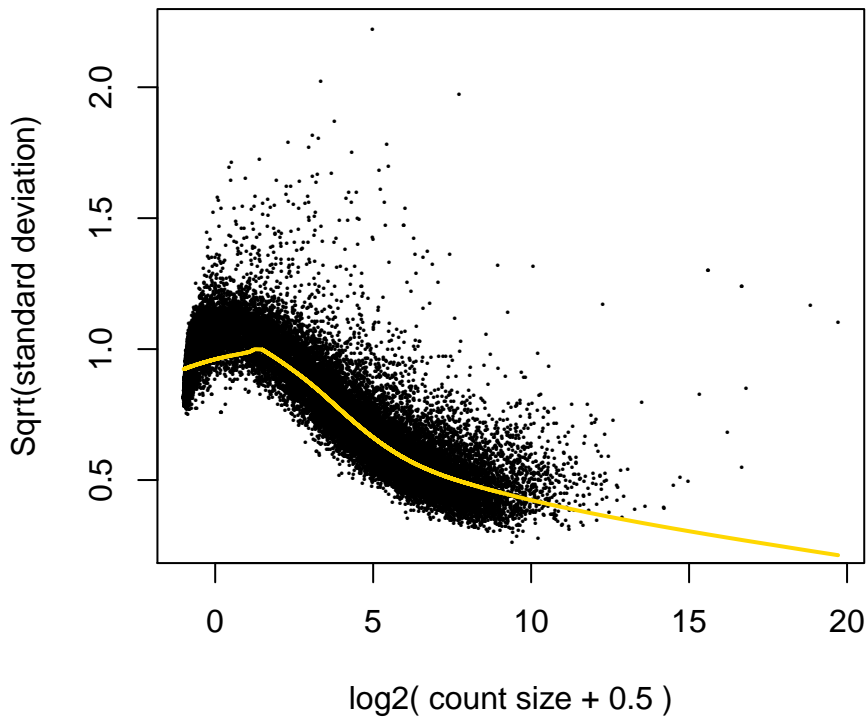


**Mean–variance trend**  
**Raw counts (gene level)**



**Mean–variance trend**  
**Filtered counts (gene level)**

