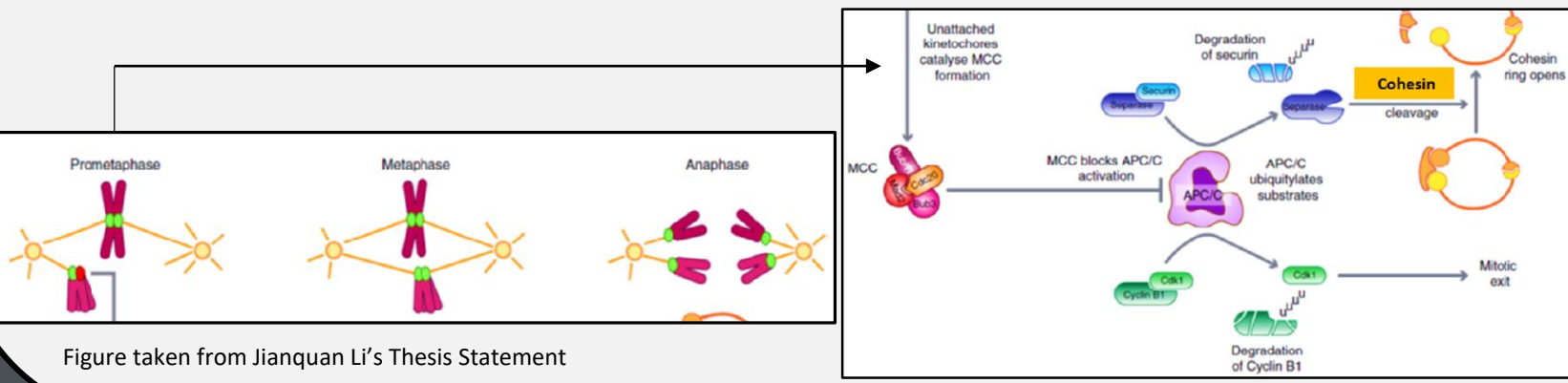


Role of p31^{COMET} Phosphorylation in Regulating the Mitotic Checkpoint Silencing in Human Cancer Cells

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Background

- The transition from metaphase to anaphase is tightly regulated by the spindle assembly checkpoint (SAC).
- SAC activation causes the inhibition of the anaphase promoting complex (APC) by a protein complex called MCC.
- p31^{COMET} is a protein complex that assists in the silencing of the SAC and can promote aneuploidy through its collaboration with Mad2.
- Phosphorylation of the p31^{COMET} weakens its interactions with Mad2.
- Conflictingly, only about half the number of p31^{COMET} are phosphorylated when SAC is turned on as a result of nocodazole treatment HeLa cells.



Aims

- To use an mGFP-p31 fusion protein to investigate the role of phosphorylation of p31^{COMET} in the SAC pathway.
- Make pCMV-AN-p31^{COMET}-mGFP plasmid DNA to express GFP tagged at the C terminus of p31^{COMET} with the aim of producing a sequenced p31^{COMET} containing flanking restriction sites.

Methods

HeLa cells were cultured and maintained in penicillin containing DMEM medium, and they were split every 2-3 days once 65-75% confluence was reached.

After splitting the cells, they were moved to an incubator at 37°C which was supplied with 5% CO₂.

Cells went through protein purification with 4 kinase inhibitors, DNase and RNase.

Samples were then run on a 10% protein gel.

Samples were transferred to a membrane from the protein gel through western blotting.

Phosphorylated and non-phosphorylated p31^{COMET} was then detected in the scanned membranes.

Figure A –

IPTG is a drug to induce p31 in the bacterial plasmid. We have tried different concentrations of it to find out which concentration can induce the most p31^{COMET}.

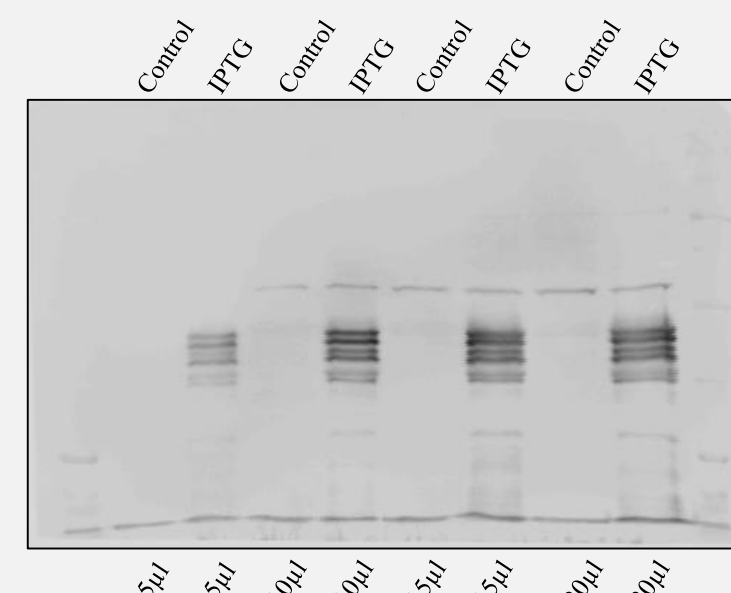


Figure B –

How p31^{COMET} phosphorylation can be detected on phos-tag gel.

Lane 1 - Control;
Lane 2 - HeLa cells treated with Nocodazole which arrests cells in mitosis;
Lane 3 - samples in Lane 2 treated with lambda Phosphatase, which could dephosphorylate p31.
GAPDH is a loading control; **Histon3S10** shows which stages the cells are. The upper band in Nocodazole treated cells is the phosphorylation form of p31. Therefore, about half of the p31 is phosphorylated in mitosis.

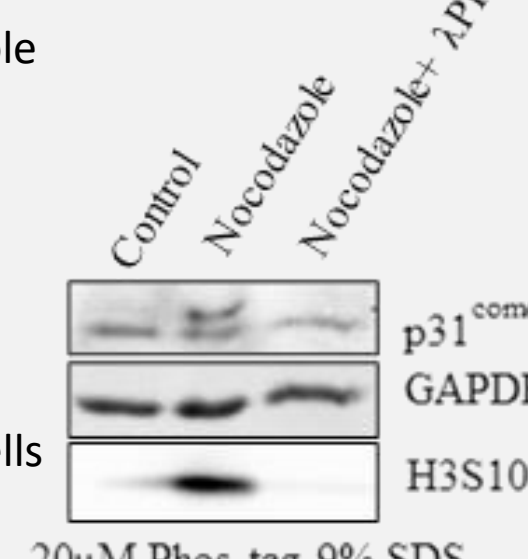


Figure C -

- We tested whether his-p31 can be phosphorylated in HeLa extract on phos-tag gel (a gel to identify phosphorylation of p31).
- We expected that if the His-p31 can be phosphorylated, the band of His-p31 should be higher than the non-phosphorylated ones.
- Lane 1** - His-p31+HeLa extract for 1 hour;
- Lane 2** - His-p31+HeLa extract +ATP for 30 min;
- Lane 3** - His-p31+HeLa extract +ATP for 1 hour;
- The left panel was stained with a p31 antibody, which showed two bands but no migration differences.
- The right panel was stained with a His antibody, which also showed no differences.
- Therefore, maybe the induced His-p31 is not the same as the endogenous p31, or His-p31 cannot be phosphorylated in HeLa extracts.

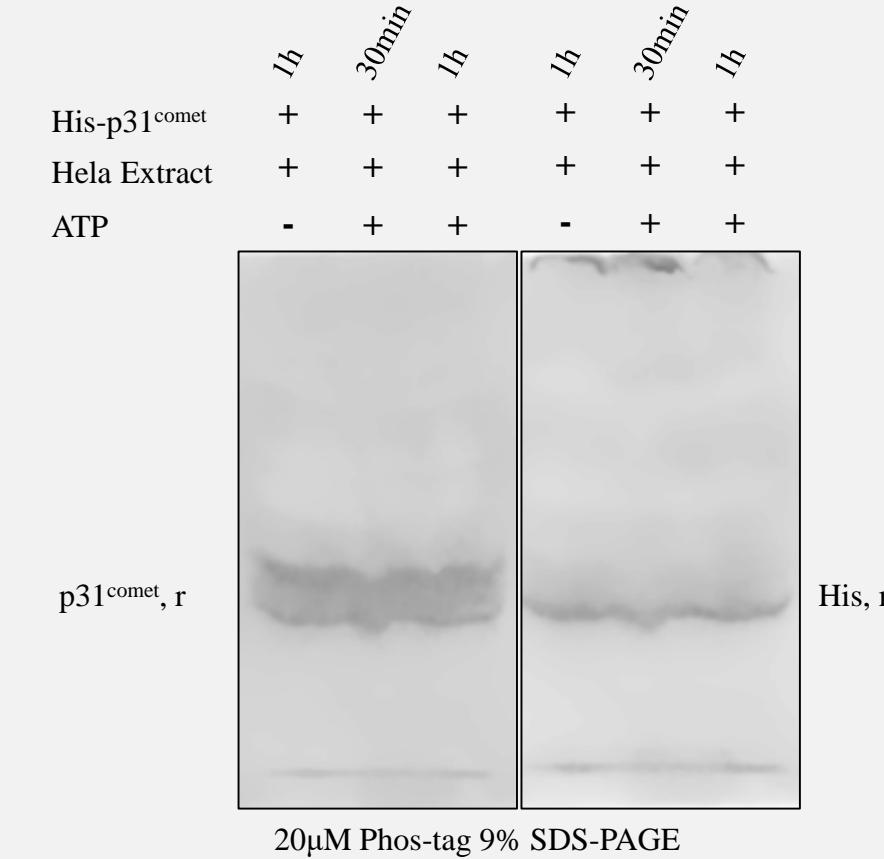
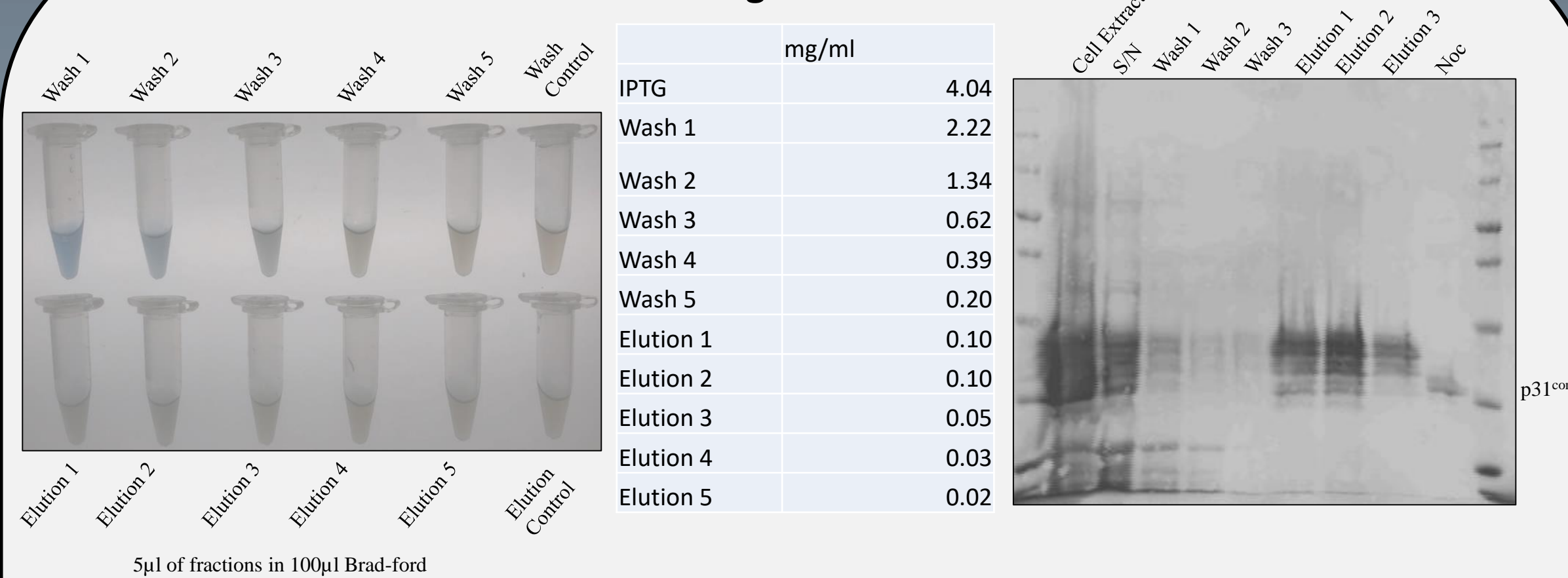


Figure D -



- (Left Picture)** – The darker the blue-ish color, the more proteins that are present.
 - First row is washed by wash buffer, which gets rid of the non-specific proteins.
 - Second row is washed with elution buffer which breaks bonds between the antibody and the beads.
- (Middle Picture)** - A table denoting the amount of p31^{COMET} protein left after every wash.
 - IPTG is used as a control
- (Right Picture)** - The samples on the left picture were run on a protein gel and these are the results that were produced.
 - The concentration of protein that is seen increases in "Elution 1" because the elution buffer is 10x more concentrated with imidazole than the wash buffer.
 - After all the washes we were able to purify the p31^{COMET} from all the other proteins.

Discussion

- The level of p31^{COMET} is low during interphase, but it starts to increase in prophase and reaches very high level at prometaphase and anaphase.
- p31^{COMET} is localized mainly in the nucleus and somewhat around the nuclear envelope during prophase.
- After the nuclear envelope breaks down, p31^{COMET} gets distributed throughout the whole cell.
- The levels of p31^{COMET} is low during interphase and high during metaphase and prometaphase and then decreases again in telophase were substantiated by western blot results.
 - Suggesting that p31^{COMET} levels are cell cycle regulated.

Figure E – Western Blot Result of elevated p31^{COMET} in HeLa Cells After Nocodazole Treatment

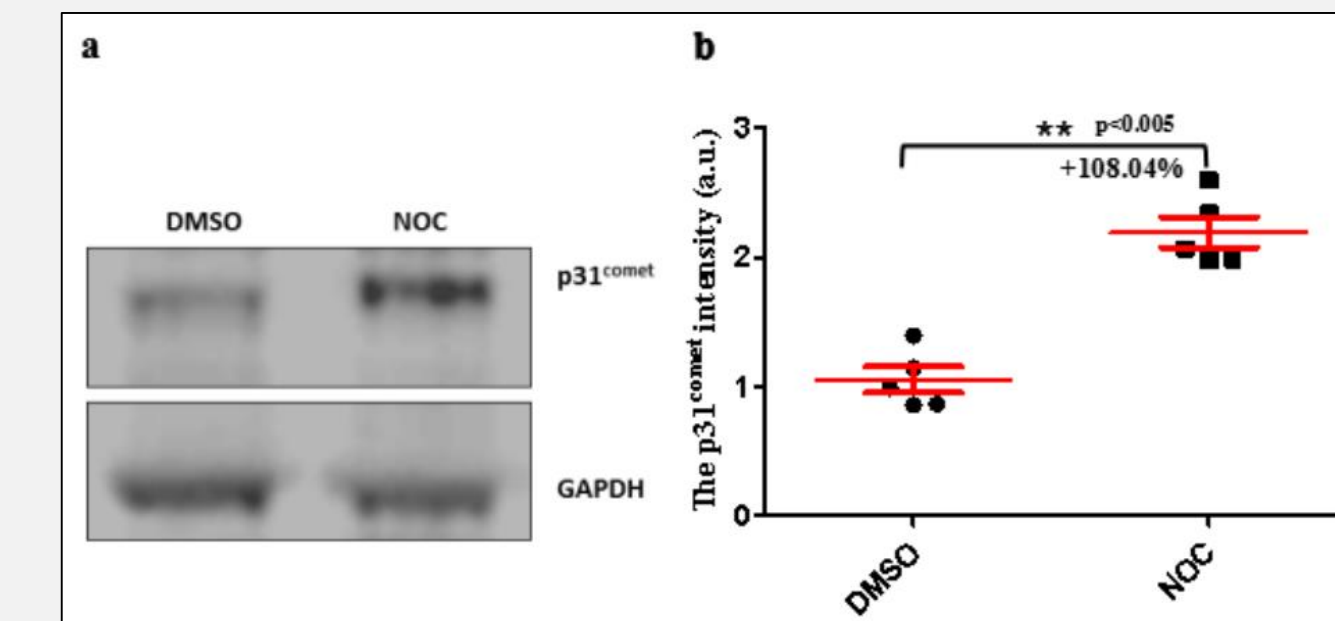


Figure taken from Jianquan Li's Thesis Statement

- After nocodazole treatment, high level of p31^{COMET} was detected during prometaphase.
- The high accumulation of the protein is caused by new protein synthesis.
 - We know this because, after nocodazole treatment, increased level of p31^{COMET} can be stopped by treating the cells with cycloheximide.
- The decline of p31^{COMET} from cells that are released from nocodazole treatment can be prevented by MG132, but not leupeptin.
 - This p31^{COMET} degradation is probably mediated by the ubiquitin-proteasome pathway.
- p31^{COMET} contains 2 degradation motifs which are the D-box and KEN-box.
 - These are signature motifs that are targeted by the APC/C E3 ligase dependent proteolysis pathway.
- This suggested that if we inhibit the APC/C ligase pathway we would get an increased amount of p31 but unfortunately it was not evident in the results.

Conclusion

- To sum it all up, more experiments need to take place to see how the APC/C contributes to p31^{COMET} degradation as they decline after nocodazole treatment release.
- The results also depict that the level of p31^{COMET} increases after cullin-1 siRNA treatment.
 - This proposes that SCF E3 ligase ubiquitin – mediated proteolysis might be the contributing factor in the degradation of p31^{COMET} as Cullin-1 is the essential core protein of SCF E3 ligase.
- Therefore, the degradation process of p31^{COMET} needs to be further investigated.

Acknowledgments

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- I would also like to thank Nanmao Dang for supervising me during intricate procedures and allowing me to work alongside her.

References:

- [1] Sharon Kaisari A et al. Role of Polo-like Kinase 1 in the Regulation of the Action of p31^{COMET} in the disassembly of MCC. 2019 | 1 - 6
- [2] Dipali A. Date et al. Phosphorylation Regulates the p31^{COMET} Mitotic Arrest- Deficient 2 Interaction to Promote SAC Activity. 2013 | 11367 – 11373
- [3] Toshiyuki Habu et al. Identification of a MAD2-Binding Protein, CMT2, and its Role in Mitosis. 2002 | 6419 – 6426
- [4] Jianquan Li. The Mechanism of Spindle Assembly Checkpoint and Mitotic Cell Death. 2018 | 1 - 99