

# Building a PDInfo Package for an Affymetrix Mapping Chip

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## 1 IMPORTANT!

Users are asked to download the packages for Affymetrix SNP chips from the BioConductor website!

The contents of this vignette explain the essence of the builder for SNP chips.

## 2 What you will need

Aside from the `pdInfoBuilder` package and its dependencies, you will need the following files in order to follow along and build a “PDInfo” package for the Affymetrix Mapping 250K NSP chip.

- `Mapping250K_Nsp.cdf`: The binary CDF file from Affymetrix containing feature-level annotation.
- `Mapping250K_Nsp_annot.csv`: Delimited text file from Affymetrix containing featureSet-level annotation.
- `Mapping250K_Nsp_probe_tab`: Delimited text file from Affymetrix containing probe sequence information for all PM probes.

## 3 Building a PDInfo package

Assuming the source files described above are in your current working directory, the following commands will produce a PDInfo package.

```
R> library("pdInfoBuilder")
R> cdfFile <- "Mapping250K_Nsp.cdf"
R> csvAnno <- "Mapping250K_Nsp_annot.csv"
R> csvSeq <- "Mapping250K_Nsp_probe_tab"
R> pkg <- new("AffySNPPDInfoPkgSeed",
             version="0.1.5",
             author="Seth Falcon", email="sfalcon@fhcrc.org",
```

```
      biocViews="AnnotationData",
      genomebuild="NCBI Build 35, May 2004",
      cdfFile=cdfFile, csvAnnoFile=csvAnno, csvSeqFile=csvSeq)
R> makePdInfoPackage(pkg, destDir=".")
R>
```

We are able to complete the above step on a dual-CPU dual-core 3GHz Xeon box with 8GB RAM in about 20 minutes. The only step that requires a significant amount of RAM is generating the coded sequence matrix. I haven't yet looked carefully at it, but I think it should run on a 4GB system.