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Erratum

Erratum to “Conservation of the heterochronic regulator Lin-28, its developmental expression and microRNA complementary sites” [Dev. Biol. 258 (2003) 432–442]☆

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Fig. 6 contained incorrect labeling and an incorrect sequence of one of the small RNAs depicted. The corrected Fig. 6 and its legend are presented here.

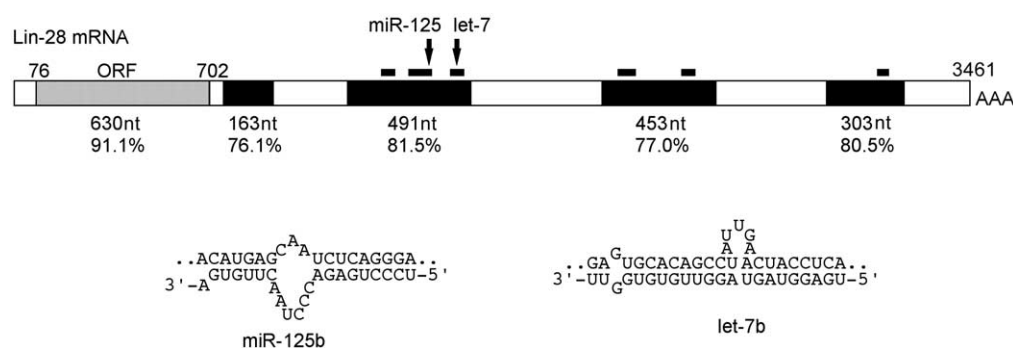


Fig. 6. Sites complementary to microRNAs miR-125 and let-7 in human and mouse Lin-28. Top, a schematic of the Lin-28 mRNA. The ORF is in gray. The numbers above the line indicate the positions of the start codon, stop codon, and poly(A) tail, respectively. The black boxes indicate regions of high sequence identity between mouse and human 3' UTRs; the lengths and percent identity of each of these regions are indicated below the schematic. The black boxes above the schematic indicate 3' UTR sequences of at least 40 nt having greater than 95% identity between mouse and human. The positions of the potential miR-125 and let-7 complementary sites are indicated by arrows. Bottom, possible secondary structures formed between the UTR and microRNAs.

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