

# test-ggplot.r

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*Thu Sep 15 01:11:47 2016*

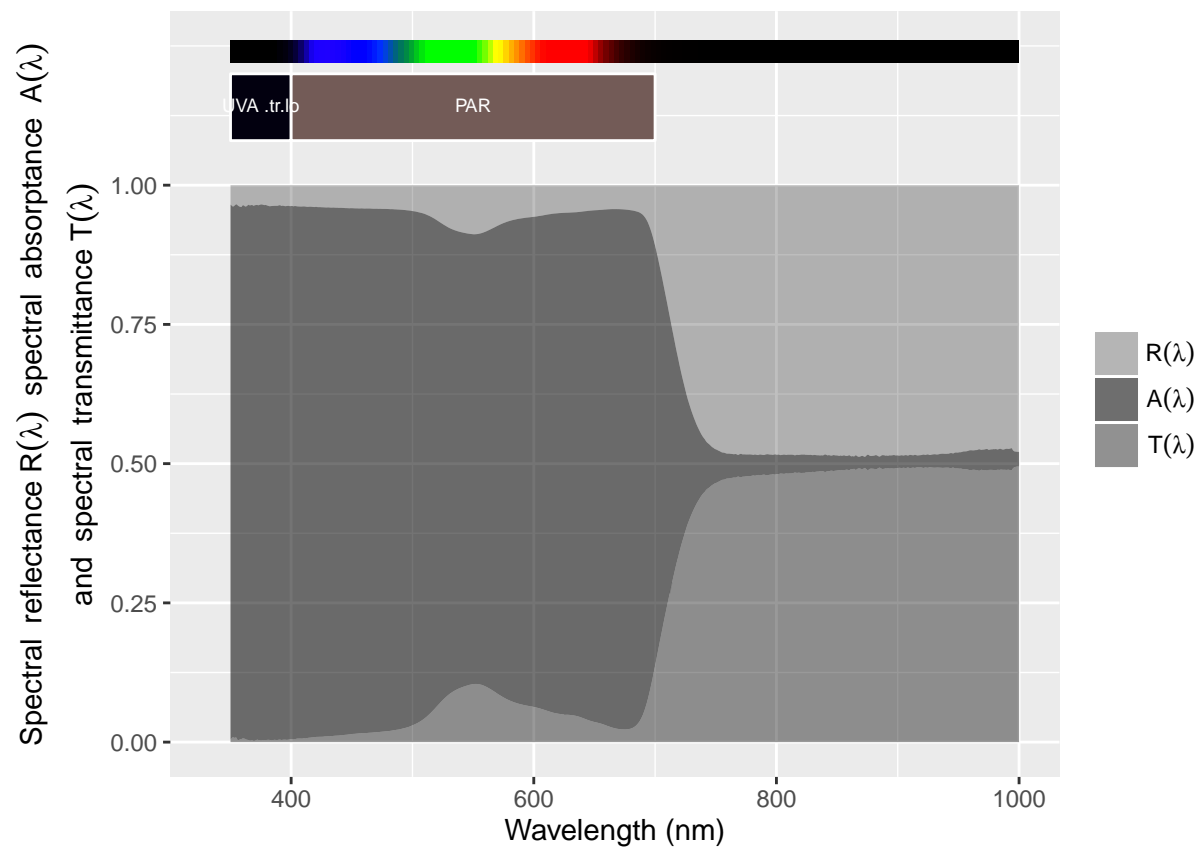
```
library(photobiologyPlants)
```

```
## Loading required package: photobiology
```

```
## Loading required package: photobiologyWavebands
```

```
library(ggspectra)
```

```
plot(Betula_ermanii.mspct$first_flush_adax)
```



```
sessionInfo()
```

```
## R version 3.3.1 (2016-06-21)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 14393)
##
## locale:
## [1] LC_COLLATE=English_United Kingdom.1252
## [2] LC_CTYPE=English_United Kingdom.1252
## [3] LC_MONETARY=English_United Kingdom.1252
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United Kingdom.1252
```

```
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] ggspectra_0.1.9.9000      photobiologyPlants_0.4.0.9001
## [3] photobiologyWavebands_0.4.1 photobiology_0.9.11
##
## loaded via a namespace (and not attached):
## [1] Rcpp_0.12.7      knitr_1.14      magrittr_1.5
## [4] munsell_0.4.3    colorspace_1.2-6 R6_2.1.3
## [7] stringr_1.1.0    plyr_1.8.4      dplyr_0.5.0
## [10] tools_3.3.1      grid_3.3.1      gtable_0.2.0
## [13] DBI_0.5-1        htmltools_0.3.5 lazyeval_0.2.0
## [16] assertthat_0.1   digest_0.6.10   tibble_1.2
## [19] ggplot2_2.1.0    tidyr_0.6.0     evaluate_0.9
## [22] rmarkdown_1.0.9013 labeling_0.3     stringi_1.1.1
## [25] scales_0.4.0
```