CORRIGENDUM

Human sat III and Drosophila hsr omega transcripts: a common paradigm for regulation of nuclear RNA processing in stressed cells

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The authors would like to apologize for the following error:
Table 1 in the article lists ‘SWI/Snf’ as one of the proteins that is associated with hsr-omega transcripts. The ‘SWI/Snf’ in Table 1 should actually read ‘Snf (Drosophila splicing necessary factor)’, which is not same as the SWI/Snf protein.

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