plot.glm

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This is a minimal example showing the diagnostic plots.

1 Effect of probability on the diagnostic statistics

This is shown in table 1.

The relationship can be shown graphically as below:

```
p \leftarrow seq(5)
h \leftarrow c(1, 3, 1.5, 3, 1)
plot(p, h, type="b",
     col="blue", axes=FALSE,
     xlab="Probability",
     ylab="Size of statistic")
axis(1, at=p, labels=c("<0.1", "0.1-0.3", "0.3-0.7", "0.7-0.9", ">0.9"))
axis(2, at=c(1, 1.5, 2, 3), labels=c("low", "low-med", "med", "high"))
dChisq <- c(2, 1.5, 2)
points(c(2, 3, 4), dChisq, type="b", col="green")
lines(x=c(1, 2), y=c(3, 2), col="green")
lines(x=c(1, 2), y=c(1, 2),col="green")
lines(x=c(4, 5), y=c(2, 1), col="green")
lines(x=c(4, 5), y=c(2, 3),col="green")
dBhat \leftarrow c(1, 3, 2, 3, 1)
points(p, dBhat, type="b", col="red")
legend(2.5, y=1.4, legend=c("h", "dChisq", "dBhat"),
       fill=c("blue", "green", "red"))
mtext("Probability vs. Size of statistic")
```

P	h	$d\chi^2$	dβ̂
< 0.1	1	l / h	1
0.1 - 0.3	h	m	h
0.3 - 0.7	l - m	l - m	m
0.7 - 0.9	h	m	h
>0.9	1	l/ h	1

Abbreviations:

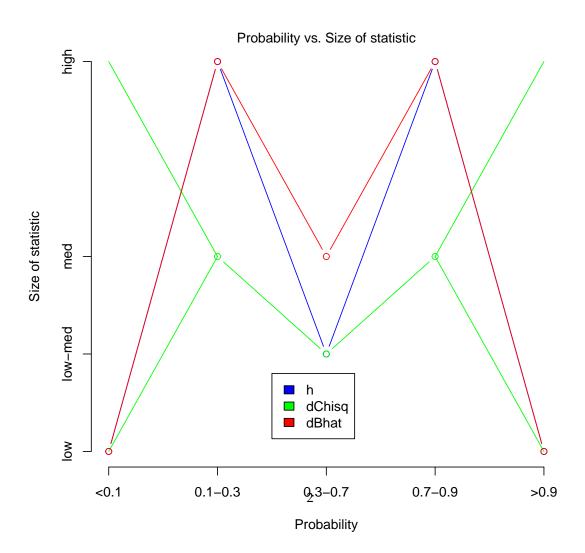
1 = low

1 - m = low to medium

m = medium

l/ h = low or high
 h = high

Table 1: Effect of probability on diagnostics



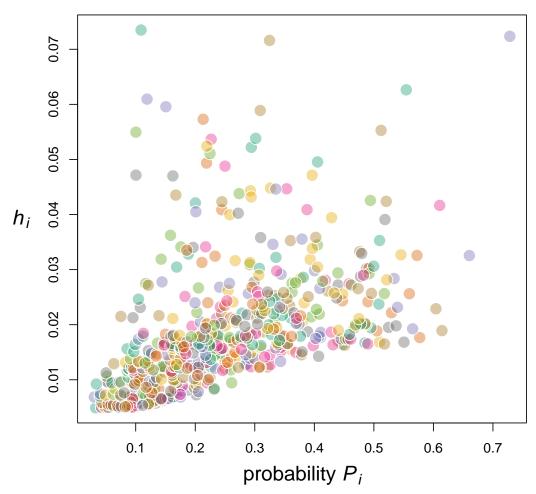
2 Sample graphical output from plot.glm

```
library("LogisticDx")
print("H&L 2nd ed. Table 4.9. Figures 5.5-5.8. Pages 177-180.")
## [1] "H&L 2nd ed. Table 4.9. Figures 5.5-5.8. Pages 177-180."
data(uis)
uis <- within(uis, {</pre>
   NDRGFP1 <- 10 / (NDRGTX + 1)
   NDRGFP2 <- NDRGFP1 * log((NDRGFP1 + 1) / 10)
})
summary(g1 <- glm(DFREE ~ AGE + NDRGFP1 + NDRGFP2 + IVHX +</pre>
              RACE + TREAT + SITE +
              AGE:NDRGFP1 + RACE:SITE,
              family=binomial, data=uis))
##
## Call:
## glm(formula = DFREE ~ AGE + NDRGFP1 + NDRGFP2 + IVHX + RACE +
     TREAT + SITE + AGE: NDRGFP1 + RACE: SITE, family = binomial,
     data = uis)
##
## Deviance Residuals:
## Min 1Q Median 3Q Max
## -1.3036 -0.7914 -0.5783 0.9902 2.6024
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -6.607764 1.177521 -5.612 2.00e-08 ***
               ## AGE
## NDRGFP1
               ## NDRGFP2 -0.461492 0.123824 -3.727 0.000194 ***
## IVHXprevious -0.633915 0.298774 -2.122 0.033862 *
             ## IVHXrecent
              ## RACEother
## TREATlong
              0.433709 0.203791 2.128 0.033320 *
              ## SITEB
## AGE:NDRGFP1 -0.015260 0.006026 -2.532 0.011329 *
## RACEother:SITEB -1.431087   0.529872   -2.701   0.006917 **
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
```

```
## Null deviance: 653.73 on 574 degrees of freedom
## Residual deviance: 597.92 on 564 degrees of freedom
## AIC: 619.92
##
## Number of Fisher Scoring iterations: 4
plot(g1, devNew=FALSE)
```

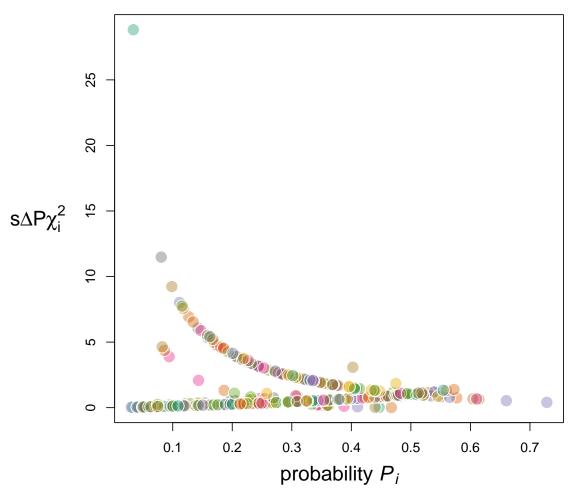
Probability $P_i \times \text{leverage } h_i$

 $0.1 < P_i > 0.9 \rightarrow h_i \sim x_i - \mu_x$ $h_i \approx$ distance of covariate pattern x_i from mean μ_x $h_i =$ diagonal of hat matrix

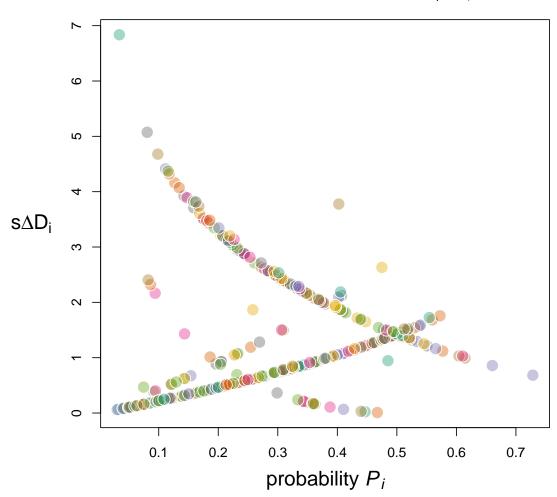


Probability $P_i \times$ scaled change in Pearson chi–sq s ΔP_i

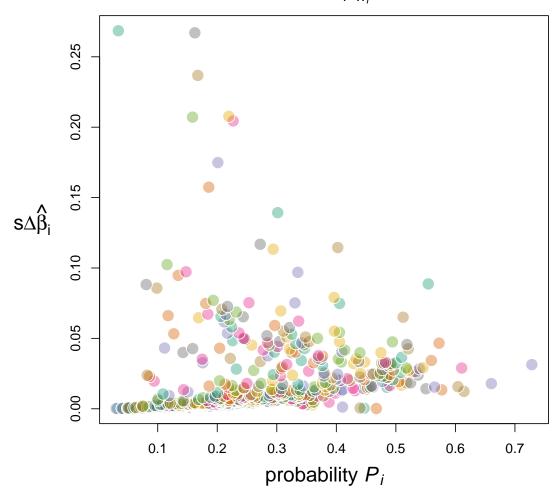
$$Pr_i = \frac{y_i - \mu_y}{\sigma_y}$$
, $s\Delta P\chi_i^2 = \frac{Pr_i}{\sqrt{1 - h_i}}$



Probability $P_i \times$ scaled change in deviance ΔD_i $dr_i = \text{sign}(y_i - \hat{y}_i) \sqrt{d_i}$, $s\Delta D_i = \frac{dr_i}{\sqrt{1 - h_i}}$

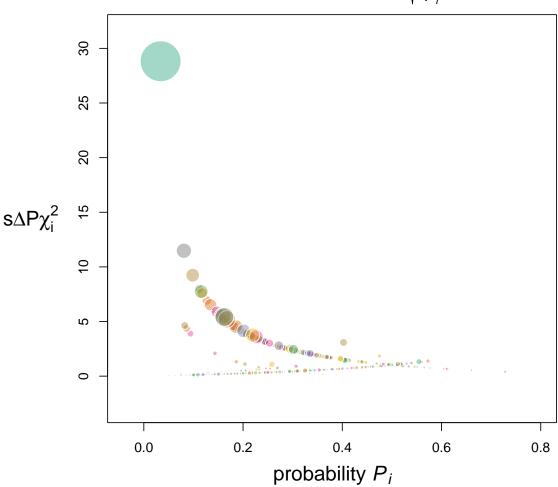


Probability $P_i \times$ scaled change in coefficients $s\Delta \hat{\beta}_i$ $s\Delta \hat{\beta}_i = \frac{sPr_i^2h_i}{1-h_i}$

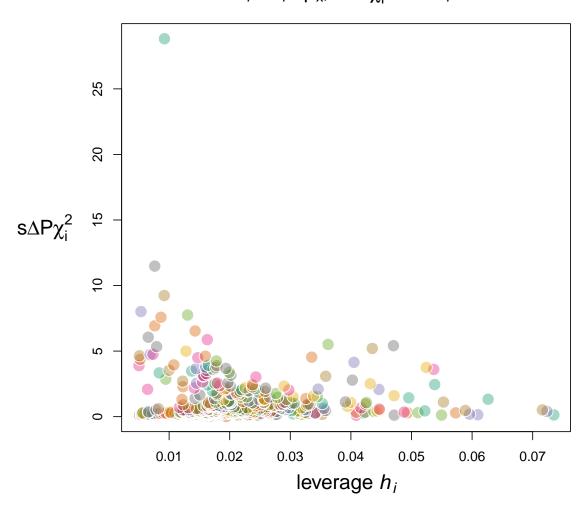


Probability $P_i \times \text{scaled change in Pearson chi-sq s}\Delta P_i$

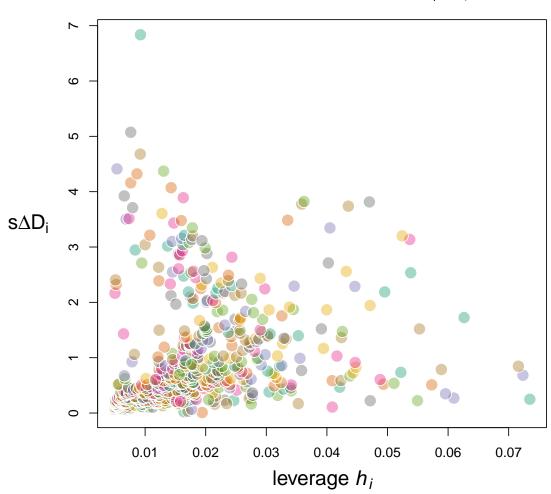
area
$$\propto s\Delta \hat{\beta}_i$$
, radius = $\sqrt{\frac{s\Delta \hat{\beta}_i}{P_i}}$



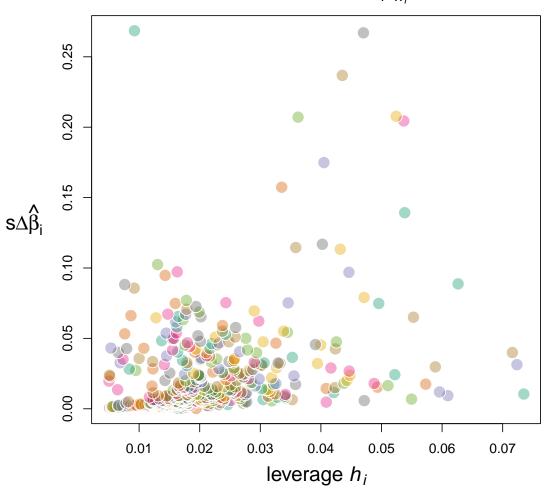
Leverage $h_i \times$ scaled change in Pearson chi–sq s $\Delta P\chi$ $h_i \approx x_i - \mu_x$, s $\Delta P\chi_i^2 = sPr_i^2$



Leverage $h_i \times$ scaled change in deviance $s\Delta D_i$ $dr_i = \text{sign}(y_i - \hat{y}_i)\sqrt{d_i}$, $s\Delta D_i = \frac{dr_i}{\sqrt{1-h_i}}$



Leverage $h_i \times$ scaled change in coefficients $s\Delta \hat{\beta}_i$ $h_i \approx x_i - \overline{x}, \ s\Delta \hat{\beta}_i = \frac{sPr_i^2h_i}{1-h_i}$



Correlation between $s\Delta P\chi_i^2,\,s\Delta D_i$ and $s\Delta {\hat \beta}_i$

