

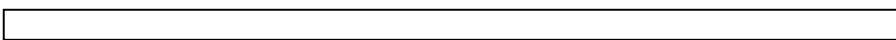
	bin 1	bin 2	...	bin m-1	bin m
gene 1					
gene 2					
gene 3					
...					
gene n					

groups

1
2
19
20

→ *quantile normalization*

average per bin



→ *linear regression*