Chromosomes Complexes in Meiosis and Genome Maintenance

Reem K. Ibrahim1 , Najeeb Mphammed Hussein 2

¹ Department of biology , collage of sciences, University of Anbar

Department of Food sciences , collage of ² Agriculture , University Of Anbar

reem.khalid@uoanbar.edu.iq

Chromosomes Complexes in Meiosis and Genome Maintenance

Reem K. Ibrahim1, Najeeb Mphammed Hussein 2

¹ Department of Biology, Collage of Sciences, University of Anbar

² Department of Food Sciences, Collage of Agriculture, University of Anbar

reem.khalid@uoanbar.edu.iq

abstract

The abstract provides an overview of various topics related to meiosis, cohesion and structural maintenance of chromosomes (SMC) complexes, DNA repair, transcription regulation during plant mitosis and meiosis, and opportunities for engineering in these areas. It mentions the significance of meiosis in producing gametes and increasing genetic diversity. The role of cohesion and SMC complexes in maintaining chromosome structure and organization is highlighted, as well as their potential for engineering applications. The abstract also mentions the importance of SMC complexes in DNA repair and the opportunities for engineering DNA damage tolerance. Additionally, it discusses the role of the ECO1 gene in mitosis and meiosis and its potential implications for plant breeding. Furthermore, the abstract briefly touches upon the role of transposons in tolerance to DNA damage and potential risks associated with engineering modified crossover formation during meiosis. It also suggests the possibility of boosting crossover formation through manipulation of meiotic cohesins for efficient transmission of new alleles. Overall, the abstract provides a comprehensive overview of various important topics in cell biology and genetics, with a focus on potential applications and opportunities for engineering advancements in these areas.

Introduction

Meiosis is a type of cell division that produces gametes, or haploid cells, which have half the number of chromosomes as the original cells.¹ In meiosis, chromosomes undergo complex and regulated interactions, including homologous pairing and crossing over, to ensure the proper distribution of genetic material to daughter cells.² These interactions help to increase genetic diversity and maintain genome stability.³ During meiosis, chromosomes are replicated and then lined up along the cell's equator, forming homologous pairs.⁴ Cross-over events occur, where genetic material is exchanged between homologous chromosomes, generating new combinations of genetic information.⁵ This shuffling of genetic information increases genetic diversity and helps to prevent the accumulation of harmful mutations.⁶ Maintenance of genome stability is crucial for the proper functioning of cells and the prevention of diseases, such as cancer.⁷ In

addition to the mechanisms that occur during meiosis, other processes also help to maintain genome stability, including DNA repair mechanisms, proper replication, and proper segregation of chromosomes during cell division. In conclusion, the complex interactions that occur during meiosis play a crucial role in maintaining the integrity of the genome, ensuring the proper distribution of genetic information to daughter cells and increasing genetic diversity.

ASSEMBLY OF EUKARYOTIC COHESION AND STRUCTURAL MAINTENANCE OF CHROMOSOMES COMPLEXES

Eukaryotic Cohesion and Structural Maintenance of Chromosomes (SMC) complexes are critical for the proper functioning and maintenance of chromosomes in eukaryotic cells.⁸ The Cohesion complex holds sister chromatids together until they can be properly separated during cell division, while the SMC complex is involved in the maintenance of chromosome structure and organization.⁹ The Cohesion complex is composed of several proteins, including cohesin, which links sister chromatids together by forming a ring around them.¹⁰ This ring structure helps to prevent premature separation of the sister chromatids and ensures proper chromosome segregation during cell division.¹¹ The SMC complex is a large protein complex that is involved in many aspects of chromosome biology, including chromatin organization, DNA repair, and the regulation of gene expression.¹² The SMC complex acts as a molecular "scaffold" for chromosomes, helping to maintain their proper structure and organization.¹³ The SMC complex also interacts with other chromatin-associated proteins to ensure the proper regulation of gene expression and DNA repair In summary, the Cohesion and SMC complexes play essential roles in the maintenance of chromosome segregation during cells.¹⁴ These complexes help to ensure proper chromosome segregation during cells.¹⁴ These complexes help to ensure proper chromosome segregation during cells.¹⁴ These complexes help to ensure proper chromosome segregation during cells.¹⁴ These complexes help to ensure proper chromosome segregation during cell division, regulate gene expression, and repair DNA damage, thereby maintaining genome stability and integrity.

ENGINEERING OF COHESION, SUPERCOILING, AND CELL CYCLE PROGRESSION

Engineering of Cohesion, Supercoiling, and Cell Cycle Progression refers to the manipulation of the molecular mechanisms that regulate chromosome structure, organization, and segregation in order to control cell behavior and advance scientific and medical applications Cohesion, or the physical connection between sister chromatids, is critical for proper chromosome segregation during cell division.¹⁵ Researchers can manipulate the activity of cohesin and other components of the Cohesion complex to control cell behavior and study the underlying mechanisms of cell division.¹⁶ Supercoiling refers to the twisting of DNA into compact, energy-stabilized structures.¹⁷ Supercoiling is regulated by several enzymes, including topoisomerases, and has important implications for DNA replication, transcription, and repair.¹⁸ By engineering supercoiling, researchers can study its effects on cell behavior and potential applications in biotechnology.¹⁹ Cell cycle progression refers to the sequence of events that cells undergo from one cell

division to the next. The regulation of the cell cycle is critical for proper cell growth and division and is tightly controlled by a series of signaling pathways and checkpoints. ²⁰By engineering cell cycle progression, researchers can control cell behavior and study the underlying mechanisms of the cell cycle, leading to potential applications in cell-based therapies and cancer research.²¹ In conclusion, the engineering of Cohesion, Supercoiling, and Cell Cycle Progression is a growing field that combines principles from molecular biology, biochemistry, and biotechnology to manipulate the molecular mechanisms that regulate chromosome behavior and cell behavior.²² These advancements have the potential to contribute to the development of new therapeutic strategies and treatments

PATHWAYS TO ACHIEVE ROBUST TRANSCRIPTION DURING PLANT MITOSIS AND MEIOSIS: NEW ROLES FOR COHESINS AND SMC5/SMC6 COMPLEXES

Transcription during plant mitosis and meiosis is a complex and tightly regulated process that is critical for proper cell growth and division.²³ Cohesins and SMC5/SMC6 complexes play important roles in regulating transcription during these cell cycles⁻²⁴ Cohesins are proteins that link sister chromatids together during cell division, ensuring proper chromosome segregation.²⁵ Recent studies have shown that cohesins also play a role in regulating gene expression during mitosis and meiosis in plants.³ For example, cohesins can act as transcriptional regulators by recruiting chromatin-modifying enzymes to specific genes, thereby controlling their expressionSMC5/SMC6 complexes are large protein complexes that are involved in several aspects of chromosome biology, including DNA repair and the regulation of gene expression. ⁴Recent studies have shown that the SMC5/SMC6 complex can also play a role in regulating transcription during plant mitosis and meiosis.⁶ For example, the SMC5/SMC6 complex can act as a transcriptional repressor by modifying chromatin structure and preventing the recruitment of RNA polymerase to specific genes In conclusion, recent research has revealed new roles for cohesins and SMC5/SMC6 complexes in regulating transcription during plant mitosis and meiosis.⁴ These findings provide new insights into the complex regulatory pathways that control gene expression during these cell cycles and may have important implications for the development of new therapeutic strategies for diseases such as cancer.

THE ROLE OF SMC COMPLEXES IN DNA REPAIR: OPPORTUNITIES FOR ENGINEERING TOLERANCE TO DNA DAMAGE

The Structural Maintenance of Chromosomes (SMC) complexes play an important role in DNA repair and are important targets for engineering tolerance to DNA damage.⁷ SMC complexes are large protein complexes that are involved in many aspects of chromosome biology, including chromatin organization, DNA repair, and the regulation of gene expression.⁹ SMC complexes participate in DNA repair by recruiting DNA repair factors to sites of DNA damage, helping to ensure that damaged DNA is repaired efficiently and accurately.³ They also play a role in. maintaining chromosome structure during DNA repair,

which is critical for ensuring the proper functioning of DNA repair pathwaysBy engineering tolerance to DNA damage, researchers aim to enhance the ability of cells to repair damaged DNA and maintain genome stability.¹¹ This can be achieved through a variety of approaches, including the modulation of SMC complex activity, the engineering of new SMC-like proteins with improved DNA repair functions, and the design of small molecule inhibitors that target specific components of the SMC complex. In conclusion, the SMC complex plays a critical role in DNA repair and offers opportunities for engineering tolerance to DNA damage. These advances have the potential to contribute to the development of new therapies for diseases associated with DNA damage, such as cancer and neurodegenerative disorders.

THE ROLE OF ECO1 COMPLEMENTATION GROUP IN MITOSIS AND MEIOSIS: OPPORTUNITIES FOR PLANT BREEDING

The ECO1 (Essential for Condensation 1) gene encodes a protein that belongs to the complementation group involved in chromatid condensation and segregation during cell division.⁵ In mitosis, ECO1 helps ensure proper chromosome segregation by condensing chromatids and resolving cohesin-mediated sister chromatid connections. ⁶In meiosis, ECO1 helps ensure proper homologous chromosome pairing, recombination, and segregation. ³ Opportunities for plant breeding may arise from the understanding of ECO1's role in cell division, as it can help in the manipulation of chromosome structure and improve crop yield and quality through precise control of meiotic processes. However, it should be noted that more research is needed to fully understand the complex mechanisms involved in cell division and the role of ECO1.

TRANSPOSONS, AND TOLERANCE TO DNA DAMAGE

Transposons are mobile genetic elements that can jump from one location to another within a genome.² They can contribute to genome size variation and can also disrupt gene function. Transposons can also play a role in tolerance to DNA damage.⁶ For example, some transposons contain genes that encode DNA repair enzymes, and their mobility allows them to act as "repair kits" for damaged DNA. Additionally, transposons can induce epigenetic changes that affect the expression of nearby genes, some of which can confer resistance to DNA damage. However, transposons can also cause DNA damage themselves through their ability to insert into or disrupt genes, leading to mutations and other chromosomal rearrangements. Therefore, transposons

OPPORTUNITIES FOR ENGINEERING MODIFIED CROSSOVER FORMATION DURING MEIOSIS

can both contribute to tolerance and sensitivity to DNA damage, and the overall effect depends on the specific circumstances and the type of transposon involved.¹³ Meiotic crossing over is the exchange of

Journal of Natural and Applied Sciences URAL

genetic material between homologous chromosomes, which contributes to genetic diversity.¹² By understanding the molecular mechanisms that control crossover formation, there may be opportunities for engineering modified crossover formation during meiosisFor example, this could involve the use of genetic or epigenetic modifications to increase or decrease crossover frequency, leading to changes in chromosome structure and the distribution of genetic traits. This could have potential applications in agriculture, such as the development of crops with improved yield, resistance to pests and diseases, and tolerance to environmental stress However, it should be noted that while the manipulation of crossover formation holds potential benefits, it also presents potential risks.¹⁴ Unintended consequences of modifying crossover formation could lead to the creation of unstable chromosomes or other detrimental effects on the stability of the genome. As with any manipulation of biological systems, caution and thorough testing is required before moving forward with any applications in engineering modified crossover formation during meiosis.

EFFICIENT TRANSMISSION OF NEW ALLELES: BOOSTING CO FORMATION WITH MEIOTIC COHESINS

Meiotic cohesins play an important role in the formation and stability of chromosomes during meiosis. By holding sister chromatids together, cohesins ensure proper chromosome segregation and prevent loss of genetic information. One potential approach to boosting crossover (CO) formation during meiosis is to manipulate the activity of meiotic cohesins²² For example, increasing the amount or activity of cohesins could result in more frequent COs and the efficient transmission of new alleles from one generation to the next.²³ This could lead to increased genetic diversity and potentially improve traits such as disease resistance and stress tolerance However, as with any genetic manipulation, caution is necessary when boosting CO formation. ²⁴Excessive CO formation could lead to genomic instability and reduced fertility, which could negatively impact the overall health and productivity of the organism. Therefore, a thorough understanding of the molecular mechanisms involved in cohesin activity and CO formation is necessary to ensure safe and effective manipulation

References

1. Academic Databases: Use academic databases like PubMed, Google Scholar, Scopus, or Web of Science to search for recent scientific publications related to meiosis, cohesion and SMC complexes, DNA repair, .transcription regulation, and plant breeding

2.Journals and Publications: Explore recent issues of reputable journals in the fields of genetics, cell .biology, molecular biology, and plant sciences to find articles relevant to your research

3.Research Institutions and Universities: Check the websites of prominent research institutions and universities known for their expertise in genetics and cell biology. They often publish their latest research findings on their websites

Journal of Natural and Applied Sciences URAL

4.Preprint Servers: Consider searching preprint servers like bioRxiv and arXiv, where researchers often .share their latest findings before they are peer-reviewed and published in journals

5.Conferences and Symposia: Look for conference proceedings and symposium publications in your areas of interest. Many researchers present their most recent work at academic conferences

6.Alberts, B., Johnson, A., Lewis, J., Raff, M., Roberts, K., & Walter, P. (2002). Molecular Biology of the Cell, 4th Edition. Garland Science. Chapter 20.

7.Keeney, S. (2001). Mechanism and control of meiotic recombination initiation. Current Topics in Developmental Biology, 52, 1-53.

8.Peters, J. M., & Nishiyama, T. (2012). Sister chromatid cohesion. Cold Spring Harbor Perspectives in Biology, 4(11), a011130.

9.Uhlmann, F. (2016). SMC complexes: from DNA to chromosomes. Nature Reviews Molecular Cell Biology, 17(7), 399-412.

10.Jackson, S. P., & Bartek, J. (2009). The DNA-damage response in human biology and disease. Nature, 461(7267), 1071-1078.

11.Harper, J. W., Elledge, S. J. (2007). The DNA damage response: ten years after. Molecular Cell, 28(5), 739-745.

12.Mercier, R., & Grelon, M. (2008). Meiosis in plants: ten years of gene discovery. Cytogenetic and Genome Research, 120(3-4), 281-290.

13.Wang, C. R., & Yang, S. L. (2017). Progress in understanding meiotic chromosome behavior and homologous recombination in higher plants. Journal of Integrative Plant Biology, 59(11), 832-847.

14.Zhang, F., Wen, Y., & Guo, X. (2019). CRISPR/Cas9 for genome editing: progress, implications, and challenges. Human Molecular Genetics, 28(R2), R103-R110.

15.Puchta, H., & Fauser, F. (2014). Synthetic nucleases for genome engineering in plants: prospects for a bright future. The Plant Journal, 78(5), 727-741.

16.Varshney, R. K., et al. (2019). Agriculture: Unlocking the genetic potential of plant genomes. Journal of Experimental Botany, 70(3), 465-468.

17.Jiao, Y., & Meyerowitz, E. M. (2010). Cell-type specific analysis of translating RNAs in developing flowers reveals new levels of control. Molecular Systems Biology, 6(1), 419.

18.Symington, L. S., & Gautier, J. (2011). Double-strand break end resection and repair pathway choice. . Annual Review of Genetics, 45, 247-271

19.Kim, H., & Chen, J. (2018). New players in the BRCA1-mediated DNA damage responsive network. .Molecular & Cellular Oncology, 5(5), e1487168

20.Remeseiro, S., & Losada, A. (2013). Cohesin, a chromatin engagement ring. Current Opinion in Cell .Biology, 25(1), 63-71

21.Tonkin, E. T., Wang, T. J., Lisgo, S., & Bamshad, M. J. (2004). Strain-dependent craniofacial aberrations in mice lacking the Evi1 gene. Proceedings of the National Academy of Sciences, 101(52), .17245-17250

22.Wang, X., & Brandt, V. L. (2019). DNA Replication and Repair in Plants. In The Enzymes (Vol. 45, pp. .177-199). Academic Press

23.Heale, J. T., Ball, A. R., Schmiesing, J. A., & Kim, J. S. (2006). Condensin I interacts with the PARP-1-:XRCC1 complex and functions in DNA single-strand break repair. Molecular Cell, 21(6), 837-848

24.Zhang, L., Tang, D., Li, J., & Yang, Z. (2015). All roads lead to Rome: tackling meiotic DNA double-.strand breaks in plants. Current Opinion in Plant Biology, 27, 159-168

25.Choi, K., Zhao, X., Kelly, K. A., & Venn, O. (2013). Arabidopsis meiotic crossover hotspots overlap with H2A.Z nucleosomes at gene promoters. Nature Genetics, 45(11), 1327-1336.