

# **Double-strand breaks and novel DNA intermediates derived from partial processing of alkylation damage in G2 cells of yeast**

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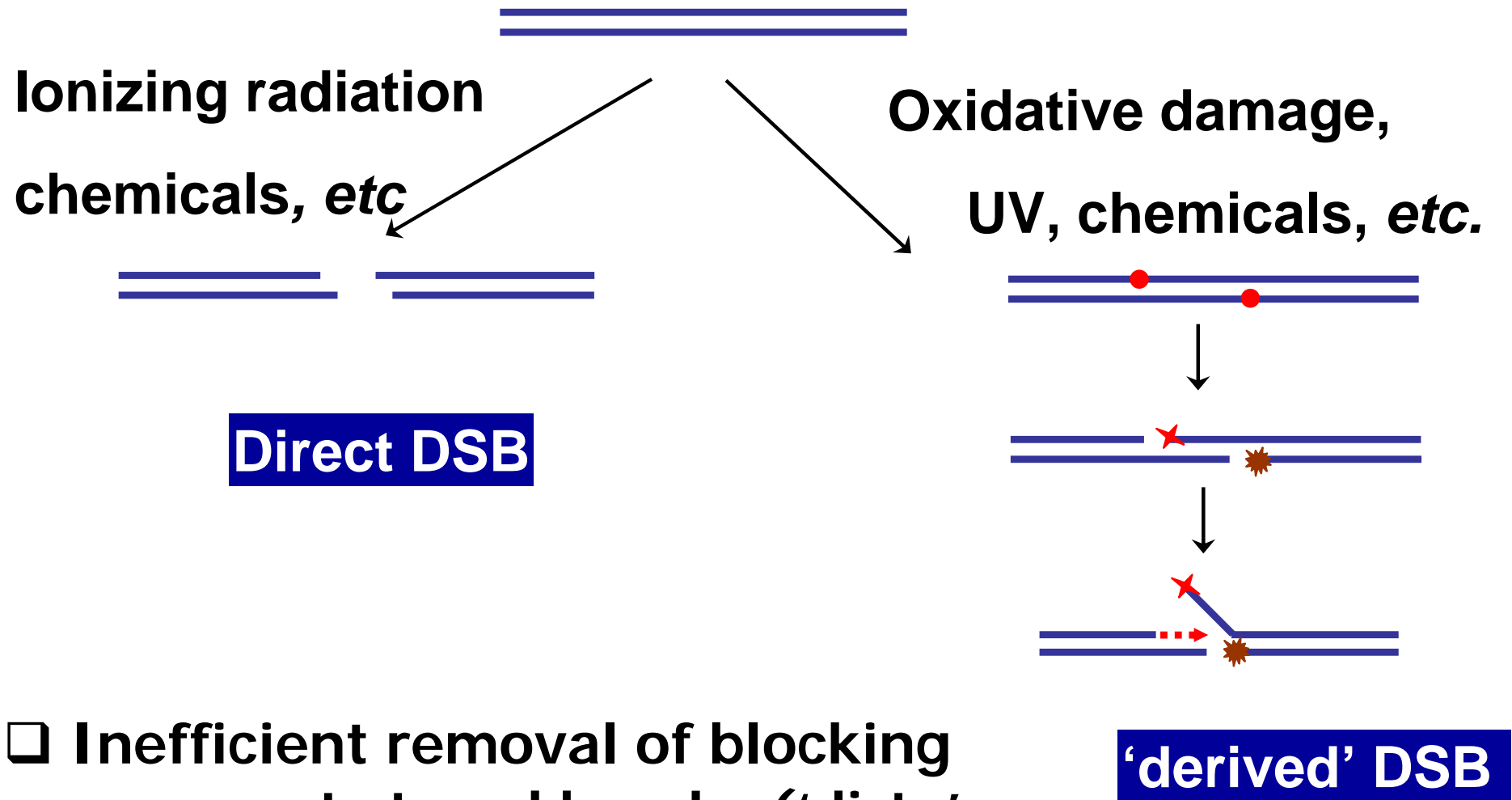
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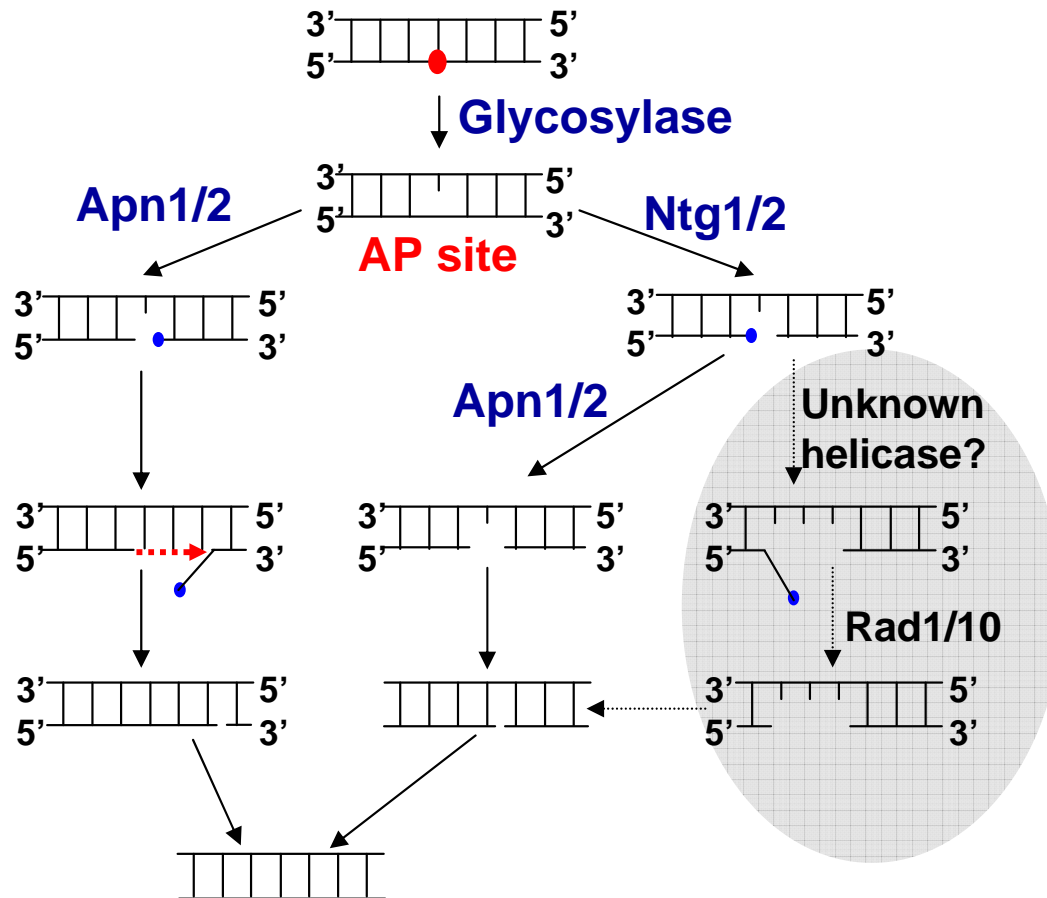
National Institute of Environmental Health Sciences (NIEHS)

# 'Derived' DSBs and 'dirty' break ends



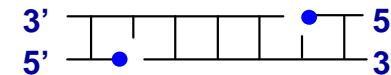
- ❑ Inefficient removal of blocking groups at strand breaks ('dirty' ends) might lead to derived DSBs

# Yeast **Base Excision Repair (BER)**, AP endonucleases and 3'-dirty ends



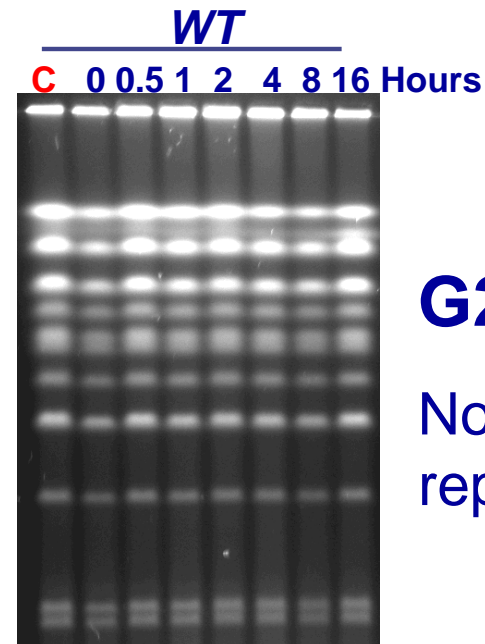
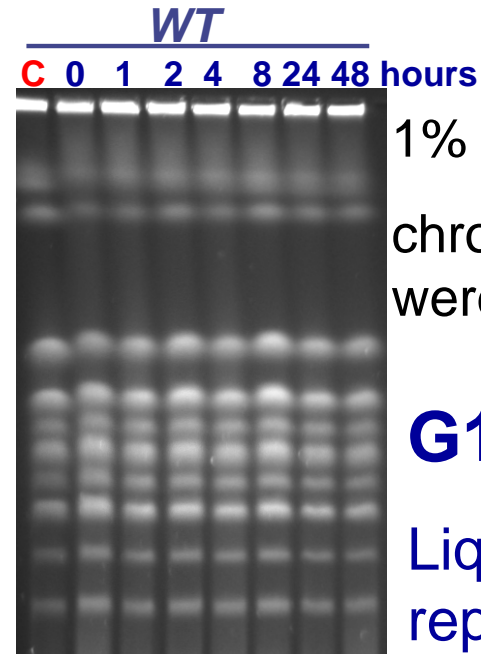
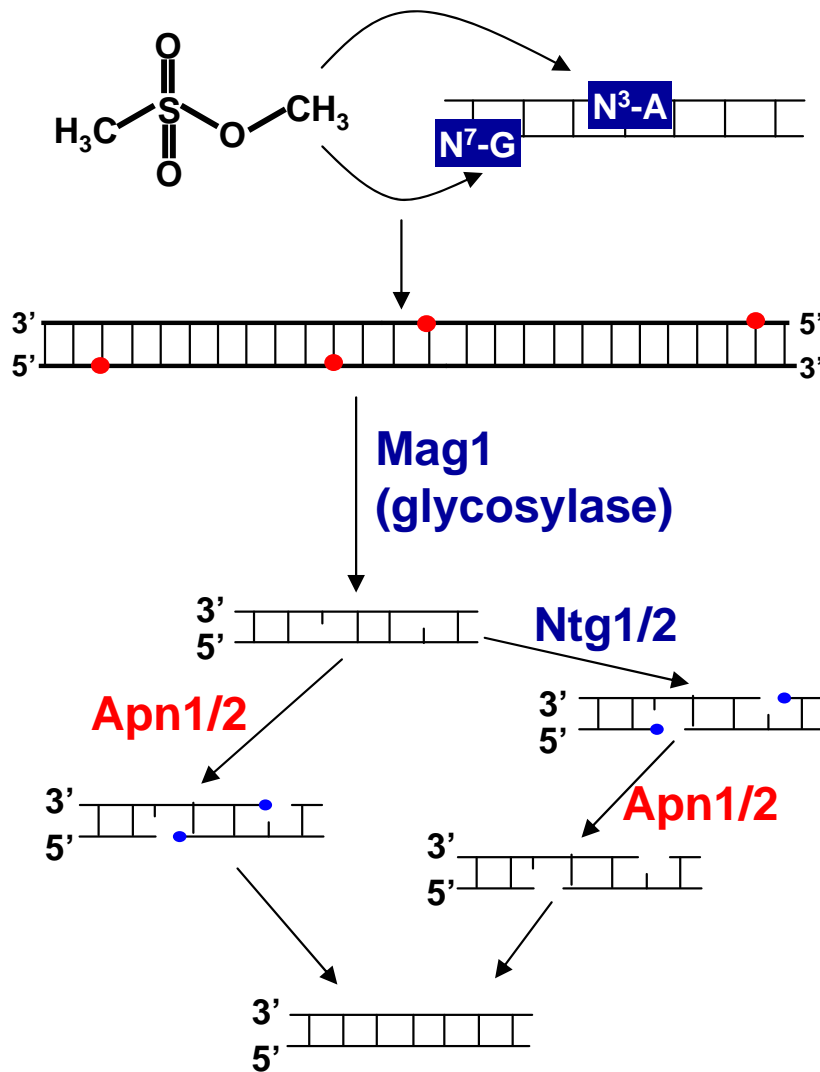
1. Can base damage generate DSBs outside of S-phase?

2. Can repair generated SSBs be converted into DSBs within the cell?

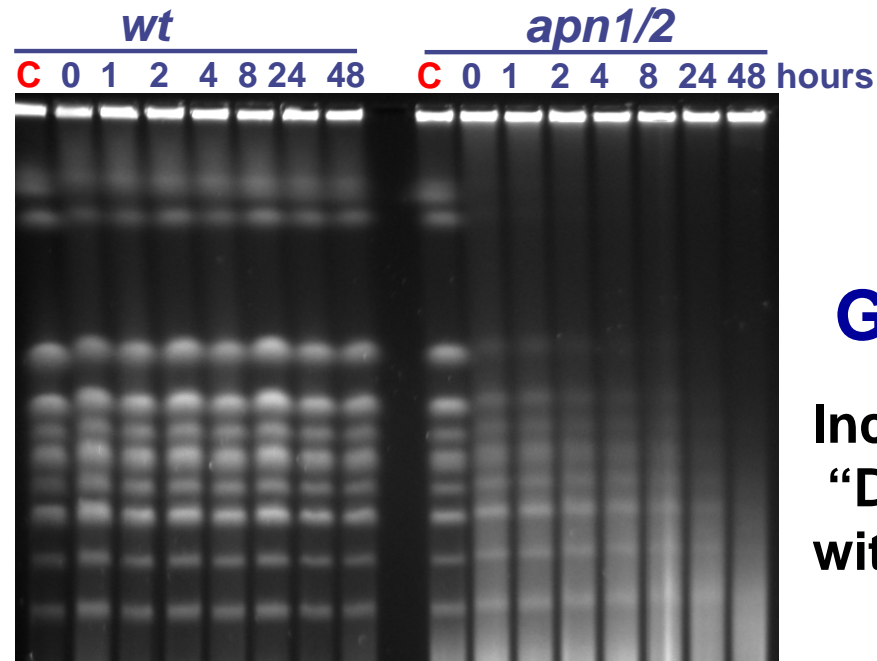
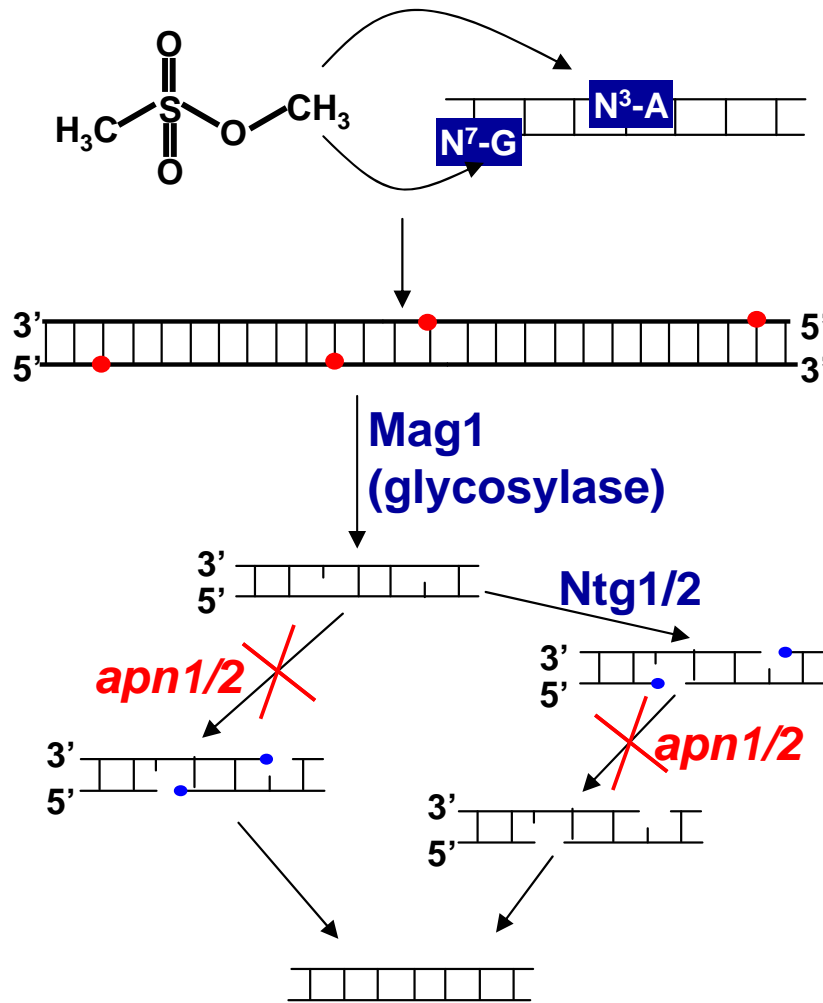


3. What are the consequences of 3'-dirty ends?

# Does MMS generate DSBs in wild type G1 or G2 cells?

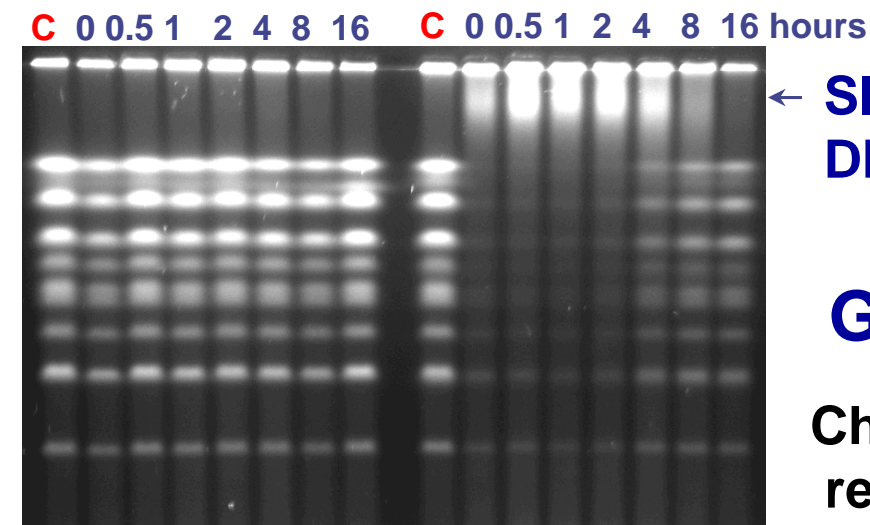


# Does partial processing of alkylation damage (*apn1*<sup>-</sup>, *apn2*<sup>-</sup>) lead to “DSBs” in G1 and/or in G2



**G1**

Increase in  
“DSBs”  
with time



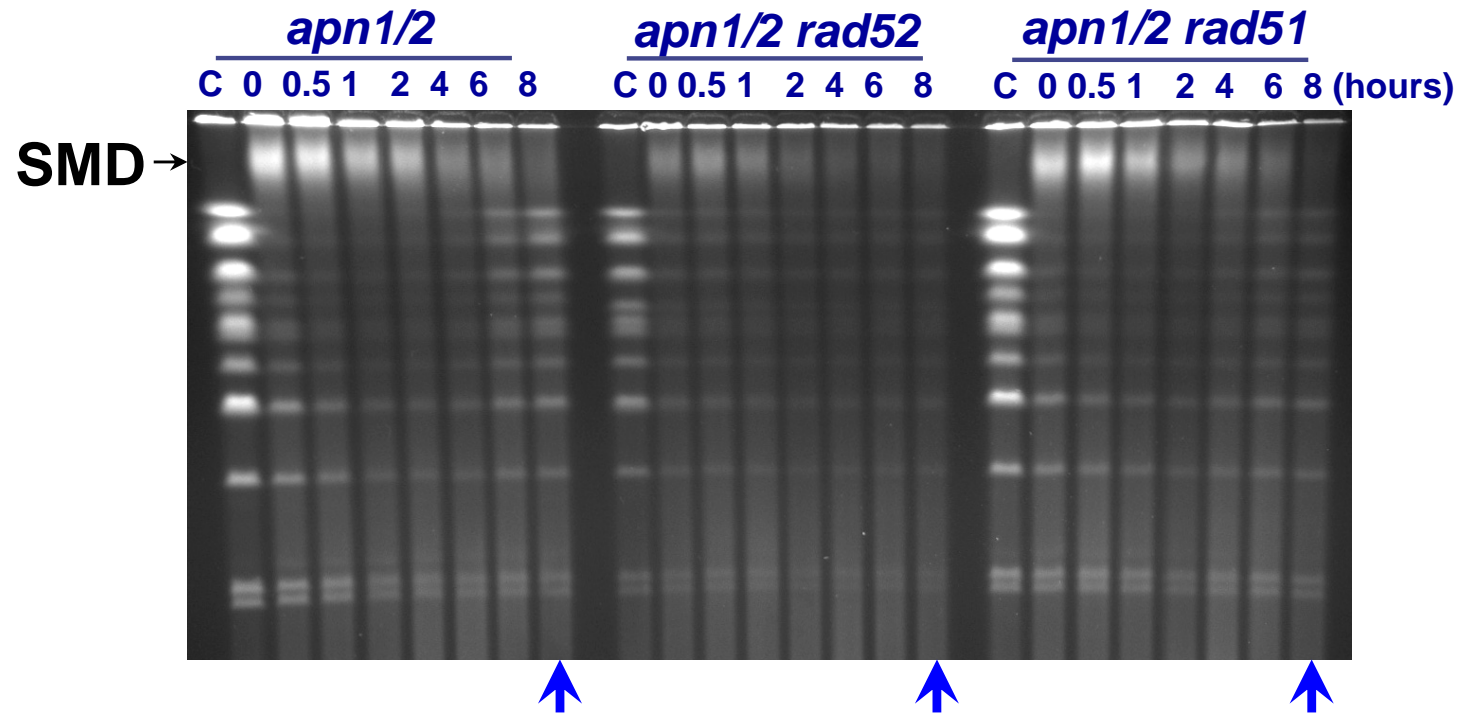
← **Slow Moving DNA (SMD)**

**G2**

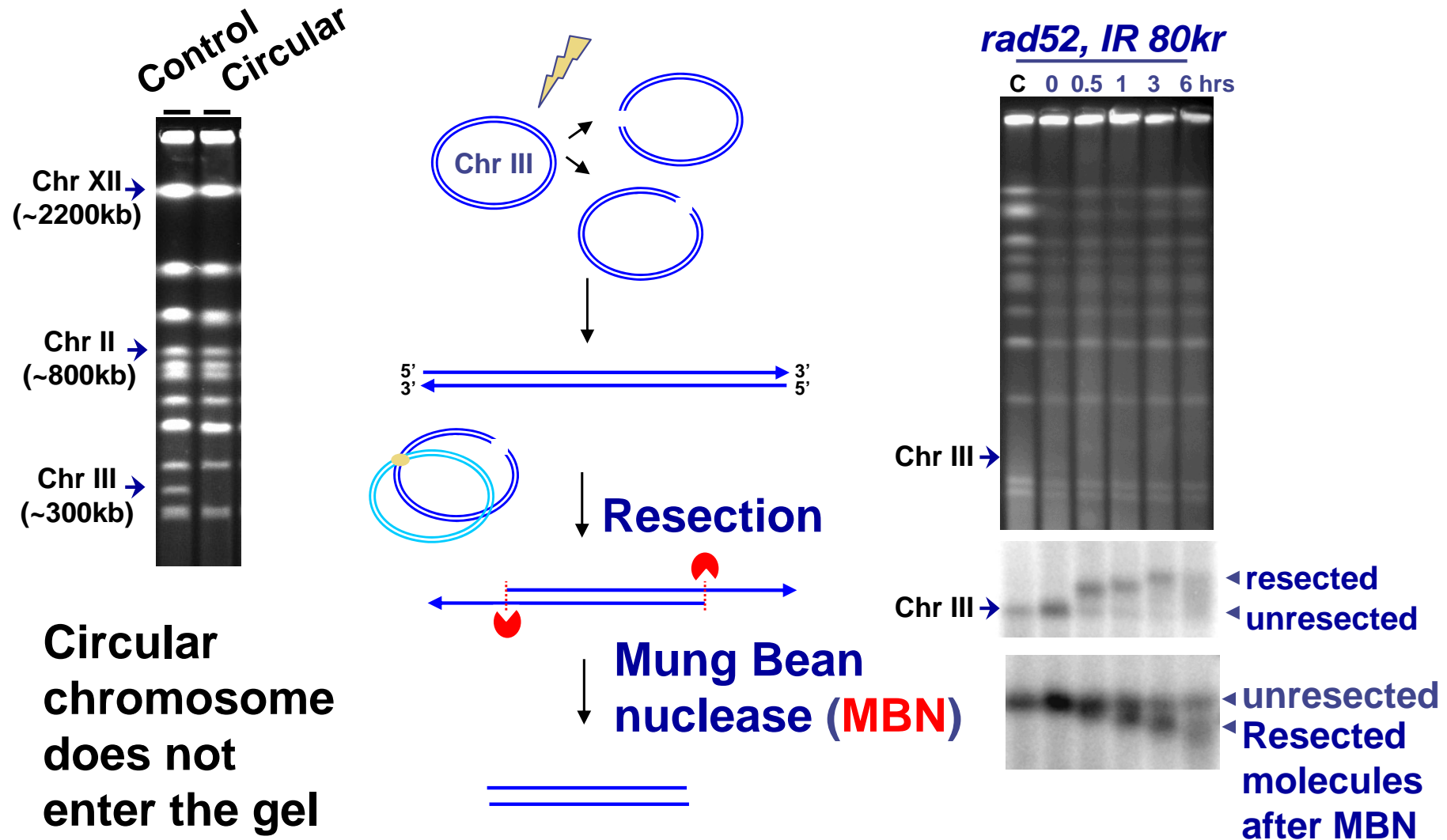
Chromosome  
restitution

G1 cells, Ma et al., NAR, 2008; G2 cells, Ma et al., PLoS Genetics, in press

# Chromosome restitution--unlike SMD--occurs through DSB recombinational repair in G2



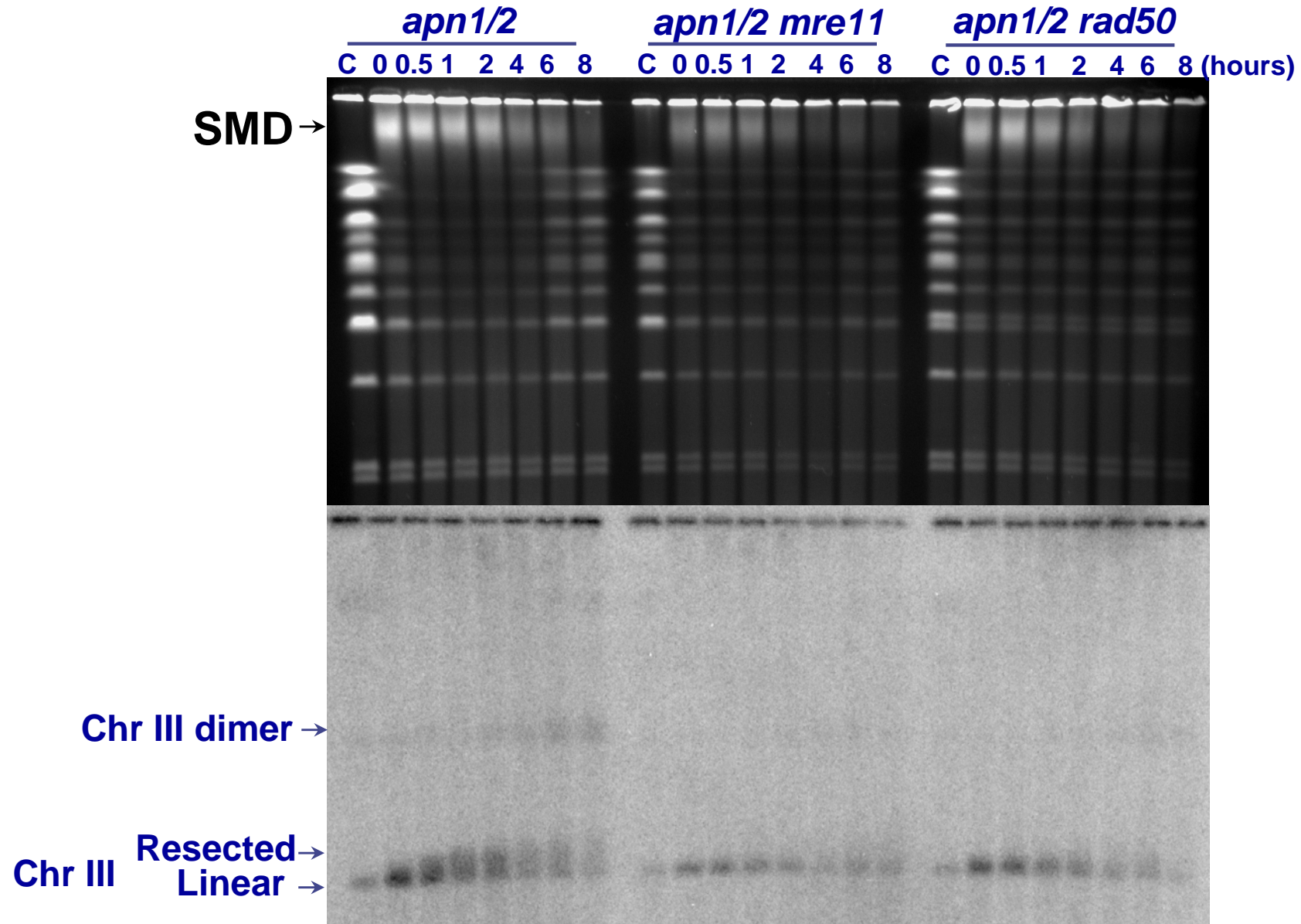
# Are the random DSBs generated by MMS in G2 cells subject to resection, an early step in DSB repair





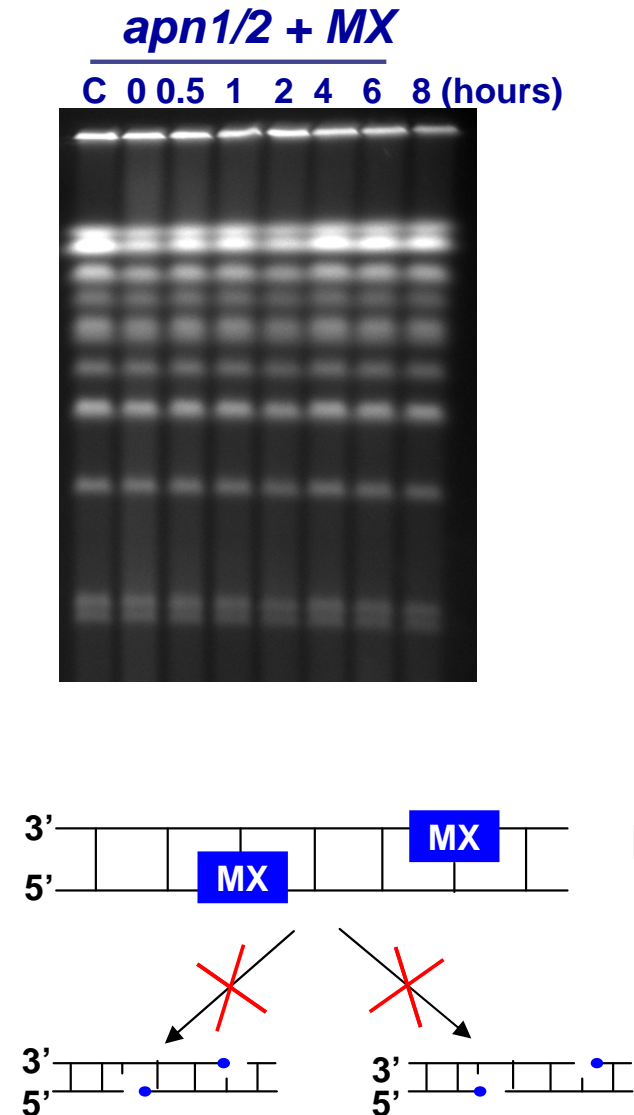
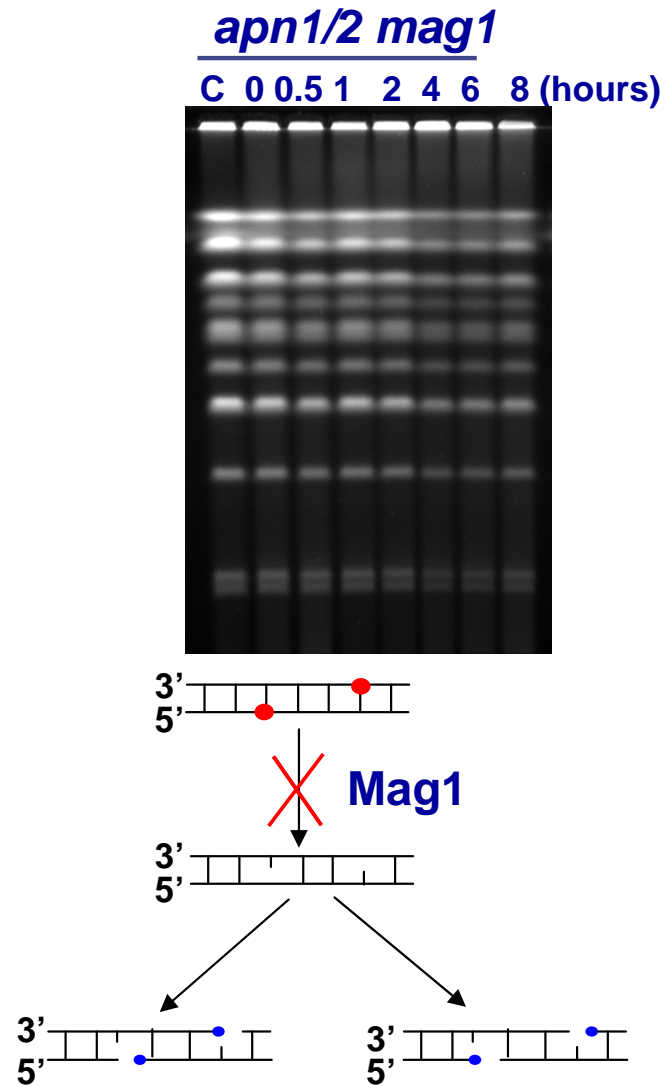
MMS-derived DSBs in G2 cells **are** resected;  
Resection depends on Rad50/Mre11 (MRX)

**Resection does not depend on Rad51 or Rad52 (not shown)**

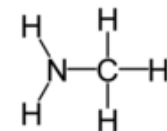




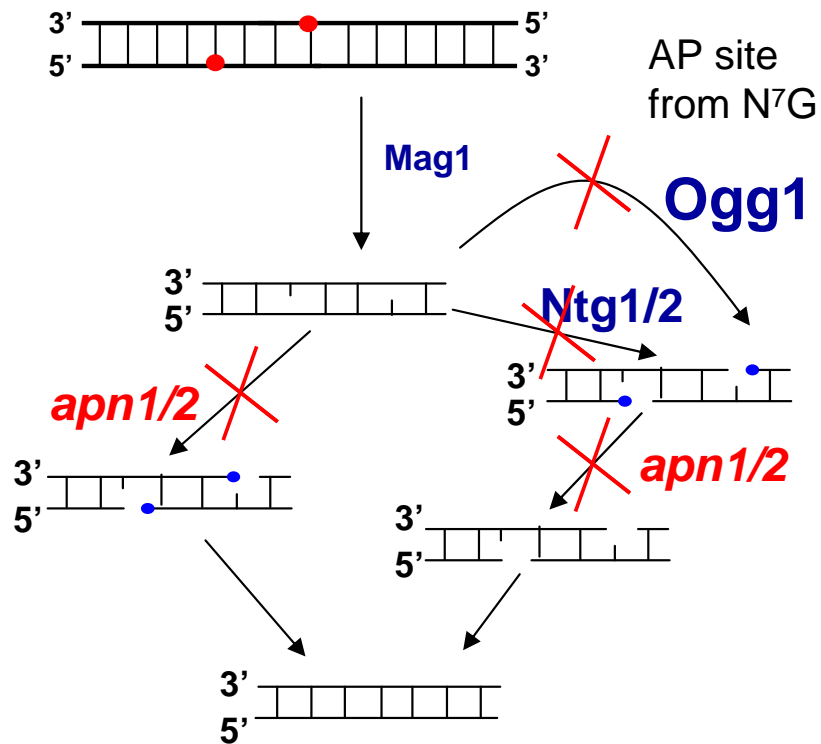
# Deletion of *MAG1* or stabilization of AP sites prevents SMD as well as DSBs



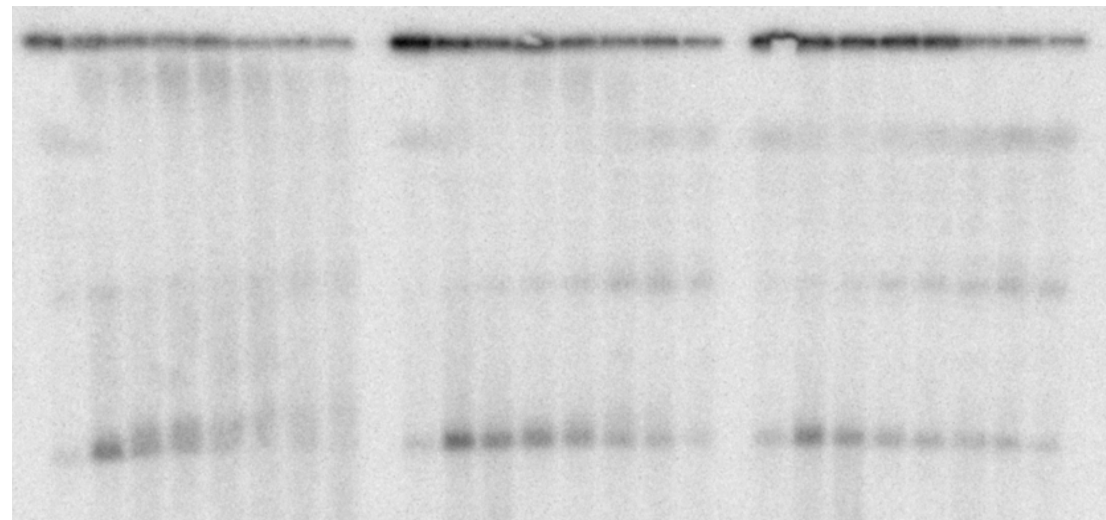
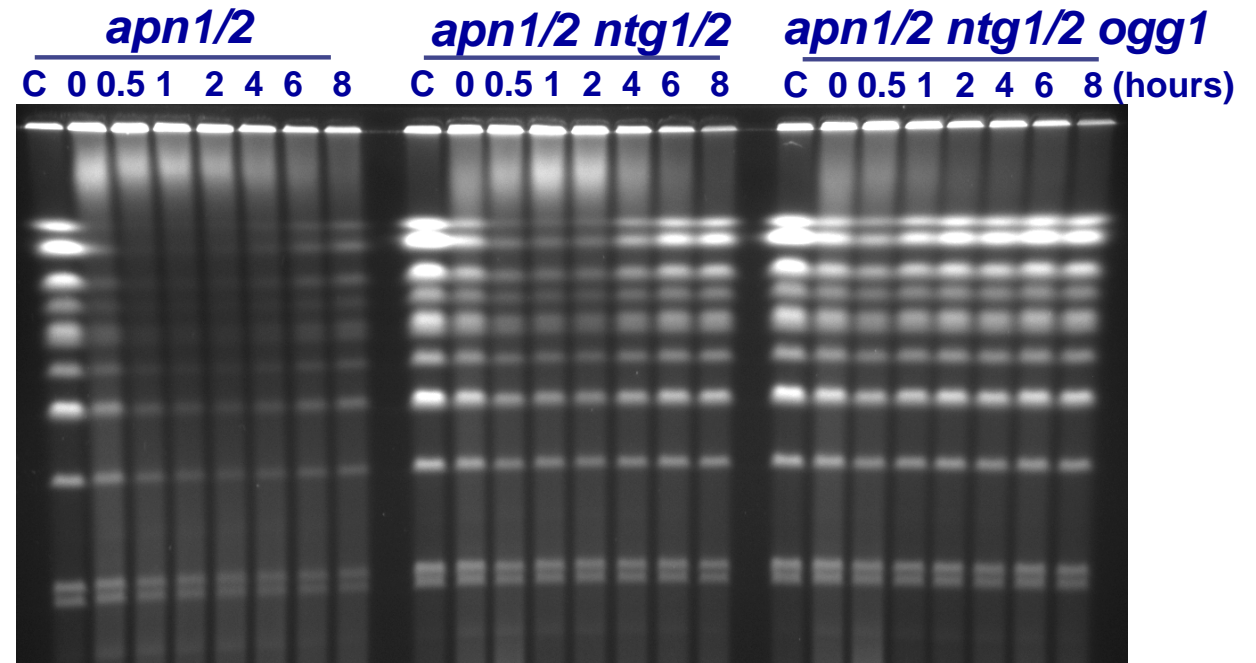
**Methoxyamine (MX)  
binds to AP sites**



# Resectable DSBs and SMD in the *apn1 apn2* mutant are due to a combination of Ntg1/2 and Ogg1



Chr III dimer →  
 Chr III Resected →  
 Chr III Linear →



# Derived DSBs and SMD intermediates resulting from partial processing of MMS lesions

