

Using samtools C libraries with *Rsamtools*

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Modified: 20 March, 2015. Compiled: October 1, 2015

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This document is meant for package developers wanting to use the [samtools](#) C libraries provided by *Rsamtools*. The instructions here are based on the 'Writing R Extensions' manual associated with R-2.13; consult the current version of the manual for up-to-date instructions.

1 Background

Rsamtools arranges to install static versions of the `libbam` and `libbcf` libraries. The idea is that other packages can use these to implement C functionality that uses these libraries. This means that the samtools libraries are available in a consistent version to users across platforms, without requiring installation of additional software.

Rsamtools takes the following approach. On installation, *Rsamtools* contains a snapshot of the samtools library source code under `src/samtools`. *Rsamtools* makes static version(s) of the samtools libraries `libbam.a` and `libbcf.a`. These static libraries are specific to the operating system on which *Rsamtools* is being installed, are found under `usrlib${R_ARCH}` in the user library location specified by the mechanism (e.g., `biocLite`, `install.packages`) used to install *Rsamtools*. At the same time, *Rsamtools* copies headers required to use the library to the location `include/samtools`.

2 Use

To use these libraries, the third party package developer needs to (1) discover the appropriate header files when their package is built, and (2) link in the libraries.

Note that in order to link correctly across platforms your package must provide *both* the respective `src/Makevars` and `src/Makevars.win` files.

2.1 Discover header files

To discover appropriate header files at package installation time, add

```
LinkingTo: Rsamtools
```

to the DESCRIPTION file, and reference the relevant include files as, for instance,

```
#include "samtools/bam.h"
```

The content of the include files can be found in the *Rsamtools* source (under `src/samtools`) or at their installed location.

2.2 Link to static libraries

Linking to the static libraries is accomplished by including the following code in `src/Makevars.win` for *Windows*:

```
SAMVARS=$(shell echo 'cat(system.file("usretc", .Platform[["r_arch"]], \
  "Rsamtools.mk", package="Rsamtools", mustWork=TRUE))' | \
  "${R_HOME}/bin/R" --vanilla --slave)
include $(SAMVARS)
```

```
PKG_LIBS=$(SAMTOOLS_LIBS)
PKG_CPPFLAGS=$(SAMTOOLS_CPPFLAGS)
```

and with the following code in `src/Makevars` for all other platforms:

```
SAMTOOLS_PATH=\
  `echo 'cat(system.file("usrlib", .Platform[["r_arch"]], \
    package="Rsamtools", mustWork=TRUE))' | \
    "${R_HOME}/bin/R" --vanilla --slave`
SAMTOOLS_LIBS="$(SAMTOOLS_PATH)/libbam.a" "$(SAMTOOLS_PATH)/libbcf.a" \
  "$(SAMTOOLS_PATH)/libtabix.a" -lz -pthread
SAMTOOLS_CPPFLAGS=-D_USE_KNETFILE -DBGZF_CACHE -D_FILE_OFFSET_BITS=64 \
  -D_LARGEFILE64_SOURCE

PKG_LIBS=$(SAMTOOLS_LIBS)
PKG_CPPFLAGS=$(SAMTOOLS_CPPFLAGS)
```

This updates the environment variables `$PKG_CPPFLAGS` and `$PKG_LIBS`; if your `Makevars/Makevars.win` modifies these also, do so by adding to the respective line, e.g.,

```
PKG_LIBS=$(SAMTOOLS_LIBS) -lfoo
PKG_CPPFLAGS=$(SAMTOOLS_CPPFLAGS) -I/path/bar
```