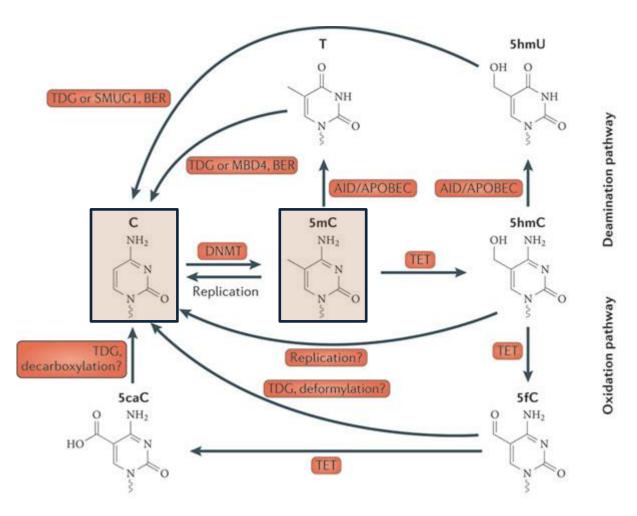
# Additional Cytosine Modifications – analysing hydroxymethylcytosine (5hmC) with oxidative BS-Seq (oxBS)

Felix Krueger felix.krueger@babraham.ac.uk January 2015



### **Cytosine modifications**

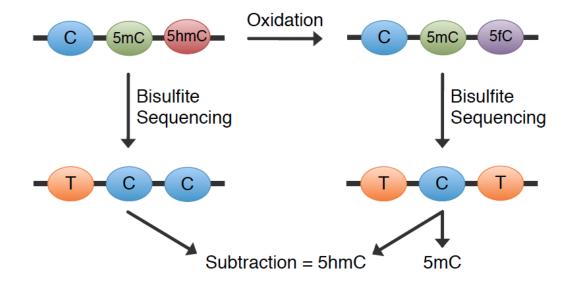


Miguel R. Branco, Gabriella Ficz & Wolf Reik Nature Reviews Genetics 13, 7-13 (January 2012)

Nature Reviews | Genetics



## Detecting 5-hydroxymethylcytosine using Oxidative Bisulfite-Seq (oxBS)

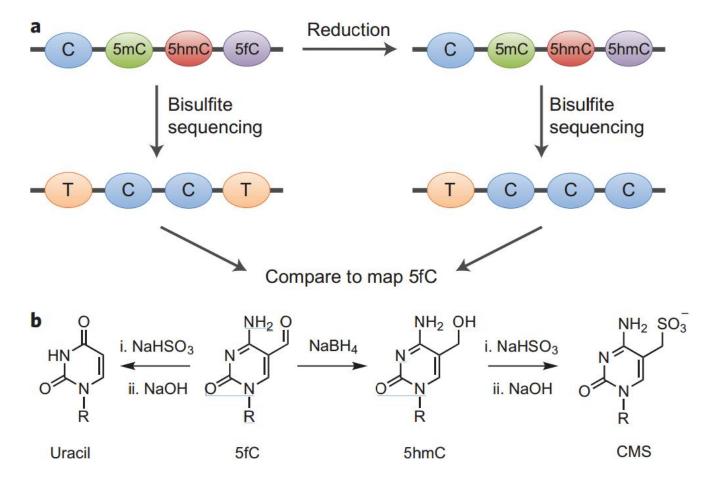


Quantitative sequencing of 5-methylcytosine and 5-hydroxymethylcytosine at single-base resolution Science, 2012 May 18;336(6083):934-7

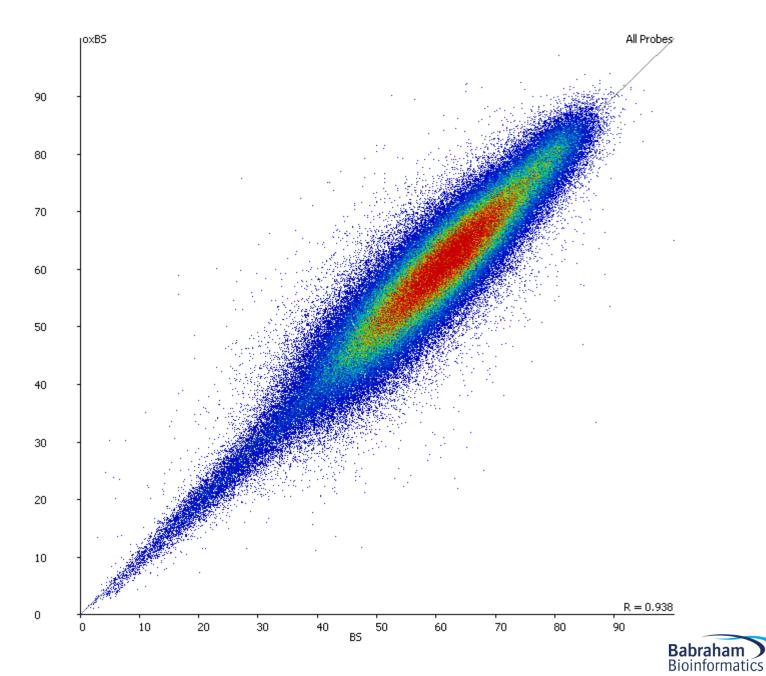




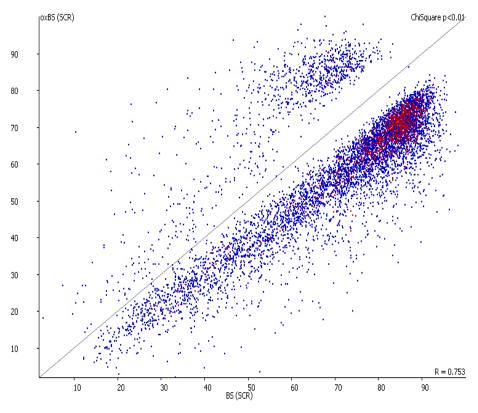
## Detecting 5-formylcytosine using Reduced Bisulfite-Seq (redBS)







#### oxBS: identifying target regions



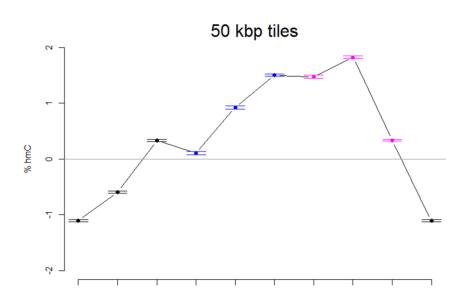
Gives an idea about false positive rate

Analysis options to gain more power:

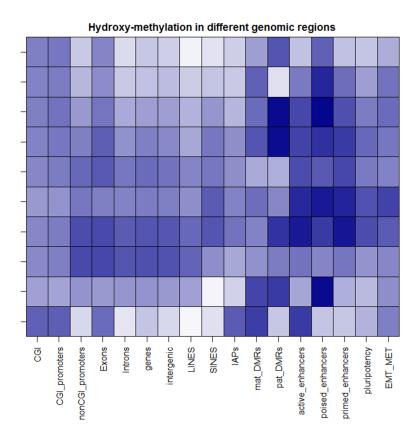
- larger running windows
- looking for sets/functional sets



#### hmC as difference BS - oxBS



genome-wide difference: sometimes negative values...



relative differences

