

## **Michael Church, Ph. D**

Postdoctoral Research Associate

### Education

October 2010 to November 2015: Department of Microbiology, Trinity College Dublin, The University of Dublin, Ireland. Awarded the degree of Ph.D. for a thesis entitled "Investigating regulation of gene transcription by the Tup1-Ssn6 co-repressor complex in *Saccharomyces cerevisiae*." Work supervised by Dr. Alastair B. Fleming.

### Publications

1. Church MC, Workman JL (2024). The SWI/SNF chromatin remodeling complex: a critical regulator of metabolism. *Biochemical Society Transactions*, *BST20231141*
2. Church MC, Price A, Li H, Workman JL (2023) The Swi-Snf chromatin remodeling complex mediates gene repression through metabolic control. *Nucleic Acids Research*, *gkad711*.
3. Lee B, Church M, Hokamp K, Al-hussain M, Bamagoos AA, Fleming AB (2023) Systematic analysis of *tup1* and *cyc8* mutants reveals distinct roles for TUP1 and CYC8 and offers new insight into the regulation of gene transcription by the yeast Tup1-Cyc8 complex. *PLoS Genet* *19(8):e1010876*.
4. Church MC, Workman JL, Sukanuma T (2021) Macrophages, Metabolites, and Nucleosomes: Chromatin at the Intersection between Aging and Inflammation. *Int. J. Mol. Sci.*, *22(19)* , 10274
5. Church MC, Fleming AB. (2017) A role for histone acetylation in regulating transcription elongation. *Transcription*. *8:1-8*.
6. Church M, Smith KC, Alhussain MM, Pennings S, Fleming AB (2017). Sas3 and Ada2 (Gcn5)-dependent histone H3 acetylation is required for transcription elongation at the de-repressed *FLO1* gene. *Nucleic Acids Res* *45(8):4413-4430*.
7. Haran J, Boyle H, Hokamp K, Yeomans T, Liu Z, Church M, Fleming AB, Anderson MZ, Berman J, Myers LC, Sullivan DJ, Moran GP (2014). Telomeric ORFs (TLOs) in *Candida* spp. encode mediator subunits that regulate distinct virulence traits. *PLoS Genet*. *10(10):e1004658*
8. Fleming AB, Beggs S, Church M, Tsukihashi Y, Pennings S. (2014). The yeast Tup1-Cyc8 (Ssn6) complex cooperates with the Hda1 and Rpd3 histone deacetylases to robustly repress transcription of the subtelomeric *FLO1* gene. *Biochim Biophys Acta GRM*. *1839(11):1242-55*