

Myb Transcription Factors: Their Role in Growth, Differentiation and Disease

Edited by
Jon Frampton



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Myb Transcription Factors Their Role In Growth Differentiation And Disease

**Bharti Mackness, Mike
Mackness, Michael Aviram, György
Paragh**

Myb Transcription Factors Their Role In Growth Differentiation And Disease:

Myb Transcription Factors: Their Role in Growth, Differentiation and Disease Jon Frampton, 2004 This volume represents the first collection of articles contributed by research leaders working on the Myb family of transcriptional regulatory proteins In more than twenty chapters the authors discuss the range of biological processes and diverse cell types in which Myb proteins operate Although concentrating on the three vertebrate Myb family members homologues from lower species are also discussed because of the light they are able to shed on the evolution and function of these proteins Individual chapters describe the involvement of Myb proteins in particular c Myb in normal and diseased development and function of many tissues including haemopoietic cells blood vessels the gastrointestinal tract and the brain Several chapters explore the mechanistic details of the action of Myb proteins especially structural features their interaction with DNA and other regulatory proteins and the variety of genes that are regulatory targets for this group of transcription factors This work will be of interest to those working directly in the field and also to the wider research community investigating the transcriptional regulation of development differentiation and growth The therapeutic potential of manipulating Myb function is also discussed making the book appealing to clinician scientists in several fields including haematology oncology and cardiology

Myb Transcription Factors Jon Frampton, 2014-09-01 **Myb Transcription Factors: Their Role in Growth, Differentiation and Disease** Jon Frampton, 2007-10-13 This volume represents the first collection of articles contributed by research leaders working on the Myb family of transcriptional regulatory proteins In more than twenty chapters the authors discuss the range of biological processes and diverse cell types in which Myb proteins operate Although concentrating on the three vertebrate Myb family members homologues from lower species are also discussed because of the light they are able to shed on the evolution and function of these proteins Individual chapters describe the involvement of Myb proteins in particular c Myb in normal and diseased development and function of many tissues including haemopoietic cells blood vessels the gastrointestinal tract and the brain Several chapters explore the mechanistic details of the action of Myb proteins especially structural features their interaction with DNA and other regulatory proteins and the variety of genes that are regulatory targets for this group of transcription factors This work will be of interest to those working directly in the field and also to the wider research community investigating the transcriptional regulation of development differentiation and growth The therapeutic potential of manipulating Myb function is also discussed making the book appealing to clinician scientists in several fields including haematology oncology and cardiology **The Paraoxonases: Their Role in Disease**

Development and Xenobiotic Metabolism Bharti Mackness, Mike Mackness, Michael Aviram, György Paragh, 2007-10-26 In September 2006 the 2nd International Conference on Paraoxonases took place in Hajd szoboszl Hungary bringing together the world s foremost experts in the field The current book is a distillation of the plenary lectures which took place at the meeting resulting in a comprehensive up to date state of the art review of current paraoxonase research The book details a

unique and novel enzyme whose physiological pathological functions are just starting to be revealed **Plant Transcription Factors** Vikas Srivastava, Sonal Mishra, Shakti Mehrotra, Santosh Kumar Upadhyay, 2022-11-09 Plant Transcription Factors Contribution in Development Metabolism and Environmental Stress provides comprehensive coverage of plant TFs and their various functions evaluating their crucial role in growth and development signaling stress management and other key plant processes Sections cover the significance of plant TFs in functional genomics the influence of phyto hormones on the modulation of plant TFs plant development and metabolism including shoot development flowering development and alkaloid biosynthesis The book's final section reviews the role of TFs in various plant stresses including temperature water and heavy metal stress Written by leading experts around the globe this book is an essential read to researchers interested in plant signaling and plant genomics Presents the latest advances in plant transcription factors and their functions Discusses the influence of phyto hormones on the modulation of plant transcription factors Highlights the relationship between plant TFs and plant development *Advances in Environmental Stress Biology and Important Agronomic Traits Improvement in Non-staple Crops* Meng Kou, Kaixuan Zhang, Jian Sun, Ivan Kreft, 2024-05-15 Non staple crops sometimes known as underutilized semidomesticated orphan and or forgotten crops usually refer to under researched grain and legume compared to staple crops such as sweetpotato buckwheat millet barley pea mung beans and adzuki beans which contain unique and beneficial nutrients that staple crops do not have Combining them with staple foods is an important guarantee for a nutrition balanced diet With the deepening of research the current research on non staple crops has gradually started to create a wide range of materials identify varieties and quality improve yield respond to environmental conditions and regulate growth and development Therefore it is an important research objective to improve the important agronomic traits of non staple crops including anthocyanins in sweetpotato rutin in buckwheat vitamins in millet glucan in barley etc through both cultivation and molecular breeding methods and to create new germplasm resources with high yield and abundant nutrients Recently notable successes have been made using genomic related approaches to uncover the genes responsible for important phenotypes in non staple crops The genetic basis of metabolomic divergence and domestication has been revealed in buckwheat and the QTLs for controlling agronomic traits such as flesh color of sweetpotato have been obtained however the function of related genes still needs further investigation In addition biotic and abiotic stresses in extreme climatic conditions change the yield and quality of crops by affecting the growth and development of crops and important metabolic regulation processes Non staple crops are often climate resilient and grown in marginal regions with low input conditions including examples for tolerance of drought stress in cowpea and buckwheat tolerance of heat in cassava and tolerance of barren in sweetpotato Investigating the mechanism of their environmental adaptability would provide new insights for breeding of not only non staple crops but also staple crops that are limited in the tolerance of a changing climate to ensure future food security It is of great theoretical significance and practical application value to

study the molecular regulatory network of non staple crops under these stress conditions Using cultivation measures plant growth regulators fertilizers and other methods to improve the environmental stress resistance and important agronomic traits in non staple crops Revealing molecular mechanisms and regulatory network under all kinds of environmental stresses in non major crops and improving stress tolerance through genetic engineering Identifying key regulatory genes of important agronomic traits in non staple crops and improving molecular breeding methods **NMR Metabolomics in Cancer**

Research Miroslava Čuperlović-Culf,2012-12-17 The application of nuclear magnetic resonance NMR metabolomics in cancer research requires an understanding of the many possibilities that NMR metabolomics can offer as well as of the specific characteristics of the cancer metabolic phenotype and the open questions in cancer research NMR metabolomics in cancer research presents a detailed account of the NMR spectroscopy methods applied to metabolomics mixture analysis along with a discussion of their advantages and disadvantages Following an overview of the potential use of NMR metabolomics in cancer research the book begins with an examination of the cancer metabolic phenotype and experimental methodology before moving on to cover data pre processing and data analysis Chapters in the latter part of the book look at dynamic metabolic profiling biomarker discovery and the application of NMR metabolomics for different types of cancer before a concluding chapter discusses future perspectives in the field Focused description of NMR spectroscopy needed by cancer biologists who are starting to use metabolomics Current overview of knowledge related to the cancer metabolic phenotype from the perspective of metabolomics applications Information about the best practices in NMR metabolomics experimentation and data preprocessing as applied to different sample types **RAS Family GTPases** Channing

Der,2007-04-29 Since 1982 Ras proteins have been the subject of intense research investigation by the biomedical research community The wide interest in Ras has been stimulated for three key reasons This book features chapters contributed by leading investigators in the field that highlight the current state of the art in Ras biochemistry structure and biology This book is an excellent reference for students in the biomedical sciences and for investigators in the field **Role of**

Transcription Factors in Gastrointestinal Malignancies Ganji Purnachandra Nagaraju,Pallaval Veera Bramhachari,2018-03-27 This book illustrates the importance of transcription factors in gastrointestinal cancer progression and metastasis with regard to understanding the mechanism and target definition in drug discovery Further it describes the complex issues associated with cancer cell growth and metastasis The respective chapters provide detailed information on the various types of transcription factors NF κB HIF 1 STAT 3 E2F1 and Sp1 and gastric associated cancers esophagus stomach colorectal liver and pancreatic cancers in connection with specific functional studies like cell cycle angiogenesis migration invasion and apoptosis These transcription factors control the expression of several signaling pathways involved in tumor growth making them ideal targets for gastrointestinal cancer therapy In closing the book provides comprehensive descriptions of the major challenges associated with gastrointestinal cancer therapy **X-ray Crystallographic Studies of**

SNAP190RcRd (Small Nuclear RNA Activating Protein) Complex and E. Coli Glycogen Synthase Fang Sheng,2008

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