

# miRNA<sub>tap</sub>.db

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miRNA<sub>tap</sub>.db

*database for miRNA<sub>tap</sub> package*

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## **Description**

This package contains data annotations (target predictions).

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## **Examples**

```
#direct targets in mouse aggregated from all sources:  
library(miRNAtap)  
targets_mouse <- getPredictedTargets('let-7a', species='mmu', method='geom')
```

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