**Annotated Bibliography of the Final Project**

1. Kim, Y. W., Lee, S., Yun, J. & Kim, A. Chromatin looping and eRNA transcription precede the transcriptional activation of gene in the β-globin locus. *Biosci. Rep.* **35,** 1–8 (2015).

2. Plank, J. L. & Dean, A. Enhancer Function: Mechanistic and Genome-Wide Insights Come Together. *Mol. Cell* **55,** 5–14 (2014).

3. Lam, M. T. Y., Li, W., Rosenfeld, M. G. & Glass, C. K. Enhancer RNAs and regulated transcriptional programs. *Trends Biochem. Sci.* **39,** 170–182 (2014).

4. Smith, E. & Shilatifard, A. Enhancer biology and enhanceropathies. *Nat. Struct. Mol. Biol.* **21,** 210–219 (2014).

5. Hsieh, C.-L. *et al.* Enhancer RNAs participate in androgen receptor-driven looping that selectively enhances gene activation. *Proc. Natl. Acad. Sci.* **111,** 7319–7324 (2014).

6. Li, W. *et al.* Functional roles of enhancer RNAs for oestrogen-dependent transcriptional activation. *Nature* **498,** 516–520 (2013).

7. Ule, J., Jensen, K., Mele, A. & Darnell, R. B. CLIP: A method for identifying protein–RNA interaction sites in living cells. *Methods* **37,** 376–386 (2005).

8. Kim, A. & Dean, A. Chromatin loop formation in the ??-globin locus and its role in globin gene transcription. *Mol. Cells* **34,** 1–5 (2012).

9. Deng, W. *et al.* Targeted Tethering of a Looping Factor. **149,** 1233–1244 (2013).

10. Lai, F. *et al.* Activating RNAs associate with Mediator to enhance chromatin architecture and transcription. *Nature* **494,** 497–501 (2013).

11. Splinter, E. *et al.* CTCF mediates long-range chromatin looping and local histone modification in the ??-globin locus. *Genes Dev.* **20,** 2349–2354 (2006).

12. Ossipow, V. & Walker, J. M. *Monoclonal Antibodies IN Series Editor*.

13. Kor, P., Antibodies, K. I. R., Lemos, J. C., Roth, C. a & Chavkin, C. Signal Transduction Immunohistochemistry. *Signal Transduct.* **717,** 197–219 (2011).