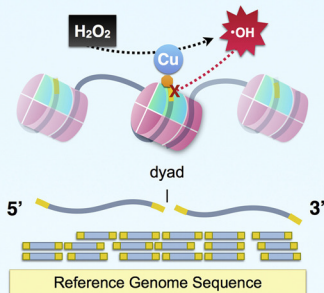


Dyad-localized  
Fenton reaction

Next-generation  
sequencing

Deconvolution and  
NCP score calculation

Genome-wide  
nucleosome map



$$W_i = f\left(\sum_j k_{(i-j)}\lambda_j + \sigma_{i1}\right); C_i = f\left(\sum_j k_{(i+j)}\lambda_j + \sigma_{i2}\right)$$

