

RNA Polymerases IV and V: evolution and function

Carlotta Porcelli, qbp693

October 8, 2017

Abstract

This is gonna be the last thing I'll write.

1 Introduction

Most of the known non-coding RNAs (ncRNAs) correspond to intergenic sequences or transcripts with unknown functions. In mammals, two of the increasingly known long ncRNAs are *Xist* and *Tsix* which are involved in the regulation of adjacent genes. In plants, small-interfering RNAs (siRNAs) guide the process of chromatin modifications. These ncRNAs are generated from dsRNAs precursors and processed by dicer (DCL) enzymes into 21-24 nucleotides long siRNAs before associating with Argonaute proteins (AGO).

2 The paradox of epigenetic control

Need for transcription in order to transcriptionally silence the same region. [10]

2.1 The RNA-mediated DNA-methylation pathway (RdDM)

3 RNA Polymerase IV

how it produces siRNAs and what do they bind to : AGO specific and dicer specific. from: [1] and from [2] picture from: [3]

3.1 The subunits

- NRPD

3.2 The products and their function

- what genes are silenced and how
- DNA methylation

[4]

4 RNA Polymerase V

* open up dna or make transcripts for binding for ago ? see: [5]

see: [6]

4.1 The subunits

[7] [8]

- NRPE

[9]

4.2 The function

two proposed models as in [4]

5 Future outlook

References

1. Zhang, X., Henderson, I.R., Lu, C., Green, P.J., and Jacobsen, S.E. (2007). Role of RNA polymerase IV in plant small RNA metabolism. *Proceedings of the National Academy of Sciences* 104, 4536–4541. URL <http://www.pnas.org/content/104/11/4536.abstract>.
2. Onodera, Y., Haag, J.R., Ream, T., Nunes, P.C., Pontes, O., and Pikaard, C.S. (2005). Plant Nuclear RNA Polymerase IV Mediates

- siRNA and DNA Methylation-Dependent Heterochromatin Formation. *Cell* *120*, 613 – 622. URL <http://www.sciencedirect.com/science/article/pii/S0092867405001510>.
3. Xu, C., Tian, J., and Mo, B. (2013). siRNA-mediated DNA methylation and H3K9 dimethylation in plants. *Protein & Cell* *4*, 656–663. URL <https://doi.org/10.1007/s13238-013-3052-7>.
 4. Pikaard, C.S., Haag, J.R., Ream, T., and Wierzbicki, A.T. (2008). Roles of RNA polymerase IV in gene silencing. *Trends in Plant Science* *13*, 390 – 397. URL <http://www.sciencedirect.com/science/article/pii/S1360138508001398>.
 5. Daxinger, L., Kanno, T., and Matzke, M. (2008). Pol V Transcribes to Silence. *Cell* *135*, 592–594. URL <https://doi.org/10.1016/j.cell.2008.10.027>.
 6. Wierzbicki, A.T., Haag, J.R., and Pikaard, C.S. (2008). Noncoding Transcription by RNA Polymerase Pol IVb/Pol V Mediates Transcriptional Silencing of Overlapping and Adjacent Genes. *Cell* *135*, 635 – 648. URL <http://www.sciencedirect.com/science/article/pii/S0092867408011926>.
 7. Wendte, J.M., Haag, J.R., Singh, J., McKinlay, A., Pontes, O.M., and Pikaard, C.S. (2017). Functional Dissection of the Pol V Largest Subunit {CTD} in RNA-Directed {DNA} Methylation. *Cell Reports* *19*, 2796 – 2808. URL <http://www.sciencedirect.com/science/article/pii/S2211124717307829>.
 8. Zhou, M. and Law, J.A. (2015). RNA Pol IV and V in Gene Silencing: Rebel Polymerases Evolving Away From Pol II's Rules. *Current opinion in plant biology* *27*, 154–164. URL <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4618083/>.
 9. Ream, T.S., Haag, J.R., Wierzbicki, A.T., Nicora, C.D., Norbeck, A.D., Zhu, J.K., Hagen, G., Guilfoyle, T.J., Paša-Tolić, L., and Pikaard, C.S. (2009). Subunit Compositions of the RNA-Silencing Enzymes Pol IV and Pol V Reveal Their Origins as Specialized Forms of RNA Polymerase II. *Molecular Cell* *33*, 192 – 203. URL <http://www.sciencedirect.com/science/article/pii/S1097276508008587>.
 10. Wierzbicki, A.T., Haag, J.R., and Pikaard, C.S. Noncoding Transcription by RNA Polymerase Pol IVb/Pol V Mediates Transcriptional Silencing of Overlapping and Adjacent Genes. *Cell* *135*, 635–648. URL <http://dx.doi.org/10.1016/j.cell.2008.09.035>.