Using samtools C libraries with *Rsamtools*

Martin Morgan

Modified: 21 March, 2011. Compiled: July 28, 2011

This document is meant for package developers wanting to usethe 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) locate the static library when their package is used.

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.

Linking to the static libraries is accomplished by including the following code in `src/Makevars`:

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

This updates the environment variables `$PKG_CPPFLAGS` and `$PKG_LIBS`; if your `Makefile` modies these also, be sure to use syntax like `PKG_LIBS+=-l...`.