

Apolipoprotein C-II : the re-emergence of a forgotten factor

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Author Correction: Comprehensive analysis of chromothripsis in 2,658 human cancers using whole-genome sequencing (Nature Genetics, (2020), 52, 3, (331-341), 10.1038/s41588-019-0576-7)

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Author Correction: Comprehensive molecular characterization of mitochondrial genomes in human cancers (Nature Genetics, (2020), 52, 3, (342-352), 10.1038/s41588-019-0557-x)

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Comprehensive analysis of chromothripsis in 2,658 human cancers using whole-genome sequencing

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

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Disruption of chromatin folding domains by somatic genomic rearrangements in human cancer

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

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The landscape of viral associations in human cancers

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