

# Summary of new features of genomation

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<code>library(devtools)</code>	1
<code>install_github("BIMSBbioinfo/genomation",build__vignettes=FALSE)</code>	1
<code>library(genomation)</code>	1

`library(devtools)`

`install_github("BIMSBbioinfo/genomation",build__vignettes=FALSE)`

`library(genomation)`

`library(devtools) load_all("../genomation") library(GenomicRanges)`

