SUPPLEMENTAL DATA

SUPPLEMENTAL TABLE I. Antibodies used in this study.

Bold letters indicate antibodies used for immunoblots in Figs 3, 4.

Target	Accession	Antibody	Epitope mapping or	Source
1	number		peptide/recombinant	
İ	number		protein sequence	
<u> </u>			segment	
Nup153	NM_005124	MAb (PF190x7A8)	aa 493-611	(1)
Tpr	U69668.1	MAb (203-37)	aa 1370-1623	(2)
Nup93	NM_014669	α-peptide polyclonal antibodies	aa 586-606	(3)
			aa 371-389	
Nup96	AAL56659	α -peptide polyclonal antibodies	aa 130-146	(3)
			aa 574-596	
			aa 880-900	
Nup98	AAL56659	α -peptide polyclonal antibodies	aa 206-219	(3)
İ			aa 596-618	
RanBP2	NM_006267	α -peptide polyclonal antibodies	aa 2180-2211	(4)
<u> </u>			aa 2553-2828	(5)
Nup107	NM_020401	α -peptide polyclonal antibodies	aa 33-51	(3)
<u> </u>			aa 437-926	(6)
Nup205	XM_371954	α -peptide polyclonal antibodies	aa 51-65	this study
İ			aa 1783-1802	(3)
Nup214	NM_005085	α -peptide polyclonal antibodies	aa 2075-2090	R.
- · • F ·				Kehlenbach
Nup133	NM_018230	α -full length protein polyclonal antibodies	#759-71 not mapped	(6)
			#756-63 not mapped	
Nup160	XP_113678	α -peptide polyclonal antibodies	aa 1059-1554	J. Koeser
Nup188	XM_497078	α -peptide polyclonal antibodies	aa 8-259	J. Koeser
11 u p100			aa 1519-1747	
Nup155	NP-705618	α -peptide polyclonal antibodies	aa 33-58	this study
			aa 719-739	
Nup75	AF514995	α -full length protein polyclonal	not mapped	J. Koeser
N. 70	CA C20410	antibodies	1.01	(2)
Nup50	CAG30419	α -peptide polyclonal antibodies	aa 1-21	(3)
			aa 126-147	.1 1
			aa 1-21 + 126-147 + 193-215	this study
Nup88	NM_002532	α-peptide monoclonal antibodies	aa 314-425	BD
	_			Bioscience
Nup62	NM_153719	α-peptide polyclonal antibodies	aa 175-187	this study
I			aa 510-522	
Nup54	NM_017426	α-peptide polyclonal antibodies	aa 310-323	this study
Nup58	AL646102	α-peptide polyclonal antibodies	aa 224-247	this study
Nup35	NP_612142	α-peptide polyclonal antibodies	aa 1-29	this study
			aa 143-166	
Nup37	NP_076962	α-peptide polyclonal antibodies	aa 76-95	this study

Nup43	AAM76708	α-peptide polyclonal antibodies	aa 365-380	this study
Seh1	NP_112493	α-peptide polyclonal antibodies	aa 342-360	this study
Sec13	NP_109598	α-peptide polyclonal antibodies	not mapped	(7)
Aladin	NP_056480	α-peptide polyclonal antibodies	aa 24-45	this study
CG1	NP_031368	α-peptide polyclonal antibodies	aa 37-59	this study
			aa 66-94	
gp210	NP_079199	α-peptide polyclonal antibodies	aa 1871-1887	this study
POM121	Q9Y2N3	α-peptide polyclonal antibodies	aa 141-153	this study
			aa 347-368	
			aa 1204-1226	
			aa 448-650	D. Goerlich

SUPPLEMENTAL RESULTS

NPC cleavage is mediated by caspases in TRAIL- and etoposide-induced apoptosis. To confirm that in TRAIL- and etoposide-induced apoptosis cleavage of individual nucleoporins is a caspase-mediated event, we made use of the pan-caspase inhibitor z-VAD-fmk. This inhibitor efficiently protected the cells both from TRAIL and etoposide-induced cell death and abolished nucleoporin cleavage. CA-074, an inhibitor of the unrelated cysteine protease cathepsin B had no or only little effect (Supplemental Fig. 1A, C). Surprisingly, in etoposide-treated cells cleavage of Nup214 was sensitive to CA-074, however this may be due to the intrinsically low level of cleavage of this protein in apoptosis. Already a small decrease in the abundance of the Nup214 fragment might drop the corresponding signal below the detection limit of the immunoblot.

LEGENDS TO SUPPLEMENTAL FIGURES

SUPPL. FIG.1 Nucleoporin proteolysis is mediated by caspases in TRAIL- and etoposide-induced apoptosis. Apoptosis was induced in HeLa cells by treatment with 300 ng/ml TRAIL for 3 h (*A*, *B*) or with 50 µM etoposide after synchronization in S-phase for 12 h (*C*, *D*) in the presence and absence of different protease inhibitors. (*A*, *C*) Whole cell extracts from control (co) and treated HeLa cells were immunoblotted with antibodies specific for the indicated nucleoporins. (*B*, *D*) The extent of apoptosis was assessed by cell staining with Hoechst-33342 and scoring for condensed nuclei. Data are means ± standard deviation from triplicate determinations.

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SUPPLEMENTAL FIG.1





