The RNA binding protein Npl3 promotes resection of DNA double-strand breaks by regulating the levels of Exo1

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<b>Strain</b>	<b>Relevant genotype</b>	<b>Source</b>
<b>JKM139</b>	MATa ho hml $\triangle$ ::ADE1 hmr $\triangle$ ::ADE1 ade1-100 leu2-3,112 lys5	$[1]$
	trp1::hisG ura3-52 ade3::GAL-HO	
<b>YLL3466</b>	JKM139 npl34::NATMX	This study
184/10A	JKM139 mec14::HIS3 sml14::KANMX	$[2]$
<b>YLL1854</b>	JKM139 MRE11-18MYC::TRP1	$\overline{3}$
DMP6178/9B	JKM139 MRE11-18MYC::TRP1 npl3A::NATMX	This study
<b>YLL3187</b>	JKM139 XRS2-3HA:: URA3	[4]
DMP6195/2A	JKM139 XRS2-3HA::URA3 npl3A::NATMX	This study
<b>YLL3501</b>	JKM139 RAD50-3HA:: URA3	$\lceil 5 \rceil$
DMP6196/1B	JKM139 RAD50-3HA::URA3 npl3Δ::NATMX	This study
<b>YLL3101</b>	JKM139 SAE2-3HA::TRP1	This study
DMP6179/8C	JKM139 SAE2-3HA::TRP1 npl3A::NATMX	This study
DMP6030/3A	JKM139 SGS1-3HA:: URA3	This study
DMP6182/7B	JKM139 SGS1-3HA::URA3 npl3A::NATMX	This study
<b>DMP5923/6A</b>	JKM139 DNA2-18MYC::TRP1	[4]
DMP6180/3A	JKM139 DNA2-18MYC::TRP1 npl3A::NATMX	This study
<b>YLL1959</b>	JKM139 EXO1-18MYC::TRP1	[4]
DMP6010/3B	JKM139 EXO1-18MYC::TRP1 npl3A::NATMX	This study
DMP6010/6C	JKM139 EXO1-18MYC::TRP1 npl34::NATMX	This study
<b>YLL1540</b>	JKM139 $exol\Delta$ ::LEU2	$\lceil 3 \rceil$
<b>YLL3287</b>	JKM139 rrp64::NATMX	[4]
DMP6293/25D	JKM139 rrp6A::NATMX npl3A::NATMX	This study
DMP6293/30A	JKM139 rrp6A::NATMX npl3A::NATMX	This study
YLL3695.1	JKM139 rrp6A::NATMX EXO1-18MYC::TRP1	This study
DMP6293/37C	JKM139 rrp6 $\triangle$ ::NATMX EXO1-18MYC::TRP1 npl3 $\triangle$ ::NATMX	This study
<b>YLL3467</b>	JKM139 NPL3-3HA::TRP1	This study
<b>YLL3012</b>	JKM139 DDC2-3HA:: URA3	[4]
DMP6009/5A	JKM139 DDC2-3HA::URA3 npl3A::NATMX	This study
DMP5991/13A	JKM139 RFA1-18MYC::TRP1	This study
DMP5991/3A	JKM139 RFA1-18MYC::TRP1 npl3A::NATMX	This study
YLL3096	JKM139 MEC1-9MYC::TRP1	[6]
DMP6238/5C	JKM139 MEC1-9MYC::TRP1 npl3A::NATMX	This study
<b>YLL3526</b>	JKM139 RFA3-3HA::TRP1	[4]
DMP6181/4A	JKM139 RFA3-3HA::TRP1 npl34::NATMX	This study
<b>YLL3222</b>	JKM139 TEL1-3HA::NAT	[6]
DMP6590/19C	JKM139 TEL1-3HA::NAT npl3Δ::NAT	This study
344-115B2	MATa his 3-513:: TRP1:: his 3-537 ura 3-52 trp1 leu2	[7]
YLL3873	$his3-513::TRPI::his3-537$ MATa $ura3-52$ trp1 leu2	This study
	$npl3\Delta$ :: $KAN$	

**Table S1.** *Saccharomyces cerevisiae* **strains used in this study**





Figure S1. Loading control of western blot in Figure 1. (A-C) Galactose was added to exponentially growing (A,B) or G2-arrested (C) YEPR cell cultures to induce HO. Protein extracts were subjected to western blot with anti-Rad53 (A,C) or anti-HA (B) antibodies or stained with Coomassie as a loading control. (D-E) YEPD G1-arrested cell cultures (αf) were released in fresh medium containing CPT (50  $\mu$ M) (D) or MMS (0,02%) (E). Protein extracts were subjected to western blot with anti-Rad53 antibodies or stained with Coomassie. (F) Phleomycin (15 μg/ml) was added to YEPD G2-arrested cell cultures in the presence of nocodazole to maintain the G2 arrest. Protein extracts were analyzed by western blot with anti-Rad53 antibodies or stained with Coomassie.



**Figure S2. High levels of RNase H1 do not suppress the checkpoint defect of cells lacking Npl3.** (A) Exponentially growing cell cultures of wild type JKM139 and an otherwise isogenic *npl3*Δ strain, both carrying a centromeric plasmid either expressing the *RNH1* gene from the *tetO* promoter or empty (vect), were arrested in G1 with  $\alpha$ -factor and plated on galactosecontaining plates (time zero) to follow microcolonies formation. (B) YEPR exponentially growing cell cultures were transferred in YEPRG to monitor Rad53 phosphorylation by western blot. The same amounts of protein extracts were separated on SDS-PAGE and stained with Coomassie as a loading control.





Figure S3. Loading control of western blot in Figure 3. Exponentially growing YEPR cell cultures were transferred to YEPRG at time zero. Protein extracts were subjected to western blot analysis with anti-Rad53 antibodies or stained with Coomassie as a loading control.



**Figure S4. Npl3 is not enriched at an HO-induced DSB.** (A) Schematic representation of *MAT* locus on chromosome III. The relative positions of the HO cleavage site and of primer pairs used for chromatin immunoprecipitation (ChIP) are shown. (B) G2-arrested YEPR wild type (no tag) and *NPL3-HA* cell cultures were transferred to YEPRG in the presence of nocodazole to maintain the G2 block and subjected to ChIP analysis with anti-HA antibodies and subsequent qPCR. Relative fold enrichment of the Npl3-HA fusion protein at the indicated distances from the HO cleavage site was determined. Plotted values are the mean values +SD  $(n=3)$ .



Figure S5. Levels of resection proteins in the absence of Npl3. Exponentially growing YEPR cell cultures of JKM139 derivative strains expressing the indicated tagged proteins were transferred in YEPRG at time zero. Protein extracts were subjected to western blot with antibodies specific for the indicated tags. The same amounts of protein extracts were separated on SDS-PAGE and stained with Coomassie as a loading control.



**Figure S6. Exo1 is not degraded by proteasome in the absence of Npl3.** Protein extracts prepared from exponentially growing YEPD cell cultures treated with the proteasome inhibitor MG132 (75  $\mu$ M) for 3 hours (+) or untreated (-) were subjected to western blot analysis with anti-MYC antibodies. The same amounts of protein extracts were separated on SDS-PAGE and stained with Coomassie as a loading control.



**Figure S7. The lack of Npl3 increases the defects of cells lacking Sae2 or Sgs1.** (A) Exponentially growing cell cultures of JKM139 derivative strains were serially diluted (1:10) before being spotted out onto YEPD plates with or without CPT. (B) Meiotic tetrads from diploid cells with the indicated genotype were dissected on YEPD plates that were incubated at 30°C for 3 days, followed by spore genotyping.



**Figure S8. Exo1 high levels do not reduce the elevated frequency of mitotic recombination caused by Npl3 lack.** Wild type and *npl3Δ* strains carrying the *his3-513::TRP1::his3-537* heteroallelic duplication on chromosome XV were transformed with either a *EXO1* 2μ plasmid or an empty vector  $(2\mu)$ . 10 independent clones for each strain were plated on complete medium plates to evaluate their viability and on plates lacking histidine to select the His+ recombinants generated by mitotic recombination at the *HIS3* locus. The number of the His+ colonies was evaluated and normalized to the viability of the strain to determine the recombina-Figure S8. Exo1 high levels do not reduce the elevent<br>since  $\frac{d}{dx}$   $\frac{d}{dx}$   $\frac{d}{dx}$ <br> $\frac{d}{dx}$ <br> $\frac{d}{dx}$ <br>Figure S8. Exo1 high levels do not reduce the elevent<br>caused by Npl3 lack. Wild type and  $npl3\Delta$  strain<br>heteroa



Figure S9. Loading control of western blot in Figure 5. Exponentially growing YEPR cell cultures were transferred in YEPRG at time zero. Protein extracts were subjected to western blot with anti-Rad53 antibodies or stained with Coomassie as a loading control.



Figure S10. Levels of checkpoint proteins in the absence of Npl3. Exponentially growing YEPR cell cultures of JKM139 derivative strains expressing the indicated tagged proteins were transferred in YEPRG at time zero. Protein extracts prepared at different time points after HO induction were subjected to western blot analysis with antibodies specific for the indicated proteins or tags. The same amounts of protein extracts were separated on SDS-PAGE and stained with Coomassie as a loading control.

![](_page_11_Figure_0.jpeg)

**Figure S11. Cell cycle arrests of cells treated with UV.** (A) Exponentially growing YEPD cell cultures of wild type JKM139 and otherwise isogenic *npl3Δ* and *exo1Δ* strains (exp) were arrested in G1 with  $\alpha$ -factor ( $\alpha$ f), UV irradiated (75 J/m<sup>2</sup>), and held in G1 in the presence of α-factor. FACS analyses of DNA content to verify the cell cycle arrest in G1. (B) Exponentially growing YEPD cell cultures of the strains in (A) (exp) were arrested in G2 with nocodazole (noc), UV irradiated  $(75 \text{ J/m}^2)$ , and held in G2 in the presence of nocodazole. FACS analyses of DNA content to verify the cell cycle arrest in G2.

## **References**

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