

# miRNA<sup>Atap</sup>.db

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miRNA<sup>Atap</sup>.db

*database for miRNA<sup>Atap</sup> 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(miRNAAtap)
targets_mouse <- getPredictedTargets('let-7a',species='mmu', method='geom')
```

# Index

miRNAAtap.db, [1](#)  
miRNAAtap.db-package (miRNAAtap.db), [1](#)