several representative nucleoporins in apoptosis. In addition to Nup153 these included RanBP2, Nup214, Tpr and p62, each of which is a constituent of a different substructure of the NPC. RanBP2, a Ran-binding protein of 358 kD, and Nup214 are XFXFG-type nucleoporins located at the short filaments attached to the cytoplasmic side of the NPC.<sup>11,23–26</sup> p62, which represents another XFXFG nucleoporin, is located at or near the center of the NPC.<sup>27–30</sup> Tpr, a protein of 267 kD which lacks XFXFG sequence motifs, is a constitutive element of thin filaments which are attached to the NPC's nucleoplasmic side and which project into the nuclear interior beyond the dimensions of the nuclear basket.<sup>31,32</sup> We found that Nup153, RanBP2, Nup214 and Tpr, but not p62, are specifically and rapidly cleaved by a distinct set of caspases during apoptosis.

We then analyzed a subset of nuclear transport factors, some of which are known to interact with Nup153, RanBP2 or Tpr. These are importin  $\beta$ , the major receptor for classical NLS-dependent import, importin  $\alpha$ , an adapter molecule which mediates the interaction between importin  $\beta$  and the NLS-bearing transport substrate, and Ran, a Ras-family GTPase which controls the binding of transport receptors to their substrates.<sup>33</sup> We discovered that Ran, importin  $\alpha$  and importin  $\beta$  are not degraded but redistributed between the nucleus and the cytoplasm in the early phase of apoptosis. Concomitantly, poly(A)<sup>+</sup>-RNA accumulated in the nucleus.

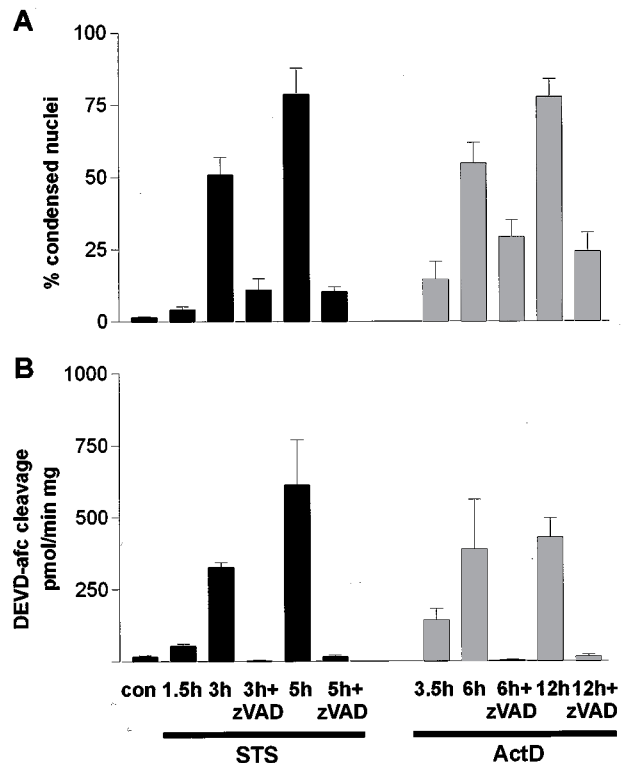
Both the relocation of transport factors and nuclear mRNA accumulation occurred prior to caspase-mediated chromatin condensation, and independently from caspase-mediated cleavage of nucleoporins. Taken together our findings suggest the existence of caspase-independent pathways that signal to the nucleus the initiation of the death programme.

## Results

### Cleavage of nucleoporins Nup153, RanBP2, Nup214 and Tpr in HeLa cells undergoing apoptosis

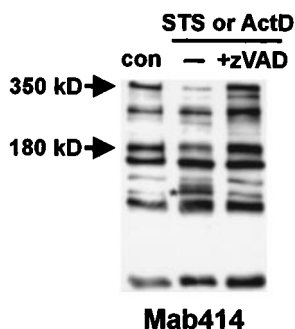
To investigate the fate of nuclear pore complex proteins in apoptosis, we treated HeLa cells with two classic apoptotic triggers: staurosporine (STS), a general protein kinase inhibitor, which induces apoptosis in many different cell types,<sup>34</sup> and actinomycin D (ActD), an inhibitor of RNA polymerases, used as a stimulus acting primarily on the nucleus.<sup>35</sup>

After 3 h of exposure to STS and 6 h of exposure to ActD, about 50% of the cells underwent apoptosis as judged by nuclear condensation. Concomitantly, we measured a 30-fold increase in the cleavage of the caspase substrate DEVD-afc, compared to control cells, which showed the efficient activation of caspase-3 like proteases (Figure 1). Both cell death and caspase activation were effectively inhibited by the poly-caspase inhibitor zVAD-fmk (20  $\mu$ M).<sup>36</sup>

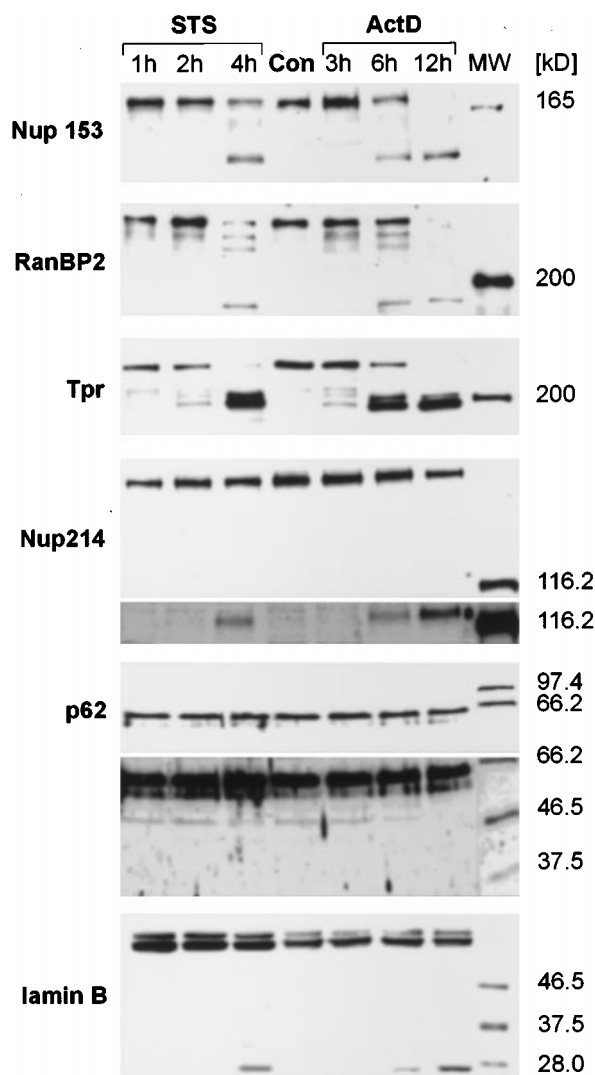


**Figure 1** Staurosporine (STS) and actinomycin D (ActD) induced cell death in HeLa cells. **(A)** Apoptosis was quantified by staining cells with H 33342 and Sytox and scoring Sytox impermeable cells with condensed nuclei. **(B)** Caspase 3 like activity was determined by DEVD-afc cleavage. Data are means  $\pm$  standard deviation from triplicate determinations. Con: control cells

Whole cell extracts from treated HeLa cells were incubated with the polyspecific antibody Mab414 which recognises several members of the XFXFG-family of nucleoporins including RanBP2, Nup153, p62 and others.<sup>9,27</sup> As shown in Figure 2, primarily two proteins with electrophoretic mobilities corresponding to Mr of 350 and 180 were degraded. Their cleavage was suppressed by preincubation with zVAD-fmk. Immunoblotting with specific antibodies identified the two cleaved nucleoporins as RanBP2 and Nup153 (Figure 3). One major degradation intermediate, recognised by different Nup153-specific antibodies, was a polypeptide of Mr 130. This fragment most likely corresponds to the Mab414-reactive 130 kD polypeptide enriched in apoptotic cells (see Figure 2). A Nup153 degradation product of 130 kD has recently also been observed by others.<sup>8</sup> Proteolytic processing of RanBP2 was monitored with an antibody specific for an epitope in the protein's C-terminal half. In this case, three cleavage products of 170, 220 and 250 kD were labelled. Cleavage of a further XFXFG-nucleoporin, Nup214, yielded a fragment of 120 kD, which became apparent only after long exposure times. In contrast, the other member of the XFXFG-family, p62, was not degraded (Figure 3), indicative that cleavage of RanBP2, Nup214 and Nup153 did not merely reflect an overall and simultaneous destruction of the nuclear pore complex.



**Figure 2** Proteolysis of XFXFG nucleoporins in apoptosis. Whole HeLa cell extracts were prepared from control cells and apoptotic cells treated with staurosporine (STS) or actinomycin D (ActD) in absence or in presence of zVAD fmk followed by immunoblotting with the polyspecific antibody Mab414. The asterisk indicates the 130 kD cleavage product of Nup153

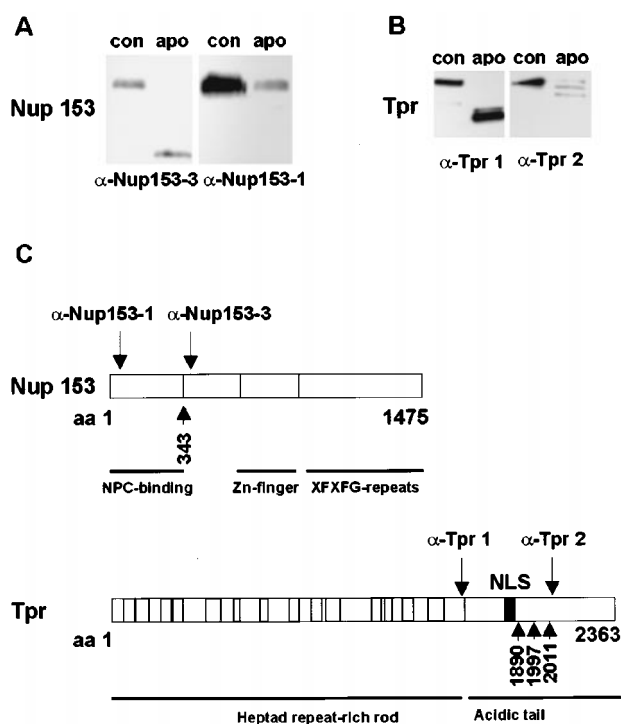


**Figure 3** Nup153, RanBP2, Tpr and Nup214 are cleaved in apoptosis. Whole cell extracts were prepared from control cells (con) and apoptotic cells treated with staurosporine (STS) and actinomycin D (ActD) at the indicated time points followed by immunoblotting with antibodies specific for the indicated nucleoporins. Long exposures of the Nup214 and p62 blots are included. Cleavage of lamin B as marker for apoptosis is also shown

To investigate whether other, non-XFXFG nucleoporins, were also targets of apoptotic degradation, we studied the fate of the NPC-associated protein Tpr. Induction of apoptosis resulted in cleavage of Tpr yielding two immunoreactive degradation products: a minor one of 200 kD and a major fragment of 190 kD.

Cleavage of Nup153, RanBP2, Nup214 and Tpr occurred 4 h after STS and 6 h after ActD treatment and coincided with the degradation of lamin B, which has been widely used as a marker for the onset of apoptotic nuclear execution.<sup>3</sup>

To narrow in the cleavage sites on Nup153 and Tpr, we then used peptide specific antibodies which were raised against different regions of these proteins. As seen in Figure 4, the Nup153 cleavage product was recognised by an antibody specific for a peptide comprising aa 391 to 404 ( $\alpha$ -Nup153-3), but not by an antibody directed against the very N-terminus of the protein ( $\alpha$ -Nup153-1), suggesting that cleavage occurred between aa 36 and 391. This region contains one sequence with homology to caspase recognition sites (DITD at position 343). Cleavage at this



**Figure 4** Mapping of Nup153 and Tpr cleavage products. (A) Whole HeLa cell extracts from control cells (con) and apoptotic cells (apo) were immunoblotted with antibodies against specific regions of the proteins. (B) Schematic representation of the Nup153 and Tpr primary structures. The tripartite structure of Nup153 consists of NPC binding domain, zinc finger region, and XFXFG repeat domain. The Tpr protein is divided into two major domains, a coiled coil dominated, heptad repeat rich rod domain, and an acidic carboxyterminal one which contains the protein's nuclear localization signal (NLS). Amino and carboxyterminal amino acids (aa), the location of putative caspase cleavage sites (short arrows) and the relative position of the epitopes recognised by the peptide specific antibodies used in this investigation (long arrows) are indicated

site would result in a polypeptide with a predicted MW of 126 kD, which fits well with the size of the fragment observed in the Western blot.

In the case of Tpr, the two cleavage products of 190 and 200 kD were recognised only by an antibody directed against a peptide stretching from aa 1622 to 1640 ( $\alpha$ -Tpr-1), whereas an antibody binding to aa 2063–2084 ( $\alpha$ -Tpr-2) failed to give a signal. The region between aa 1640 and 2063 contains three putative caspase cleavage sites at positions D<sub>1890</sub>, D<sub>1997</sub> and D<sub>2011</sub>. While cleavage at position 1890 may give rise to the fragment of 190 kD, the second fragment could originate either by cleavage at aa 1997 or 2011. The size difference between the major and the minor fragment would be in both cases 12–14 kD, which agrees with the separation between the two Tpr-specific bands observed in the Western blot.

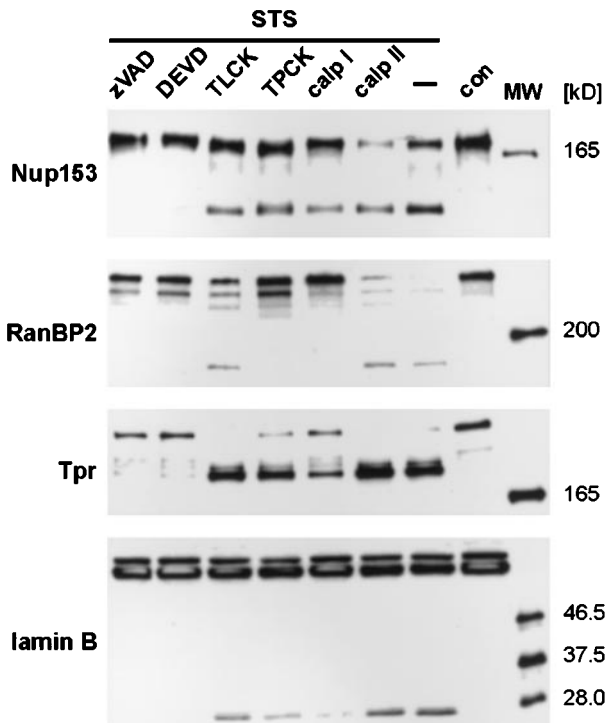
A variety of specific protease inhibitors were then tested for their ability to inhibit nucleoporin cleavage. In the STS model, cleavage of Nup153 and Tpr was most effectively, if not exclusively, inhibited by zVAD-fmk and DEVD-CHO, indicating that cleavage of these nucleoporins is dependent on caspase activity (Figure 5). Identical results were obtained also for Nup214 (not shown). Caspase inhibitors prevented processing also of RanBP2 to the 220 kD and 170 kD, but not to the 250 kD fragment. Of the other protease inhibitors, only calpain inhibitor I inhibited cleavage of RanBP2. This may reflect the broad inhibition

spectrum of this inhibitor or the possibility that besides caspases, calpains may also be involved in RanBP2 proteolysis during apoptosis.

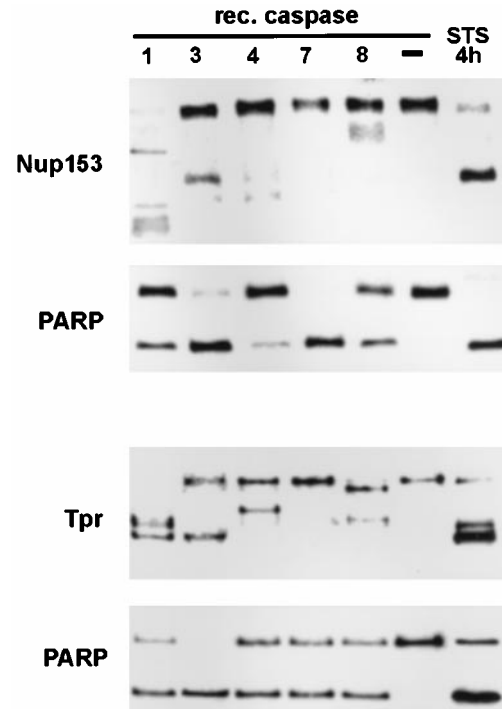
To confirm that caspases are responsible for the cleavage of Tpr and Nup153 we incubated permeabilised cells in the presence of different purified recombinant caspases. As shown in Figure 6, the Nup153 signature fragment at 130 kD was found only after incubation with caspase 3. Also other caspases cleaved Nup153 in the *in vitro* assay, but these gave rise to fragments clearly different from those seen *in vivo*: Caspase 1 consistently degraded Nup153 almost to completion, while caspase 8 yielded a weak band at 150 kD.

The *in vivo* cleavage pattern of Tpr, with the two proteolytic fragments at 190 and 200 kD was reproduced *in vitro* only after incubation with caspase 1, thus identifying Tpr as a substrate for this caspase in apoptosis. Caspase 3 cleaves Tpr only to the 190 kD fragment, possibly explaining why, *in vivo*, this fragment is more abundant compared to the 200 kD fragment, since in cells, Tpr may be cleaved by both caspases concomitantly.

Taken together, our results show that Nup153, RanBP2, Nup214 and Tpr are cleaved by caspases in the course of apoptosis. This cleavage is independent from the apoptotic stimulus, since it occurs in two distinct models of cell death and thus is likely to be a constitutive part of the apoptotic execution programme.



**Figure 5** Nucleoporin cleavage is prevented by caspase inhibitors. HeLa cells were treated with staurosporine (STS) for 4 h in the presence or absence of different protease inhibitors. Whole cell extracts of STS treated and control cells (con) were then analyzed by immunoblotting with nucleoporin specific antibodies. Inhibition of lamin B cleavage is shown as reference



**Figure 6** *In vitro* cleavage of Nup153 and Tpr by recombinant caspases. HeLa cells were permeabilised with digitonin as described and incubated with different purified caspases. After immunoblotting with  $\alpha$  Nup153 and  $\alpha$  Tpr antibodies, the filters were reprobbed with  $\alpha$  PARP antibody as a control for caspase activity. As a reference, a comparable amount of whole cell extract from cells treated with staurosporine for 4 h was loaded on the same gel

### Nuclear transport factors are abnormally distributed in apoptotic cells

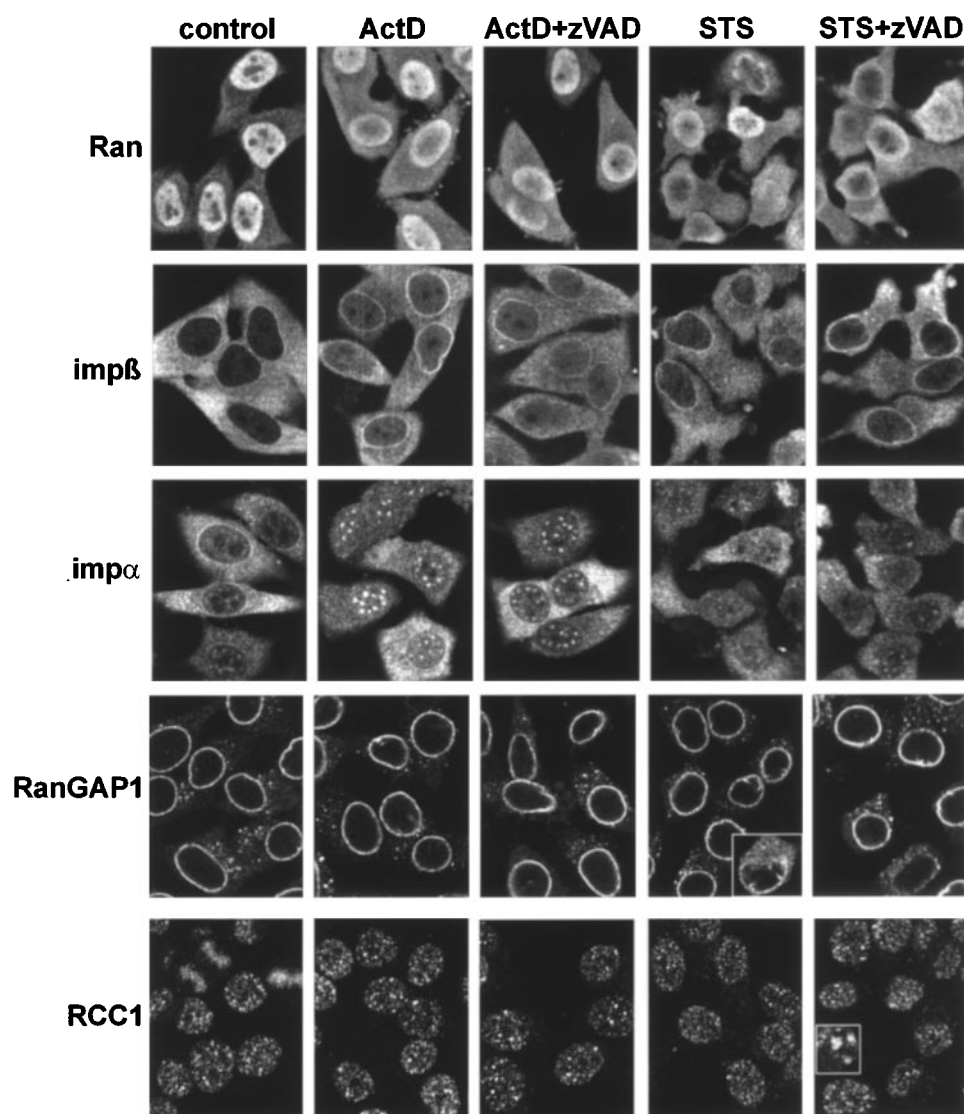
After having analyzed the cleavage of Nup153, RanBP2, Nup214 and Tpr in apoptosis, we studied the fate of shuttling nuclear transport factors that reportedly bind to these nucleoporins, i.e. importin  $\alpha$ , importin  $\beta$  and Ran.<sup>14,15,37-39</sup>

We also studied the components of the Ran cycle, RanGAP1, a GTPase activating protein for Ran<sup>40</sup> and RCC1, its nucleotide exchange factor.<sup>41</sup> In non-apoptotic interphase cells, these proteins exhibit a characteristic subcellular distribution: the majority of Ran is located within the nuclear interior, while the importins are found mainly in the cytoplasm. A small fraction of importin  $\beta$  is also known to occur bound to the NPCs.<sup>42,43</sup> RanGAP1 occurs in two fractions, one cytoplasmic and the other, which is modified by a small ubiquitin like peptide, is attached at the nuclear pore via

binding to RanBP2.<sup>44-47</sup> RCC1 is a nuclear protein which binds to chromatin.<sup>48</sup>

Immunofluorescence analysis of Ran in control cells showed the expected nucleoplasmic localisation of the protein with exclusion of the nucleoli (Figure 7). Upon treatment with STS and ActD the protein redistributed from the nuclear compartment to the cytosol. This occurred in cells with no visible sign of chromatin condensation. Although Ran still accumulated in the nucleus, the concentration gradient over the nuclear membrane was visibly diminished and nucleolar exclusion was lost.

Under control conditions, importin  $\beta$  and importin  $\alpha$  localised mainly to the cytoplasm and to a small fraction at the nuclear membrane. After treatment with both STS and ActD, the importin  $\beta$ -specific signal in the nucleus visibly increased, indicating a redistribution of the protein, while nuclear rim staining was retained. Redistribution of importin



**Figure 7** Ran, importin  $\beta$  and importin  $\alpha$  are relocated during apoptosis. HeLa cells fixed 1.5 h after staurosporine (STS) and 3.5 h after actinomycin D (ActD) stimulation in the absence or presence of the caspase inhibitor zVAD fmk were stained with the specified antibodies as described. The inset in the panel with RanGAP1 immunostaining of STS treated cells shows a late apoptotic cell with increased cytoplasmic RanGAP1 staining

$\alpha$  was clearly detectable only in STS-treated cells, where nuclear rim staining was completely lost, whereas after ActD treatment, importin  $\alpha$  appeared in speckled structures in the nucleus. Relocalisation of both importins occurred prior to chromatin condensation.

In contrast, localisation of RanGAP1 at the nuclear rim was not affected until the nuclei were completely condensed (see insert in Figure 7). In these late apoptotic cells, the RanGAP1 signal in the cytoplasm clearly increased, indicating dissociation of the protein from the NPC. Similarly, RCC1 distribution was not influenced by treatment with apoptosis inducers until the stage of chromatin condensation. In late apoptotic nuclei, RCC1 staining changed from a finely punctate pattern to a clumpy appearance reflecting its association with the condensed DNA.

To ensure that changes in fluorescence distribution were not caused by proteolytic degradation of the factors themselves, we examined their integrity by immunoblotting. As shown in Figure 8, Ran, RanGAP1, RCC1, importin  $\beta$  and importin  $\alpha$  were present in constant amounts in HeLa cells undergoing apoptosis. We could not detect any sign of proteolytic cleavage, confirming that the altered pattern of immunofluorescence observed in apoptotic cells reflected a true change in the localisation of the proteins.

Most importantly, redistribution of Ran, importin  $\beta$  and importin  $\alpha$  occurred already at 1.5 h after STS and 3.5 h after ActD treatment, i.e., when caspases were not yet active (see Figure 1) and nuclei did not exhibit any sign of

chromatin condensation. At these time points, cytochrome c release, an indicator for the activation of the caspase 9 pathway,<sup>49</sup> had not yet occurred (not shown). Additionally, redistribution was not affected by zVAD-fmk at concentrations that protected from apoptosis and completely abolished nucleoporin cleavage.

This implies, that nucleoporin cleavage was not causally linked to the observed relocalisation of Ran and importins, and that caspase-independent signals are directed to the nuclear transport machinery early during the execution of the apoptotic programme.

### The permeability of the nuclear pore is altered at an early stage of apoptosis

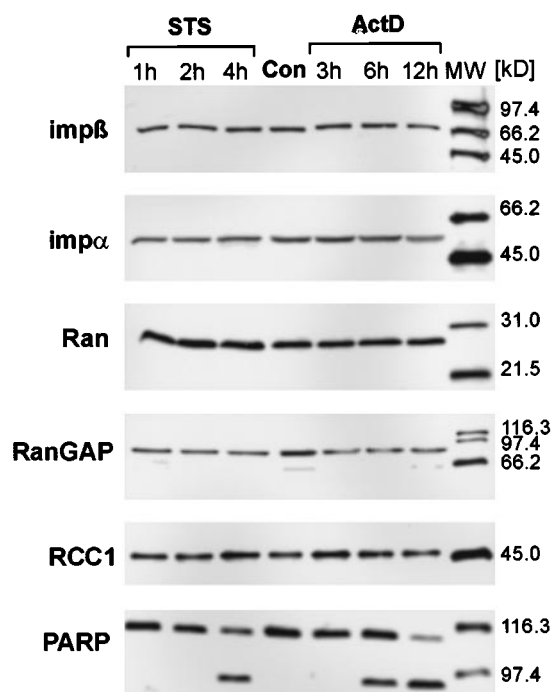
The redistribution of nuclear transport factors in early apoptotic cells may have resulted from a loss of selectivity of the nuclear permeability barrier, allowing for unrestrained passage of these factors through the nuclear pore. We therefore decided to probe the permeability of the nuclear barrier during apoptosis by microinjecting fluorescent dextrans into the cytoplasm of HeLa cells and following their distribution after apoptotic stimulation. To this purpose we tried dextrans with three different molecular weights: 40 kD being in the range of the exclusion limit of the nuclear pore, 70 kD and 150 kD clearly exceeding the exclusion limit, but differing in their diffusion rates.<sup>50</sup>

The 40 kD dextran equilibrated between the nucleus and the cytoplasm during the recovery of the cells after microinjection, whereas the 150 kD did not enter the nucleus until caspases were active and nucleoporins and lamins were cleaved (not shown). In untreated cells, the 70 kD dextran was also, as expected, efficiently excluded from the nucleus and its diffusion was not rate-limiting.

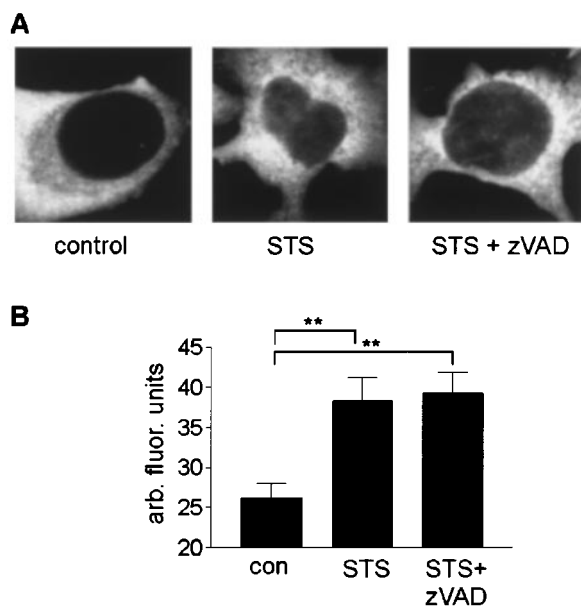
Nuclear penetration of 70 kD dextran was detected 1.5 h after stimulation with STS (Figure 9). In addition, zVAD-fmk did not prevent 70 kD dextran permeation into the nucleus, showing that modulation of the diffusion diameter of the NPC is also independent from caspases.

### poly(A)<sup>+</sup>-RNA accumulates in nuclei of cells committed to apoptosis

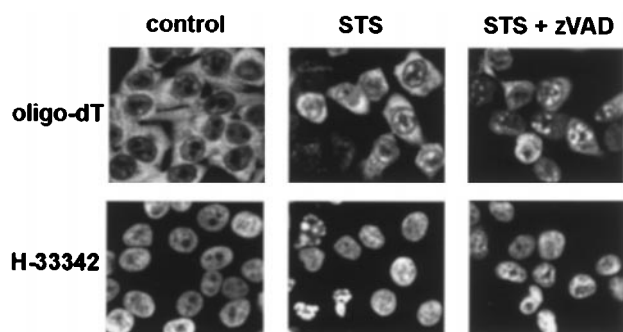
Based on the observation that early in apoptosis the permeability of the nuclear envelope increases and nuclear transport factors are relocalised, we investigated whether macromolecular trafficking across the nuclear membrane may be altered concomitantly. To this end, we studied the distribution of mRNA in non-apoptotic and stimulated HeLa cells by *in situ* hybridisation with a biotinylated oligo-dT 45-mer. As shown in Figure 10, control HeLa cells showed a high cytoplasmic poly(A)<sup>+</sup>-RNA content compared to the nucleus, which exhibited a faint staining with nucleolar exclusion. Treatment with STS for 1.5 h induced an increase in intranuclear labelling and the appearance of intranuclear speckles of intense fluorescence, indicating a high local concentration of mRNA. Notably, this occurred in nuclei that had a normal chromatin distribution as assessed by Hoechst counterstaining. Cells with condensed nuclei were not labelled by oligo-dT, indicating that at this stage the mRNA



**Figure 8** Nuclear transport factors are not degraded during apoptosis. While HeLa cell extracts were prepared from control cells and apoptotic cells treated with staurosporine (STS) and actinomycin D (ActD) at the indicated time points followed by immunoblotting with antibodies specific for importin  $\beta$ , importin  $\alpha$ , Ran, RanGAP1 and RCC1. Cleavage of PARP as marker for apoptosis is shown as reference



**Figure 9** The nucleus of cells undergoing apoptosis becomes permeable to 70 kD fluorescent dextran. **(A)** Confocal images of HeLa cells after cytoplasmic microinjection of 70 kD dextran and stimulation with staurosporine (STS) in the absence or presence of zVAD fmk. **(B)** Evaluation of nuclear fluorescence. The average pixel intensity over the nuclear area was calculated using the MCID image analysis software. Data were collected from three different experiments. 15–20 nuclei were scored in each experiment. Statistical significance was calculated using the Student's *t* test.  $**P < 0.05$



**Figure 10** Intranuclear poly(A)<sup>+</sup> RNA accumulation in apoptosis. HeLa cells were stimulated with staurosporine (STS) in the absence or presence of zVAD fmk. After 1.5 h they were fixed and subjected to *in situ* hybridisation with a biotin oligo d(T)<sub>45</sub> mer. The upper panel shows the distribution of the poly(A)<sup>+</sup> RNA, the lower panel the DNA stained with Hoechst 33342

had been degraded. Preincubation with zVAD-fmk, which prevented nuclear fragmentation as well as nucleoporin cleavage (see Figures 1 and 3), had no effect on the redistribution of the mRNA-specific fluorescence signal. This excludes the possibility that the Nup153 and Tpr fragments produced in apoptosis are responsible for nuclear mRNA accumulation.

## Discussion

The caspase family of cysteine proteases executes apoptosis in many cell systems.<sup>51</sup> While death may in some cases

intervene also in caspase-inhibited cells,<sup>52</sup> caspase-mediated proteolysis is responsible for the classical morphology of the apoptotic demise. In particular, the relevant changes, occurring in the nucleus of apoptotic cells, are believed to require caspase activity. This has supported the idea that nuclear alterations are situated downstream to the major signalling and controlling pathways of the apoptotic programme.

Here we show that factors which control nucleocytoplasmic transport, redistribute between the nucleus and the cytoplasm prior to caspase activation, as measured by DEVD-afc cleavage, and independently from caspase-mediated cleavage of nuclear proteins. Furthermore, we also show that early in apoptosis permeability of the nuclear pore complex changes, rendering it more permeable for larger-sized dextran molecules normally excluded in non-apoptotic cells. At the same time, we find that mRNA accumulates into the nucleus. This implies that a cytoplasmic/nuclear signalling system may induce early nuclear dysfunction upstream or independently from the pathways leading to caspase activation.

## Nucleoporin cleavage by caspases

Our results show that the peripheral nucleoporins Nup153, RanBP2, Nup214 and Tpr are cleaved by caspase as part of the proteolytic degradation of the nucleus. Two of these nucleoporins, namely Nup153 and Tpr are located on the nucleoplasmic side of the nuclear pore where they represent components of the nuclear basket and the pore-attached intranuclear filaments, respectively.<sup>11,53</sup> They both have been shown to bind importin  $\beta$ , although with different binding characteristics, and have been reported to be involved in mRNA export.<sup>13,39,54</sup> Nup153 has also been shown to bind Ran and to play a role in multiple types of nuclear import and export.<sup>15,55</sup>

RanBP2 is localised to the filaments which emanate from the nuclear pore on its cytoplasmic side<sup>23,24</sup> and associates with RanGAP1. RanBP2 constitutes probably the principal site where transport receptor/Ran–GTP complexes are disassembled and Ran-bound GTP is hydrolysed, which allows export receptors to release their cargo and import receptors to bind another import substrate (reviewed in<sup>33</sup>). Nup214 is also located on the cytoplasmic side of the nuclear pore and is essentially required for nucleocytoplasmic transport.<sup>56</sup> Nup214 has been shown to bind the export receptor exportin 1 in a Ran–GTP dependent manner.<sup>57</sup> Therefore, it seems conceivable, that cleavage of these nucleoporins may shut down or alter the efficiency of Ran- and importin/exportin dependent nuclear transport as part of the nuclear execution programme. In this respect, it is interesting to notice that the Nup153, 130 kD fragment still contains the Ran and importin  $\beta$  binding regions, but lacks the NPC binding domain, and therefore will likely dissociate from the NPC.

In contrast, it is unknown which part of Tpr represents the binding site for importin  $\beta$ <sup>39</sup> and consequently, whether this part is proteolytically removed from the region containing the protein's nuclear pore complex binding domain. The latter is known to reside within the 190 kD

caspace cleavage product<sup>54,58</sup> (Cordes, unpublished data), meaning that this fragment may remain associated with the NPC.

The apparent discrepancy between the electrophoretic mobility of the two Tpr fragments of 190 kD and 200 kD and the calculated molecular mass of the putative cleavage products (cleavage sites: D<sub>1890</sub> and D<sub>1997</sub> or D<sub>2011</sub>) being 210 kD and 222 or 223 kD, may be due to an abnormal mobility of these polypeptides in SDS gel electrophoresis. However, we cannot exclude that Tpr may have additional cleavage sites that do not fit the known consensus sequences recognised by caspases.

Nup153 and Tpr are cleaved *in vitro* by caspase 3 and caspase 1, respectively, yielding an identical cleavage pattern to that observed in cells. While caspase 3 is known to process a wide variety of substrates, there is only one other nuclear substrate cleaved by caspase 1 besides Tpr. This is the transcriptional factor NF-E2, whose biological role is unclear.<sup>59</sup>

Another nucleoporin, p62, which is also considered to play an important role in nucleocytoplasmic transport, was not degraded in apoptosis, although it contains at least two consensus sequences for caspase recognition in its carboxyterminal domain. Whether this protein is protected from degradation by caspases due to its central location within the NPC and its complex formation with other nucleoporins<sup>60,61</sup> remains unknown.

### **Increase of nuclear pore permeability and redistribution of transport factors in cells undergoing apoptosis**

Prior to the cleavage of the four nucleoporins we observed an increased permeability of the nuclear envelope to 70 kD dextran, which was insensitive to caspase inhibition. Variations of the diffusion limit of the nuclear pore may be reversible and subjected to regulation. This is suggested by the observation that changes in the pore diffusion limit to gold-coated particles occur in cells that transit from a quiescent to a proliferating or to a transformed state.<sup>62,63</sup> In addition, intracellular signals such as changes in nuclear calcium levels seem to modulate the passage of intermediate sized dextrans (10–70 kD) through the nuclear pore.<sup>64</sup> The pathophysiological implications of changes in nuclear permeability remain unclear. One possible effect of increasing the pore radius in apoptotic cells may be to facilitate the transfer to the nucleus of relevant signals prior to the activation of the caspase execution machinery, and independently of the possible alterations in nucleocytoplasmic transport.

The observed redistribution of Ran, importin  $\alpha$  and importin  $\beta$  may indeed be indicative that the import and export mechanisms controlling the differential localisation of these proteins may be disturbed in cells undergoing apoptosis. Although, since Ran still accumulates in the nucleus and RanGAP1 correctly localises at the NPC in STS-treated cells, at least this part of the Ran system may still be functional. A major disturbance of nuclear import may instead occur in ActD treated cells, where importin  $\alpha$  appears to be sequestered in dense nuclear bodies, from

where it is most likely not accessible for its own export receptor CAS.<sup>65</sup> Block of importin  $\alpha$  export from the nucleus may negatively affect the import of classical-NLS dependent substrates, since they would lack their interaction partner in the cytosol.

### **poly(A)<sup>+</sup>-RNA accumulation in cells undergoing apoptosis**

In cells stimulated to undergo apoptosis, mRNA accumulated into the nucleus independently from caspase activation and prior to detectable chromatin rearrangements (Figure 10). Several mechanisms could be responsible for intranuclear mRNA accumulation. Among them are the modification or degradation of essential export factors like TAP<sup>66</sup> or the disturbance of the maturation and transport of hnRNP, which are the major mRNA carriers in eukaryotic cells.<sup>67</sup> Since pre-mRNA needs to be properly spliced before export may occur,<sup>68</sup> it may also be possible that the increase in the poly(A)<sup>+</sup>-RNA-specific signal observed in preapoptotic nuclei reflects an accumulation of improperly spliced transcripts. This assumption would also be consistent with the observation that the poly(A)<sup>+</sup>-RNA-specific signal is largely localised in discrete intranuclear spots, which might represent pre-mRNA retained by the splicing machinery.

Independently of the underlying mechanisms, the early nuclear accumulation of poly-(A)<sup>+</sup>-RNA after apoptotic stimulation may itself contribute to the propagation of the apoptotic signal, for example, by suppressing the synthesis of short lived repressors of the apoptotic pathway. This may represent an additional caspase-independent amplification system that would reinforce mitochondrial and/or cytoplasmic signals.

### **Relevance of caspase-dependent and -independent nuclear alterations in apoptosis**

Two independent alterations that may affect nuclear function can be suggested by our findings: (i) an early, caspase-independent redistribution of transport factors and nuclear accumulation of mRNA and (ii) a secondary caspase-mediated disruption of the nuclear pore itself, which may eventually result in the irreversible collapse of nucleocytoplasmic trafficking.

Active import seems to be required to translocate into the nucleus some, but not other, pro-apoptotic factors. For example, the nuclear translocation of the mitochondrial protein AIF seems to require transport through the pore.<sup>17</sup> Several other factors, like caspases themselves, as well as caspase-activated molecules like Acinus, CAD and DEDD, have been suggested to exert their effects following import into the nucleus.<sup>18,22,69</sup> In view of our findings, it is unlikely that factors that are produced as a consequence of caspase activation would need active nuclear import, possibly at difference with AIF, whose nuclear effects are caspase-independent. Four major nucleoporins that are essential for nuclear import are in fact degraded simultaneously and concomitantly to the cleavage of other caspase substrates.

On the other hand, our finding that nuclear transport factors and poly-(A)<sup>+</sup>-RNA redistribute well before caspase



activation when the permeability of the NPC is also altered, strongly suggests that relevant signals may be exchanged between the nucleus and other cellular compartments in a caspase-independent fashion early during apoptosis. This may constitute an important and previously unrecognised aspect of the apoptotic programme.

## Materials and Methods

### Cell culture and apoptosis induction

HeLa 229 cells were grown in DMEM medium containing 5% foetal calf serum at 37°C in a humidified atmosphere, and passaged every 3 days by fivefold dilution into fresh medium. For induction of apoptosis, staurosporine (Sigma, Taufkirchen) and actinomycin D (Sigma, Taufkirchen) were added to a final concentration of 0.5 µM and 0.5 µg/ml respectively. For protease inhibitor assays, cells were preincubated with the inhibitor (100 µM for all inhibitors except z-VAD-fmk (N-benzyloxycarbonyl-Val-Ala-aspartyl-fluoromethylketone) (20 µM) and DEVD-CHO (N-acetyl-Asp-Glu-Val-Asp-aldehyde) (200 µM)) for 30 min before induction. All inhibitors were from Bachem Biochemica, Heidelberg, Germany.

### Viability assay

HeLa cell cultures were stained with a mixture of the membrane permeant DNA dye H-33342 (500 ng/ml) and the membrane impermeant DNA dye SYTOX (500 nM) (Molecular Probes, Eugene, OR, USA). Cells with intact plasma membrane and characteristically condensed or fragmented nuclei were scored as apoptotic. Data were collected from a minimum of three independent experiments.

### Measurement of caspase activity

Caspase-3-like activity (measured by DEVD-afc (N-acetyl-Asp-Glu-Val-aspartyl-aminofluoro-methylcoumarine) cleavage) was assayed essentially as described before.<sup>70,71</sup> Cells were lysed in 25 mM HEPES pH 7.5, 5 mM MgCl<sub>2</sub>, 1 mM EGTA, 0.5% Triton X-100. The fluorimetric assay was carried out in microtiter plates with a substrate concentration of 40 µM and a total protein amount of 3–4 µg. Cleavage of DEVD-afc was followed in reaction buffer (50 mM HEPES pH 7.5, 10 mM DTT, 1% sucrose, 0.1% CHAPS) over a period of 20 min at 37°C with λ<sub>ex</sub>=390 nm and λ<sub>em</sub>=505 nm, and the activity was calibrated with afc-standard solutions. Measurements were run in triplicate and repeated at least three times.

### Antibodies

Mab414 (BAbCO, Richmond, CA, USA) α-lamin B (Calbiochem, Schwalbach), α-PARP (Clone C2-10, Pharmingen/Becton Dickinson, Hamburg, Germany), α-RCC1 (Santa Cruz Biotechnology, CA, USA) and α-importin β (Alexis, San Diego, CA, USA) were obtained commercially. α-CAN/Nup214 was a generous gift from R Kehlenbach.

Synthetic peptides for immunisation were synthesised by t-Boc chemistry and coupled via carboxy- or aminoterminal cysteine residues to maleimide-activated keyhole limpet hemocyanin as described.<sup>53</sup> Nup153 peptides correspond to amino acids (aa) 21–36 (α-Nup153-1) and aa 391–404 (α-Nup153-3) of the human sequence.<sup>72</sup> The RanBP2 peptide corresponds to aa 2285–2314 of human RanBP2/Nup358.<sup>24</sup> Human Tpr peptides correspond to aa 1622–1640 (α-Tpr-1) and to aa 2063–2084 (α-Tpr-2) of the 2363-aa

Tpr sequence. Immunisation of guinea-pigs and affinity purification of antibodies was as reported earlier.<sup>53</sup> The mouse monoclonal antibody PF190x7A8 reactive with Nup153 has been described.<sup>10</sup> α-RanGAP1 antibodies were raised in rabbits against the purified recombinant human protein as described for the α-importin α and α-Ran antibodies.<sup>43</sup>

### SDS gel electrophoresis and immunoblotting

For the analysis of nuclear pore proteins, SDS–PAGE was performed according to Thomas and Kornberg.<sup>73</sup> Proteins were blotted onto nitrocellulose using a wetblot chamber (Biorad Trans-Blot Cell) and the filters then incubated overnight at 4°C in TNT (50 mM Tris pH 8.0, 150 mM NaCl, 0.05% Tween 20) containing 5% milk-powder. Incubation with primary antibody was overnight at 4°C. Filter washings were in TNT.

For lamin and PARP immunoblots, proteins were separated on 12% Laemmli-polyacrylamide gels and then blotted onto nitrocellulose in a Bio-Rad Semidry Blotter. Incubations and washings were as above.

### In vitro cleavage of nuclear pore proteins

For *in vitro* cleavage of Nup153, frozen permeabilised cells, prepared as described by Görlich *et al.*,<sup>74</sup> were washed in reaction buffer (50 mM HEPES pH 7.5, 10 mM DTT, 1% sucrose, 0.1% CHAPS). For caspase 1 cleavage the reaction buffer contained 100 mM HEPES pH 7.5, 10% sucrose, 10 mM DTT and 0.1% CHAPS. 4–6 × 10<sup>4</sup> cells (corresponding to 80–100 µg of total protein) were mixed with 200 ng of purified caspase 1, 3, 4, 7 or 8 in 20 µl of reaction buffer. After 1 h incubation at 37°C, the cells were pelleted, resuspended in sample buffer and analyzed by SDS–PAGE.

For the analysis of Tpr cleavage, washing and incubation steps had to be minimised due to the intrinsic instability of the protein. Cells were permeabilised in 96-well plates on ice for 5 min with 40 µg/ml digitonin. The digitonin containing solution was removed and replaced by 20 µl of reaction buffer containing recombinant caspases. After 30 min at 37°C, the plate was briefly centrifuged at 4°C, the supernatant was removed and the reaction stopped by addition of hot sample buffer.

Caspase 1 and 8 were kindly provided by DW Nicholson, caspase 4 and 7 by R Talanian and caspase 3 by FO Fackelmeyer.

### Immunocytochemistry

HeLa cells grown on coverslips were fixed in 4% freshly prepared paraformaldehyde in PBS (10 min). Cells were then washed with 50 mM NH<sub>4</sub>Cl in PBS, followed by two washes in PBS, permeabilised in 0.1% Triton X-100 in PBS for 5 min and again washed twice in PBS. For Ran immunostaining, permeabilisation was in 0.1% Triton+0.02% SDS.

Incubations with primary antibodies were for 30 min, followed by extensive washes in PBS. Incubations with Alexa 594 coupled secondary antibodies (Molecular Probes, Eugene, OR, USA) were for 20 min. Nuclei were counterstained with Hoechst-33342 at 125 ng/ml final concentration. Coverslips were mounted in Glycerol/DABCO/PBS and analyzed with a Leica TCS confocal microscope.

### In situ hybridisation

Fluorescence *in situ* hybridisation of HeLa cells employing a biotinylated oligo-dT 45mer (MWG Biotech, Ebersberg) was carried out according to Huang *et al.*<sup>75</sup> Cells grown on coverslips were fixed

for 15 min at room temperature in 3% paraformaldehyde in PBS. After three 15 min washes in PBS, the cells were permeabilised for 5 min on ice in pre-cooled 0.5% Triton X-100 in PBS. After three 5 min washes in PBS, coverslips were inverted onto 50  $\mu$ l of hybridisation mix and incubated overnight at 42°C in a humidified chamber. The hybridisation mix consisted of 2 $\times$ SSC containing 1 mg/ml *E. coli* tRNA, 10% dextran sulphate, 25% formamide and 50  $\mu$ g/ml biotin-oligo dT(45). After hybridisation, the cells were washed three times for 15 min at 42°C in 2 $\times$ SSC and once for 15 min at 42°C, in 0.5 $\times$ SSC. For detection, cover slips were first incubated with 50  $\mu$ l blocking solution (4 $\times$ SSC, 30 mg/ml BSA (Roche Biochemicals) 0.1% Tween 20) for 30 min at room temperature in a humidified chamber, followed by incubation with detection solution (4 $\times$ SSC, 10 mg/ml BSA, 0.1% Tween 20, 10  $\mu$ g/ml FITC-Avidin (Vector Laboratories)) under the same conditions. After three 5 min washes in 4 $\times$ SSC/0.1% Tween 20 at 40°C, cover slips were stained with 125 ng/ml Hoechst-33342 and mounted on a drop of Glycerol/DABCO/PBS. Fluorescent images were recorded on a Leica TCS confocal microscope.

### Microinjection of fluorescent dextran

Microinjection was performed with a semi-automated system (Eppendorf-Netheler-Hinz GmbH, Hamburg, Germany). HeLa cells were grown onto glass-bottomed petri dishes. Prior to injection, the culture medium was replaced by fresh culture medium buffered with 25 mM HEPES, pH 7.0. TexasRed-labelled dextran (Molecular Probes Eugene, OR, USA) was microinjected into the cytoplasm, the cells were then allowed to recover for 2 h in fresh medium and stimulated with 0.5  $\mu$ M STS to undergo apoptosis. For caspase inhibition, cells were incubated for 30 min with 20  $\mu$ M z-VAD-fmk prior to stimulation. One and a half hours after the addition of STS, images were recorded using a Leica TCS confocal microscope. All images were taken at similar laser intensity and instrument settings.

### Evaluation of nuclear fluorescence

Images of microinjected cells were evaluated using the MCID image analysis software (Imaging Research Inc., Canada). A total of 50–60 nuclei from three independent experiments was scored. All images were background corrected and for each cell, the average fluorescence level was measured in a region of interest corresponding to the area of the nucleus. Statistical significance was calculated using the Student's *t*-test.

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