# Power of Wald test for interaction

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## 1 Power of Wald test:

#### 1.1 For one parameter:

If we consider using Wald test to test the null hypothesis  $H_0: \theta = \theta_0$ , the test statistic will be

$$\sqrt{\frac{n}{I^{-1}(\hat{\theta})}} \left( \hat{\theta} - \theta_0 \right)$$

where  $\hat{\theta}$  is the MLE of  $\theta$  and  $I^{-1}(\hat{\theta})$  is inverse of the fisher information evaluated at the MLE.

Under null hypothesis, this test statistic follows a standard normal distribution. If the alternative hypothesis  $H_a: \theta = \theta_1 \neq \theta_0$ , then the power of our test can be computed as:

$$1 - \Phi(\Delta + z_{a/2}) + \Phi(\Delta - z_{a/2})$$

where 
$$\Delta = \sqrt{\frac{n}{I^{-1}(\hat{\theta})}}(\theta_0 - \theta_1)$$
.

The important information above is that the power of Wald test will depend on several things at the same time:

- The difference of  $\theta_0 \theta_1$
- The sample size n
- The true information's inverse  $I^{-1}(\theta_1)$

In general, higher power is achieved when  $\Delta$  is large. That implies, we have higher power if we have larger  $|\theta_0 - \theta_1|$ , n or smaller  $I^{-1}(\theta_1)$ .

#### 1.2 General Expression and Non-centrality parameter:

In general for a vector of parameter  $\beta \in \mathbb{R}^p$ , if we are interested in testing the null hypothesis of linear combination of  $\beta$ :  $L\beta = 0$ , then the test statistic will be

$$(L\hat{\beta})^T L I_n^{-1}(\hat{\beta}) L^T (L\hat{\beta})$$

which follows a chi-square distribution with 1 degree of freedom under the null hypothesis. The matrix  $I_n^{-1}(\hat{\beta})$  is the inverse of the fisher information matrix of the whole sample, evaluated at  $\beta = \hat{\beta}$ .

Specifically, if we want to test the case where L = (0, 0, ..., 1, 0..., 0), then our test statistic reduced to

$$I_n^{-1}(\hat{\beta})_{[i,i]}(\hat{\beta}_i)^2$$

where  $\beta_i$  is the i-th component of  $\beta$ ,  $I_n^{-1}(\hat{\beta})_{[i,i]}$  is the i-th diagonal term of  $I_n^{-1}(\hat{\beta})$ . Under the alternative hypothesis that  $\beta_i = \beta_{i1} \neq 0$ , the limiting distribution will be a non-central chi-square distribution with non-centrality parameter being

$$I_n^{-1}(\beta)_{[i,i]}\beta_{i1}$$

here  $I_n^{-1}(\beta)$  is inverted fisher information matrix evaluated at the true parameter vector  $\beta$ .

Note that to compute  $I_n^{-1}(\beta)$ , we may need more than just  $\beta_{i1}$ . Even though we are only interested in testing  $\beta_i$ , that does not imply we can compute  $I_n^{-1}(\beta)$  just using the true value of  $\beta_i$ . We will discuss this problem in the next section in details.

# 2 When will other parameters matter for our power?

#### 2.1 Wald test for lm:

To answer this question, we need to know when will  $I_n^{-1}(\beta)$  depend on parameters other than  $\beta_i$ . It turns out that if the true underlying model of our data is a linear regression model,  $I_n^{-1}(\beta)$  will not depend on any parameter. However, if the true underlying model is not an ordinary linear regression model, but a generalized linear regression model, then  $I_n^{-1}(\beta)$  will depend on all the regression parameters.

If we consider the linear regression model:

$$y = \beta_0 + \beta_G G + \beta_E E + \beta_{GE} G \times E + \epsilon$$

where  $\epsilon \sim N(0, \sigma^2)$ 

Then the fisher information matrix at  $\beta$  can be computed as

$$XX^T/\sigma^2$$

Note that this matrix is only affected by the variance parameter  $\sigma^2$ , and not affected by any regression parameter  $\beta$ . Therefore, the power function of Wald test for  $\beta_{GE} = 0$  will only depend on the magnitude of the true value of  $\beta_{GE}$ .

### 2.2 Wald test for glm:

On the other hand, if the true data generating model is

$$\mu = g^{-1}(\beta_0 + \beta_G G + \beta_E E + \beta_{GE} G \times E)$$

where g(.) is a specified link function connecting the linear predictor with the mean of y.

In this case, the information matrix can be written as

$$XW(\beta)X^T$$

where the matrix  $W(\beta)$  is a diagonal matrix with each term depends on all the regression parameter  $\beta$ .

For a specific example, if we are using probit regression model, then the matrix  $W(\beta)$  will be

$$W(\beta) = diag \left\{ \frac{\phi^2(s_i)}{\Phi(s_i)(1 - \Phi(s_i))} \right\}$$

where  $s_i = \beta_0 + \beta_G G_i + \beta_E E_i + \beta_{GE} G_i \times E_i$  and  $\phi$ ,  $\Phi$  are the density and cdf of standard normal distribution respectively.

## 3 Computational Experiment:

Here we illustrate the phenomenon above using some computational experiment, and compare the empirical power and the theoretical power:

## 3.1 Assume data generated from ordinary linear model

In this section, we consider continuous trait generated from a linear regression model:

First case, we consider the true  $\beta_{GE}=0.1$  and true main effect  $\beta_E=1$ :

```
set.seed(100)
N <- 1000
G \leftarrow sample(c(0,1,2), size = N, replace = T, prob = c(0.16,0.48, 0.36))
E <- rnorm(N)
beta0 <- -1
betaG <- 0.3
betaE <- 1
betaGE <- 0.1
ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)</pre>
mod <- glm(ylat~G+E+I(G*E), family = gaussian(link = "identity"))</pre>
#### Get the design matrix:
X <- cbind(rep(1,N),mod$model[,-1])</pre>
### Compute the weight matrix W:
beta <- c(beta0,betaG,betaE,betaGE)</pre>
### The true information matrix
I <- as.matrix(t(X)) %*% as.matrix(X)</pre>
#### Invert to get the true covariance matrix
V <- solve(I)</pre>
### Compute the power function
### Assume d = beta0 - beta1, where beta0 = 0 is what we are testing as null:
delta \leftarrow sqrt(1/V[4,4])*(0-beta[4])
alpha \leftarrow 0.05
Power <- 1- pnorm(delta - qnorm(alpha/2)) + pnorm(delta + qnorm(alpha/2))
```

## [1] 0.6030156

```
#### Illustrate that this power is correct:
set.seed(100)
p1 <- c()
for (i in 1:1000) {
   ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)
   mod <- glm(ylat~G+E+I(G*E), family = gaussian(link = "identity"))
   p1[i] <- summary(mod)$coefficient[4,4]
}
emp_power <- mean(p1 <= alpha)
emp_power</pre>
```

Second case, we consider the true  $\beta_{GE} = 0.1$  fixed but change the true main effect  $\beta_E$  from 1 to 0:

```
set.seed(100)
N <- 1000
G \leftarrow sample(c(0,1,2), size = N, replace = T, prob = c(0.16,0.48, 0.36))
E <- rnorm(N)
beta0 <- -1
betaG <- 0.3
betaE <- 0
betaGE <- 0.1
ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)</pre>
mod <- glm(ylat~G+E+I(G*E), family = gaussian(link = "identity"))</pre>
#### Get the design matrix:
X <- cbind(rep(1,N),mod$model[,-1])</pre>
### Compute the weight matrix W:
beta <- c(beta0,betaG,betaE,betaGE)</pre>
### The true information matrix
I <- as.matrix(t(X)) %*% as.matrix(X)</pre>
#### Invert to get the true covariance matrix
V <- solve(I)</pre>
### Compute the power function
### Assume d = beta0 - beta1, where beta0 = 0 is what we are testing as null:
delta \leftarrow sqrt(1/V[4,4])*(0-beta[4])
alpha <- 0.05
Power <- 1- pnorm(delta - qnorm(alpha/2)) + pnorm(delta + qnorm(alpha/2))
```

```
#### Illustrate that this power is correct:
set.seed(100)
p1 <- c()
for (i in 1:1000) {
   ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)
   mod <- glm(ylat~G+E+I(G*E), family = gaussian(link = "identity"))
   p1[i] <- summary(mod)$coefficient[4,4]
}
emp_power <- mean(p1 <= alpha)
emp_power</pre>
```

We can see that both the theoretical power and empirical power are not changed at all.

### 3.2 Assume data generated from probit regression model

Here, we consider the binary trait generated from a probit regression model:

First case, we consider the true  $\beta_{GE} = 0.1$  and true main effect  $\beta_E = 1$ :

```
set.seed(100)
N <- 1000
G \leftarrow sample(c(0,1,2), size = N, replace = T, prob = c(0.16,0.48, 0.36))
E \leftarrow rnorm(N)
beta0 <- -1
betaG <- 0.3
betaE <- 1
betaGE <- 0.1
ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)</pre>
y \leftarrow ifelse(ylat >=0, 1, 0)
mod <- glm(y~G+E+I(G*E), family = binomial(link = "probit"))</pre>
#### Get the design matrix:
X <- cbind(rep(1,N),mod$model[,-1])</pre>
### Compute the weight matrix W:
beta <- c(beta0,betaG,betaE,betaGE)</pre>
#beta <- as.numeric(mod$coefficients)</pre>
W \leftarrow c()
for (i in 1:N) {
  si <- as.numeric(as.numeric(X[i,]) %*% beta)</pre>
  w[i] <- (dnorm(si)^2)/(pnorm(si)*(1-pnorm(si)))
### The true information matrix
I <- as.matrix(t(X)) %*% diag(w,nrow = N,ncol = N) %*% as.matrix(X)</pre>
```

```
#### Invert to get the true covariance matrix
V <- solve(I)

### Compute the power function

### Assume d = beta0 - beta1, where beta0 = 0 is what we are testing as null:
delta <- sqrt(1/V[4,4])*(0-beta[4])
alpha <- 0.05
Power <- 1- pnorm(delta - qnorm(alpha/2)) + pnorm(delta + qnorm(alpha/2))
Power</pre>
```

```
#### Illustrate that this power is correct:
set.seed(100)
p1 <- c()
for (i in 1:1000) {
   ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)
   y <- ifelse(ylat >=0, 1, 0)
   mod <- glm(y~G+E+I(G*E), family = binomial(link = "probit"))
   p1[i] <- summary(mod)$coefficient[4,4]
}
emp_power <- mean(p1 <= alpha)
emp_power</pre>
```

### ## [1] 0.167

Second case, we consider the true  $\beta_{GE} = 0.1$  fixed but change the true main effect  $\beta_E$  from 1 to 0:

```
set.seed(100)
N < -1000
G \leftarrow sample(c(0,1,2), size = N, replace = T, prob = c(0.16,0.48, 0.36))
E <- rnorm(N)
beta0 <- -1
betaG <- 0.3
betaE <- 0
betaGE <- 0.1
ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)</pre>
y <- ifelse(ylat >=0, 1, 0)
mod <- glm(y~G+E+I(G*E), family = binomial(link = "probit"))</pre>
#### Get the design matrix:
X <- cbind(rep(1,N),mod$model[,-1])</pre>
### Compute the weight matrix W:
beta <- c(beta0,betaG,betaE,betaGE)</pre>
#beta <- as.numeric(mod$coefficients)</pre>
```

```
w <- c()
for (i in 1:N) {
    si <- as.numeric(as.numeric(X[i,]) %*% beta)
    w[i] <- (dnorm(si)^2)/(pnorm(si)*(1-pnorm(si)))
}

### The true information matrix
I <- as.matrix(t(X)) %*% diag(w,nrow = N,ncol = N) %*% as.matrix(X)

#### Invert to get the true covariance matrix
V <- solve(I)

### Assume d = beta0 - beta1, where beta0 = 0 is what we are testing as null:
delta <- sqrt(1/V[4,4])*(0-beta[4])
alpha <- 0.05
Power <- 1- pnorm(delta - qnorm(alpha/2)) + pnorm(delta + qnorm(alpha/2))
Power</pre>
```

```
#### Illustrate that this power is correct:
set.seed(100)
p1 <- c()
for (i in 1:1000) {
   ylat <- beta0 + betaG*G + betaE*E + betaGE*G*E + rnorm(N)
   y <- ifelse(ylat >=0, 1, 0)
   mod <- glm(y~G+E+I(G*E), family = binomial(link = "probit"))
   p1[i] <- summary(mod)$coefficient[4,4]
}
emp_power <- mean(p1 <= alpha)
emp_power</pre>
```

### ## [1] 0.35

We can see that both the theoretical power and empirical power increased greatly.