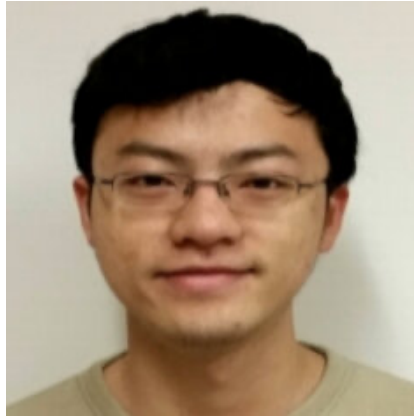


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PUBLICATIONS

- Wu H, **Wu X**, Shen L, Zhang Y (2014). [Single-base resolution analysis of active DNA demethylation using methylase-assisted bisulfite sequencing](#). *Nat Biotechnol.* 32(12):1231-1240.
- Wu H, **Wu X**, Zhang Y (2016). [Base-resolution profiling of active DNA demethylation using MAB-seq and caMAB-seq](#). *Nature Protocols* 11(6):1081-1100.
- Wu X**, Inoue A, Suzuki T, Zhang Y (2017) [Simultaneous mapping of active DNA demethylation and sister chromatid exchange in single cells](#). *Genes Dev.* 31(5):511-523
- Chen R, **Wu X**, Jiang L, Zhang Y (2017) [Single-Cell RNA-Seq Reveals Hypothalamic Cell Diversity](#). *Cell Reports* 18(13):3227-3241
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