



# 5 years of community annotation using Wikipedia

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**Rfam** is a database of non-coding RNA families, each represented by a multiple sequence alignment and a covariance model (CM)<sup>1</sup>. Our multiple sequence alignments and secondary structure annotations are generated manually by specialist biocurators. However, since June 2007 (Release 8.1) Rfam has been successfully coordinating community annotation via the online encyclopedia Wikipedia to provide textual annotation for these families<sup>2</sup>.

The last Rfam release 10.1 (June 2011) contains 1,973 families which are linked to around 900 Wikipedia articles. New edits to these articles are downloaded on a daily basis and displayed on our website along side our specialist information.

## Each Rfam family has a link to one Wikipedia article

Wikipedia article is chosen by curator at time of family build



If no suitable article is available, a curator may:

- Create a new article
- Link to a generic RNA article, e.g. miRNA, ncRNA, snoRNA
- Each article has a link back to Rfam

This article mapping is maintained by Rfam

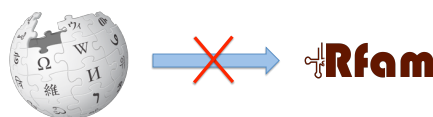


## Wikipedia article is mirrored on the Rfam website alongside our specialist information

Rfam specific data tabs

Rfam infobox

## Wikipedia content on rfam.sanger.ac.uk is not live



Rfam articles requiring approval

Unapproved articles

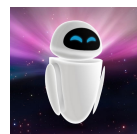
Title	Last editor	Last edit date	Current category	Approved	Reverted	Reverted
let-7 family	let-7 family	2011-06-17	let-7 family	1	0	0
let-7 family	let-7 family	2011-06-17	let-7 family	1	0	0
let-7 family	let-7 family	2011-06-17	let-7 family	1	0	0

Pre-approved articles

Title	Last editor	Last edit date	Current category	Approved	Reverted	Reverted
let-7 family	let-7 family	2011-06-17	let-7 family	1	0	0
let-7 family	let-7 family	2011-06-17	let-7 family	1	0	0
let-7 family	let-7 family	2011-06-17	let-7 family	1	0	0

The Rfam wiki approval app allows us to approve content before displaying on our website.

## The Rfambot



Wikipedia allows for automation using bots

Rfambot automates Rfam-related routine tasks e.g. Rfam infobox, release & image updates

Many non-Rfam specific bots edit & improve our mapped articles

## Rfam specific information is not open to Wikipedia editing



Text annotation

Rfam infobox



EVERYTHING ELSE

e.g. Sequence alignments, Covariance models, Secondary structures, Types, Names, Article mapping

## Wikipedia page edits and page views

Since June 2007

16,607 edits have been made by 2,213 editors

~351 edits were reverted – mainly by external editors

~600 articles have been edited > 10 times

15 articles have received > 120 edits

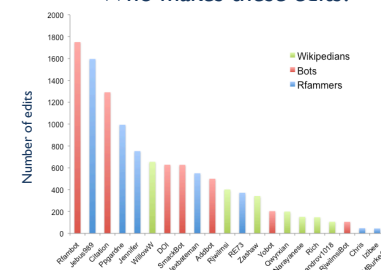
February 2012

Each article was viewed > 10

tRNA, microRNA, Ribosomal RNA and Ribozyme articles were each viewed > 1000 times

15% of Rfam web access came via Wikipedia

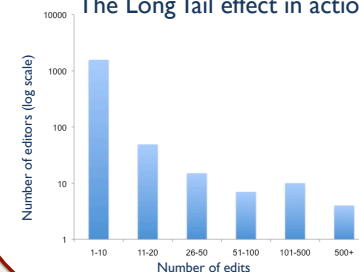
## Who makes these edits?



Editors ranked by number of edits made since June 2007. Only the top editors are shown

Edits to our articles of interest are made by many people and bots not associated with Rfam

## The Long Tail effect in action



Distribution of the number of edits each editor makes

Most editors make only a few edits but we have lots of editors

## Why Wikipedia works for Rfam

- Stable editing infrastructure
- Huge user community – scientific and non-scientific editors – excellent quality and vandalism policing
- The “Google factor”
- “Chores” can be automated by bots
- Rfam specialist data is untouched
- Editor contributions exceed Rfam manpower!

References:

- 1 Rfam: Wikipedia, clans and release Gardner PP, Daub J, Tate J, Moore BL, Osuch IH, Griffiths-Jones S, Finn RD, Nawrocki EP, Kolbe DL, Eddy SR, Bateman A. Nucleic Acids (2011) doi: 10.1093/nar/gkq1129
- 2 The RNA WikiProject: community annotation of RNA families. Daub J, Gardner PP, Tate J, Ramsköld D, Manske M, Scott WG, Weinberg Z, Griffiths-Jones S, Bateman A. RNA (2008) 12:2462-2464

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