

Posttranscriptional Regulation

- Control of gene expression usually involves the control of transcription initiation.
- But gene expression can be controlled after transcription, with mechanisms such as:
 - RNA interference
 - alternative splicing
 - RNA editing
 - mRNA degradation

Posttranscriptional Control Of Gene Expression

Ghedira Kais



Posttranscriptional Control Of Gene Expression:

Post-transcriptional Control of Gene Expression ,2003 Post-Transcriptional Control of Gene Expression

John E.G. McCarthy, Mick F. Tuite, 2013-06-29 The last ten years have witnessed a remarkable increase in our awareness of the importance of events subsequent to transcriptional initiation in terms of the regulation and control of gene expression In particular the development of recombinant DNA techniques that began in the 1970s provided powerful new tools with which to study the molecular basis of control and regulation at all levels The resulting investigations revealed a diversity of post transcriptional mechanisms in both prokaryotes and eukaryotes Scientists working on translation mRNA stability transcriptional anti termination or other aspects of gene expression will often have met at specialist meetings for their own research area However only rarely do workers in different areas of post transcriptional control regulation have the opportunity to meet under one roof We therefore thought it was time to bring together leading representatives of most of the relevant areas in a small workshop intended to encourage interaction across the usual borders of research both in terms of the processes studied and with respect to the evolutionary division prokaryotes eukaryotes Given the breadth of topics covered and the restrictions in size imposed by the NATO workshop format it was an extraordinarily difficult task to choose the participants However we regarded this first attempt as an experiment on a small scale intended to explore the possibilities of a meeting of this kind Judging by the response of the participants during and after the workshop the effort had been worthwhile

Post-transcriptional Control of Gene Expression Celine Sin, 2016 Gene expression describes the process of making functional gene products e.g proteins or special RNAs from instructions encoded in the genetic information e.g DNA This process is heavily regulated allowing cells to produce the appropriate gene products necessary for cell survival adapting production as necessary for different cell environments Gene expression is subject to regulation at several levels including transcription mRNA degradation translation and protein degradation When intact this system maintains cell homeostasis keeping the cell alive and adaptable to different environments Malfunction in the system can result in disease states and cell death In this dissertation we explore several aspects of gene expression control by analyzing data from biological experiments Most of the work following uses a common mathematical model framework based on Markov chain models to test hypotheses predict system dynamics or elucidate network topology Our work lies in the intersection between mathematics and biology

The Control of Gene Expression at the Posttranscriptional Level Richard Lit, 1991

Post-Transcriptional Gene Regulation Erik Dassi, 2016-08-23 This volume aims to provide the most recent advances in techniques for studying gene expression regulation at the post transcriptional level Post Transcriptional Gene Regulation Second Edition is organized in six sections describing bioinformatics approaches for studying post transcriptional regulation various expression profiling approaches protein RNA interaction and non coding RNA investigation techniques methods for profiling RNA modifications and other techniques such as alternative translation initiation or polyadenylation sites

determination Written for the Methods in Molecular Biology series chapters include introductions to their respective topics lists of the necessary materials and reagents step by step readily reproducible laboratory protocols and tips on troubleshooting and avoiding known pitfalls Authoritative and practical Post Transcriptional Gene Regulation Second Edition is a versatile resource to researchers studying post transcriptional regulation introducing the most recent techniques and providing a comprehensive guide to their implementation *Post-transcriptional Regulation of Gene Expression* Mariano A. Garcia-Blanco, 2005 *Post-transcriptional Control of GAP-43 Gene Expression* Kao-chung Francis Tsai, 1996

Transcriptional and Post-transcriptional Control of Therapeutic Gene Expression During Disease Activity Hodan Hassan Ahmed Mohamed, 2015 *Post-transcriptional Control of Gene Expression* Laura Ghisolfi, 2008 *Transcriptional and Post-transcriptional Regulation of Gene Expression* Jun Ding, 2016 Regulation of gene expression includes a variety of mechanisms to increase or decrease specific gene products Gene expression can be regulated at any stage from transcription to post transcription and it is essential to almost all living organisms as it increases the versatility and adaptability by allowing the cell to express the needed proteins In this dissertation we comprehensively studied the gene regulation from both transcriptional and post transcriptional points of view Transcriptional regulation is by which cells regulate the transcription from DNA to RNA thereby directing gene activity Transcriptional factors TFs play a very important role in transcriptional regulation and they are proteins that bind to specific DNA sequences regulatory elements to regulate the gene expression Current studies on TF binding are still very limited and thus it leaves much to be improved on understanding the TF binding mechanism To fill this gap we proposed a variety of computational methods for predicting TF binding elements which have been proved to be more efficient and accurate compared with other existing tools such as DREME and RSAT peaks motif On the other hand studying only the transcriptional gene regulation is not enough for a comprehensive understanding Therefore we also studied the gene regulation at the post transcriptional level MicroRNAs miRNAs are believed to post transcriptionally regulate the expression of thousands of target mRNAs yet the miRNA binding mechanism is still not well understood In this dissertation we explored both the traditional and novel features of miRNA binding and proposed several computational models for miRNA target prediction The developed tools outperformed the traditional microRNA target prediction methods e.g miRanda and TargetScan in terms of prediction accuracy precision recall and time efficiency *Transcriptional and Post-transcriptional Regulation* Kais Ghedira, 2018 This book focuses on the transcriptional and post transcriptional gene regulations and presents a detailed portrait of many novel aspects related to highlighting the importance of key TFs in some vital biological processes the role of certain TFs to control some infectious diseases the role of non coding RNAs in controlling mRNA expression the involvement of these non coding RNAs in diseases and the interplay between TFs and microRNAs as key players for gene expression regulation giving a complete picture of how genes are regulated at the cellular level The editor embarked upon this writing project entitled Transcriptional and Post transcriptional

Regulation to make pertinent contributions accessible to the scientific community Hopefully a large audience will enjoy reading and benefit from the chapters of this book

Deciphering the Gene Expression Control in Epigenetic, Post-transcriptional and Translational Regulation Jianan Lin, 2019 Regulation of gene expression through which cells increase or decrease the gene products is an essential part of development which not only determines the cellular differentiation but also responds to the environmental changes A wide range of gene regulation mechanisms are involved in two major processes from Deoxyribonucleic acid DNA to Ribonucleic acid RNA and from RNA to protein also known as transcription and translation respectively Though the regulation of gene expression is not fully understood this complex process has been characterized to several major steps which includes epigenetic regulation transcriptional regulation post transcriptional regulation and translational regulation The present study covers three main projects related to three of the aforementioned steps of gene regulation First we study the DNA methylation dynamics in the bovine early embryos To understand the epigenetic reprogramming and regulation in the embryonic development we characterize the methylation process at the single base level in early embryos Second we develop a novel method called Protein RNA Association Strength PRAS to predict the functional targets of RNA Binding Proteins RBPs that play important roles in the regulation of gene expression in the post transcriptional process The development of various Cross linking and immunoprecipitation with high throughput sequencing CLIP seq data makes it possible to investigate the transcriptomic binding sites of RBPs We aim to fill the gap between the peak calling methods and the interpretation of RBPs biological functions based on CLIP seq data Third we study the regulation of c MYC on the mRNA translation The oncogenic c MYC MYC transcription factor has broad effects on gene expression and cell behavior We study how MYC affects the global translation of mRNAs and the translation start site usage in the human lymphoma cell line In sum we perform data analysis and methodology development in these three specific projects related to the regulation of gene expression which will help us better understand the central dogma of biology

Post-transcriptional control of copia retrotransposon gene expression Susan J. Thomson, 1996 **Roles of Cofactors in Transcriptional and Post-transcriptional Regulation of Gene Expression** Hongwei Li, 2003 Stochastic Modeling of Post-transcriptional Regulation of Gene Expression in the Limit of Strong Interactions Robert Barr, 2018

Post-transcriptional Regulation of Gene Expression by the Polypyrimidine Tract Binding Protein Kirsty J. Sawicka, 2010

Regulation of Post-transcriptional Gene Expression by CPEB4 and RNF219 Fabian Poetz, 2020 **Computational**

Analysis of Transcriptional and Post-transcriptional Regulation of Gene Expression Andreas J. Gruber, 2017 **Post Transcriptional Regulation of Gene Expression by a Nuclear Polyadenylated RNA Binding Protein** Ronald Earl Hector, 2000 *Transcriptional and Post-transcriptional Regulation* Ghedira Kais, 2018-10-10 This book focuses on the transcriptional and post transcriptional gene regulations and presents a detailed portrait of many novel aspects related to highlighting the importance of key TFs in some vital biological processes the role of certain TFs to control some infectious

diseases the role of non coding RNAs in controlling mRNA expression the involvement of these non coding RNAs in diseases and the interplay between TFs and microRNAs as key players for gene expression regulation giving a complete picture of how genes are regulated at the cellular level The editor embarked upon this writing project entitled Transcriptional and Post transcriptional Regulation to make pertinent contributions accessible to the scientific community Hopefully a large audience will enjoy reading and benefit from the chapters of this book

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