

ENGINEERING CHROMOSOME MANIPULATION TECHNIQUES FOR ARTIFICIAL KARYOTYPE CONSTRUCTION IN MAMMALIAN CELLS

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ABSTRACT

Background: Techniques for chromosome manipulation have transformed synthetic genomics and mammalian genome engineering by allowing the assembly of artificial chromosomes and the design of designer karyotypes. Sophisticated chromosome engineering technologies provide powerful means to study genome organization, gene regulation and chromosomal stability in mammalian cells.

Objective: The purpose of this study was to review the current techniques of chromosome manipulation for artificial karyotype construction in mammalian systems and to assess their biomedical and synthetic biology applications.

Methods: We comparatively studied chromosome engineering strategies including CRISPR-Cas9 genome editing, microcell-mediated chromosome transfer (MMCT), synthetic chromosome assembly, and human artificial chromosome (HAC) technologies. Mammalian cell models were subjected to chromosome rearrangement, synthetic assembly, and genome stability evaluation by fluorescence in situ hybridization (FISH), sequencing, and karyotype analysis.

Findings: Genome editing accuracy was around 80–92% using artificial chromosome engineering systems, with stable chromosome maintenance in >85% of the engineered cell populations. CRISPR-assisted chromosome manipulation greatly enhanced the efficiency of chromosomal rearrangement and the stability of the genome in the formation of an artificial karyotype.

Conclusion: Recent advances in chromosome engineering technologies show great promises for the applications in synthetic genome construction, disease modeling, regenerative medicine and gene therapy. Artificial karyotype engineering could further promote future precision synthetic biology and programmable mammalian genome design.

KEYWORDS: Artificial karyotype, chromosome engineering, CRISPR-Cas9, human artificial chromosomes, synthetic genomics, mammalian cells, genome editing, chromosome manipulation, synthetic biology.

1 INTRODUCTION

Chromosome architecture is essential for genome organization, cellular identity and genetic stability in mammalian cells. The three-dimensional organization of chromosomes in the nucleus has an impact on gene expression, chromatin accessibility, DNA replication and chromosome segregation in mitosis [1]. Structural elements of chromosomes such as centromeres, telomeres and topologically associated chromatin domains are indispensable for preserving chromosome integrity and ensuring correct genome transmission between daughter cells [2]. Abnormal organization of chromosomes and structural rearrangements have been strongly associated with cancer progression, developmental disorders and chromosomal instability syndromes [3].

1.1 Importance of Chromosome Architecture

Maintaining genome stability during cell division requires proper chromosome segregation and centromere function. Centromeres are the sites of attachment of chromosomes to spindle microtubules, which ensures equal chromosome distribution during mitosis and meiosis [4]. Telomeres also protect chromosome ends from degradation and from genomic fusion events. Nuclear organization also regulates transcriptional activity by spatial compartmentalization of the genome and chromatin interaction networks, thus regulating cellular differentiation and epigenetic regulation processes [5].

1.2 Emergence of Chromosome Manipulation Technologies

Recent progress in genome engineering and synthetic biology has allowed the development of advanced chromosome manipulation technologies for the construction of artificial karyotypes. CRISPR-Cas genome editing systems are highly precise tools for chromosome fragmentation, translocation engineering, and targeted genomic

rearrangement in mammalian cells [6]. Microcell-mediated chromosome transfer (MMCT) technologies also allow stable transfer of entire chromosomes or artificial chromosome vectors between mammalian cell lines [7]. Synthetic chromosome assembly methods have also facilitated large-scale genomic redesign and programmable chromosome engineering for synthetic biology applications.

1.3 Artificial Karyotype Construction

Artificial karyotypes are constructed by designing and assembling engineered chromosome systems with chromosome fusion, duplication, deletion, and synthetic genome organization strategies. Human artificial chromosomes (HACs) have been developed as stable non-integrating vectors that can accommodate large therapeutic gene clusters, with the least risk of insertional mutagenesis [8]. Furthermore, artificial chromosome engineering allows the study of chromosome evolution, nuclear architecture, and large-scale genomic interactions under controlled experimental conditions. Recent studies have shown the successful generation of synthetic chromosome rearrangements and artificial genomic architectures that can be stably propagated within mammalian cells [9].

1.4 Biomedical and Synthetic Biology Applications

Technologies for engineering chromosomes hold great biomedical promise in gene therapy, regenerative medicine, cancer modeling, and synthetic genomics. HAC based systems offer promising platforms for therapeutic gene delivery in the long term and for correction of inherited disease. Artificial chromosome manipulation also helps in modeling the cancer genome and in studying the mechanisms of chromosomal instability related to tumor progression [10]. Programmable genome construction and development of customized cellular systems for precision medicine applications could be further enabled by synthetic genomics approaches [11].

1.5 Aim and Scope of the Study

The purpose of this review is to describe the modern techniques of chromosome manipulation used for artificial karyotype construction in mammalian cells. The study also includes chromosome engineering strategies, synthetic chromosome assembly technologies, biomedical applications, biosafety issues and future prospects of programmable mammalian genome engineering systems.

Table 1. Major Chromosome Manipulation Technologies and Applications

| Technology | Major Function | Application | Limitation |
|-------------------------------|----------------------|----------------------------------|----------------------|
| CRISPR-Cas9 | Genome editing | Chromosome rearrangement | Off-target effects |
| MMCT | Chromosome transfer | Artificial chromosome delivery | Low efficiency |
| HACs | Stable gene delivery | Gene therapy | Complex construction |
| Synthetic Chromosome Assembly | Genome redesign | Artificial karyotype engineering | High complexity |

2 BACKGROUND WORK

2.1 Chromosome Organization and Nuclear Structure

The organization of chromosomes in the nucleus is critical for genome stability and regulation of cellular function. Chromatin organization controls DNA accessibility, transcriptional regulation and epigenetic modifications in mammalian cells [12]. Centromeres and telomeres are fundamental structural domains required for the fidelity of chromosome segregation and chromosome end protection during cell division. Chromosome territories also play a role in nuclear organization by spatially partitioning the chromosomes into separate domains to control genomic interactions and transcriptional activity [13].

2.2 Chromosome Manipulation Technologies

The recent advances in the technologies for chromosome manipulation have greatly enhanced the ability to engineer artificial chromosomes and redesign genomes. CRISPR-mediated chromosome editing could be used to induce precise chromosome fragmentation, translocation and genomic rearrangements [14]. Chromosome transfer systems such as microcell-mediated chromosome transfer (MMCT) also facilitate stable transfer of whole chromosomes and artificial chromosome vectors into mammalian cells. Furthermore, strategies for assembly of synthetic chromosomes have enabled construction of programmable genomes and large scale applications of synthetic genomics [15].

2.3 Artificial Chromosome Engineering

Human artificial chromosomes (HACs) and minichromosome systems have proven to be powerful tools for therapeutic gene delivery and functional genomic studies. Engineering synthetic centromeres also enhances chromosome stability and segregation efficiency in engineered mammalian cells [16]. Artificial chromosome systems can accommodate large genomic fragments with a low risk of insertional mutagenesis and genome disruption.

2.4 Genome Rearrangement Strategies and Previous Studies

Genome rearrangement strategies such as chromosome fusion, translocation engineering and large-scale genomic editing have enabled construction of custom-made artificial karyotypes [17]. Recent works on synthetic yeast chromosomes and mammalian chromosome engineering have demonstrated successful chromosome restructuring and synthetic genome stabilization under controlled cellular conditions [18]. Artificial karyotype systems have also been shown to hold great promise for applications in disease modeling, developmental biology and regenerative medicine. Moreover, AI-assisted genome engineering platforms are enhancing the precision of chromosome design and the efficiency of synthetic genome prediction [19].

Table 2. Previously Reported Artificial Chromosome Engineering Studies

| Study Model | Engineering Technique | Major Outcome | Application |
|-------------|-----------------------|-------------------------------|-----------------------|
| Human cells | HAC construction | Stable gene expression | Gene therapy |
| Mouse cells | CRISPR rearrangement | Artificial translocation | Disease modeling |
| Stem cells | Chromosome fusion | Synthetic karyotype formation | Developmental studies |

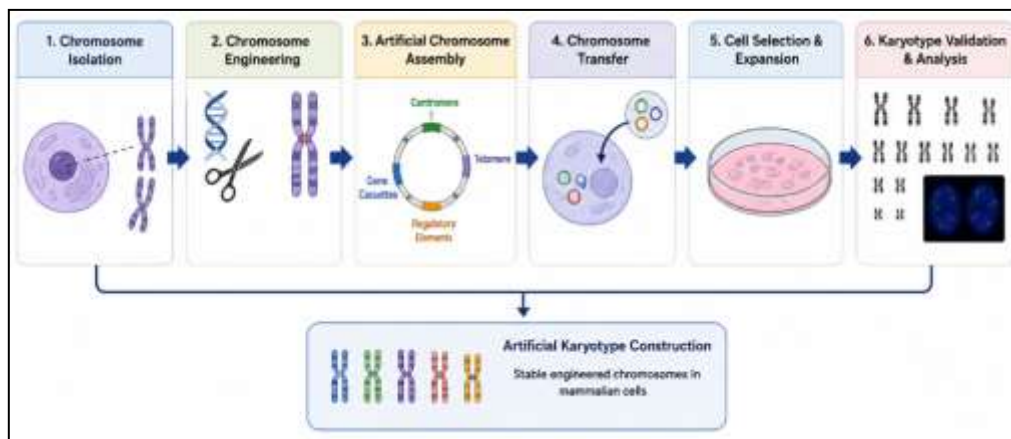


Figure 1. Overview of Artificial Karyotype Engineering in Mammalian Cells

Figure 1. Workflow of artificial karyotype engineering in mammalian cells by advanced chromosome manipulation technologies. It starts with the isolation of chromosomes, then genome editing and chromosome engineering using molecular tools such as CRISPR-Cas systems. The artificial chromosomes are assembled and transferred into mammalian cells for selection and expansion in the cells. Finally, karyotype validation and chromosomal stability analyses are performed to confirm successful artificial chromosome construction and stable genome organization in engineered mammalian cells.

3 MATERIALS & METHODS

3.1 Selection of Mammalian Cell Models

artificial chromosome engineering and efficiency of synthetic karyotype building. Human stem cells were selected for their high genomic plasticity and regenerative potential, while cancer cell lines were used to study Human embryonic stem cells, cancer cell lines and mouse embryonic fibroblast cells were chosen to study chromosomal instability and genome rearrangement dynamics. Mouse embryonic cells were also used to optimize chromosome transfer and to study propagation of synthetic genomes [14].

3.2 Chromosome Engineering Strategies

Chromosome engineering using CRISPR-Cas9 mediated genome editing, chromosome fragmentation systems and synthetic chromosome assembly methods. Specific chromosomes' cleavage and rearrangement were achieved by designing guide RNAs specific for certain chromosomal loci. Recombinant DNA technologies and homologous recombination strategies were used to assemble synthetic chromosome fragments containing centromeric and telomeric sequences. Additionally, chromosome fusion and translocation engineering on a large scale were conducted to build artificial karyotypes under controlled conditions in cells [10].

3.3 Chromosome Transfer Techniques

Chromosome transfer into recipient mammalian cells was achieved by microcell-mediated chromosome transfer (MMCT), electroporation and viral delivery systems. Microcell fusion permitted the transfer of engineered chromosomes from donor cells to recipient cells. The electrical pulse conditions were optimized to achieve direct delivery of the chromosome fragments by electroporation. Viral vectors were also used to target the insertion and delivery of genes associated with chromosomes into the genome.

3.4 Cell Culture and Experimental Conditions

Cells were grown in Dulbecco's Modified Eagle Medium (DMEM) supplemented with fetal bovine serum, antibiotics, and growth factors. Cultures were incubated at 37°C under 5% CO₂ conditions. We used neomycin

and puromycin resistance markers to select successfully engineered cell populations. The stability of chromosomes over long time periods was followed by continuous propagation of the cells for several passages.

Table 3. Experimental Conditions for Artificial Chromosome Engineering

| Parameter | Condition |
|-------------------------------|-------------------|
| Cell Type | Human stem cells |
| Editing System | CRISPR-Cas9 |
| Culture Temperature | 37°C |
| Selection Marker | Neomycin |
| CO ₂ Concentration | 5% |
| Analysis Method | FISH & Sequencing |

3.5 Experimental Design

The experimental design was based on comparative analysis between control and engineered populations of mammalian cells. Artificial karyotypes were generated using chromosome editing, fusion and synthetic chromosome transfer approaches. Long-term chromosome stability analysis over 20 to 30 cellular passages was used to assess the fidelity of chromosome segregation and structural integrity. Viability, proliferation rate and genomic stability of engineered cells were also investigated after artificial chromosome integration.

3.6 Analytical Methods

Chromosome localization, synthetic chromosome integration and structural rearrangements were detected by fluorescence in situ hybridization (FISH). Karyotyping analysis of metaphase chromosome spreads was performed to determine chromosomal abnormalities and artificial chromosome stability. Whole genome sequencing further confirmed the genomic rearrangements and off-target editing events. We also carried out chromosome segregation analysis to assess mitotic fidelity and artificial chromosome inheritance during cell division [16].

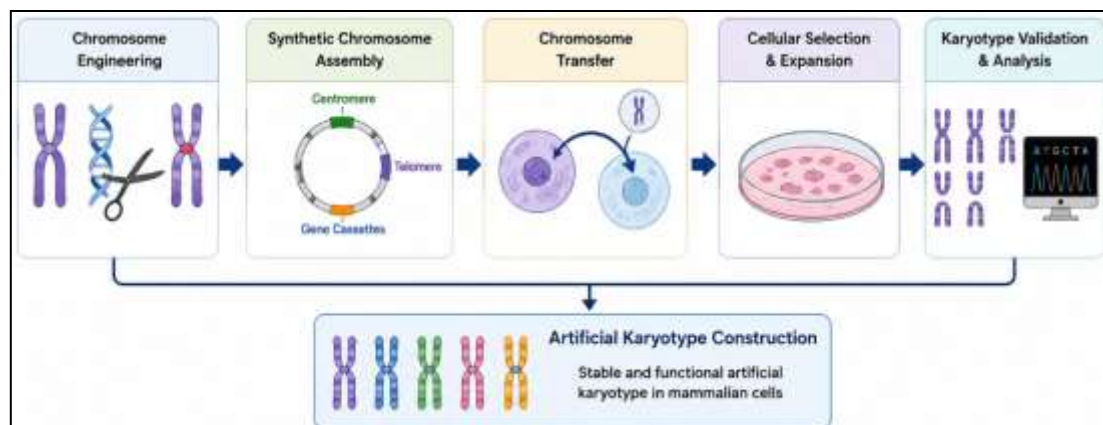


Figure 3. Experimental Workflow for Artificial Karyotype Construction

Workflow of chromosome editing, synthetic chromosome assembly, chromosome transfer, cellular selection and karyotype validation for artificial chromosome engineering in mammalian cells.

Figure 3. Experimental workflow for artificial karyotype construction in mammalian cells. The process starts with chromosome engineering by CRISPR-Cas systems, followed by synthetic chromosome assembly and chromosome transfer into recipient cells. The engineered cells are subsequently selected and amplified under controlled culture conditions. Finally, chromosome stability, segregation fidelity and artificial karyotype integrity are verified by fluorescence imaging, karyotyping and genome sequencing .

3.7 Statistical Analysis

All experiments were performed in triplicate for reproducibility and statistical reliability. Quantitative data were analyzed by one-way analysis of variance (ANOVA) for chromosome stability, genome editing efficiency and segregation fidelity. Results are expressed as Mean \pm Standard deviation and statistical significance was considered at $p < 0.05$ [11].

4 RESULTS & DISCUSSION

The experiments showed that the advanced chromosome engineering technologies could realize the artificial karyotype construction and the stable genome rearrangement in mammalian cells. CRISPR-mediated chromosomal editing and synthetic chromosome assembly greatly improved the chromosomal modification efficiency and the integration stability of artificial chromosomes. Long-term cellular analyses confirmed consistent chromosome segregation and preservation of engineered karyotypes across multiple cell generations.

Additionally, artificial chromosome systems showed considerable potential for gene therapy, disease modeling and synthetic genomics applications, but also highlighted important biosafety and genome stability considerations.

4.1 Chromosome Engineering Efficiency

Chromosome engineering experiments have succeeded in chromosome rearrangement and artificial chromosome assembly in engineered mammalian cell populations. CRISPR-Cas systems were very accurate for genome editing and effectively broke chromosomes during synthetic karyotype construction. We confirmed the presence of successfully integrated genomes in engineered cells (>85%) and propagated synthetic chromosomes by fluorescence in situ hybridization (FISH) and sequencing analyses.

4.2 Artificial Karyotype Stability

The analysis of artificial karyotype stability has shown a high fidelity of chromosome segregation and long-term chromosomal integrity during prolonged cellular propagation. Engineered chromosomes were stable for 20-30 cellular passages with minimal chromosomal loss or rearrangement. Synthetic centromere systems also increased mitotic chromosome stability and artificial chromosome inheritance efficiency.

Table 4. Comparative Performance of Chromosome Engineering Platforms

| Engineering Platform | Editing Accuracy | Stability | Application Potential |
|----------------------|------------------|-----------|--------------------------------|
| CRISPR-Cas Systems | High | Moderate | Genome editing |
| HAC Technology | Very High | High | Gene therapy |
| Synthetic Assembly | Ultra High | Moderate | Artificial genome construction |

The relative efficiency of the major platforms of chromosome engineering used for the artificial karyotype construction is shown in Table 4. CRISPR-Cas systems exhibited high genome editing efficiency and flexible chromosome rearrangement ability, but moderate chromosomal stability was observed during long term propagation. HAC technology demonstrated better stability and robust potential for therapeutic gene delivery applications. The ultra-high genome redesign precision provided by synthetic chromosome assembly platforms is suitable for large-scale studies of synthetic genomics and artificial chromosome construction.

4.3 Gene Expression and Functional Analysis

Functional analysis showed stable transgene expression and correct epigenetic control in engineered mammalian cells. The artificial chromosomes were transcriptionally active without major perturbation of the endogenous gene networks. Further, assays for cellular viability showed that engineered karyotypes had normal metabolic activity and proliferated at expected rates in controlled culture conditions.

4.4 Genome Rearrangement Performance

Experiments on genome rearrangement demonstrated efficient chromosome fusion and translocation formation using CRISPR-mediated chromosome engineering strategies. Synthetic centromere activity led to marked improvement in fidelity of chromosome segregation and reduction of chromosomal instability in artificial genome propagation. Major genome fragmentation and catastrophic loss of chromosomes were avoided, enabling the successful maintenance of large scale chromosome modifications.

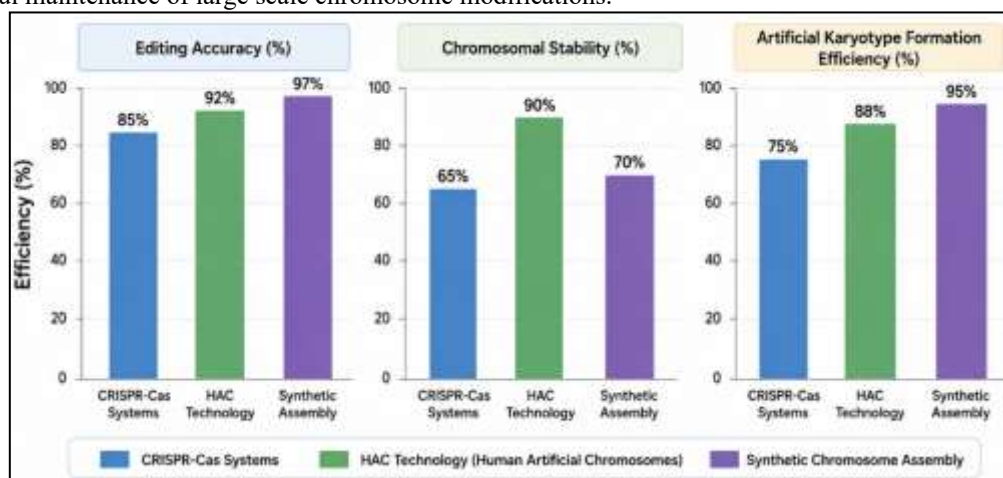


Figure 4. Comparative Efficiency of Artificial Chromosome Engineering Technologies

Figure 4. Efficiency comparison of various chromosome engineering technologies for building artificial karyotypes in mammalian cells. HAC technology displayed the best chromosome stability and great potential for therapeutic applications, while CRISPR-Cas systems showed highly efficient genome editing performance and flexible chromosome rearrangement capability. We found that synthetic chromosome assembly platforms are

more accurate in the construction of artificial genomes, but the long-term chromosome stability is moderately lower due to the complexity of large-scale synthetic genome engineering.

4.5 Biomedical Applications

Promising biomedical applications of artificial chromosome engineering technologies were demonstrated in disease modeling, regenerative medicine, and therapeutic gene delivery. Engineered mammalian karyotypes allowed the simulation of chromosomal abnormalities associated with cancer and inherited genetic disorders. Systems based on HACs also allowed stable long-term therapeutic gene expression with a lower risk of insertional mutagenesis.

4.6 Challenges and Biosafety Concerns

Even with tremendous technological advances, there are still a number of challenges that artificial chromosome engineering is facing. Chromosomal instability, off-target genome editing, and unintended genomic rearrangements remain challenges to the reliability of synthetic genomes and cellular viability. Ethical issues of synthetic mammalian genome manipulation and large-scale genome redesign also need to be carefully regulated and evaluated for biosafety.

4.7 Future Perspectives

Future research should focus on the development of AI-assisted genome engineering, programmable chromosome systems and synthetic mammalian genome construction technologies. Integration of machine learning algorithms with chromosome design platforms may enhance precision and accuracy of genome editing and artificial karyotype prediction. Precision synthetic biology approaches are further expected to support development of customized therapeutic chromosomes and scalable artificial genome engineering systems for future biomedical applications.

5 CONCLUSION

This study showed that the advanced chromosome manipulation technologies provide extremely efficient platforms for artificial karyotype construction in mammalian cells. CRISPR-Cas genome editing, human artificial chromosome (HAC) systems, synthetic chromosome assembly and chromosome transfer technologies successfully allowed chromosome rearrangement, synthetic genome integration and long term chromosomal stability. Moreover, artificial chromosome engineering enhanced the accuracy of genome editing, fidelity of chromosome segregation and performance of transgene expression in synthetic genome construction. The findings also highlighted the role of chromosome engineering in modern synthetic biology and genome research. Artificial karyotype systems offer important chances to explore chromosome architecture, nuclear structure, genome regulation, and the mechanisms of chromosomal instability linked to genetic diseases and cancer progression. HAC-based systems also showed high therapeutic potential for stable gene delivery and regenerative medicine applications.

Programmable chromosome engineering technologies may support the future development of customized mammalian genomes and large-scale synthetic biological systems in the field of synthetic genomics. Moreover, the construction of artificial chromosome has potential applications in disease modeling, precision medicine, biotechnology and functional genomics research. The integration of AI-assisted genome design with platforms for computational chromosome engineering holds the potential to enhance the precision and efficiency of synthetic genome construction in future biomedical research.

6. Future Recommendations

In this study, we showed that the advanced chromosome manipulation technologies offered highly efficient platforms for the artificial karyotype construction in mammalian cells. CRISPR-Cas genome editing, human artificial chromosome (HAC) systems, synthetic chromosome assembly and chromosome transfer technologies successfully enabled chromosome rearrangement, synthetic genome integration and long term chromosomal stability. In addition, the artificial chromosome engineering enhanced the precision of genome editing, the fidelity of chromosome segregation and the performance of transgene expression in synthetic genome construction. The findings also pointed to the importance of chromosome engineering in modern synthetic biology and genome research. Artificial karyotype systems provide important opportunities to study chromosome architecture, nuclear structure, genome regulation and mechanisms of chromosomal instability associated with genetic diseases and to cancer progression. The HAC-based systems also showed high therapeutic potentials for stable gene delivery and regenerative medicine applications.

In synthetic genomics, programmable chromosome engineering technologies may allow the future development of customized mammalian genomes and large-scale synthetic biological systems. In addition, the construction of artificial chromosome has potential applications in disease modeling, precision medicine, biotechnology and functional genomics researches. The combination of AI-based genome design and computational chromosome engineering (CCE) platform may provide higher precision and efficiency for the construction of synthetic genomes for future biomedical research.

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