NUAK1 coordinates growth factor-dependent activation of mTORC2 and Akt signaling.

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Supplementary Information

Figure S1. NUAK1 inhibition does not impact on mTOR-Rictor association.

A. Immunoblot (IB) of the Immunoprecipitation (IP) of endogenous mTOR and Co-IP of endogenous Rictor after 5 minutes of EGF stimulation in MDA-MB-231 cells. **B-C.** IB of IP of endogenous mTOR and Co-IP of endogenous Rictor after 60 minutes of EGF stimulation in MDA-MB-231 (B) and U87 (C) cells.



Figure S2. NUAK1 effect on mTOR subcellular distribution.

A. Representative confocal images of mTOR in MDA-MB-231 cells expressing inducible shRNAs for NUAK1. Inducible models were treated with or without doxycycline by 24 hours and then serum-starved overnight followed by stimulation with EGF for 60 minutes Green, mTOR; Red, RFP; Blue, Nuclei. **B.** Representative confocal images of mTOR and Rab5 in MDA-MB-231 cells serum-starved overnight followed by 90 minutes of pretreatment with DMSO or HTH-01-015 (10 μ M) before stimulation with EGF for 60 minutes. Red, Rab5; Green, mTOR; Blue, Nuclei. **C.** Quantification of mTOR/Rab5 co-localization from B. Each bar represents the mean \pm SD, Student t test.



Figure S3. NUAK1 inhibition does not affect early endosomes distribution.

Representative confocal images of Rab5 and Lamp1 under NUAK1 inhibition in MDA-MB-231 after EGF stimulation for 60 minutes. Left, merge; Right, Rab5 and Lamp1 + Hoechst images. Cells borders were marked with a boundary. Green, Rab5; Red, Lamp1; Blue, nuclei



Figure S4. NUAK1 effect on Akt signaling under different cellular conditions.

A. IB of the effect of NUAK1 inhibition on Akt signaling in normal growth conditions. MDA-MB-231 cells were treated with DMSO or HTH-01-015 (10 μM) for 2, 6, 12 and 24 hours. β-actin was used as loading control. **B.** IB of the effect of NUAK1 inhibition on the Akt signaling upon oxidative stress. MDA-MB-231 cells were pretreated with DMSO or HTH-01-015 (10 μM) by 1-hour before treatment with H₂O₂ by 0, 30 and 120 minutes. GAPDH was used as loading control. **C-D.** IB of Akt signaling under NUAK1 inhibition in U87 (C) and SW480 (D) cells serum-starved overnight followed by 1-hour of pretreatment with DMSO or HTH-01-015 (10 μM) before stimulation with EGF. **E.** IB of Akt signaling under NUAK1 inhibition in MDA-MB-231 cells serum-starved overnight followed by 1-hour of pretreatment with DMSO or HTH-01-015 (10 μM) before stimulation with EGF.



Figure S5. NUAK1 co-localize with endogenous Rab5.

A. IF of MDA-MB-231 cells after 0 (-EGF) and 15 (+EGF) minutes of EGF stimulation. Representative confocal images for HA-NUAK1 WT and endogenous Rab5. Red, HA-Tagged-NUAK1; Green, endogenous Rab5 (early-endosome marker); Blue, nuclei. **B.** Quantification of HA-NUAK1/Rab5 co-localization from A. Each bar represents the mean ± SD, not significant (ns), Student t test.

Α	cDNA 188611-014_E07_p0f_NUAK_K84A_6_5p0A8E.ab1	CAGTEGEETGEGGGATETAGETTAGEEACEATGGAETAAAGAEGATGAEGATAAGEAA	8 60
	cDNA	GAATTCGAAGGGCCGCCGCCGCCTGTGG	28
	180611-014_E07_pBf_NUAK_KB4A_6_SpBABE,ab1	GGGATCCACTAGTAACGGCCGCCAGTGTGCTCGGAATTCGAAGGGGCCGCCGCCGCCTGTGG	120
	CDNA	C66666ACC6CCCC6ACTT6666CT66666CGCC666CTCTCCCC6A6A66C66T66C66	88
	188611-014_E07_p8f_NUAK_K84A_6_SpBABE,ab1	C66666ACC6CCC6ACTT6666CT66666C6C666CTCTCCCC6A6A66C66T66C66	18#
	CDNA	GGECGACTGCAGCCCTGGAGCCCAGGAAGCCGCACGGGGTGAAGCGGCATCACCACAAGC	148
	188611-014_E07_p0f_NUAK_K84A_6_SpBABE,ab1	GGECGACTGCAGCCCTGGAGCCCAGGAAGCCGCACGGGGTGAAGCGGCATCACCACAAGC	240
	cDNA	ACAACTTGAMGCACCGCTACGAGCTGCAGGAGACCCTGGGCAAAGGCACCTACGGCAAAG	208
	180511-014_E07_p8f_NUAK_K84A_6_SpBABE.ab1	ACAACTTGAMGCACCGCTACGAGCTGCAGGAGACCCTGGGCAAAGGCACCTACGGCAAAG	300
	cDNA	TCAAGCGGGCCACCGAGAGGTTTTCTGGCCGAGTGGTTGCTATAAAATCCATTCGTAAGG	268
	180611-014_E07_p0f_NUAK_K84A_6_Sp0A0E,ab1	TCAAGCGGGCCACCGAGAGGTTTTCTGGCCGAGTGGTTGCTATAGCCTCCATTCGTAAGG	368
	cDNA	ACAMAATTAMEGATGAACAAGACATEGTTCACATCAGACGAGAGATTGAGATCATGTCAT	328
	180611-014_E07_p0f_NUAK_K84A_6_Sp0ADE,ab1	ACAMAATTAAGGATGAACAAGACATEGTTCACATCAGACGAGAGATTGAGATCATGTCAT	420
	cDNA	CTCTCAACCATCCTCATATCATCAGTATTTATGAAGTGTTTGAGAACAAAGATAAGATTG	388
	188611-014_E07_p0f_NUAK_K84A_6_SpBA8E.ab1	CTCTCAACCATCCTCATATCATCAGTATTTATGAAGTGTTTGAGAACAAAGATAAGATTG	480
	cona	TGATCATCATGGAATATGCCAGCAAAGGGGAGCTGTACGATTACATCAGTGAGCGGCGAC	448
	180611-014_007_007_NUAK_K84A_6_SpBABE,ab1	TGATCATCATGGAATATGCCAGCAAAGGGGGGGCTGTACGATTACATCAGTGAGCGGCGAC	540
	c0NA	GCCTCAGTGAGAGGGAGACCCGGCACTTCTTCCGGCAGATCGTCTCTGCTGTGCACTATT	588
	180611-014_E07_p0f_NUAK_K84A_6_Sp0A8E,ab1	GCCTCAGTGAGAGGGGAACCCGGCACTTCTTCCGGCAGATCGTCTCTGCTGTGCACTATT	600
	cDNA	GTCACAAGAACGGTGTGGTCCACCGGGACTTGAAGCTGGAAAATATACTGCTCGATGACA	568
	180511-014_E07_p0f_NUAK_K84A_6_SpBADE.ab1	GTCACAAGAACGGTGTGGTCCACCGGGACTTGAAGCTGGAAAATATACTGCTCGATGACA	668
	cDNA	ACTGCAATATTAAGATTGCTGACTTTGGGGCTTTCCAACCTGTACCAGAAGGATAAGTTCT	628
	180611-014_E07_p8f_NUAK_K84A_6_SpBABE,ab1	ACTGCAATATTAAGATTGCTGACTTTGGGGCTTTCCAACCTGTACCAGAAGGATAAGTTCT	720
	cDNA	TACAAACGTTTTGT666AGTCCACTCTATGCATCTCCTGAGATTGTCAAT666AGACCTT	683
	180611-014_E07_p0f_NUAK_K84A_6_SpBA8E,ab1	TACAAACGTTTTGT666AGTCCACTCTATGCATCTCCTGAGATTGTCAAT666AGACCTT	788
	cDNA	ACCGAGGGCCAGAGGTGGACAGCTGGGCCCTGGGTGTGTTGCTTTACACTCTTGTTTATG	748
	180511-014_007_p0f_NUAK_K84A_6_5p0A00,ab1	ACCGAGGGCCAGAGGTGGACAGCTGGGCCCTGGGTGTGTTGCTTTACACTCTTGTTTATG	840
	c0NA	GAACAATGCCCTTCGATGGTTTCGATCACAAAAACCTCATTCGGCAAATCAGCAGCGGAG	888
	180611-014_E07_p8f_NUAK_K84A_6_SpBA8E.ab1	GAACAATGCCCTTCGATGGTTTCGATCACAAAAACCTCATTCGGCAAATCAGCAGCGGAG	988
	cdNA	AGTACCGGGAGCCAACACAGCCCTCAGATGCTCGAGGACTCATACGGTGGATGCTGATGG	863
	188511-814_E87_p8f_NUAK_K84A_6_Sp8A8E,ab1	AGTACCGGGAGCCAACACAGCCCCTCAGATGCTCGAGGACTCATACGGTGGATGCTGATGG	968



Figure S6. Validation of NUAK1 Kinase Dead mutant (K84A).

A. Alignment of human NUAK1 WT (cDNA) with the sequence obtained by sequencing of human NUAK1 K84A. The Alignment validated the mutation at Lysine 84 (K84) by Alanine (A). **B.** IB of the activity of FLAG-NUAK1 WT and FLAG-NUAK1 K84A. phospho-MYPT1 at Ser-445 was used as a positive control. NUAK1 K84A mutant does not induced MYTP1 phosphorylation.



Figure S7. Long-term inhibition of NUAK1 dramatically induces morphological changes. Representative confocal images of MDA-MB-231 cells treated with DMSO, HTH-01-015 (5 μ M) or Palbociclib (1 μ M) (used as a positive control) for 4 days. Red, α -tubulin; Green, Phalloidin (F-actin); Blue, nuclei (n=3).