

# Guide to the ngram Package

An n-gram Babbler

## Guide to the **ngram** Package

AN N-GRAM BABBLER

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#### 1 Introduction

An n-gram is an ordered sequence of n "words" taken from a body of text. For example, consider the string A B A C A B B. This is the "blood code" for the video game Mortal Kombat for the Sega Genesis, but you can pretend it's a biological sequence or something boring if you prefer. If we examine the 2-grams (or bigrams) of this sequence, they are:

```
A B, B A, A C, C A, A B, B B
```

or without repetition:

```
AB, BA, AC, CA, BB
```

That is, we take the input string and group the "words" 2 at a time (because n=2). If we form all of the n-grams and record the next "words" for each n-gram (and their frequency), then we can generate new text which has the same statistical properties as the input.

The **ngram** package is an R package for constructing n-grams and generating new text as described above. It also contains a few preprocessing utilities to aid in this process. Additionally, the C code underlying this library can be compiled as a standalone shared library.

#### 2 Installation

In this section, we will describe the various ways that one can install the **ngram** package.

#### 2.1 Installing from Source

The sourcecode for this package is available (and actively maintained) on GitHub. To install an R package from source on Windows, you will need to first install the Rtools package. To install an R package from source on a Mac, you will need to install the latest Xcode, which you can get from the App store.

The easiest way to install **ngram** from GitHub is via the **devtools** package by Hadley Wickham. To install **ngram** using **devtools**, simply issue the command:

```
library(devtools)
install_github(repo="ngram", username="wrathematics")
```

from R. Alternatively, you could download the sourcecode from github, unzip this archive, and issue the command:

```
R CMD INSTALL ngram-master
```

from your shell.

#### 2.2 Installing from CRAN

The usual

```
install.packages("ngram")
```

from an R session should do it.

#### 3 Using the Package

#### 3.1 Background

The input to the n-gram processor must be a single string (character vector of length 1). To aid in what could be a repetitive task, the package offers the concat() function. For example:

So if data is coming from multiple files, the simplest way to merge them together would be to call

```
1 x <- readLines("file1")
2 y <- readLines("file2")
3
4 str <- concat(x, y)</pre>
```

The ngram() function (which does the processing and forms the n-grams) always splits words at a space. You can preprocess the string with R's regular expression utilities, such as gsub(), or use the preprocess() utility in the ngram package to be able to split at non-spaces for the purpose of n-gram generation (by inserting your own beforehand).

#### 3.2 Package Use and Example

The general process goes

1. Prepare the input string; you may find concat() and preprocess() useful (see the previous subsection).

- 2. Process with ngram().
- 3. Generate new text with babble(), and/or
- 3.5 Extract pieces of the processed ngram data with the get.\*() functions.

Let us return to the example sequence of letters from Section 1. If we store this string in x:

```
1 x <- "A B A C A B B"
```

then the next step is to process with ngram():

```
1 library(ngram)
2 ng <- ngram(x, n=2)</pre>
```

We can then inspect the sequence:

```
1 > ng
2 [1] "An ngram object with 5 2-grams"
```

If you don't have too many n-grams, you may want to print all of them by calling print() directly, with option full=TRUE:

```
> print(ng, full=TRUE)
  C A
  B {1} |
  B A
  C {1} |
  ВВ
  NULL {1} |
10
  A C
11
  A {1} |
12
13
  A B
14
  A {1} | B {1} |
```

Here we see each 3-gram, followed by its next possible "words" and each word's frequency of occurrence (occurrence following the given n-gram). So in the above, the first n-gram printed C A has B as a next possible word, because the sequence C A is only ever followed by the "word" B in the input string. On the other hand, A B is followed by A once and B once. The sequence B B is terminal, i.e. followed by nothing; we treat this case specially.

Next, we might want to generate some new strings. We for this, we use babble():

```
babble(ng, 10)
| [1] "A C A B B B A C A B "
```

```
3 > babble(ng, 10)
4 [1] "B B C A B A C A B A "
5 babble(ng, 10)
6 [1] "A C A B A C A B A C "
```

This generation includes a random process. For this, we developed our own implementation of MT19937, and so R's seed management does not apply. To specify your own seed, use the **seed**= argument:

#### 3.3 Important Notes About the Internal Representation

The entirety of the interesting bits of the **ngram** package take place outside of R (completely in C). Observe:

```
> str(ng)
Formal class 'ngram' [package "ngram"] with 6 slots
..@ str_ptr:<externalptr>
..@ strlen : int 13
..@ n : int 2
..@ ng_ptr :<externalptr>
..@ ngsize : int 5
..@ wl_ptr :<externalptr>
```

So everything is wrangled up top as an S4 class, and underneath the data is stored as 2 linked lists, outside the purview of R. This means that, for example, that you cannot save the n-gram object with a call to save(). If you do and you shut down and restart R, the pointers will no longer be valid.

Extracting a the data into a native R data structure is not currently possible. Full support is planned for a later release. Some pieces can be extracted. At this time, get.ngrams() and get.string() are implemented, but get.nextwords() is not.