
Rat mRNA&IncRNA Epitranscriptomic microarray databse

References

1. Cunningham, F., M. R., et al., Ensembl 2015. *Nucleic Acids Res*, 2015. 43(Database issue): D662-669.
2. Kozomara A., miRBase: annotating high confidence microRNAs using deep sequencing data. *Nucleic Acid Res*, 2014. 42(Database issue): p. D68-73.
3. Harrow J., et al., GENCODE: the reference human genome annotation for the ENCODE project, 2012. 22(9): p. 1760-74.
4. Amaral, P.P., et al., lncRNADB: a reference database for long noncoding RNAs. *Nucleic Acids Res*, 2011. 39(Database issue): p. D146-51.
5. Pruitt, K.D., T. Tatusova, and D.R. Maglott, NCBI Reference Sequence (RefSeq): a curated nonredundant sequence database of genomes, transcripts and proteins. *Nucleic Acids Res*, 2005. 33(Database issue): p. D501-4.
6. Bejerano, G., et al., Ultraconserved elements in the human genome. *Science*, 2004. 304(5675): p. 1321-5.
7. Rinn, J.L., et al., Functional demarcation of active and silent chromatin domains in human HOX loci by noncoding RNAs. *Cell*, 2007. 129(7): p. 1311-23.
8. Mercer, T.R., et al., Specific expression of long noncoding RNAs in the mouse brain. *Proc Natl Acad Sci U S A*, 2008. 105(2): p. 716-21.
9. Guttman, M., et al., Chromatin signature reveals over a thousand highly conserved large non-coding RNAs in mammals. *Nature*, 2009. 458(7235): p. 223-7.
10. Benson, D.A., et al., GenBank: update. *Nucleic Acids Res*, 2004. 32(Database issue): p. D23-6.