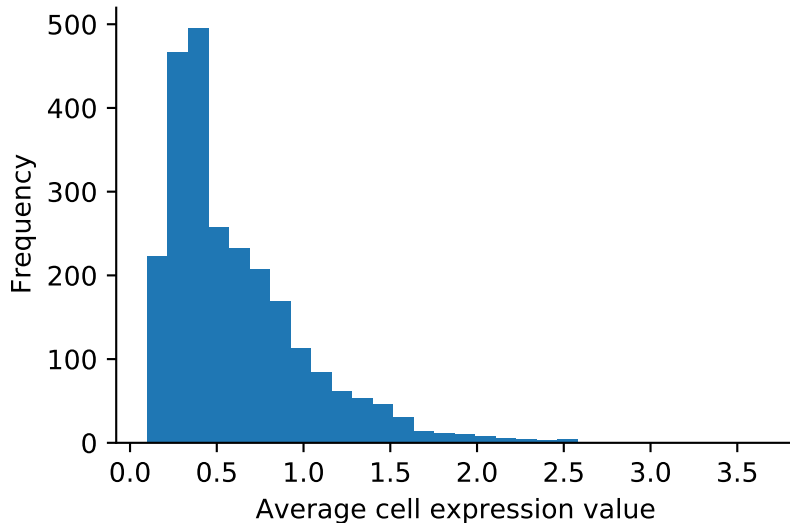


**(a) Average cell expression value
in Q Bladder**



**(b) Average expression value in cells
predicted in the wrong cluster**

