

gradient

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1 Global epistatsis model

We assume that genotypes are stored in a $N \times L$ matrix, where N is the number of individuals and L is the number of loci. The model has the following parameters:

β , vector of effect sizes for different loci

α , power term that controls linearity

σ^2 , variance of the Gaussian error term (which has mean 0)

μ , population-wide mean

We define the mean for individual i as

$$\mu_i = \text{sgn}(b_i)|b_i|^\alpha + \mu,$$

where $b_i = \beta^T X_i = \sum_j \beta_j X_{ij}$. Alternatively, we write the phenotype Y_i for individual i as

$$Y_i = \text{sgn}(b_i)|b_i|^\alpha + \mu + \epsilon_i,$$

where ϵ_i is the Gaussian error for individual i .

In this model, the loglikelihood is

$$\begin{aligned} LL(\theta) &= \sum_{i=1}^n \log P(Y_i | \theta) \\ &= \sum_{i=1}^n \log \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(Y_i - \mu_i)^2}{2\sigma^2}\right) \\ &= -\frac{n}{2} \log 2\pi\sigma^2 - \frac{1}{2\sigma^2} \sum_{i=1}^n (Y_i - \mu_i)^2. \end{aligned}$$

1.0.1 gradients

First, the gradient for the effect size β_l . Defining operator $D := \frac{\delta}{\delta\beta_l}$,

$$\begin{aligned}
DLL(\theta) &= D \left(-n \log \sqrt{2\pi\sigma^2} - \frac{1}{2\sigma^2} \sum_{i=1}^n (Y_i - \mu_i)^2 \right) \\
&= -\frac{1}{2\sigma^2} \sum_{i=1}^n D(Y_i - \mu_i)^2 \\
&= \frac{1}{2\sigma^2} \sum_{i=1}^n 2(Y_i - \mu_i) D\mu_i
\end{aligned}$$

We have

$$D\mu_i = D[sgn(b_i)|b_i|^\alpha]$$

where $b = \sum_{j=1}^k \beta_j G_{ij}$ and $sgn(x)$ is the sign function. We have $Dsgn(b_i) = 0$, so, by the product rule,

$$\begin{aligned}
D\mu_i &= sgn(b) D(|b_i|^\alpha) \\
&= sgn(b_i) \alpha |b_i|^{\alpha-1} D|b_i| \\
&= sgn(b_i) \alpha |b_i|^{\alpha-1} sgn(b_i) G_{il} \\
&= \alpha |b_i|^{\alpha-1} G_{il}
\end{aligned}$$

And thus

$$= DLL(\theta) = \frac{\alpha}{\sigma^2} \sum_{i=1}^n (Y_i - \mu_i) \left| \sum_{j=1}^k \beta_j G_{ij} \right|^{\alpha-1} \cdot G_{il}.$$

1.0.2 Now we need to calculate with respect to μ , σ^2 and α .

For α Defining $D := \frac{\delta}{\delta\alpha}$, we again have

$$DLL(\theta) = \frac{1}{2\sigma^2} \sum_{i=1}^n 2(Y_i - \mu_i) D\mu_i \tag{1}$$

(2)

We can calculate

$$\begin{aligned}
D\mu_i &= D(sgn(b)|b|^\alpha + \mu) \\
&= sgn(b)|b|^\alpha \log |b|
\end{aligned}$$

where again $b = \sum_{j=1}^k \beta_j G_{ij}$. (And all the log's are base e .) Thus

$$\frac{\delta}{\delta\alpha} LL(\theta) = \frac{1}{\sigma^2} \sum_{i=1}^n (Y_i - \mu_i) sgn(b_i) |b_i|^\alpha \log |b_i|.$$

For μ Defining $D := \frac{\delta}{\delta\mu}$, we again have

$$DLL(\theta) = \frac{1}{2\sigma^2} \sum_{i=1}^n 2(X_i - \mu_i) D\mu_i \quad (3)$$

$$(4)$$

We can calculate

$$\begin{aligned} D\mu_i &= D(\text{sgn}(b)|b|^\alpha + \mu) \\ &= 1 \end{aligned}$$

and thus

$$\frac{\delta}{\delta\mu} LL(\theta) = \frac{1}{\sigma^2} \sum_{i=1}^n (X_i - \mu_i)$$

For σ^2 (remembering to ignore the squared, since it's just part of the parameter's name) Defining $D := \frac{\delta}{\delta\sigma^2}$, we again have

$$DLL(\theta) = D \left(-n \log \sqrt{2\pi\sigma^2} - \frac{1}{2\sigma^2} \sum_{i=1}^n (X_i - \mu_i)^2 \right) \quad (5)$$

$$= -\frac{n}{2\sigma^2} + \frac{1}{2\sigma^2} \sum_{i=1}^n (X_i - \mu_i)^2 \quad (6)$$

2 checking the gradient numerically

```
In [69]: import numpy as np
          from scipy.optimize import approx_fprime
          import numpy.random as npr
```

```
In [89]: k = 10
          G = npr.choice([-1,0,1], size = k, replace = True)
          beta = npr.uniform(-1,1, size = k)
          mu_0 = 4.5
          alpha = 4
```

```
In [90]: def mu_i(betas):
          b = np.sum(betas*G)
          absb = np.abs(b)
          sgnb = np.sign(b)
          return sgnb * absb**alpha + mu_0
```

```
In [91]: def Dmu_i(betas):
          absb = np.abs(np.sum(betas*G))
          return alpha*(absb**(alpha-1.0))*G
```

```
In [92]: mu_i(beta)
```

```
Out[92]: -8.932148169982433
```

```
In [93]: approx_fprime(beta, mu_i, 1e-9)
```

```
Out[93]: array([ 0.          , -28.06526389,  28.06526389, -28.06526389,  
                -28.06526389,  28.06526389,  28.06526389, -28.06526389,  
                -28.06526389,  28.06526389])
```

```
In [94]: Dmu_i(beta)
```

```
Out[94]: array([ 0.          , -28.06526173,  28.06526173, -28.06526173,  
                -28.06526173,  28.06526173,  28.06526173, -28.06526173,  
                -28.06526173,  28.06526173])
```

Checks out!