

- Dykstra, B., Olthof, S., Schreuder, J., Ritsema, M. & de Haan, G. Clonal analysis reveals multiple functional defects of aged murine hematopoietic stem cells. *J. Exp. Med.* 208, 2691–2703 (2011).
- Pang, W. W. et al. Human bone marrow hematopoietic stem cells are increased in frequency and myeloid-biased with age. Proc. Natl Acad. Sci. USA 108, 20012–20017 (2011).
- 23. Ghezraoui, H. et al. Chromosomal translocations in human cells are generated by canonical nonhomologous end-joining. *Mol. Cell* **55**, 829–842 (2014).
- 24. Wyatt, D. W. et al. Essential roles for Polymerase θ-mediated end joining in the repair of chromosome breaks. *Mol. Cell* **63**, 662–673 (2016).
- García-Rubio, M. L. et al. The Fanconi anemia pathway protects genome integrity from R-loops. PLoS Genet. 11, e1005674 (2015).
- Schwab, R. A. et al. The Fanconi anemia pathway maintains genome stability by coordinating replication and transcription. Mol. Cell 60, 351–361 (2015).
- Jaber, S., Toufektchan, E., Lejour, V., Bardot, B. & Toledo, F. p53 downregulates the Fanconi anaemia DNA repair pathway. Nat. Commun. 7, 11091 (2016).
- Oberbeck, N. et al. Maternal aldehyde elimination during pregnancy preserves the fetal genome. Mol. Cell 55, 807–817 (2014).
- Kawashima, N. et al. Aldehyde dehydrogenase-2 polymorphism contributes to the progression of bone marrow failure in children with idiopathic aplastic anaemia. Br. J. Haematol. 168, 460–463 (2015).

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Author Contributions J.I.G., G.P.C. and K.J.P. conceived the study and wrote the manuscript. J.I.G. conducted the majority of the experiments and analysed the data. G.P.C. assisted in the characterization of genomic instability in $Aldh2^{-/-}$ Fancd2 $^{-/-}$ mice and single HSC transplantation. F.L. analysed the survival of chicken DT40 cells, performed western blotting and assisted with the analysis of micronucleus samples. L.M. assisted with single cell transplantation and performed the BigBlue *in vivo* point-mutation analysis. S.L. and F.Y. performed the M-FISH karyotyping of mouse metaphases. N.P. performed validations of indels by targeted deep sequencing. G.G., S.R., S.N.-Z. and M.R.S. provided assistance with the analysis and interpretation of sequencing data.

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