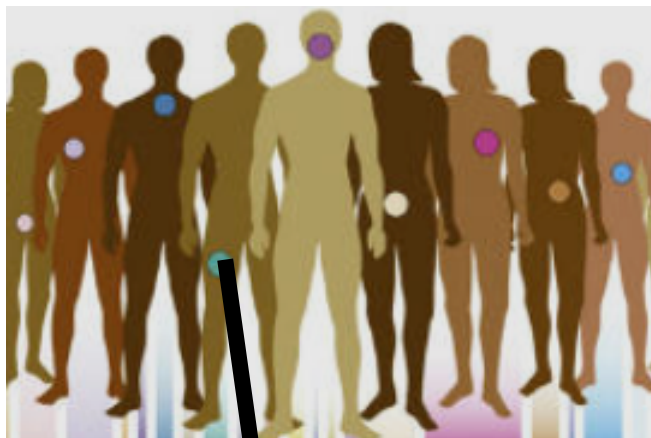


**Condition 1**

**Condition 2**

**Samples/  
Individuals**

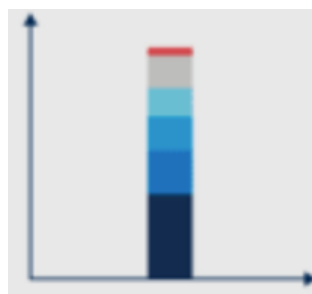


For each **sample**

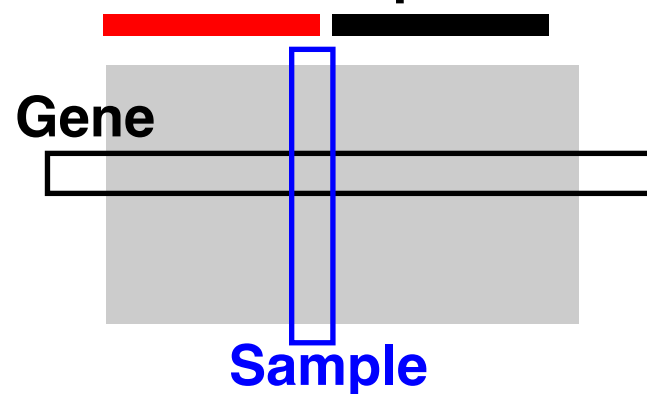
**Cells from part  
of the tissues**



**Bulk RNASeq:  
Mixture of cells**

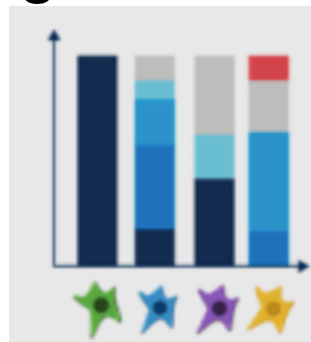


**Bulk RNASeq data matrix**



**Single cell RNASeq**

**Gene  
expressions**



**scRNASeq data matrix**

