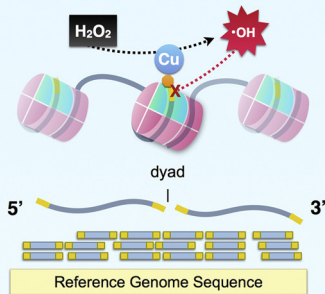


Dyad-localized
Fenton reaction

Next-generation
sequencing

Deconvolution and
NCP score calculation

Genome-wide
nucleosome map



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

