Soil Database Interface

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```
suppressMessages(library(soilDB))
data(loafercreek)
## aggregate major horizon types over 1cm intervals
# categorize major horizon types
hz.tab <- rev(sort(table(loafercreek$hzname)))</pre>
hz.tab[hz.tab > 5]
##
##
   A Bt1 Bt2 Cr Bt3 Oi R Crt
## 47 45 44 25 22 21 15 13
# add generalized hz name
loafercreek$hz <- rep("other", times = nrow(loafercreek))</pre>
# generalize horizons
loafercreek$hz[grep("0", loafercreek@horizons$hzname)] <- "0"</pre>
loafercreek$hz[grep("A", loafercreek@horizons$hzname)] <- "A"</pre>
loafercreek$hz[grep("Bt", loafercreek@horizons$hzname)] <- "Bt"</pre>
loafercreek$hz[grep("Cr", loafercreek@horizons$hzname)] <- "Cr"</pre>
loafercreek$hz[grep("R", loafercreek@horizons$hzname)] <- "R"</pre>
# convert generalized hz to factor
loafercreek$hz <- factor(loafercreek$hz)</pre>
loafercreek.hz.agg <- slab(loafercreek, fm = ~hz)</pre>
# wide->long format
loafercreek.hz.agg.long <- melt(loafercreek.hz.agg,</pre>
    id.var = c("top", "bottom", "contributing_fraction",
        "all.profiles", "variable"), variable_name = "horizon")
```

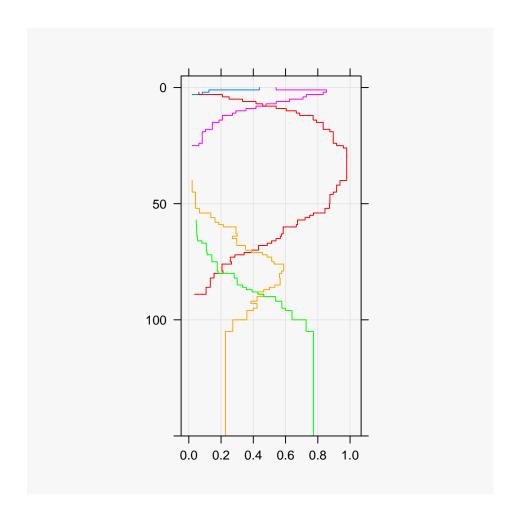


Figure 1: : horizon proportions

```
# plot horizon type proportions
p1 <- xyplot(top ~ value, groups = horizon, data =
loafercreek.hz.agg.long,
   ylim = c(150, -5), type = c("S", "g"), horizontal = TRUE,
   subset = value > 0 & horizon != "other", asp = 2, ylab = "",
   xlab = "")
```