



(REVIEW ARTICLE)



## Proteins in the kinetochore generate forces to drive chromosomes movement

Amer Alasadi <sup>1,\*</sup>, Basim Turki Alyousif <sup>2</sup> and Noor Fadhil <sup>3</sup>

<sup>1</sup> Department of Medical Basic Science, College of Nursing, University of Thi-Qar- Iraq.

<sup>2</sup> Marshes Research Center, University of Thi-Qar- Iraq.

<sup>3</sup> Al Karama Teaching hospital, Wassit Health Office, Iraq.

GSC Biological and Pharmaceutical Sciences, 2022, 20(02), 046–051

Publication history: Received on 26 June 2022; revised on 31 July 2022; accepted on 02 August 2022

Article DOI: <https://doi.org/10.30574/gscbps.2022.20.2.0313>

### Abstract

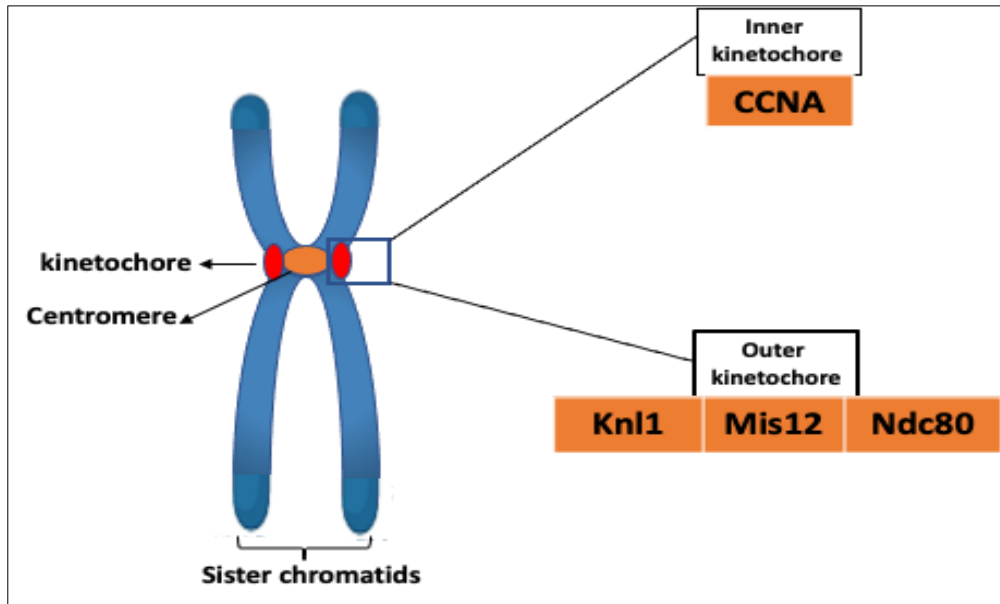
To accomplish the precise segregation of chromosomes, the kinetochore must form incentive interaction with motor proteins. Any error happens in chromosome segregation led to aneuploidy that consider as main factor to cause the cancers and many birth defects. Kinetochore is a proteinous complex structure consist of a hundred of proteins. It's performed many functions such as it connects movement of chromosomes to microtubule, it acts as observer for chromosomes bioorientation, beside the main role in chromosomes segregation. In this review we described the proteins content of kinetochore and the important role of these proteins in generating forces to drive the movement of chromosomes, also we clarified some experiments that used to identify these proteins and support their functions.

**Keywords:** Kinetochore; Chromosomes; Aneuploidy; KMN; FRAP

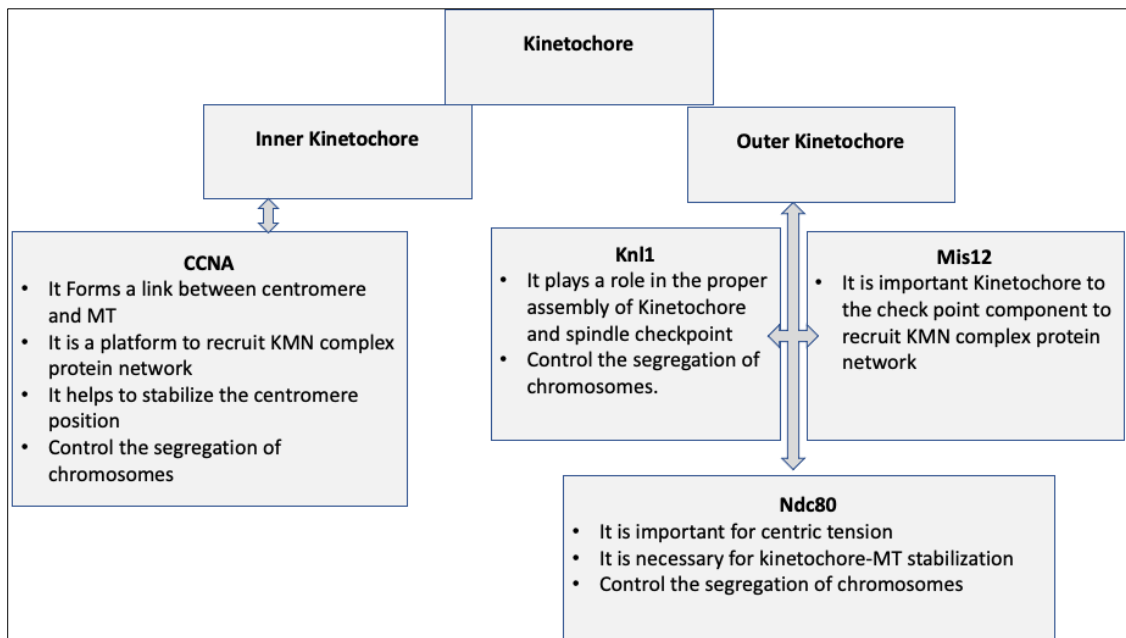
### 1. Introduction

Kinetochore is multi protein's structure which forms several connections between the centromere and kinetochore microtubules. The inner kinetochore contains an essential protein that are existed in the centromere called constitutive centromere-associated network (CCAN) which present through cell cycle, while the outer kinetochore proteins assemble at beginning of prophase and leave at the end of mitosis [1]. The outer kinetochore comprises from three complexes (Knl1, Misl2 and Ndc80), which all these named KMN network as indicated in Figure 1. These proteins create a stable attachment site with microtubule, and they have a controlling role in the polymerization state of kinetochore microtubule (KMT) as described here in Figure 2 [2].

\* Corresponding author: Amer Alasadi  
Department of Medical Basic Science, College of Nursing, University of Thi-Qar- Iraq.



**Figure 1** The structure of the kinetochore proteins. The schematic diagram illustrates the structure of the chromosome and the proteins structure of kinetochore



**Figure 2** Functions of kinetochore proteins. The above graphic figure shows the functions of each protein in the kinetochore protein complex

## 2. Forces to drive the chromosome movement

Three main forces are believed to be an important to drive the chromosome movement. The first forces come from kinetochore proteins that consider the major forces. It pulls KT and associated chromosome to the spindle pole; KT moves poleward when the depolarization happen at the plus end of MT. The second forces generate by MT flux and its present in some types of cells. The polar injection is the third forces, and the chromosome movement occurs when interpolar MT interacts with Kinesin family that are present in the chromosome arm [3].

Kinetochores generate the forces by three ways, either by coupling of KT to depolarized MT, or from the tugging action of minus end motor protein (dynein) and, from the motor enzymes have an important role in tethering KT to MT [4]. Additionally, studies in *Drosophila* proposed the reduction of KT dynein this will lead to reduce chromosomal movement rate at anaphase. The reduction process in the *Drosophila* was happened by two ways, the first way by making mutation in *Drosophila* ZW10 and rod to disrupt RZZ components and the second way by inhibition cytoplasmic dynein via using p50 or by anti DHC antibody, also the confocal microscopy was showed the KTs alignment were not fully completed and unattached and set outside the mitotic spindle in the absences of the dynein [5-7].

### 3. Approaches to identify the protein contents of kinetochore

Three major kinetochore proteins were identified by different approaches as summarized in the Table 1, using human antibodies, the combination of genetic, RNA interference (RNAi) and biochemical methods were used to identify others kinetochore proteins that have an important role in chromosome segregation [8-13]. In last decade a large of kinetochore proteins were identified by using mass spectrometry in model organisms [14]. Other researchers used the Delta Assay to investigate the subunits that formed the Ndc80 complex, which is consider the main MT binding components of outer kinetochore. The results depended on the average of separation between two kinetochores inner and outer [15&16].

Additionally, a new technique named Elegant Fluorescence Microscopy used for showing the whole composition of kinetochore and it concluded the kinetochore is a tri-complex assembly and outer kinetochore has an attachment site with microtubule [17]. In addition, some researchers were used the mitotic arrest cells for identification some particles in kinetochore that have ability to bind with MT via multiple binding site [18].

There are many an important protein that play an impressive role in cell cycle events including EBI, CLIP170, and MAP215. These are plus end direct proteins and have a controlling role in the polymerization state of KT-MT attachment site and their functions are accomplished in budding yeast by Dam1 complex [19]. EBI has a binding and stabilizing role in MT seam [20]. CLIP170 helps in creating the primary connection between MT and kinetochore [21]. MAP250 acts to enhance the MT polymerization that is necessary for chromosome segregation [22&23].

Recently, GFP-tagged fusion model used to identify a new protein in human kinetochore named Ska1. Ska1 plays a several roles in coupling of chromosome movement, depolarization of MT and chromosome segregation [24&25]. Other researchers were used high pressure freezing method to study kinetochore proteins depend on preparing the samples for electron microscopy and it was better than chemical fixation because it has ability to preserve ultrastructure of the samples. This study proved the poleward chromosome movement result due to forces that arise in kinetochore [26&27].

Fluorescent Recovery after Photo bleaching (FRAP) technique was applied to investigate the dynamic property of MT by observing a bleached spot made on the lattice of fluorescent MT of Integral spindle. The searchers showed in the epithelial cells of kidney the photo bleached section stayed fixed on fluorescent MT of KT while the chromosome continued and crossed the bleached region, and these events proposed the MT depolarization happens at the KT [28]. Another study was revealed the KT contains a small members of motor proteins such as dynein, ZW10 and CENP-E and these motor proteins have an important role in chromosome movement to poleward [29 & 30].

Cells free approach were used to detect the role of the Ndc80 and dynein/dyactin, the inhibition of Ndc80 complex lead to delay dynein/dyactin rapid expression through prometaphase that depend on the KT poleward movement. Thus, dynein/dyactin movement continue without completing chromosome alignment and segregation at anaphase because the Ndc80 complex was absent [31]. A recent study showed the Ndc80 complex in the KT may act as a forcer transducer and it's stretched along the KT axis [32]. Similarly, the researchers described the Dam1 complex in budding yeast and their results clarified Dam1 complex role in the enhance the Ndc80 ability to mediate KT-MT attachment [33].

Conversely, in *Drosophila* the researchers were used Fluorescent Speckles Microscopy (FSM) technique to clarify the role of microtubule influx in producing the forces that drive chromosome movement to poleward. FSM is a useful method to following MT influx movement in the polymer, and their result showed the MT flux is the main forces for chromosome movement and this conclusion came by monitoring the speckles movement of injected labeled tubulin along MT [34]. Similarly, in the spermatocyte of insect, the results revealed the MT flux of kinetochore is the main forces to initiate the movement of chromosome [35]. Also, in fission yeast *Schizosaccharomyces pombe* the studies found it contains a member of motors proteins and none of these are necessary or involved for viability, so the chromosomes movement accomplish without motor proteins [36].

**Table 1** Approaches used to identify the protein contents of kinetochore. Table shows the different methods can be used to identify the proteins structure the kinetochore, and the principle of each method summarizes here:

Name of Assay	Principle
Delta Assay	Measure the distance of florescence spot between proteins
Human Antibody	Specific Antibody to recognize the target protein
Genetic, biochemical and RNAi	Depend on the inhibition or activation specific protein
Mass Spectrometry	Measure the ionized content of the protein
Fluorescence Microscopy	Detect the fluorescence illuminated by protein
GFP-Tagged fusion	Mark a protein with specific tag
Fluorescent Recovery After Photobleaching (FRAP)	Measure the motility of fluorescence protein
Electron Microscopy (High Pressure Freezing)	Detect the electron contents of a protein
Cells Free	Depend on Protein-Protein interaction
Fluorescent Speckles Microscopy (FSM)	Measure the movement and the dynamic of a protein

---

#### 4. Conclusion

From all above we concluded there were many models or mechanisms, which have ability to control the movement of chromosome during cell cycle. Chromosomes movement may be happened by forces generated from the depolarization of the MT minus end at the spindle poles and the source of energy emanate from hydrolysis of GTP through MT assembly, or from the KT motor proteins and these used the ATP as source of energy to support the mechanical forces to complete the movement of chromosomes, or by supporting roles of spindle matrix in the interaction between the motor protein through the generation of forces.

---

#### Compliance with ethical standards

##### *Acknowledgments*

We would like to thank the Department of Medical Basic Science-College of Nursing-University of Thi-Qar for their support and scientific cooperation.

##### *Disclosure of conflict of interest*

The authors declare no conflicts of interest.

---

#### References

- [1] Rago F, Cheeseman IM. Review series: The functions and consequences of force at kinetochores. *J Cell Biol* 2013;4;200(5):557-65.
- [2] Tooley J, Stukenberg PT. The Ndc80 complex: integrating the kinetochore's many movements. *Chromosome Res* 2011;19(3):377-391.
- [3] Akiyoshi B, et al. Tension directly stabilizes reconstituted kinetochore-microtubule attachments. *Nature* 2010; 468:576–9.
- [4] Vorozhko VV, Emanuele MJ, Kallio MJ, Stukenberg PT, Gorbsky GJ .Multiple mechanisms of chromosome movement in vertebrate cells mediated through the Ndc80 complex and dynein/dynactin. *Chromosoma* 2008;117:169–179.
- [5] Savoian, M.S., Goldberg, M.L., Rieder, C.L. The rate of poleward chromosome motion is attenuated in *Drosophila* zw10 and rod mutants. *Nat Cell Bio* 2000 ; 2 (12). pp. 948–952.

- [6] Wu J, Misra G, Russell RJ, Ladd AJ, Lele TP, Dickinson RB. Effects of dynein on microtubule mechanics and centrosome positioning. *Mol Biol Cell* 2011 ;22(24):4834-41.
- [7] Tanaka, T.U., and Desai, A. Kinetochore-microtubule interactions: the means to the end *Curr. Opin. Cell Biol* 2008; 20, 53–63.
- [8] Earnshaw, W. C. & Rothfield, N. Identification of a family of human centromere proteins using autoimmune sera from patients with scleroderma. *Chromosoma* 1985; 91, 313–321.
- [9] Cheeseman IM, Desai A: Molecular architecture of the kinetochore–microtubule interface. *Nat Rev Mol Cell Biol* 2008; 9:33-46.
- [10] Chan GK, Schaar BT, Yen TJ. Characterization of the kinetochore binding domain of CENP-E reveals interactions with the kinetochore proteins CENP-F and hBUBR1. *J Cell Biol* 1998 Oct 5;143(1):49-63.
- [11] Przewloka MR, Zhang W, Costa P, Archambault V, D'Avino PP, Lilley KS, Laue ED, McAinsh AD, Glover DM. Molecular analysis of core kinetochore composition and assembly in *Drosophila melanogaster*. *PLoS One* 2007 May 30;2(5):e478.
- [12] Hamilton GE, Davis TN. Biochemical evidence for diverse strategies in the inner kinetochore. *Open Biol* 2020 Nov;10(11):200284.
- [13] Raaijmakers JA, Tanenbaum ME, Maia AF, Medema RH. RAMA1 is a novel kinetochore protein involved in kinetochore-microtubule attachment. *J Cell Sci* 2009 Jul 15;122(Pt 14):2436-45.
- [14] Meraldi, P., McAinsh, A.D., Rheinbay, E. et al. Phylogenetic and structural analysis of centromeric DNA and kinetochore proteins. *Genome Biol* 7 2006; 7, R23.
- [15] Wan X, O'Quinn RP, Pierce HL, Joglekar AP, Gall WE, DeLuca JG, Carroll CW, Liu ST, Yen TJ, McEwen BF, Stukenberg PT, Desai A, Salmon ED. Protein architecture of the human kinetochore microtubule attachment site. *Cell*. 2009 May 15;137(4):672-84.
- [16] Foltz, D. R. et al. The human CENP-A centromeric nucleosome-associated complex. *Nature Cell Biol* 2006; 8, 458–469.
- [17] Aussie Suzuki, Sarah K Long, Edward D Salmon. An optimized method for 3D fluorescence co-localization applied to human kinetochore protein architecture. *eLife* 2018; 7:e32418.
- [18] Johnston K, et al. Vertebrate kinetochore protein architecture: protein copy number. *J Cell Biol* 2010; 189:937–43.
- [19] Nogales, E. Architecture and flexibility of the yeast Ndc80 kinetochore complex. *J. Mol. Biol* (2008); 383, 894–903.
- [20] Sandblad, L. et al. The *Schizosaccharomyces pombe* EB1 homolog Mal3p binds and stabilizes the microtubule lattice seam. *Cell* 2006;127, 1415–1424.
- [21] Tanenbaum, M. E., Galjart, N., van Vugt, M. A. & Medema, R. H. CLIP-170 facilitates the formation of kinetochore–microtubule attachments. *EMBO J* 2006; 25, 45–57
- [22] Kline, S.L., Cheeseman, I.M., Hori, T., Fukagawa, T., and Desai, A. The human Mis12 complex is required for kinetochore assembly and proper chromosome segregation. *J. Cell Biol* 2006; 173, 9–17.
- [23] He X, Rines DR, Espelin CW, Sorger PK. Molecular analysis of kinetochore-microtubule attachment in budding yeast. *Cell* 2001 Jul 27;106(2):195-206.
- [24] Gard, D. L., Becker, B. E. & Josh Romney, S. Mapping the eukaryotic tree of life: structure, function, and evolution of the MAP215/Dis1 family of microtubule associated proteins. *Int. Rev. Cytol* 2004; 239, 179–272.
- [25] McEwen BF, Hsieh CE, Mattheyses AL, Rieder CL. A new look at kinetochore structure in vertebrate somatic cells using high-pressure freezing and freeze substitution. *Chromosoma*1998;107(6-7):366-75.
- [26] J. B. Rattner and D. P. Bazett-Jones. Kinetochore Structure: Electron Spectroscopic Imaging of the Kinetochore. *The Journal of Cell Biology* 1989; Vol 108, 1209-1219.
- [27] Gorbsky GJ, Sammak PJ, Borisy GG. Microtubule dynamics and chromosome motion visualized in living anaphase cells. *J Cell Biol*1988 ;106(4):1185-92.
- [28] Watanabe R, Hirano Y, Hara M, Hiraoka Y, Fukagawa T. Mobility of kinetochore proteins measured by FRAP analysis in living cells. *Chromosome Res* 2022 Mar;30(1):43-57.

- [29] Sharp DJ, Rogers GC, Scholey JM. Cytoplasmic dynein is required for poleward chromosome movement during mitosis in *Drosophila* embryos. *Nat Cell Biol.* 2000 Dec;2(12):922-30.
- [30] Joglekar AP, Bloom K, Salmon ED. In vivo protein architecture of the eukaryotic kinetochore with nanometer scale accuracy. *Curr Biol.* 2009 Apr 28;19(8):694-9.
- [31] Tien, J.F., N.T. Umbreit, D.R. Gestaut, A.D. Franck, J. Cooper, L. Wordeman, T. Gonen, C.L. Asbury, and T.N. Davis. 2010. Cooperation of the Dam1 and Ndc80 kinetochore complexes enhances microtubule coupling and is regulated by aurora B. *J. Cell Biol* 2010;189:713–723.
- [32] Warren JD, Orr B, Compton DA. A comparative analysis of methods to measure kinetochore-microtubule attachment stability. *Methods Cell Biol* 2020; 158:91-116.
- [33] James R. LaFountain, Jr., Christopher S. Cohan, Alan J. Siegel, and Douglas J. LaFountain. Direct Visualization of Microtubule Flux during Metaphase and Anaphase in Crane-Fly Spermatocytes. *Molecular Biology of the Cell* 2004; Vol. 15, 5724–5732,
- [34] Poleward microtubule flux is a major component of spindle dynamics and anaphase a in mitotic *Drosophila* embryos. 2002. Maddox P, Desai A, Oegema K, Mitchison TJ, Salmon ED. *Curr Biol* 2002;1;12(19):1670-4.
- [35] Grishchuk, E.L., and J.R. McIntosh. 2006. Microtubule depolymerization can drive poleward chromosome motion in fission yeast. *EMBO J* 2006; 25:4888-4896.
- [36] Waterman-Storer CM, Desai A, Bulinski JC, Salmon ED. Fluorescent speckle microscopy, a method to visualize the dynamics of protein assemblies in living cells. *Curr Biol* 1998 Nov 5;8(22):1227-30.