**PatMaN**

RRID:SCR_011821  
Type: Tool

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**Proper Citation**

PatMaN (RRID:SCR_011821)

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**Resource Information**

**URL:** [https://bioinf.eva.mpg.de/patman/](https://bioinf.eva.mpg.de/patman/)

**Description:** Software that searches for short patterns in large DNA databases, allowing for approximate matches.

**Resource Name:** PatMaN

**Proper Citation:** PatMaN (RRID:SCR_011821)

**Resource Type:** Resource, software resource

**Keywords:** c++

**Resource ID:** SCR_011821

**References:** PMID: 18467344

**Availability:** GNU General Public License, v3

**Website Status:** Last checked up

**Alternate IDs:** OMICS_00997

**Abbreviations:** PatMaN

**Mentions Count:** 40

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Ratings and Alerts
No rating or validation information has been found for PatMaN.

No alerts have been found for PatMaN.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics

We found 40 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch Infrastructure.


Taller D, et al. (2018) Expansion of Capsicum annuum fruit is linked to dynamic tissue-
specific differential expression of miRNA and siRNA profiles. PloS one, 13(7), e0200207.


Cognat V, et al. (2017) The nuclear and organellar tRNA-derived RNA fragment population in Arabidopsis thaliana is highly dynamic. Nucleic acids research, 45(6), 3460-3472.
