

## Getting SEER, A-bomb Survivor, and Human Mortality Database Data

**SEERaBomb** uses ASCII text SEER data available from <https://seer.cancer.gov/data/treatment.html>, which includes chemo and radiation information. Please unzip this data into `~/data/SEER` such that the folder `populations` holds person-year-at-risk files and the folder `incidence` holds cancer files. The A-bomb survivor data files `lsshempy.csv` and `lssinc07.csv` obtained from the Radiation Effects Research Foundation (RERF) by following the links “*The incidence of leukemia, lymphoma and multiple myeloma among atomic bomb survivors: 1950-2001*” and “*Life Span Study Cancer Incidence Data, 1958-1998*” at [http://www.rerf.jp/library/dl\\_e/index.html](http://www.rerf.jp/library/dl_e/index.html) should similarly be placed into `~/data/abomb`. For mortality relative risk time course calculations, it is also necessary to obtain a human mortality database username and password from <http://www.mortality.org/>. You should then run this:

```
install.packages("demography") #if not already installed
library(demography)
mrt=hmd.mx("USA", "username", "password")$rate # sub in your info
if(!dir.exists("~/data/usMort")) dir.create("~/data/usMort")
save(mrt, file="~/data/usMort/mrt.RData")
```

at the R prompt. Use of these file locations will simplify running scripts.