In silico reconstitution of spindle assembly: speed and error of search and capture

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#### Different stages of mitosis

#### Search and Capture (S&C)



#### **Dynamic Instability of MT**



MT dynamics described by 4 parameters:

Basic question that we will address

What is the Fast & Accurate method of capturing all chromosomes ?

Part-I: minimal capture time

Part-II: capture accuracy

# Efficient Search & Capture requires MT's dynamic instability (Holy & Leibler, PNAS, 1994)

Dynamical instability of MTs are essential for efficient search and capture.

#### 1 MT searching for 1 KT

Search time is Minimum if :

- 1 Average length of MT = distance of the KT
- 2 MTs are not rescued when they undergo catastrophe.



Efficient chromosome capture requires a biased MT dynamics (Wollman et al. Current Biol., 2005)

Many MTs and many KTs

Unbiased versus biased S&C

- High RanGTP inside the nucleus stabilize MTs.
- MTs catastrophe is high outside the nucleus compared to inside.



# Efficient chromosome capture requires a biased S&C (Wollman et al. Current Biol., 2005)



Avg. cap. time is small for biased capture.
Capture time is logarithmic in Chromosome number.
Assumption: MTs are not obstructed by Chromosomes

Previous analysis considered...

## Chromosomes are transparent

### Chromosomes are static

#### Finite volume effect of chromosome



## Visibility decreases drastically with increasing chromosome number



### Part-I: Optimal capture time ?

#### **Two different S&C scenarios**



Static positioning of chromosomes - chromosomes are fixed at their initial locations.



Dynamic positioning of chromosomes – chromosomes are

fixed at their initial locations.

# Realistic to assume finite volume of chromosomes





#### Two different S&C scenarios

#### Random walk of chromosomes



Static positioning of chromosome - chromosomes are fixed at their initial locations.





Dynamic positioning of chromosomes – chromosomes are fixed at their initial locations.

## Compare capture time between model static & dynamic positioning of chromosomes



#### Pathways of S&C

Are the centrosomal and chromosomal pathways of MT dynamics integrated and coordinated for accurate spindle assembly ?



#### Longer the KT-fiber, smaller the capture time



## Biased Search & Capture DOES THE JOB! even for slow chromosomal jiggling



## So far I have shown you

- Static positioning of chromosome is not efficient.
- Capture process is way more efficient for the dynamic positioning of the chromosomes.
- Biased microtubule dynamics required for fast capture.

### Part-II: Accuracy

#### Incorrect attachment & consequence



#### Probability of different attachments



#### Probability of different attachments



#### Chromosome rotates after initial capture



#### Probability of different attachments



#### Probability of different attachments



## Long rotation decrease correct attachments



#### Predicting Error-statistics for multipolar cells





## Summary

 Meretolic attachment is most likely to occur for flexible MT-kinetochore attachments.

Alignment of KT axis along the MT and its stabilization rescues the captured KT to be meretolically attached. Therefore amphitelic attachment is most probable in this case.

### Thank you

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For details see: PNAS 106, 15708-15713 (2009)