

Correction: Genome-wide meta-analysis identifies novel loci conferring risk of acne vulgaris

Teder-Laving, Maris; Kals, Mart; Reigo, Anu; **Ehin, Riin**; Objärtel, Telver; Vaht, Mariliis; Nikopensius, Tiit; Metspalu, Andres; Kingo, Külli European Journal of Human Genetics 2024 / art. 1192 <https://doi.org/10.1038/s41431-023-01366-0> [Journal metrics at Scopus](#) [Article at Scopus](#) [Journal metrics at WOS](#) [Article at WOS](#)

Cytological characterization and allelism testing of anther developmental mutants identified in a screen of maize male sterile lines

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Deciphering the omics of plant-microbe interaction: perspectives and new insights

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Editorial : regulatory networks in genome stability pathways

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Engineered microbes for pigment production using waste biomass

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An epigenetic alphabet of crop adaptation to climate change : review article

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Gene expression profiling of puberty-associated genes reveals abundant tissue and sex-specific changes across postnatal development

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Genome-wide meta-analysis identifies novel loci conferring risk of acne vulgaris

Teder-Laving, Maris; Kals, Mart; Reigo, Anu; **Ehin, Riin**; Objärtel, Telver; Vaht, Mariliis; Nikopensius, Tiit; Metspalu, Andres; Kingo, Külli European Journal of Human Genetics 2024 / p. 1136–1143 <https://doi.org/10.1038/s41431-023-01326-8> [Journal metrics at Scopus](#) [Article at Scopus](#) [Journal metrics at WOS](#) [Article at WOS](#)

Insights on engineered microbes in sustainable agriculture : biotechnological developments and future prospects

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Investigating the impact of vitrification on bovine ovarian tissue morphology, follicle survival, and transcriptomic signature

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Legislation of direct-to-consumer genetic testing in Europe : a fragmented regulatory landscape

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Lynch syndrome mutations shared by the Baltic States and Poland

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A meta-analysis of gene expression signatures of blood pressure and hypertension

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Oviduct as a sensor of embryo quality: deciphering the extracellular vesicle (EV)-mediated embryo-maternal dialogue

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Partial deletion of TCF4 in three generation family with nonsyndromic intellectual disability, without features of Pitt-Hopkins syndrome

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Phylogenetic insight into ABCE gene subfamily in plants

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Role of bacterial-fungal consortium for enhancement in the degradation of industrial dyes

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The birth prevalence of spinal muscular atrophy: a population specific approach in Estonia

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A two-cohort RNA-seq study reveals changes in endometrial and blood mirnome in fertile and infertile women

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uCARE Chem Suite and uCAREChemSuiteCLI : tools for bacterial resistome prediction

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