

Epigenetic regulation of transposable elements drives plant speciation

German Martinez¹, Philip Wolff¹, Zhenxing Wang¹, Jordi Moreno-Romero¹, Hua Jiang¹, Juan Santos-González¹, Keith Slotkin², Claudia Köhler¹

¹Department of Plant Biology, Uppsala BioCenter, Swedish University of Agricultural Sciences and Linnean Center for Plant Biology, Uppsala, Sweden

²Department of Molecular Genetics and Center for RNA Biology, The Ohio State University, Columbus, Ohio, USA

Polyloidization is a widespread phenomenon among plants and is considered a major speciation mechanism. Polyloid plants have a high degree of immediate post-zygotic reproductive isolation from their progenitors, as backcrossing to either parent will produce mainly nonviable progeny. This reproductive barrier is called triploid block and it is caused by malfunction of the endosperm. Our work revealed that paternal epigenetically activated small interfering RNAs (easiRNAs) are responsible for the establishment of the triploid block-associated seed abortion in *Arabidopsis thaliana*. Paternal loss of the plant-specific RNA polymerase IV suppressed easiRNA formation and rescued triploid seeds by restoring small RNA-directed DNA methylation at transposable elements, correlating with reduced expression of paternally expressed imprinted genes. Our data suggest that easiRNAs form a quantitative signal for chromosome number and their balanced dosage is required for post-fertilization genome stability and seed viability. Our data reveal a striking analogy of easiRNAs in establishing the triploid block with Piwi-interacting RNAs in hybrid dysgenesis in flies. In both models, TE-derived small RNAs transmit epigenetic information transgenerationally, pointing to a conserved role of TE-derived small RNAs in assessing gamete compatibility.