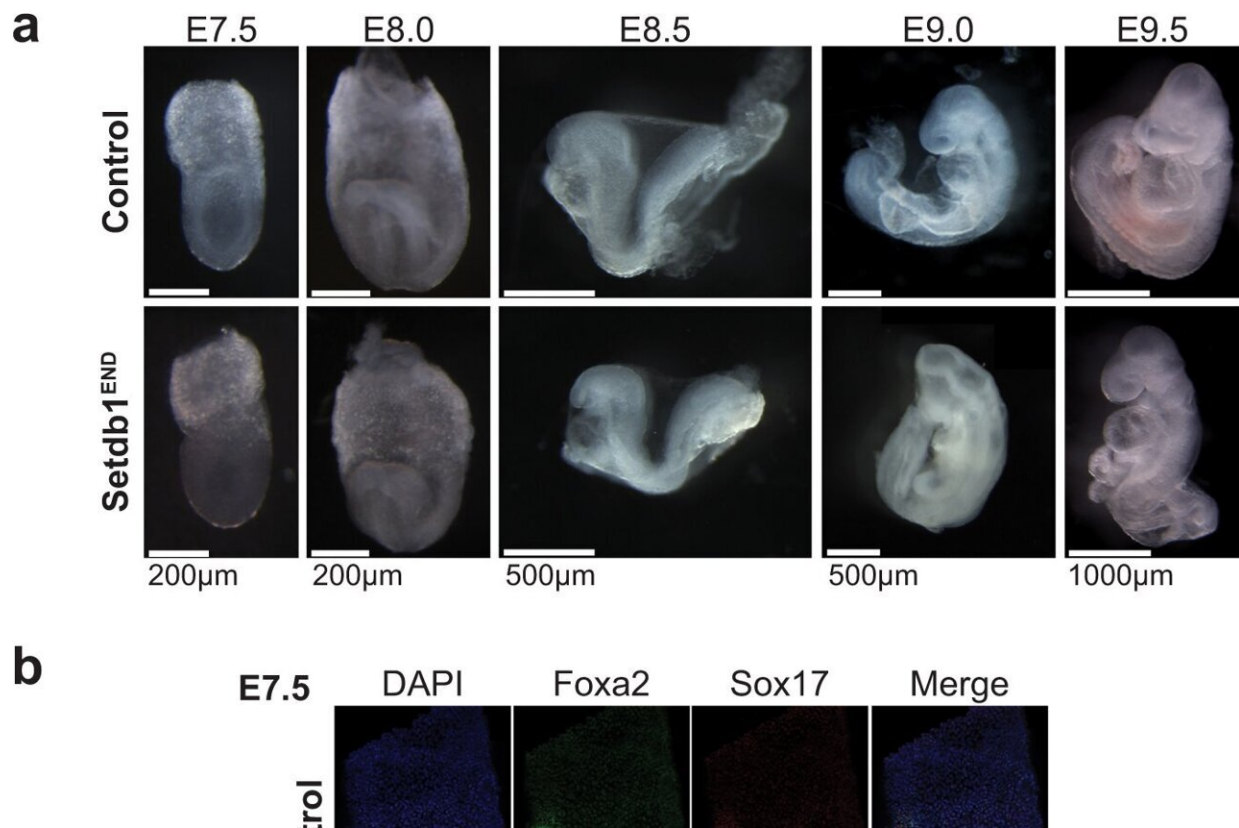


DNA modification important for silencing of endogenous retroviruses

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Loss of Setdb1 in endoderm leads to strong developmental defects during embryogenesis. **a** Lateral view of control and Setdb1^{END} embryos at E7.5 (late bud stage), E7.75 (head fold stage), E8.5 (~6 somite) and E9. No visible developmental defects can be detected in Setdb1^{END} embryos from E7.5 to E8.0. Setdb1^{END} embryos show an axis turning defect which manifests from E8.5 and leads to strong posterior truncation at E9.0. Representative images from n = 3 per genotype and stage. **b** Lateral view of E7.5 control and Setdb1^{END} embryos stained with Foxa2 and Sox17 antibodies (anterior to the left). The presence of

both markers indicates that endoderm cells could be formed in *Setdb1*^{END} embryos. Representative images from n = 3 per genotype and stage. c Hematoxylin/Eosin staining of transverse sections of E8.5 and E9.0 control and *Setdb1*^{END} embryos. The approximate positions of the sections are indicated in the schematic. The black rectangle marks the region used for magnification. Red arrows indicate the hindgut region. The black arrowheads mark the neural tube. Representative images from n = 3 per genotype and stage. Credit: *Nature Communications* (2022). DOI: 10.1038/s41467-022-32978-7

LMU molecular biologist Gunnar Schotta has investigated which mechanisms prevent endogenous retroviruses from becoming overly active.

Endogenous retroviruses (ERVs) have established themselves in the [human genome](#) in the course of evolution and play a major role in normal gene regulation. Excessive ERV activity, however, can lead to diseases such as autoimmunity and cancer. And so cells have developed mechanisms for recognizing and silencing [endogenous retroviruses](#).

The silencing is accomplished by packaging the corresponding DNA sections into a less accessible structure. This process is facilitated by modifications to the histone proteins that package the DNA (H3K9me3) and a modification to the DNA itself (DNA methylation). It has not been clear before now, however, which of these modifications are really important for ERV silencing.

The team—led by Gunnar Schotta from LMU's Biomedical Center Munich—has found that in certain [cell types](#), DNA methylation is decisive and the presence of H3K9me3 is not sufficient to silence ERVs. This discovery furnishes an important basis, for example, for the investigation of aberrantly activated ERVs in the context of diseases such as cancer and autoimmunity.

The study was published in *Nature Communications*.

More information: Zeyang Wang et al, Dominant role of DNA methylation over H3K9me3 for IAP silencing in endoderm, *Nature Communications* (2022). [DOI: 10.1038/s41467-022-32978-7](https://doi.org/10.1038/s41467-022-32978-7)

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