

```
###The following packages need to be installed to convert Bruker Flex
format files to mzml using R
##
install.packages("MALDIquant")
install.packages("MALDIquantForeign")
##
###load the following packages
##
library("MALDIquant")
library("MALDIquantForeign")
##
###import your Bruker Flex files
##
spectra<-importBrukerFlex(path="###Unconverted file location here###")
##
###export to mzml format
##
exportMzMl(spectra, path="###Location for exported data###", force=FALSE)
```