

# KIF18A MUTATIONS LEAD TO CELL DIVISION ERRORS : INSIGHT INTO DIMINISHED FERTILITY IN YOUNG WOMEN

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## BACKGROUND

- Female fertility depends on coordinated organ systems, hormonal regulation, and genetics.
- Egg quality—especially chromosome number (euploidy)—is a key factor that declines with maternal age.
- Currently, maternal age is the only clinical predictor of egg euploidy, with quality decreasing in the mid 30s and fertility nearly lost by the mid-40s
- Elevated egg aneuploidy (EEA) drives reproductive aging, but its mechanisms remain under investigation
- The kinesin motor protein KIF18A plays a critical role in oocyte meiosis and ensuring euploidy
- Mutations of KIF18A, including S126L and T273A, have been linked to increased EEA in both animal models and young women undergoing infertility treatment (1, 2).

## OBJECTIVE

To determine how KIF18A mutations affect the following functions of the protein:

- Chromosomal alignment
- Spindle length
- Spindle polarity
- Mitotic index

## METHODS

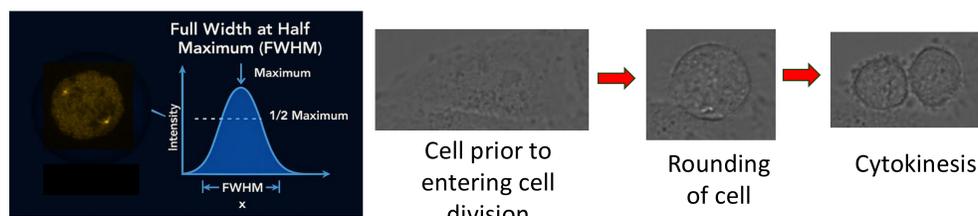
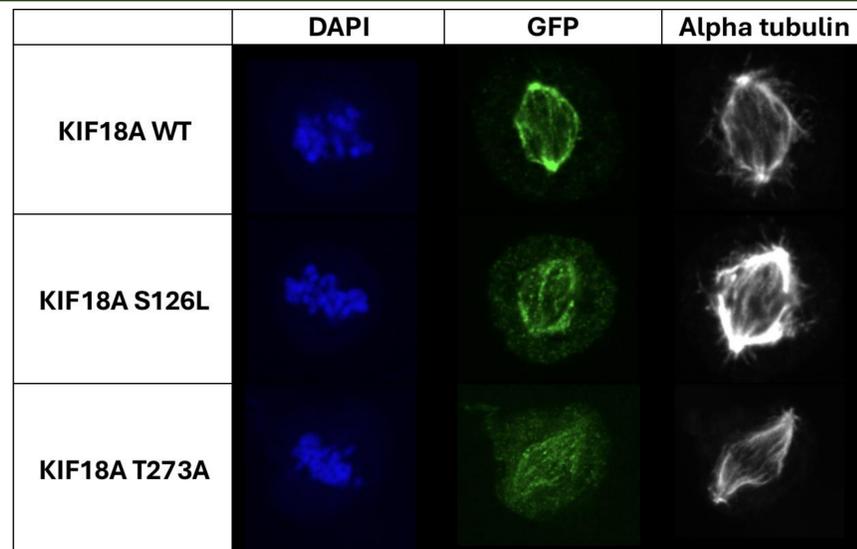
- HeLa Kyoto cell lines with GFP tagged KIF18A S126L and T273A were created using transformation and transfection via lipofectamine and electroporation techniques
- Doxycycline was used to drive expression of GFP tagged KIF18A WT, KIF18A S126L, and KIF18A T273A
- siRNA was used to knockout endogenous KIF18A
- Cells were fixed and stained for fluorescence microscopy
- Mitotic timing was assessed via 24-hour live imaging at 40x
- Alignment of chromosomes, spindle length, mitotic index, and polarity were assessed via fixed fluorescence microscopy at 40x and 60x
- To quantify mitotic timing, the time-point at which cells underwent rounding and cytokinesis was determined and the difference was calculated.
- To assess chromosomal alignment, a line spanning the spindle poles was used to measure kinetochore distribution with the Plot Profile tool in ImageJ/Fiji. Signal intensity was normalized to its maximum and plotted along the pole-to-pole axis using a custom MATLAB macro. Gaussian fitting was applied, and the Full Width at Half Maximum (FWHM) and spindle length were reported for each cell.

## ACKNOWLEDGEMENTS AND CONTACTS

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## VARIANT EXPRESSION IN MITOTIC CELLS



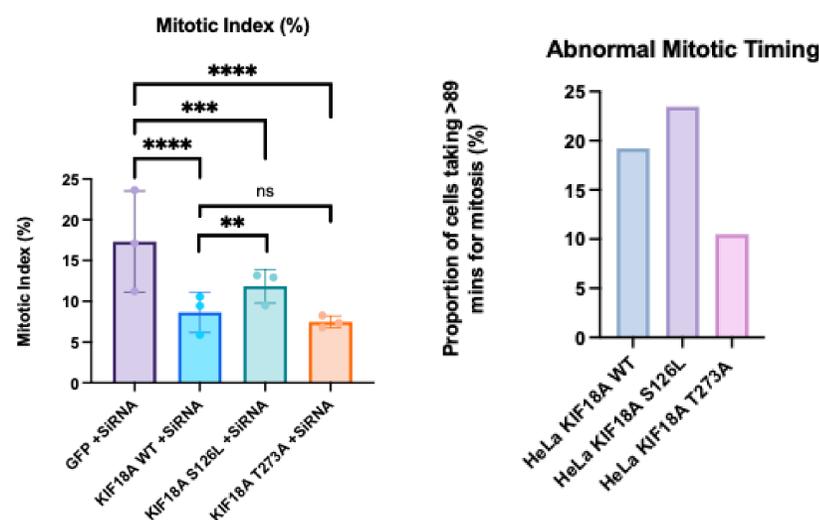
## RESULTS

### Mitotic Index

The mean mitotic index was 17.3% for GFP control, 8.6% for WT ( $p < 0.0001$ ), 11.86% for S126L ( $p = 0.0006$ ), and 7.47% for T273A ( $p < 0.0001$ ). Compared to WT, S126L had a statistically higher mitotic index ( $p = 0.003$ ), indicative of mitotic delay caused by loss of KIF18A.

### Mitotic Timing

Mean percentage of cells with abnormal mitotic timing were 19.2% for WT, 23.5% for S126L, and 10.5% for T273A.



## RESULTS

### Spindle Polarity

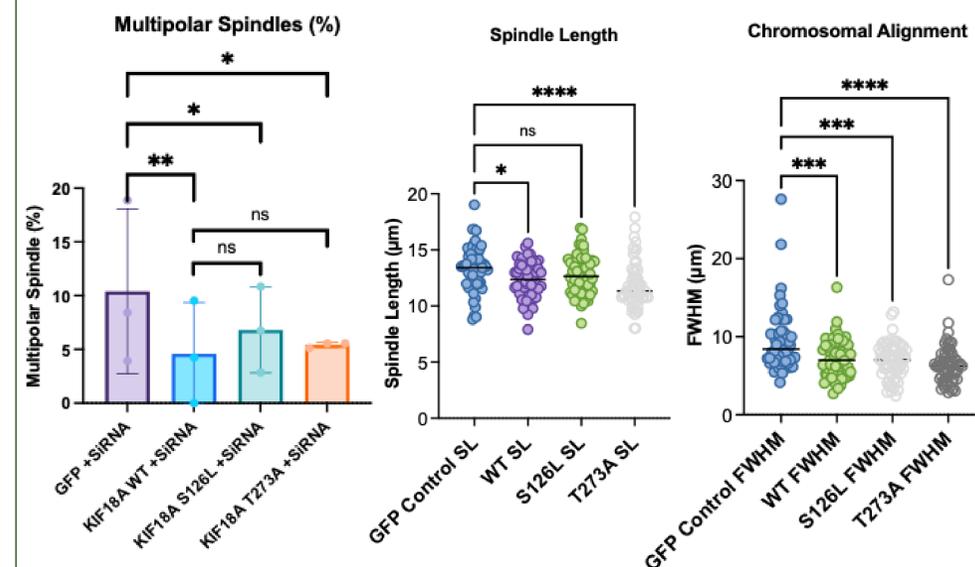
Mean percentage of cells with multipolar spindles were 10.4% for GFP control, 4.6% for WT ( $p = 0.002$ ), 6.8% for S126L ( $p = 0.048$ ), and 5.4% for T273A ( $p = 0.02$ ). Differences between WT and the variants were not statistically significant.

### Spindle Length

Median spindle lengths were 13.4  $\mu\text{m}$  for GFP control, 12.4  $\mu\text{m}$  for WT, 12.7  $\mu\text{m}$  for S126L, and 11.3  $\mu\text{m}$  for T273A. Differences were statistically significant between GFP control and both WT and T273A ( $p < 0.0001$ ).

### Chromosomal Alignment

Chromosome alignment, measured by determining the full width at half maximum (FWHM) for chromosomes in metaphase, was 8.4  $\mu\text{m}$  for GFP control, 7.0  $\mu\text{m}$  for WT, 7.0  $\mu\text{m}$  for S126L, and 6.2  $\mu\text{m}$  for T273A. FWHM was significantly lower in all KIF18A variants vs GFP control: WT ( $p = 0.0008$ ), S126L ( $p = 0.0002$ ), and T273A ( $p < 0.0001$ ).



## CONCLUSIONS

- Mitotic index, spindle length, and mitotic timing were significantly affected by the S126L mutation compared to WT.
- Spindle bipolarity and chromosomal alignment were not significantly affected by either mutation.
- These findings indicate that S126L but not T273A significantly reduces KIF18A function in promoting mitotic progression and spindle length control.
- These results suggest a mechanism to explain how infertility associated KIF18A mutations can lead to egg aneuploidy and diminished fertility in younger women.

## REFERENCES

Biswas L, Tyc KM, Aboelenain M, et al. Maternal genetic variants in kinesin motor domains prematurely increase egg aneuploidy. Proc Natl Acad Sci U S A, 2024. 121(45): p. e2414963121.  
Wu T, Luo Y, Zhang M, et al. Mechanisms of minor pole-mediated spindle bipolarization in human oocytes. Science, 2024. 385(6711): p. eado1022.