

## **SelectSNPProbes Documentation**

**Description:** Filter a file to retain only SNP probes.

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### **Summary**

#### **Parameters**

Name	Description
input file (required)	The input file. It can be a file in CN, XCN, SNP, or BSNP format.
output prefix (required)	Prefix for the output file name.

## **Output Files**

<output.prefix>.snpsonly.<input.file\_extension>
 A file filtered to only contain data for the SNP probes.

# **Platform Dependencies**

Module type: SNP Analysis

CPU type: any
OS: any
Language: Perl