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Oifferential transcript usage analysis



In 1 collection

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Protocol status: Working We use this protocol and it's working

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ABSTRACT

Differential transcript usage analysis from Kilfeather, Khoo et al., 2024



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Protocol

Differential transcript usage was performed using DRIMSeq (v1.24.0). To consider only genes containing more than 1 isoform, with a minimum level of expression, the following parameters were used:

min_samps_gene_expr = n * 0.75, min_gene_expr = 10, min_samps_feature_expr = 10, min_feature_expr = 10, min_samps_feature_prop = 10, min_feature_prop = 0.1.