

ERRATUM

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Erratum to: 'MicroRNA target prediction using thermodynamic and sequence curves'

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Unfortunately, the original version of this article [1] contained an error. Figures 6 and 7 were incorrect with the wrong captions. The figures have been corrected in the original article and is also included correctly below.

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Reference

1. Ghoshal A, Shankar R, Bagchi S, Grama A, Chaterji S. MicroRNA target prediction using thermodynamic and sequence curves. *BMC Genomics*. 2015;16:999.

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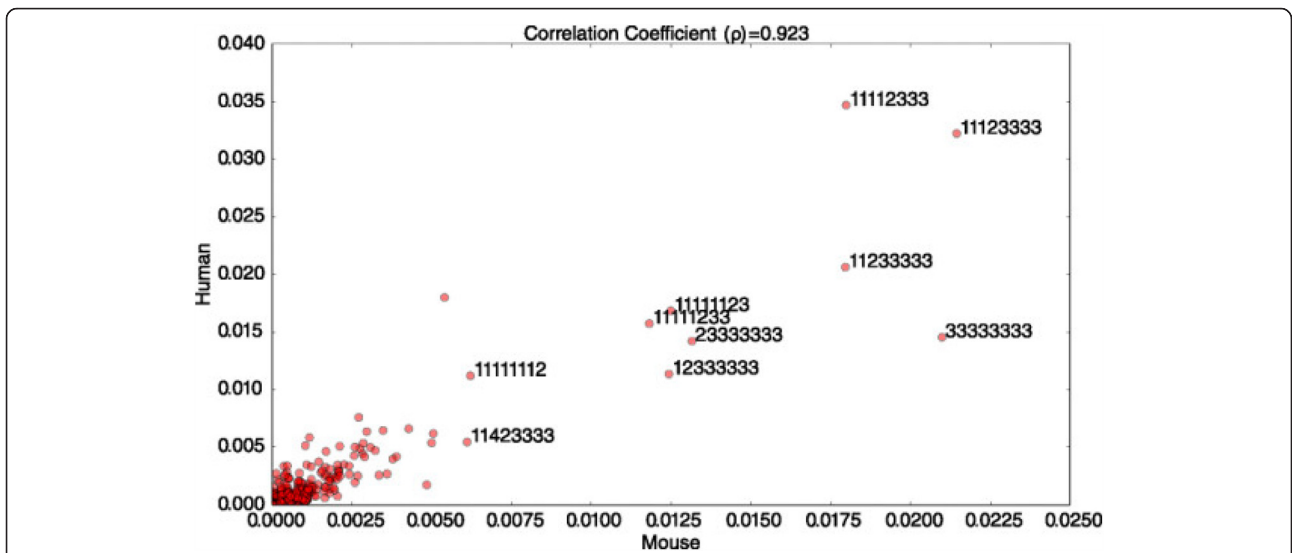


Fig. 6 Scatter plot of frequencies of various types of seed alignment patterns in set of positive miRNA-mRNA interactions for Mouse (x-axis) and Human (y-axis). Among the top-10 most frequently occurring patterns, only two, namely, the 6-mer and 7-mer, are canonical seed match patterns. In the labels for the top-10 most frequently occurring patterns, 1 indicates a match, 2 indicates a mismatch, 3 a gap, and 4 indicates a GU wobble

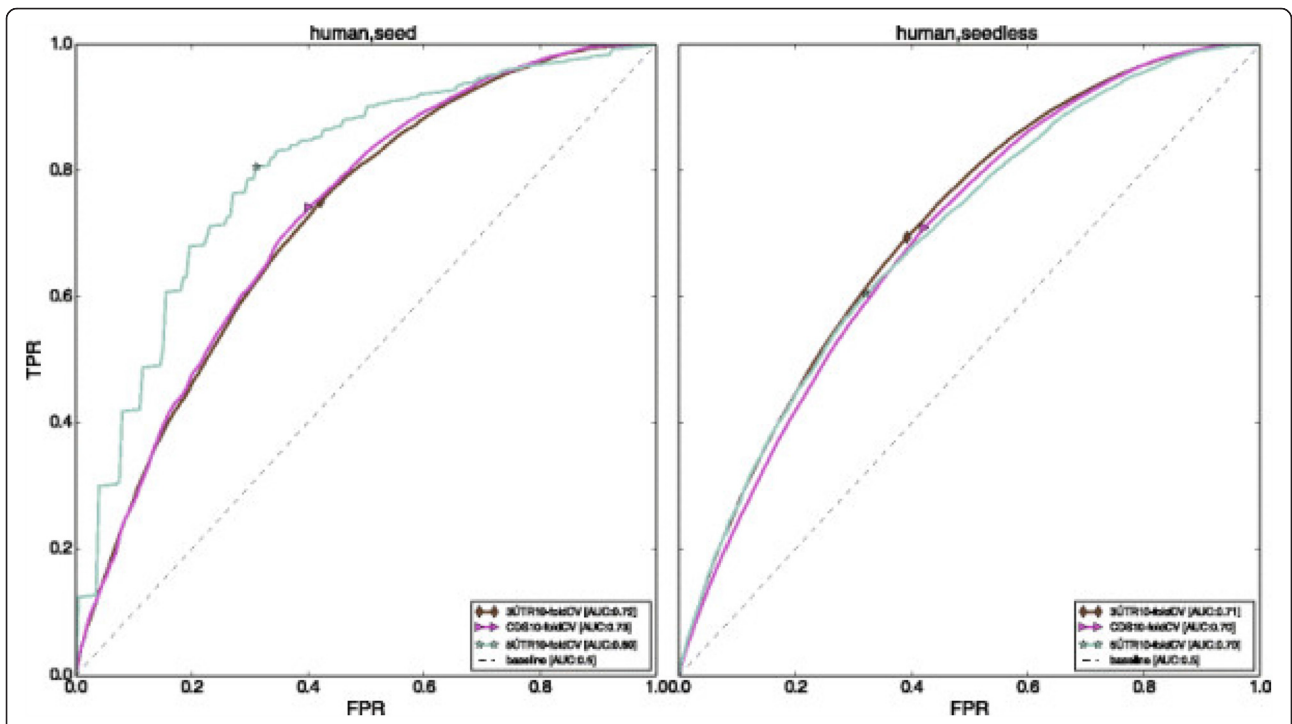


Fig. 7 ROC curves for 10-fold cross-validation performance of *Avishkar* in different regions of the gene in the human dataset