

# Examples to Genetic Analysis Package (gap)

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## Contents

This package is a spin-off of the package `gap`, whereas this vignette is a skeleton to be expanded in the near future.

After installation, you will be able to issue `library(help=gap.examples)` or to view the list within a web browser via `help.start()`. To load the package, issue `library(gap.examples)`, and you can use `lsf.str("package:gap.examples")` and `data(package="gap.examples")` to generate a list of datasets.

A PDF version of this file can be viewed with command

```
vignette("gap.examples", package="gap.examples").
```

Several other vignettes have also been moved here.