

Erratum: More Comprehensive Forensic Genetic Marker Analyses for Accurate Human Remains Identification Using Massively Parallel DNA Sequencing[BMC Genomics. 2016;17 Suppl 9:750.] DOI:10.1186/s12864-016-3087-2

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Massively parallel sequencing-enabled mixture analysis of mitochondrial DNA samples

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Parsing apart the contributors of mitochondrial DNA mixtures with massively parallel sequencing data

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