Expectation of R^2 with stratified phenotypes and genotypes

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Let Y_i , $i=1,\ldots,n$ be the stratification-adjusted phenotype centered around 0 and X_i , $i=1,\ldots,n$ be the stratification-adjusted genotype centered around 0.

We have

$$E[X_i] = 0 \forall i$$

$$E[Y_i] = 0 \forall i$$

$$Var[X_i] = \sigma_{X,i}^2$$

$$Var[X_i] = \sigma_{Y,i}^2$$

The Pearson correlation coefficient is

$$\hat{\rho} = \frac{\sum_{i=1}^{n} (X_{i} - \bar{X}) (Y_{i} - \bar{Y})}{\sqrt{\sum_{i=1}^{n} (X_{i} - \bar{X})^{2}} \sqrt{\sum_{i=1}^{n} (Y_{i} - \bar{Y})^{2}}}$$

$$\hat{\rho} = \frac{\sum_{i=1}^{n} X_{i} Y_{i}}{\sqrt{\sum_{i=1}^{n} (X_{i})^{2}} \sqrt{\sum_{i=1}^{n} (Y_{i})^{2}}}$$

$$R^{2} = \frac{(\sum_{i=1}^{n} X_{i} Y_{i})^{2}}{\sum_{i=1}^{n} (X_{i})^{2} \sum_{i=1}^{n} (Y_{i})^{2}}$$

$$= \frac{(\sum_{i=1}^{n} X_{i} Y_{i})^{2}}{\sum_{i=1}^{n} Var(X_{i}) \sum_{i=1}^{n} Var(Y_{i})}$$

$$E(R^{2}) = E\left[\frac{(\sum_{i=1}^{n} X_{i} Y_{i})^{2}}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}\right]$$

$$= \frac{E\left[(\sum_{i=1}^{n} X_{i} Y_{i})^{2}\right]}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{Var\left[\sum_{i=1}^{n} X_{i} Y_{i}\right] + E\left[\sum_{i=1}^{n} X_{i} Y_{i}\right]^{2}}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{Var\left[\sum_{i=1}^{n} X_{i} Y_{i}\right]}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{\sum_{i=1}^{n} Var(X_{i}Y_{i})}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{\sum_{i=1}^{n} \left[E(X_{i}^{2}) E(Y_{i}^{2}) - E(X_{i})^{2} E(Y_{i})^{2} \right]}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{\sum_{i=1}^{n} \left[E(X_{i}^{2}) E(Y_{i}^{2}) \right]}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{\sum_{i=1}^{n} \left[Var(X_{i}) Var(Y_{i}) \right]}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

$$= \frac{\sum_{i=1}^{n} \left[\sigma_{X,i}^{2} \sigma_{Y,i}^{2} \right]}{\sum_{i=1}^{n} \sigma_{X,i}^{2} \sum_{i=1}^{n} \sigma_{Y,i}^{2}}$$

For a binary phenotype and a diploid genotype we can express this as

$$E\left(R^{2}\right) = \frac{\sum_{i=1}^{n} \left[2\hat{p}_{X,i}\left(1-\hat{p}_{X,i}\right)\hat{p}_{Y,i}\left(1-\hat{p}_{Y,i}\right)\right]}{\sum_{i=1}^{n} 2\hat{p}_{X,i}\left(1-\hat{p}_{X,i}\right)\sum_{i=1}^{n} \hat{p}_{Y,i}\left(1-\hat{p}_{Y,i}\right)}$$