- 70. Cramer, P., Pesce, C. G., Baralle, F. E., and Kornblihtt, A. R. (1997) Functional association between promoter structure and transcript alternative splicing. *Proceedings of the National Academy of Sciences of the United States of America* **94**, 11456-11460
- 71. Merkhofer, E. C., Hu, P., and Johnson, T. L. (2014) Introduction to cotranscriptional RNA splicing. *Methods Mol Biol* **1126**, 83-96
- 72. Hasson, P., Muller, B., Basler, K., and Paroush, Z. (2001) Brinker requires two corepressors for maximal and versatile repression in Dpp signalling. *EMBO J* **20**, 5725-5736
- 73. Paroush, Z., Finley, R. L., Jr., Kidd, T., Wainwright, S. M., Ingham, P. W., Brent, R., and Ish-Horowicz, D. (1994) Groucho is required for Drosophila neurogenesis, segmentation, and sex determination and interacts directly with hairy-related bHLH proteins. *Cell* **79**, 805-815
- 74. Grbavec, D., and Stifani, S. (1996) Molecular interaction between TLE1 and the carboxylterminal domain of HES-1 containing the WRPW motif. *Biochem Biophys Res Commun* 223, 701-705
- 75. Adelman, K., and Lis, J. T. (2012) Promoter-proximal pausing of RNA polymerase II: emerging roles in metazoans. *Nat Rev Genet* **13**, 720-731
- 76. Gaertner, B., and Zeitlinger, J. (2014) RNA polymerase II pausing during development. *Development* **141**, 1179-1183
- 77. Bartkowiak, B., Liu, P., Phatnani, H. P., Fuda, N. J., Cooper, J. J., Price, D. H., Adelman, K., Lis, J. T., and Greenleaf, A. L. (2010) CDK12 is a transcription elongation-associated CTD kinase, the metazoan ortholog of yeast Ctk1. *Genes Dev* **24**, 2303-2316
- 78. Lee, J. S., and Shilatifard, A. (2007) A site to remember: H3K36 methylation a mark for histone deacetylation. *Mutation research* **618**, 130-134
- 79. Wagner, E. J., and Carpenter, P. B. (2012) Understanding the language of Lys36 methylation at histone H3. *Nature reviews. Molecular cell biology* **13**, 115-126