Single-cell RNA-sequencing: variance stabilizing transformations

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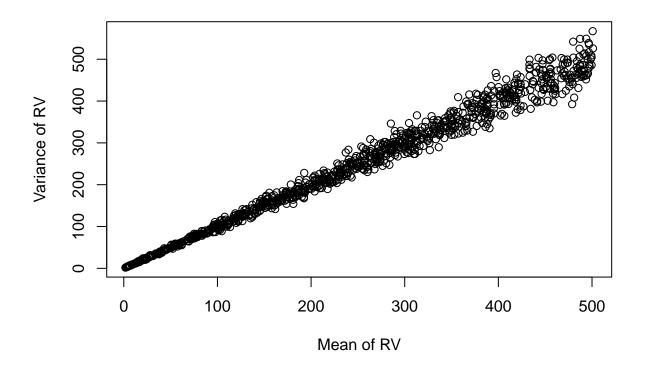
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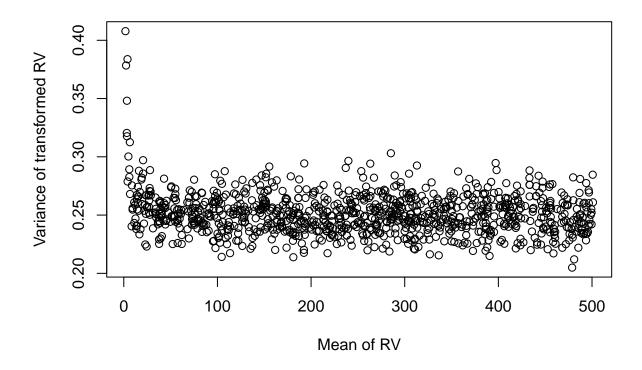
1 Approximating the variance stabilizing transformation of a Poisson random variable

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- A random variable $Y \sim Poi(\mu)$ has $Var(Y) = E(Y) = \mu$.
- We are looking for a variance stabilizing transformation (VST) f(Y) such that Var(f(Y)) = c, with c any constant. In particular, we need Var(f(Y)) to be independent of μ .
- A first-order Taylor series gives us $f(Y) \approx f(\mu) + (Y \mu)f'(\mu)$.
- Rearranging gives us $\left\{f(Y) f(\mu)\right\}^2 = (Y \mu)^2 f'(\mu)^2$.
- Which may be written as $Var(f(Y)) = Var(Y)f'(\mu)^2$.
- This shows us that we want a transformation such that $f'(\mu)^2 = 1/\mu$ because then $\forall \mu : Var(f(Y)) = 1!$ Let's find out.
- The last bullet point can be written as $f'(\mu) = \frac{1}{\sqrt{\mu}}$ and therefore $f(\mu) = \int \frac{1}{\sqrt{\mu}} d\mu = 2\mu^{1/2}$.
- Finally, this shows us that the transformation $f(Y) = 2Y^{1/2}$ ensures Var(f(Y)) = 1. Similar, as is often written in the scientific literature, the transformation $f(Y) = Y^{1/2}$ ensures Var(f(Y)) = 1/4.
- Note that these derivations all rely on the first-order Taylor expansion to be a good approximation. The VST will therefore work better for random variables with a high mean μ as the distribution will be more discrete for random variables with a low mean. We show this using simulation below.





Question. Why is the first plot heteroscedastic?

Answer.

Remember that $Var(\hat{\mu}) = \frac{\hat{\mu}}{n}$. Since for all random variables in our simulation n is equal, in our case we have that $Var(\hat{\mu}) = c\hat{\mu}$. The variance on estimating the mean thus increases with the mean.