## 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	1 / 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:

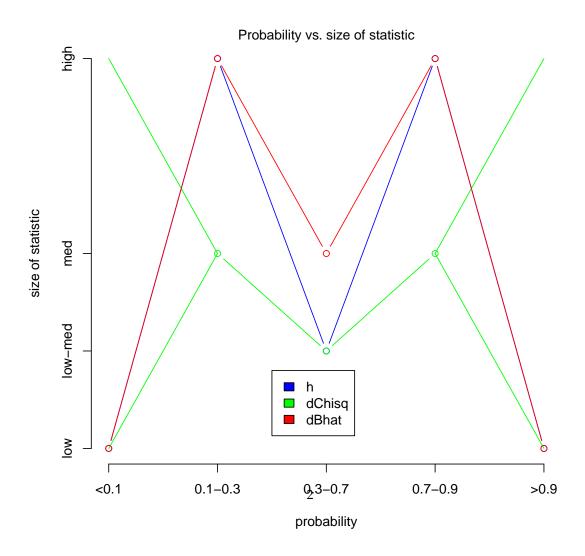
1 = low

1 - m = low to medium

m = medium

l/ h = low or high
 h = high

Table 1: Effect of probability on diagnostics



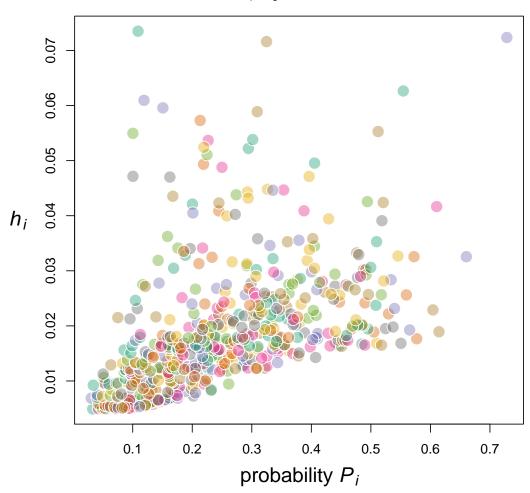
#### 2 Sample graphical output from plot.glm

```
library("LogisticDx")
## 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
## -1.304 -0.791 -0.578 0.990
                                2.602
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -6.60776 1.17752 -5.61 2.0e-08 ***
                ## AGE
                ## NDRGFP1
## NDRGFP2
               ## IVHXprevious -0.63391 0.29877 -2.12 0.03386 *
## IVHXrecent -0.70526 0.26163 -2.70 0.00703 **
## RACEother
              0.68626 0.26419 2.60 0.00939 **
0.43371 0.20379 2.13 0.03332 *
## TREATlong
                ## SITEB
## AGE:NDRGFP1 -0.01526 0.00603 -2.53 0.01133 *
## RACEother:SITEB -1.43109 0.52987 -2.70 0.00692 **
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 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: 619.9
```

```
##
## Number of Fisher Scoring iterations: 4
plot(g1, devNew=FALSE)
```

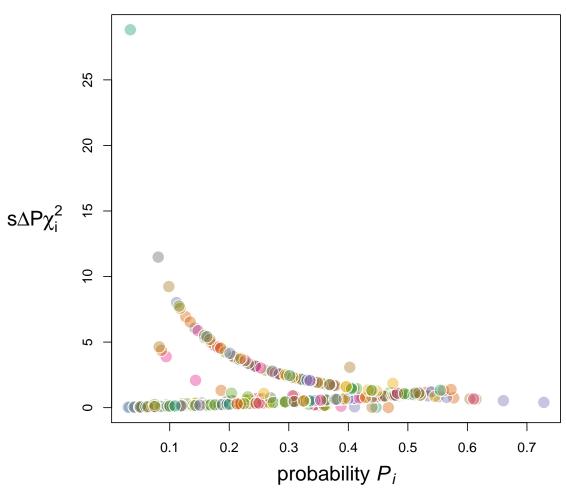
## Probability $P_i \times \text{leverage } h_i$

 $\begin{array}{ccc} 0.1 <\!\!P_i\!\!>\!\!0.9 & \to & h_i \!\!\simeq\!\! x_i\!\!-\!\!\mu_x \\ h_i \!\!\approx\!\! & \text{distance of covariate pattern } x_i \text{ from mean } \mu_x \\ h_i \!\!=\!\! & \text{diagonal of hat matrix} \end{array}$ 

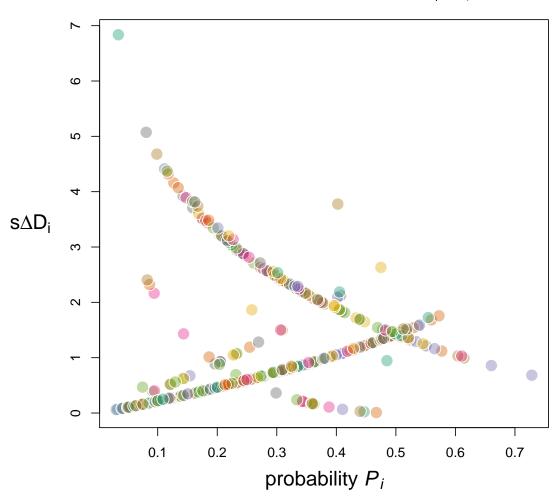


Probability  $P_i \times$  scaled change in Pearson chi–sq s $\Delta P_i$ 

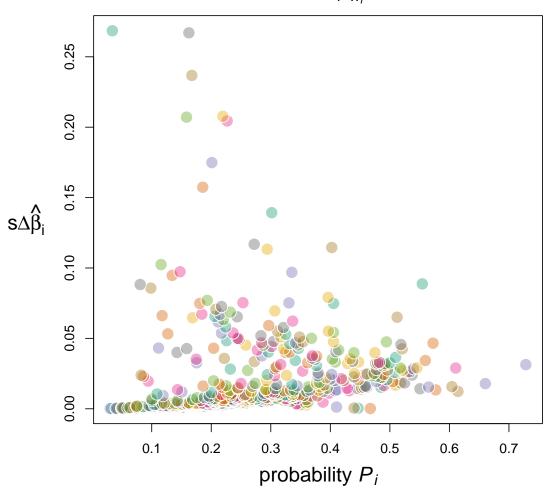
$$Pr_i = \frac{y_i - \mu_y}{\sigma_y}$$
,  $s\Delta P\chi_i^2 = \frac{Pr_i}{\sqrt{1 - h_i}}$ 



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

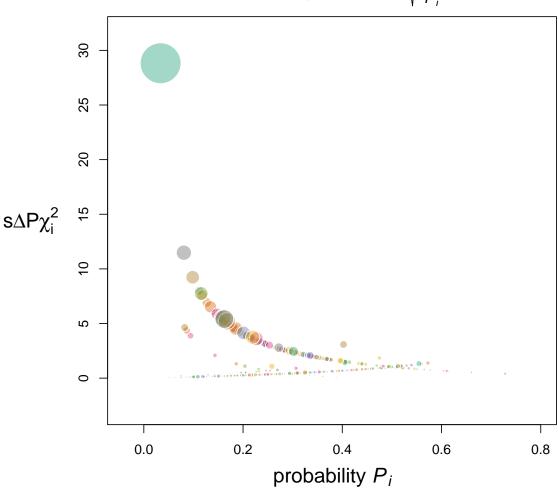


Probability  $P_i \times$  scaled change in coefficients  $s\Delta \hat{\beta}_i$   $s\Delta \hat{\beta}_i = \frac{sPr_i^2h_i}{1-h_i}$ 

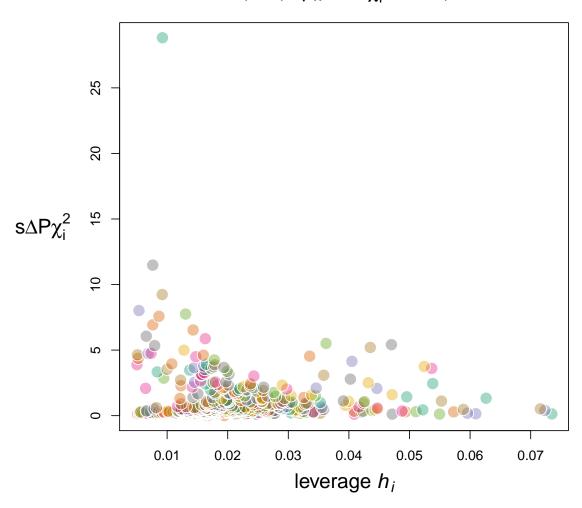


Probability  $P_i \times \text{scaled change in Pearson chi-sq s}\Delta P_i$ 

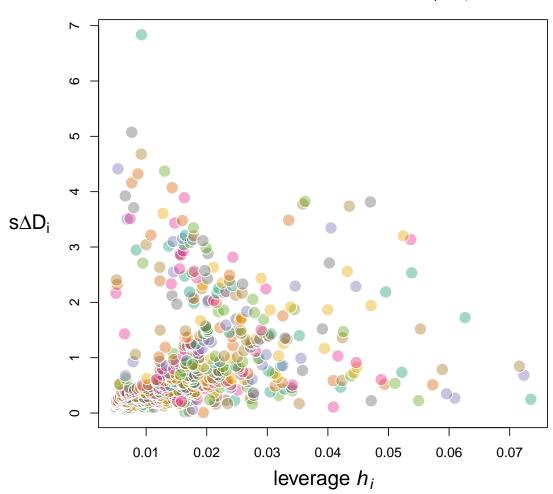
area 
$$\propto s\Delta \hat{\beta}_i$$
, radius =  $\sqrt{\frac{s\Delta \hat{\beta}_i}{P_i}}$ 



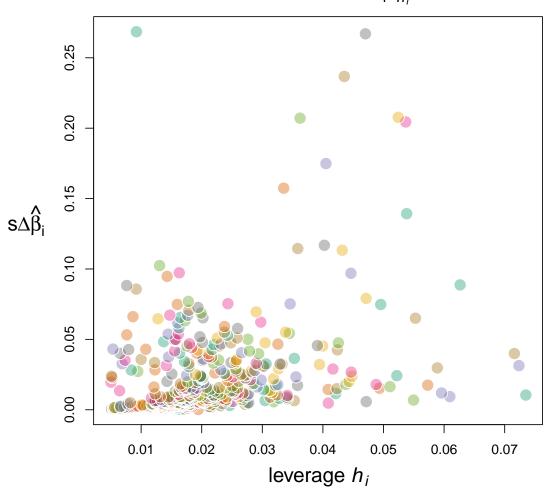
## Leverage $h_i \times$ scaled change in Pearson chi–sq s $\Delta P\chi$ $h_i \approx x_i - \mu_x$ , s $\Delta P\chi_i^2 = sPr_i^2$



## 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 $s\Delta {\hat \beta}_i$

