

A novel PHD-finger protein 14/KIF4A complex overexpressed in lung cancer is involved in cell mitosis regulation and tumorigenesis

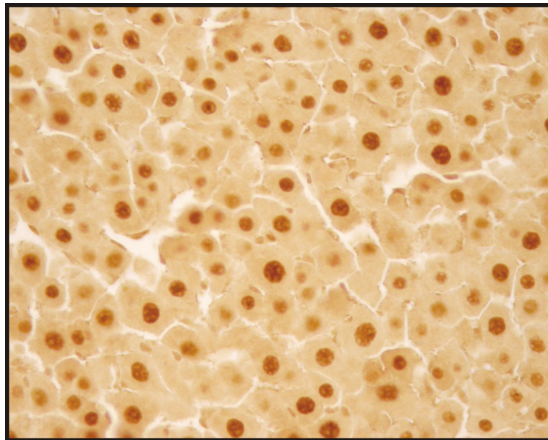
SUPPLEMENTARY DATA

SUPPLEMENTARY METHODS

***In vitro* binding assay**

GST-PHF14 (N-terminal 1-160 a.a) and GST were purified from BL21 bacteria and eluted from glutathione-Sepharose 4B beads (GE Healthcare, UK) as suggested by

the manufacturer. For binding assays, pEGFP-C1 KIF4A full length was incubated with equivalent amounts of GST or GST-PHF14 (N-terminal 1-160 a.a) fusion proteins. After incubation, bead-bound proteins were separated by SDS-PAGE and analyzed by western blotting.

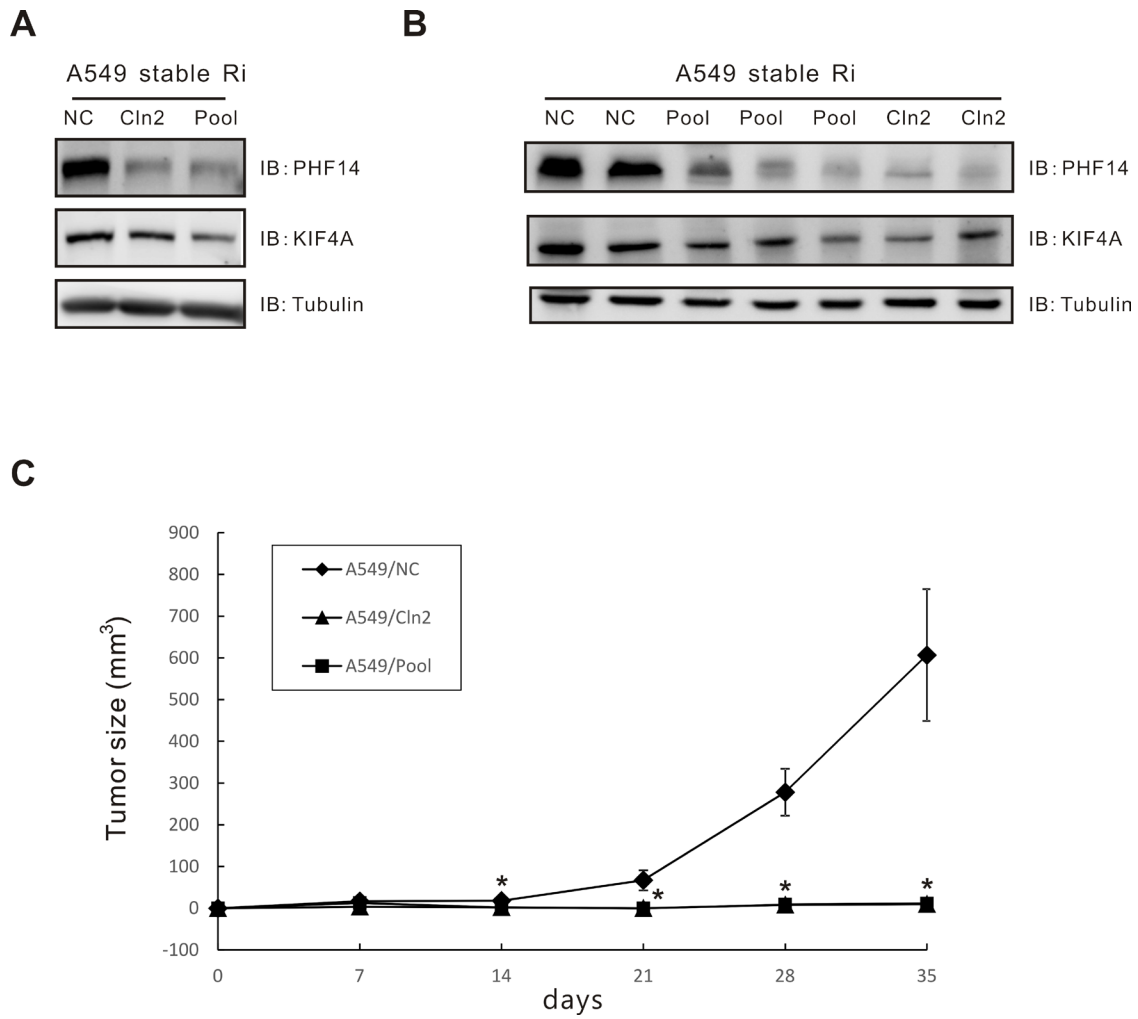


- Antigen

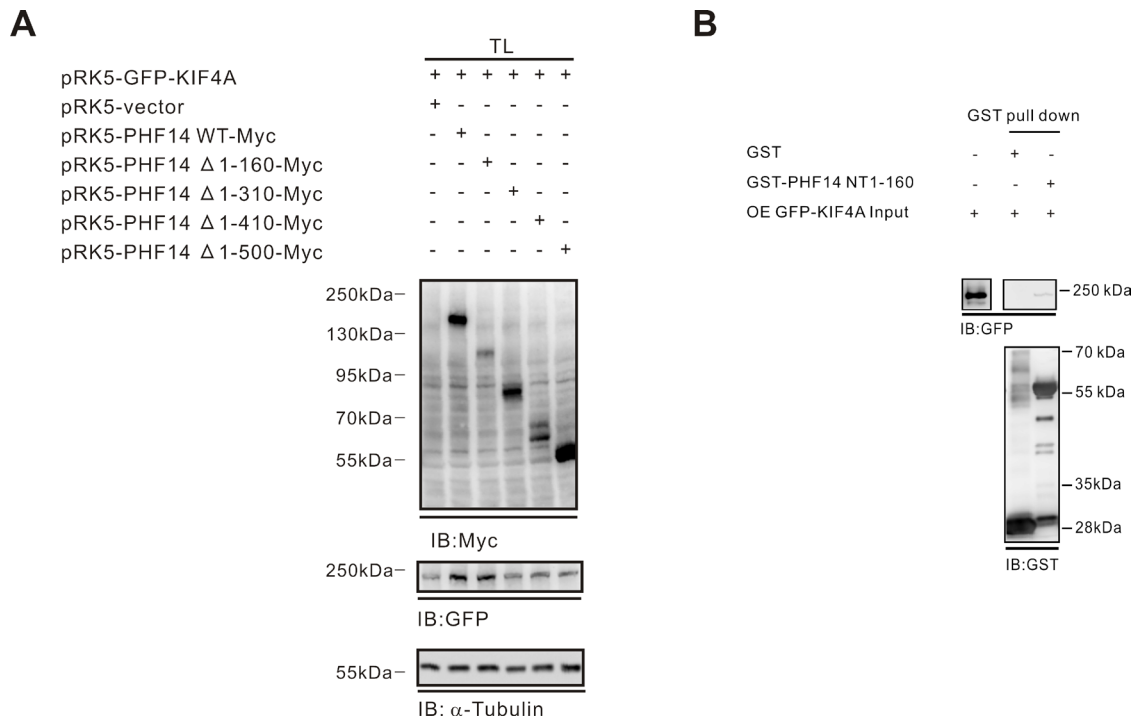


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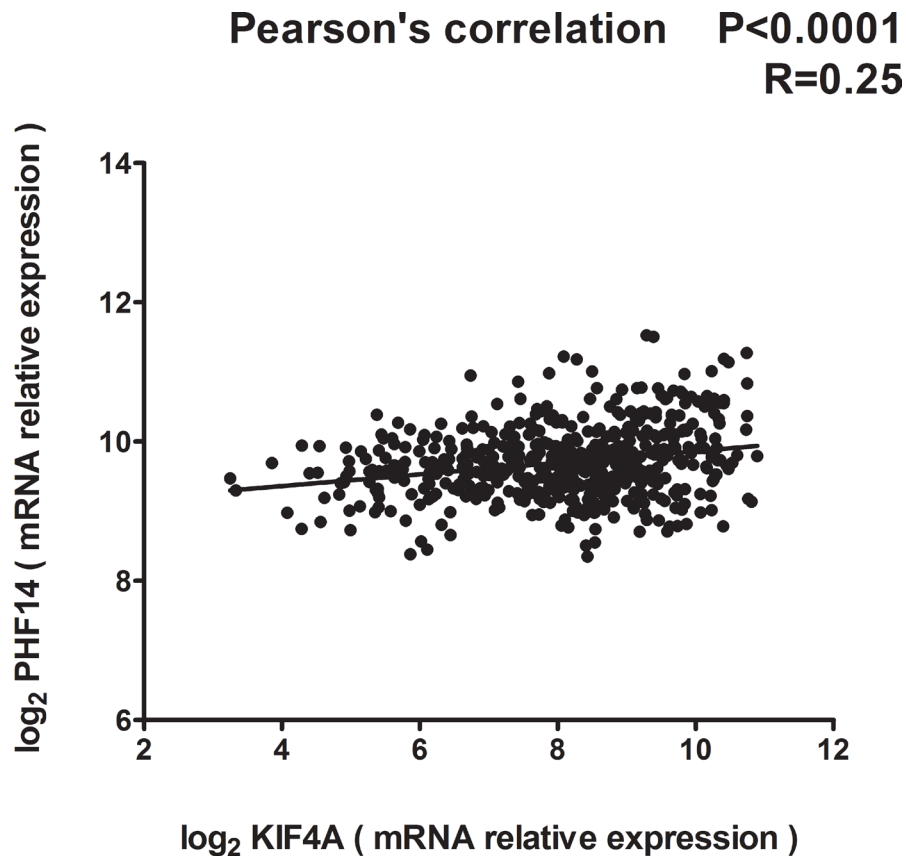
Supplementary Figure 2: The specificity of the PHF14 polyclonal antibody. Immunohistochemical analysis of mouse liver showing PHF14 expression using the anti-PHF14 polyclonal antibody without (upper) or with (bottom) pre-incubation with the antigen.



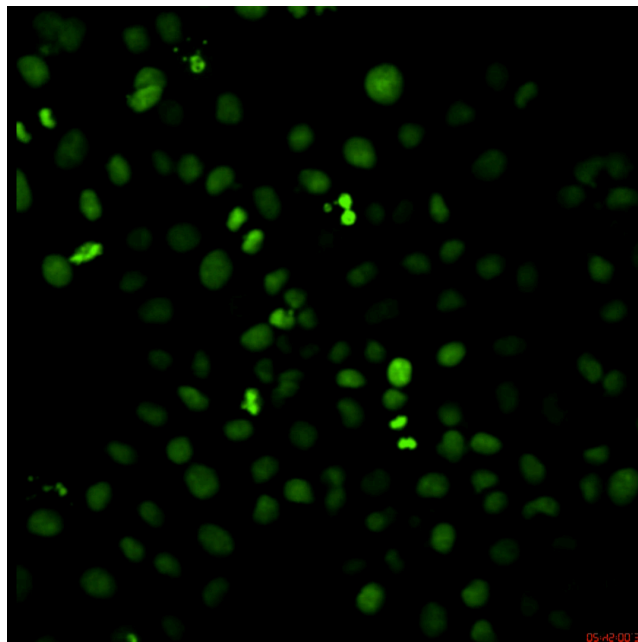
Supplementary Figure 3: PHF14/KIF4A expression and tumor formation in nude mice. **A.** PHF14/KIF4A expression in A549 control cells (NC) or PHF14 knockdown-cells (A549/Cln2 or PHF14-knockdown pool cells). **B.** PHF14/KIF4A expression in A549 NC, Cln2 or pool cells - formed tumors six weeks after inoculation. **C.** Average tumor size in the mice injected with different cells as indicated. The mean value \pm s.d. was calculated from 4-6 samples in each group. * $P < 0.05$.



Supplementary Figure 4: PHF14 interacts with KIF4. **A.** Co-overexpression of myc-tagged PHF14 or its mutants with GFP-tagged WT KIF4A. Different constructs were transfected into the HCT-116 cells. The exogenous PHF14 and KIF4A were detected using an anti-Myc/anti-GFP antibody respectively. α -Tubulin was used as the loading control. **B.** GST pull-down assay. Purified GST-tagged fragments were incubated with 293T cell lysates containing GFP-tagged KIF4A WT.

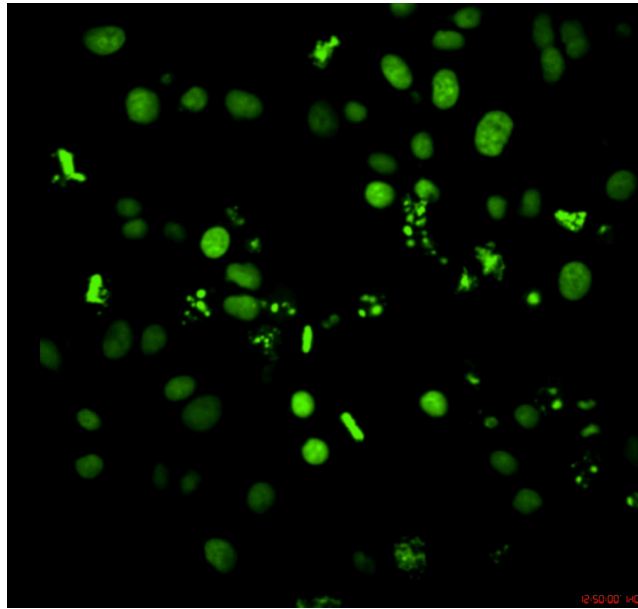


Supplementary Figure 5: PHF14 co-overexpressed with KIF4A. The correlation between PHF14 and KIF4A mRNA expression levels in paired NSCLC samples. RNAseq gene expression profiles were retrieved from TCGA database, $n=516$, $R=0.246$, $P < 0.0001$.



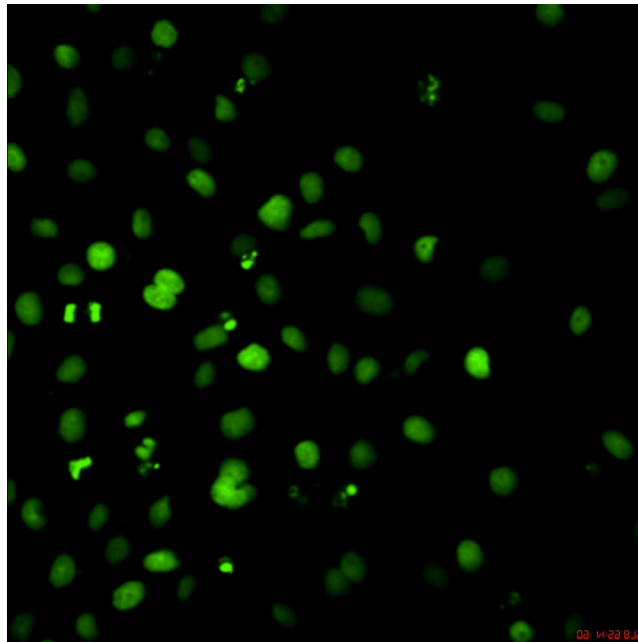
Supplementary Movie 1: Control HeLa cells undergoing mitosis. HeLa cells stably expressing histone H₂B-GFP (green) and transfected with control siRNA. Images were captured every 5 min for about 12h.

See Supplementary Movie 1



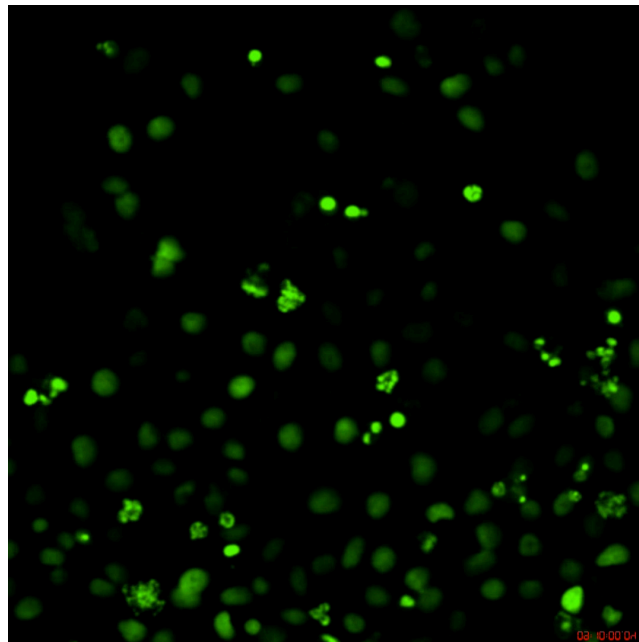
Supplementary Movie 2: Abnormal mitosis in PHF14-knockdown cell. HeLa cells stably expressing histone H₂B-GFP (green) were transfected with siRNA targeting PHF14. Images were captured every 5 min for about 12h.

See Supplementary Movie 2



Supplementary Movie 3: Abnormal mitosis in KIF4A-knockdown cell. HeLa cells stably expressing histone H₂B-GFP (green) were transfected with siRNA targeting KIF4A. Images were captured every 5 min for about 12h.

See Supplementary Movie 3



Supplementary Movie 4: Abnormal mitosis in PHF14/KIF4A double-knockdown cell. HeLa cells stably expressing histone H₂B-GFP (green) were transfected with siRNA targeting PHF14 and KIF4A. Images were captured every 5 min for about 12h.

See Supplementary Movie 4

Supplementary Table 1: Mutations of PHF14 detected in lung squamous cell carcinoma and lung adenocarcinoma from TCGA dataset

| Disease Type | Participants ID | Gene Name | Sample Name | CDS Mutation | AA Mutation | domain location |
|------------------------------|-----------------|-----------|------------------------------|--------------|-------------|----------------------|
| Lung Squamous Cell Carcinoma | TCGA-33-4532 | PHF14 | TCGA-33-4532-01A-01D-1267-08 | c.2030G>A | p.S677N | CC2 |
| Lung Squamous Cell Carcinoma | TCGA-34-5239 | PHF14 | TCGA-34-5239-01A-21D-1817-08 | c.2313-6A>T | p.? | PHD3 |
| Lung Adenocarcinoma | TCGA-44-2656 | PHF14 | TCGA-44-2656-01A-02D-0969-08 | c.202G>T | p.E68* | before CC domain1 |
| Lung Adenocarcinoma | TCGA-44-2656 | PHF14 | TCGA-44-2656-01A-02D-0969-08 | c.1437A>T | p.S479S | PHD2 |
| Lung Adenocarcinoma | TCGA-44-2659 | PHF14 | TCGA-44-2659-01A-01D-0969-08 | c.1373G>T | p.C458F | PHD2 |
| Lung Squamous Cell Carcinoma | TCGA-46-3769 | PHF14 | TCGA-46-3769-01A-01D-0983-08 | c.205G>T | p.E69* | before CC domain1 |
| Lung Squamous Cell Carcinoma | TCGA-46-3769 | PHF14 | TCGA-46-3769-01A-01D-0983-08 | c.2004A>G | p.L668L | CC2 |
| Lung Adenocarcinoma | TCGA-49-6742 | PHF14 | TCGA-49-6742-01A-11D-1855-08 | c.1342C>T | p.R448C | PHD2 |
| Lung Adenocarcinoma | TCGA-50-6590 | PHF14 | TCGA-50-6590-01A-12D-1855-08 | c.1318G>T | p.E440* | PHD2 |
| Lung Squamous Cell Carcinoma | TCGA-66-2758 | PHF14 | TCGA-66-2758-01A-02D-1522-08 | c.1520A>G | p.K507R | between PHD2 and CC2 |
| Lung Adenocarcinoma | TCGA-67-3771 | PHF14 | TCGA-67-3771-01A-01W-0928-08 | c.881C>T | p.S294F | between CC1 and PHD1 |

Supplementary Table 2: Mutations of PHF14 detected in lung adenocarcinoma

| Disease Type | Gene Name | Sample Name | CDS Mutation | AA Mutation | domain location |
|---------------------|------------------|--------------------|---------------------|--------------------|------------------------|
| Lung Adenocarcinoma | PHF14 | 136 | c.1718C>T | p.L569F | between PHD2 and CC2 |
| Lung Adenocarcinoma | PHF14 | 203 | c.1818C>T | p.S602F | between PHD2 and CC2 |
| Lung Adenocarcinoma | PHF14 | 1516 | c.1818C>T | p.S602F | between PHD2 and CC2 |
| Lung Adenocarcinoma | PHF14 | 1574 | c.1818C>T | p.S602F | between PHD2 and CC2 |
| Lung Adenocarcinoma | PHF14 | 1165 | c.2797ATG>TGA | p.MK928NE | after PHD4 |