

NEIL3-mediated mitotic base excision repair of oxidative lesions at telomeres prevents senescence in hepatocellular carcinoma

Helge Gad^{1,3*}, Zhao Zhenjun², Carlos Benitez-Buelga¹, Kumar Sanjiv¹, Hua Xiangwei⁴, He Kang², Feng Mingxuan², Zhao Zhicong², Ulrika Warpman Berglund¹, Xia Qiang², Thomas Helleday^{1,3}

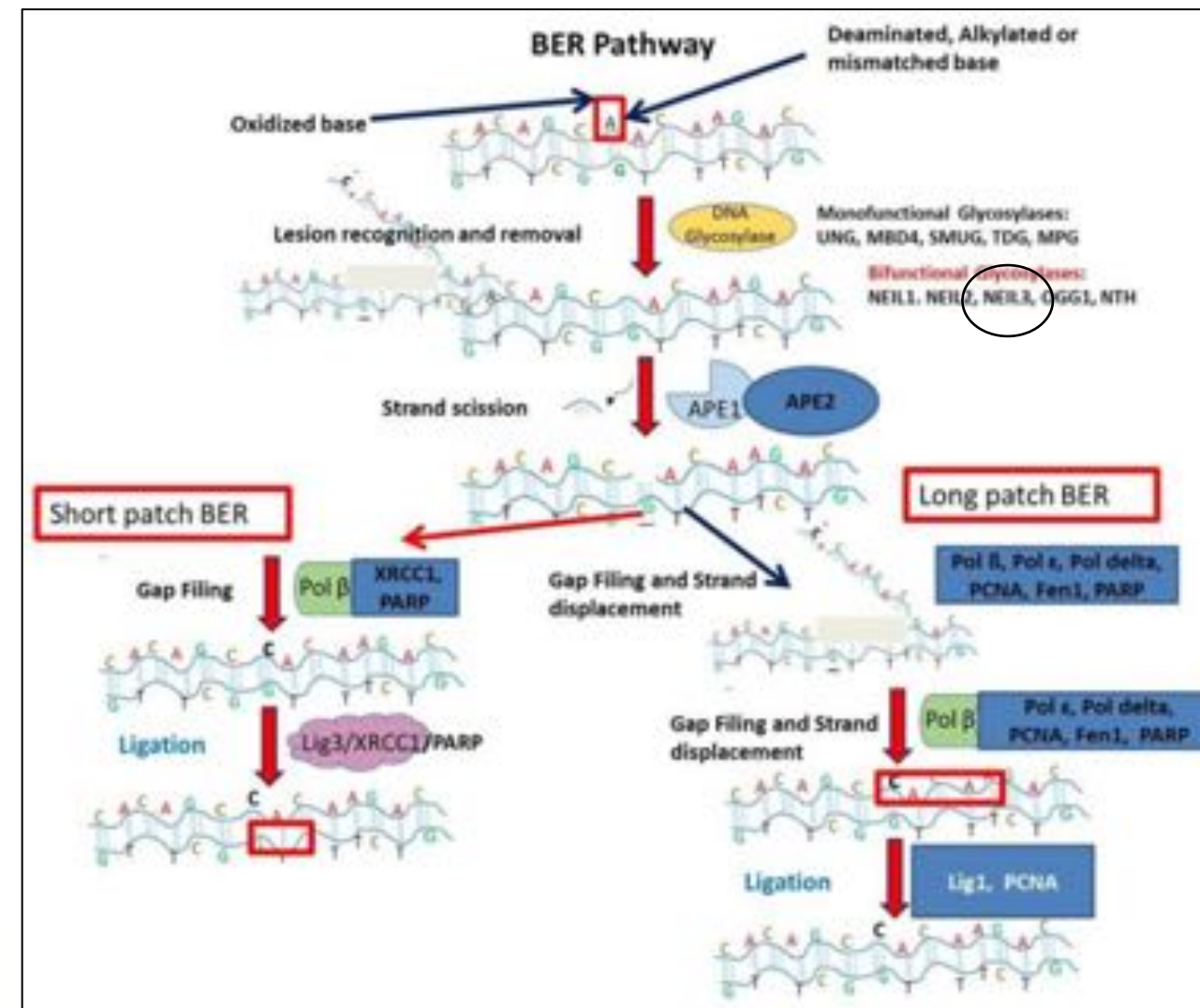
1) Science for Life Laboratory, Department of Oncology-Pathology, Karolinska Institutet, S-171 76 Stockholm, Sweden. 2) Department of Liver Surgery, Renji Hospital, School of Medicine, Shanghai Jiao Tong University, Shanghai, China. 3) Weston Park Cancer Centre, Department of Oncology and Metabolism, University of Sheffield, Sheffield S10 2RX, UK. 4) Organ Transplantation Center, the Affiliated Hospital of Qingdao University, Qingdao, China

Introduction

Hepatocellular carcinoma (HCC) is the most common type of primary liver cancer and the third leading cause of cancer death worldwide. Chronic liver diseases contributes to accumulation of reactive oxygen species (ROS) and inflammation, leading to cirrhosis and eventually HCC. Although many approaches have been suggested to treat HCC (e.g., surgery, transarterial chemoembolization (TACE), immunotherapy and targeted therapies), effective drugs and non-surgical treatment for HCC patients remain very limited

ROS production increases during G2/M-phase and induction of oxidative damage in this phase arrests cells in pro-metaphase, suggesting that the level of endogenous DNA damage varies during the cell cycle and have more severe effects in certain phases.

It is well established that there is differential DNA repair at telomeres. Telomeres are sensitive to oxidative damage, resulting in cell senescence, chromosome fusion and apoptosis. While BER pathways are described to be active at telomeres, information about these processes or potential differential roles of glycosylases are generally lacking.



Summary

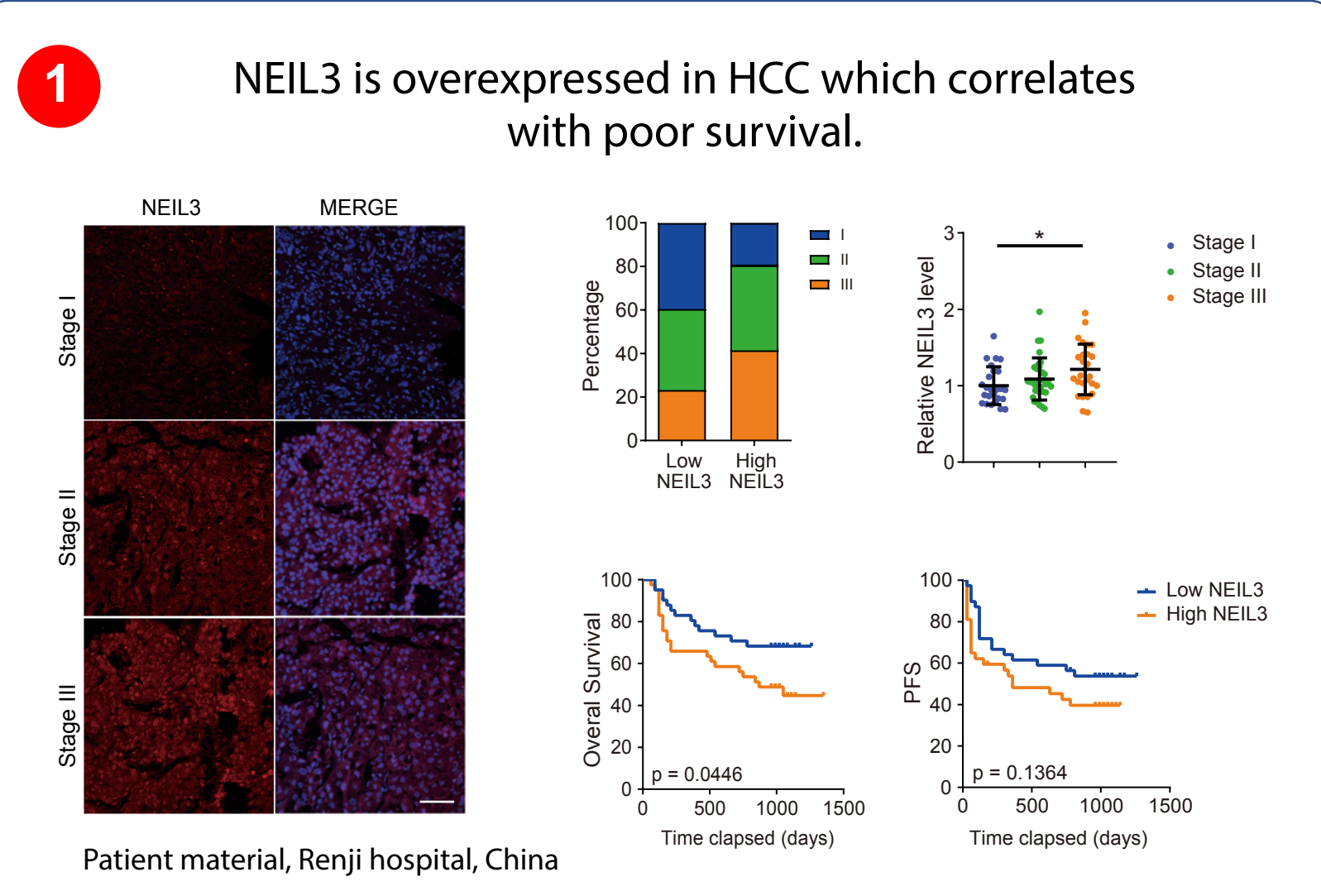
NEIL3 is overexpressed in HCC which correlates with poor survival. NEIL3 depleted HCC cell lines accumulate oxidative DNA lesions specifically at telomeres, resulting in Telomere dysfunctional foci (TIFs) and 53BP1 foci formation.

NEIL3 relocates to telomeres following oxidative DNA damage during mitosis and recruits apurinic endonuclease 1 (APE1), indicating activation of base excision repair.

NEIL3, but not NEIL1 or NEIL2, is required to initiate base excision repair at oxidized telomeres that is dependent on APE1 and Polβ.

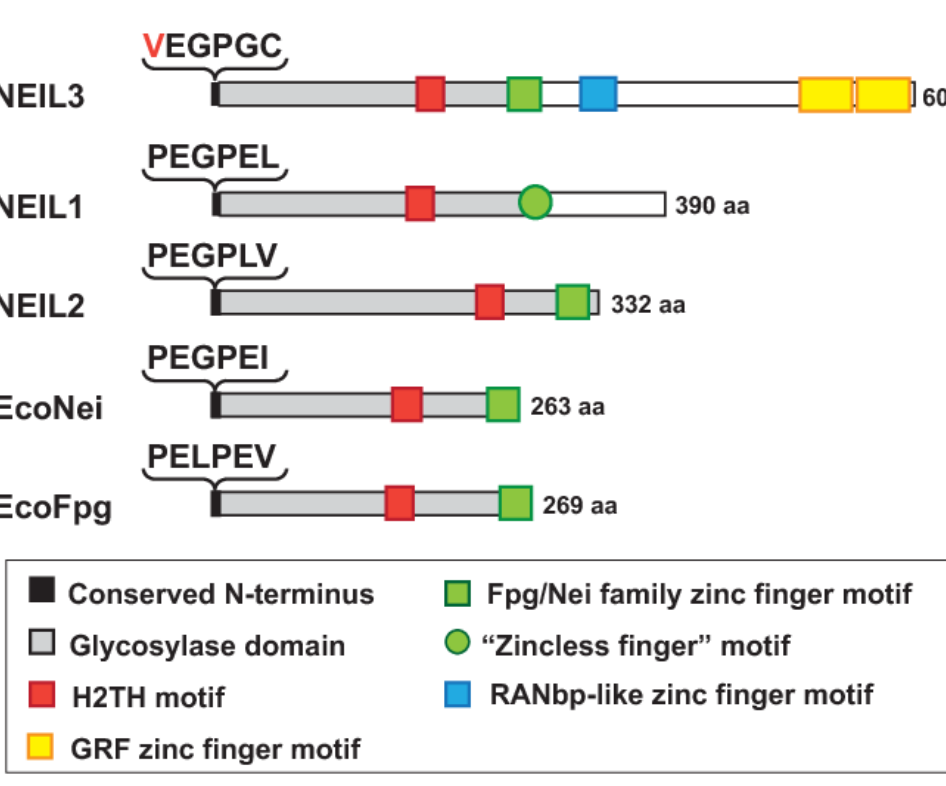
Repetitive exposure of oxidizing damage in NEIL3 depleted cells induced chromatin bridges and damaged telomeres.

These data suggest NEIL3 could be a target for therapeutic intervention of HCC, and perhaps a combination treatment with a NEIL3 inhibitor and oxidizing compounds (e.g., elesclomol) could prove to be a good strategy to induce ROS and prevent its repair in the tumour.

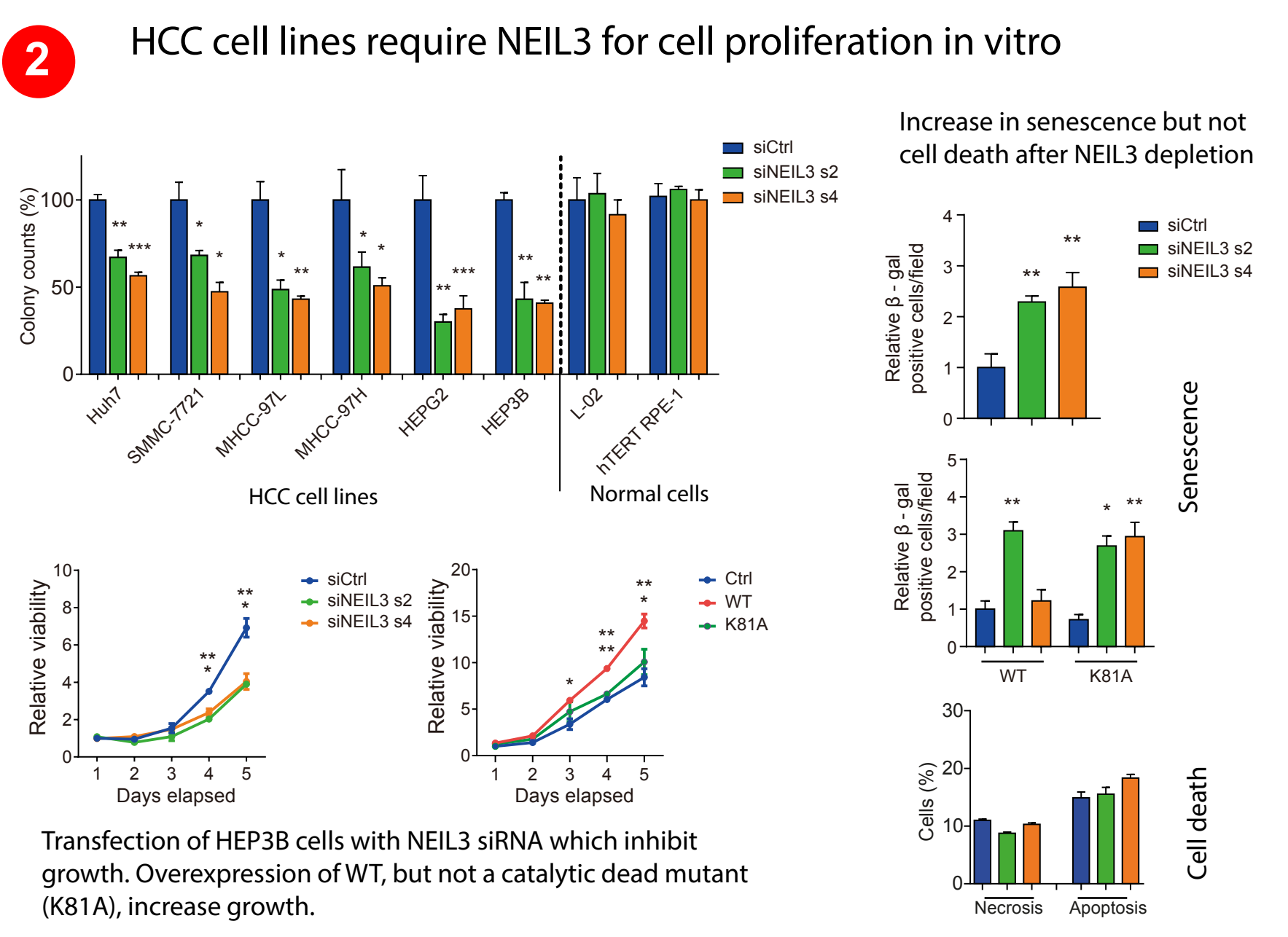


Oxidative damaged bases in DNA are repaired by Base Excision Repair (BER). NEIL3 is a multi-functional glycosylase removing hydantoins (spiroiminodihydantoin (Sp) and guanidinohydantoin (Gh)) and thymine glycol (Tg) from the DNA. NEIL3 has been implicated in repair of inter-strand crosslinks, replication associated damage and telomere damage. NEIL3 has an unstructured C-terminal domain that can interact with TRF1, FEN-1, PCNA and APE-1.

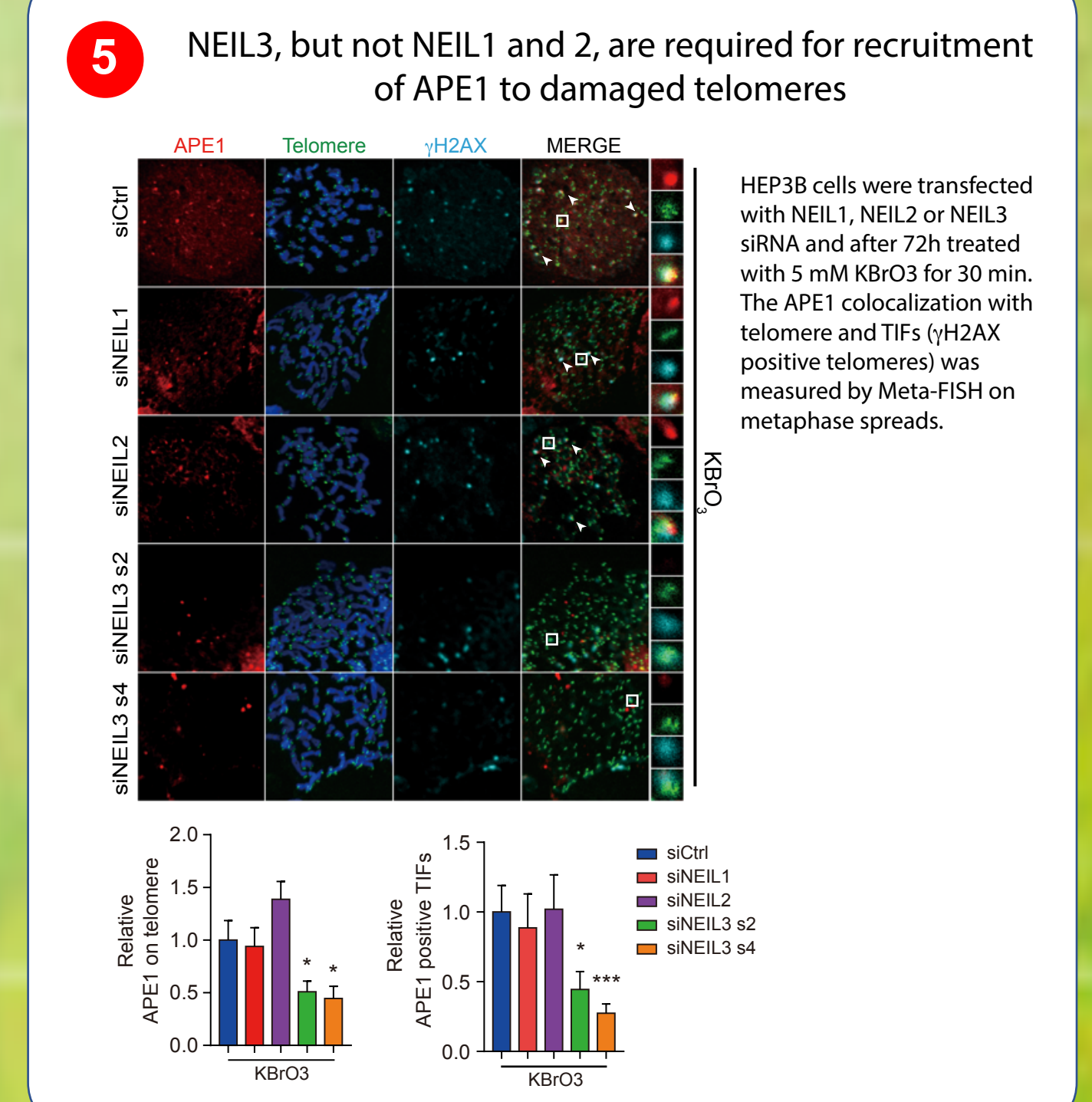
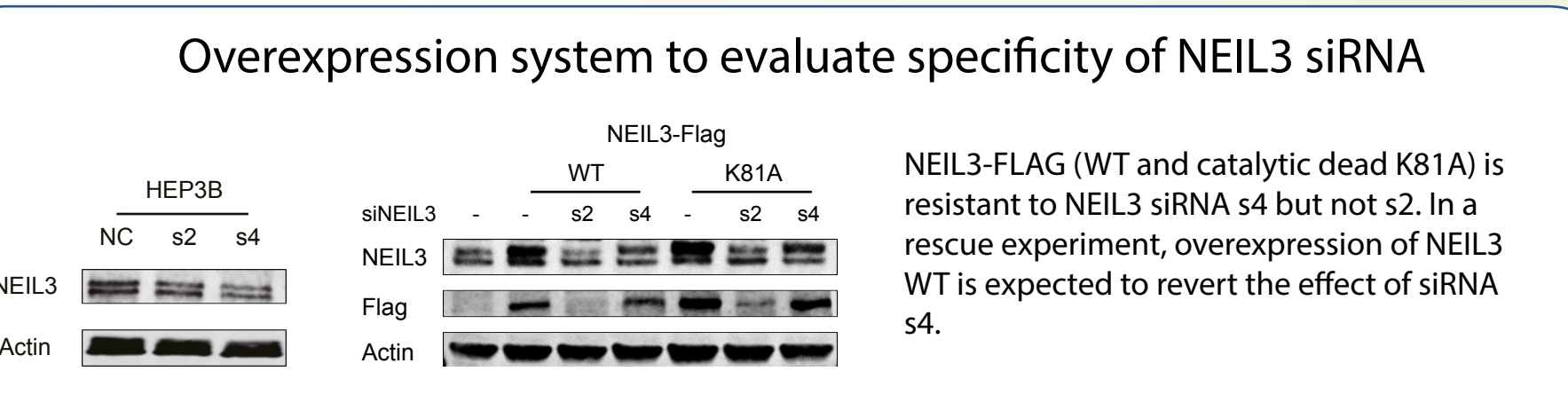
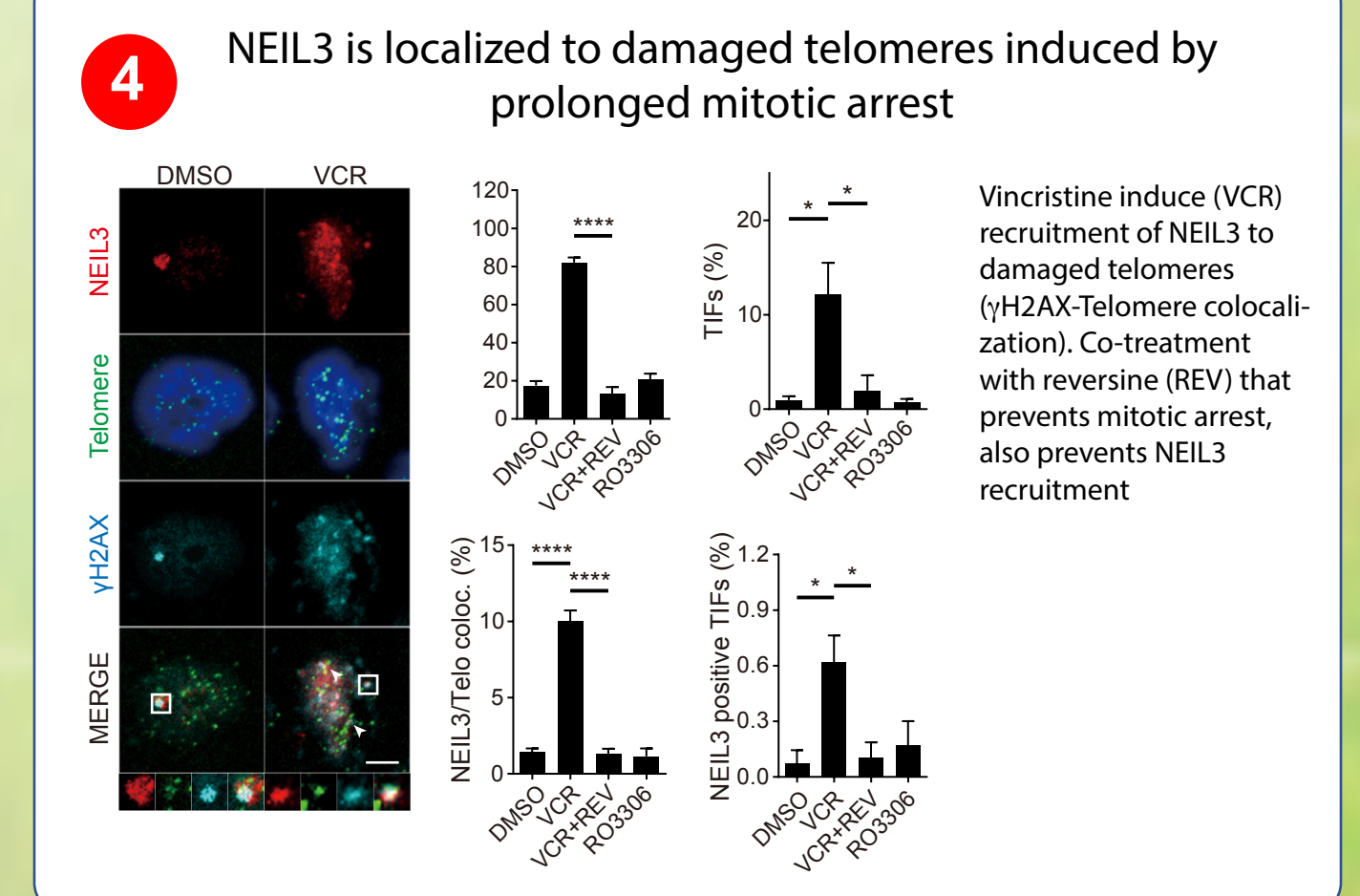
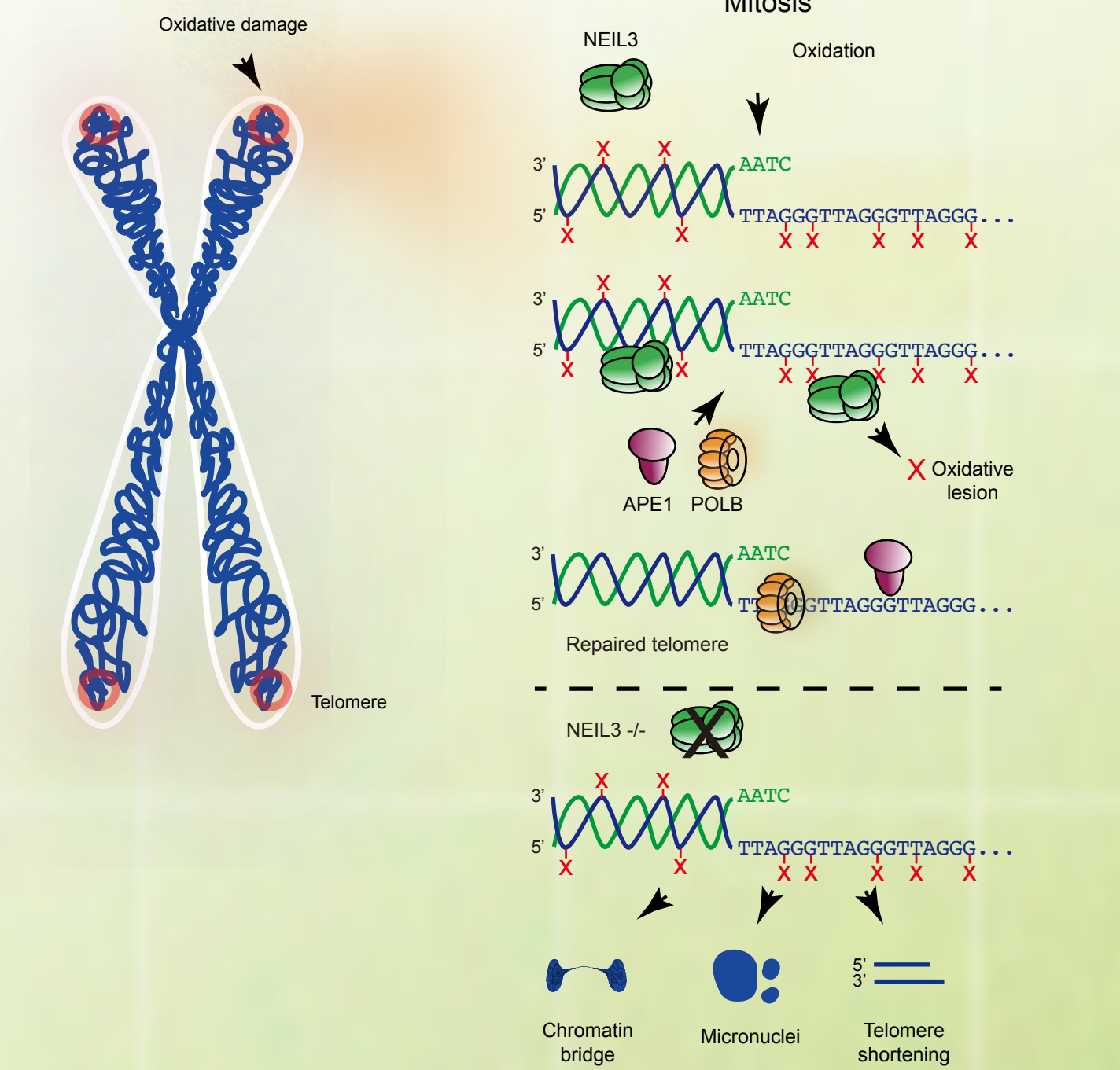
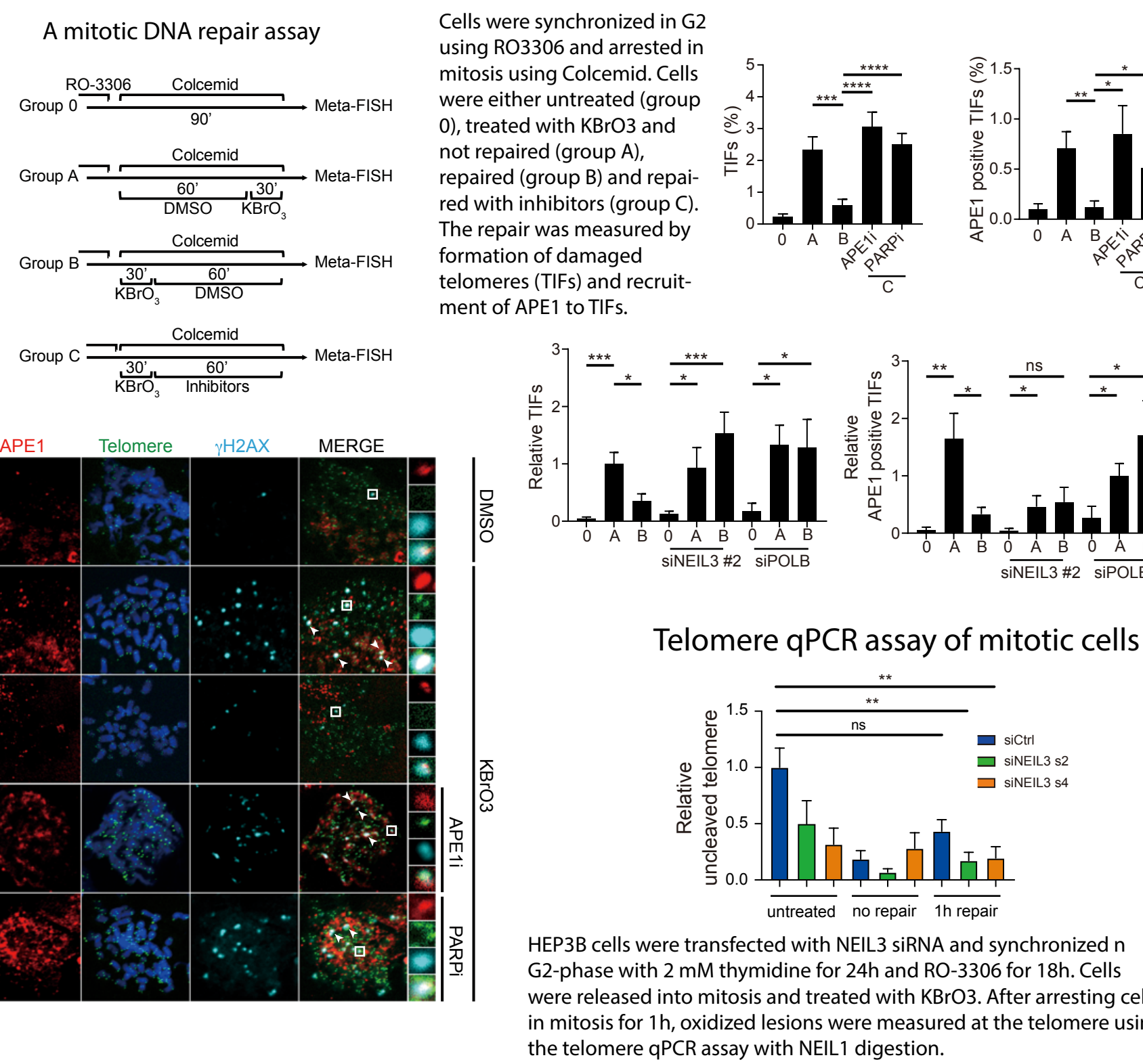
NEIL3 has an unique C-terminus with several protein binding motifs



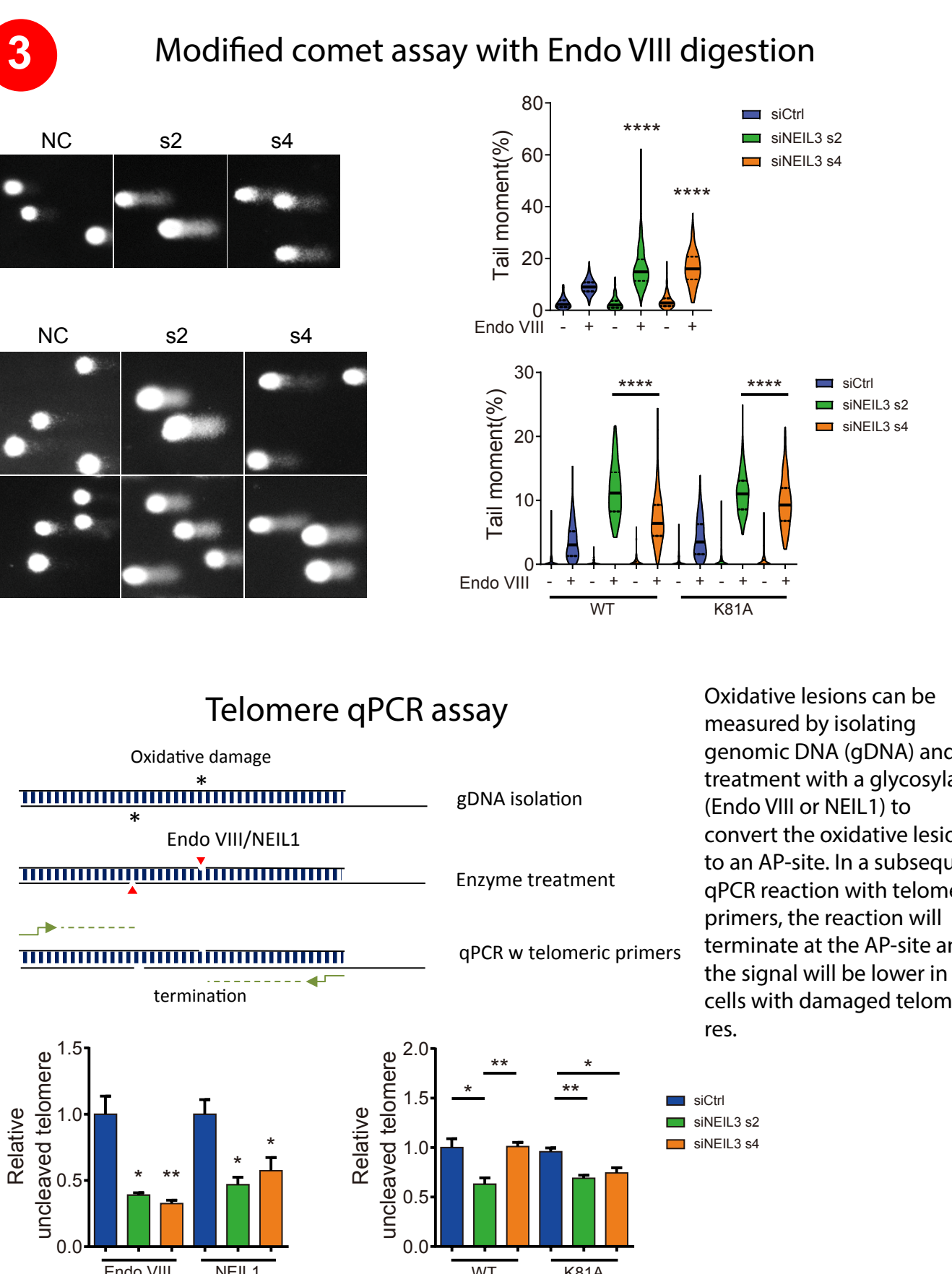
The increase of NEIL3 expression in G2/M phase and the observation that oxidative stress has profound effects in mitosis, suggests that NEIL3 has a crucial role in DNA repair in this phase.



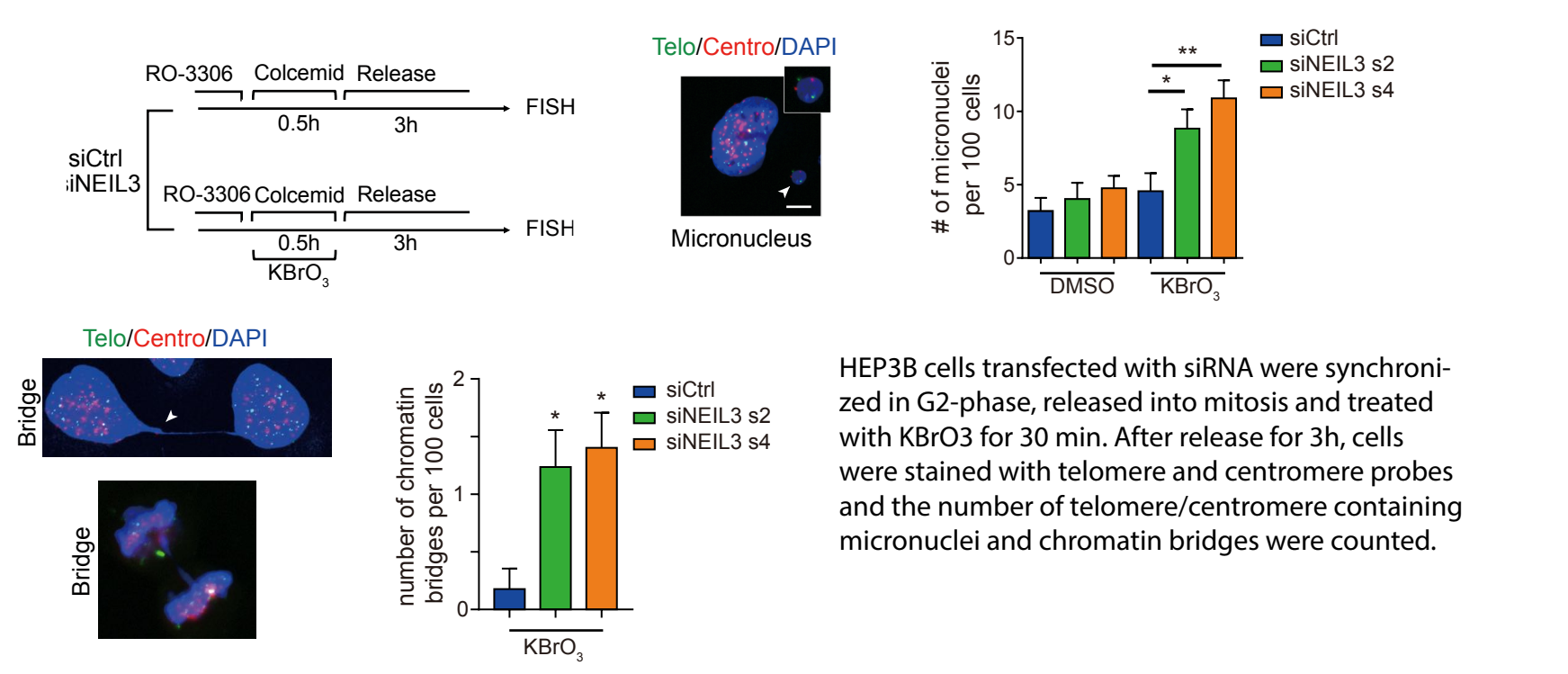
6 NEIL3 recruits APE1 to damaged telomeres and promotes repair of oxidized lesions in mitosis



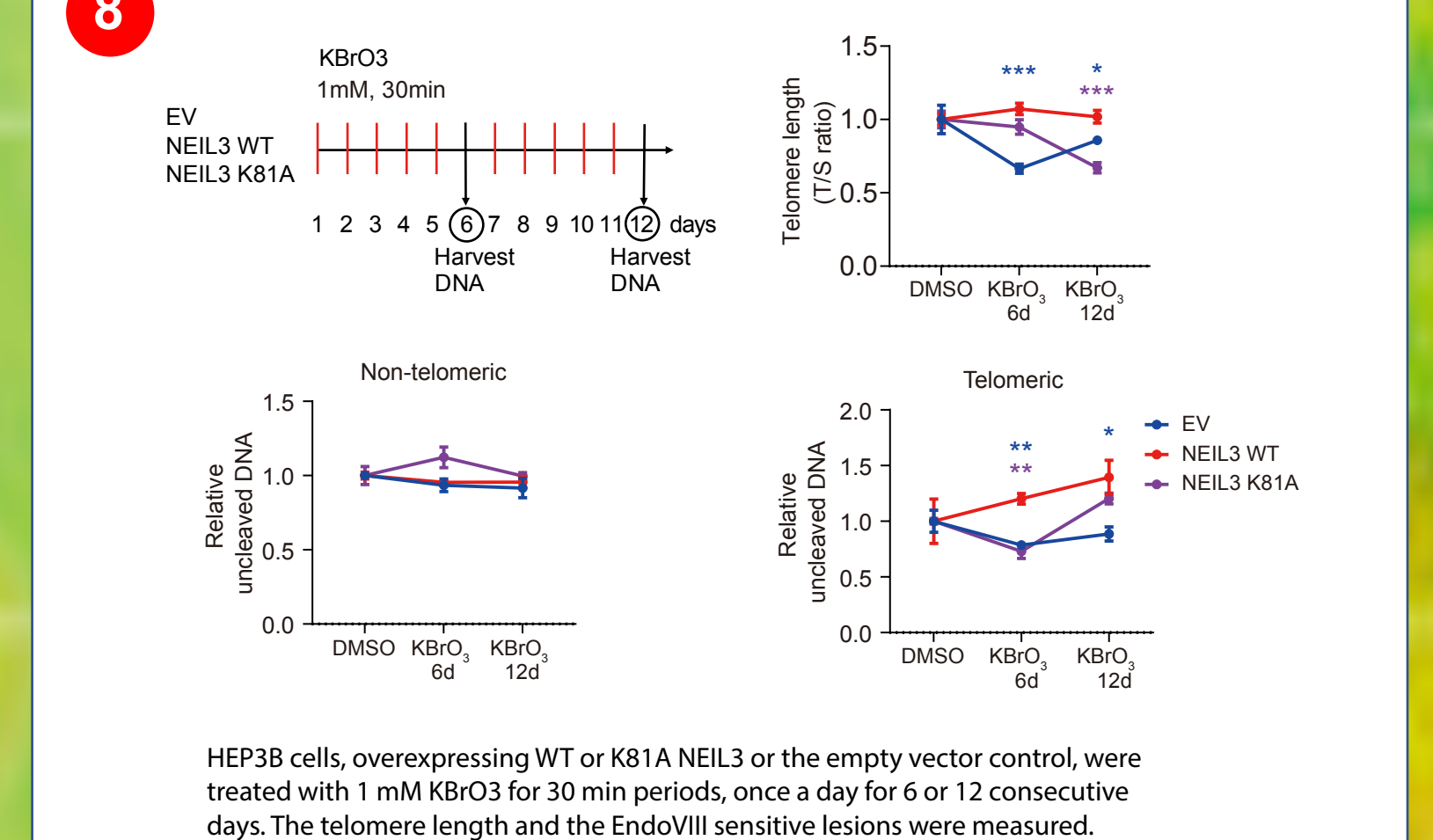
3 NEIL3 silencing induces oxidative damage at telomeres



7 NEIL3 protects HEP3B cells from genomic instability induced by oxidative damage



8 Overexpression of NEIL3 protects cells from long-term effects of oxidative stress



The University of Sheffield.

SciLifeLab



Karolinska Institutet