

# NECAB3 Promotes Activation of Hypoxia-inducible factor-1 during Normoxia and Enhances Tumorigenicity of Cancer Cells

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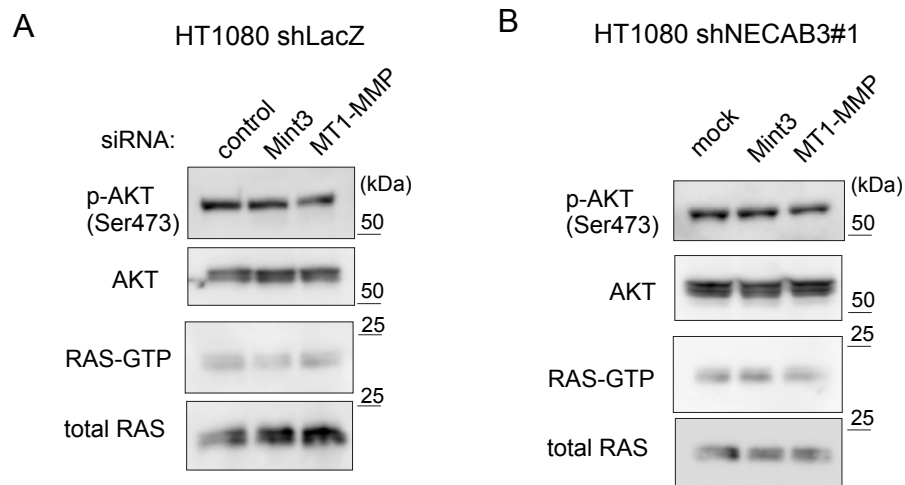
## Supplementary Materials and Methods

**shRNA sequences.** shRNAs against Mint3 had sequence 5'-gaugauggcgguggugacgggacgaaucccgucaccaccgcca-3' (#1) and 5'-gcggguucuugguccuguaugacgaaucauacaggaccaagaaccgc-3' (#2).

**Primer sequences.** Primers specific for *NECAB1* (sense, 5'-ttcctcgacatactgaggag-3' and antisense, 5'-tcacatagctcttctgtgc-3'), *NECAB2* (sense, 5'-cgacagaaccacatcaaacc-3' and antisense, 5'-atcctccgtggcagatggaa-3'), *NECAB3* (sense, 5'-ccgcagagcagacaagaatg-3' and antisense, 5'-ccagtgctctgagaagtag-3') were used for real-time PCR.

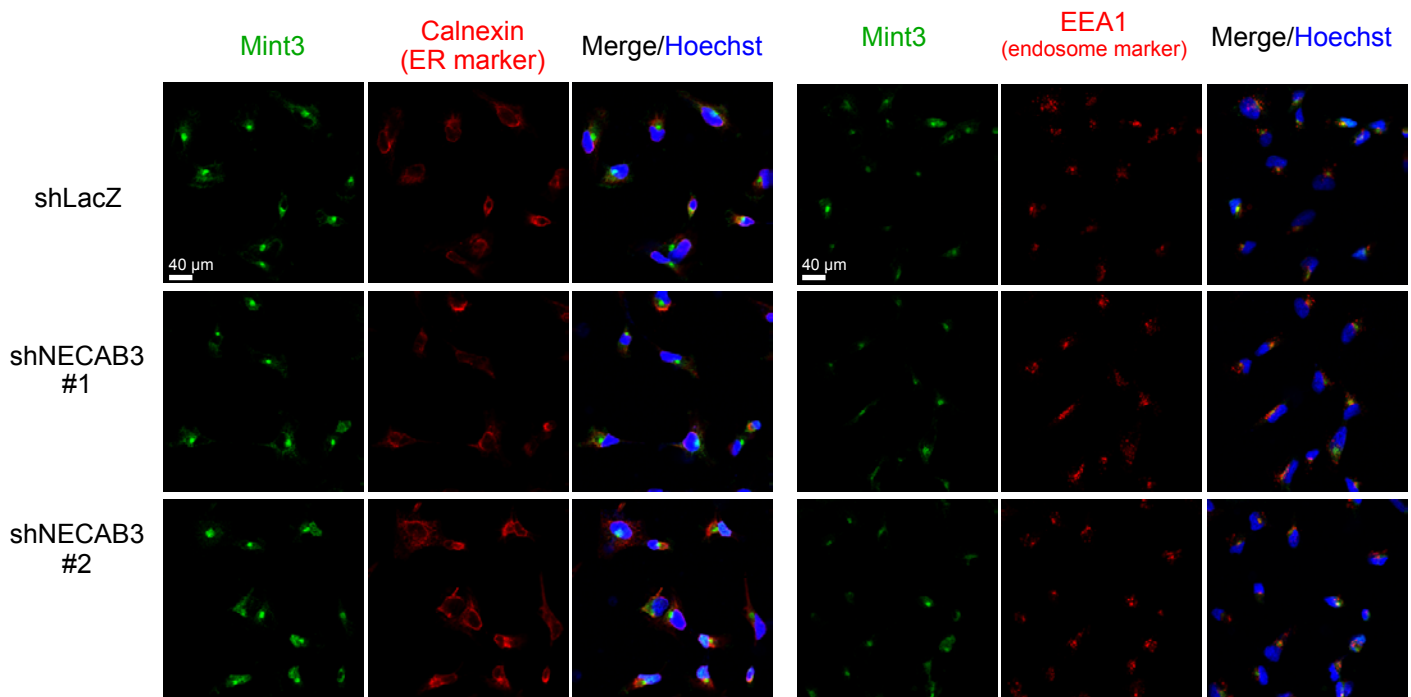
**Antibodies.** Rabbit antibodies against calnexin (Abcam, Cambridge, U.K.) and EEA1 (Cell Signaling Technology, Danvers, MA) were used for immunostaining.

## Supplementary Figures.

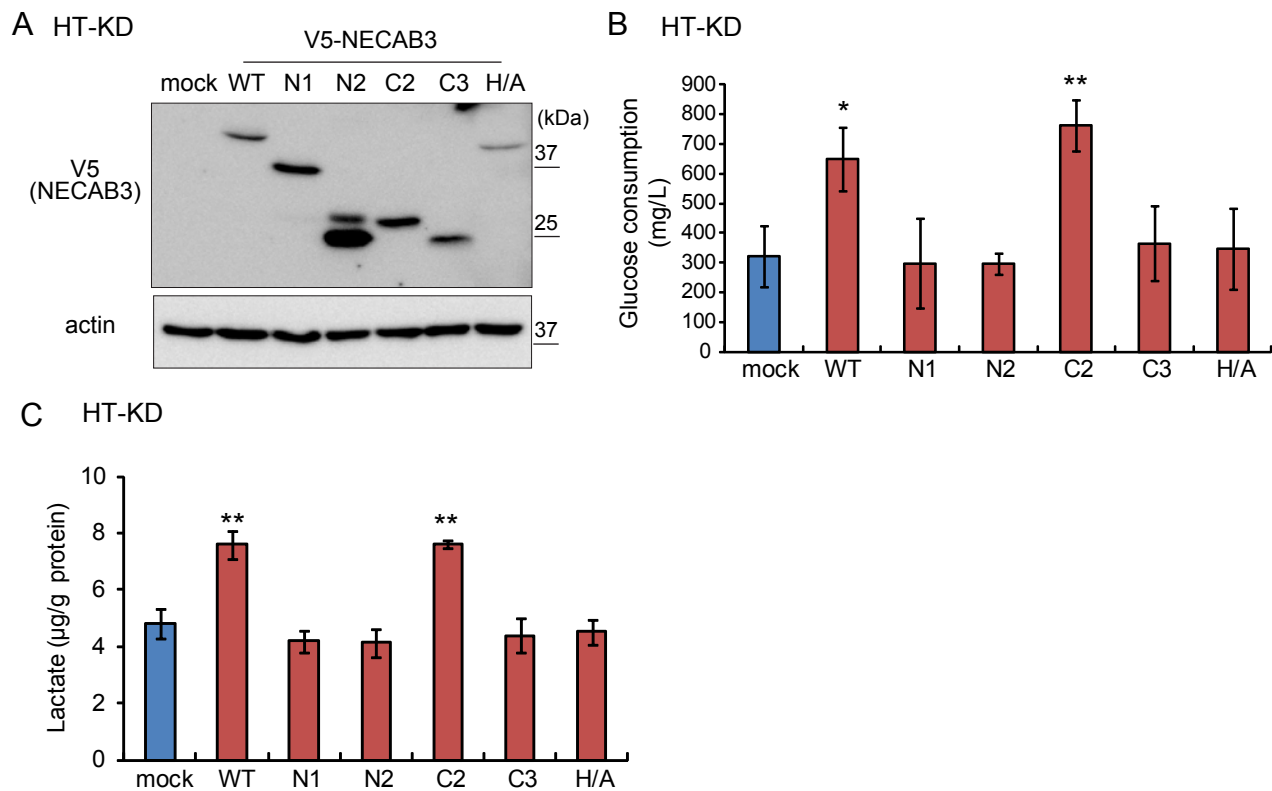


**Figure S1. Knockdown or overexpression of MT1-MMP and Mint3 did not affect AKT phosphorylation and Rac activation.**

(A-B) Immunoblotting of phospho-AKT (Ser473), AKT, active Ras (Ras-GTP), and total Ras in whole cell lysates from control (HT1080shLacZ, **A**) or NECAB3-depleted cells (HT1080shNECAB3#1, **B**) transfected with siRNAs against MT1-MMP, Mint3, or control sequences.



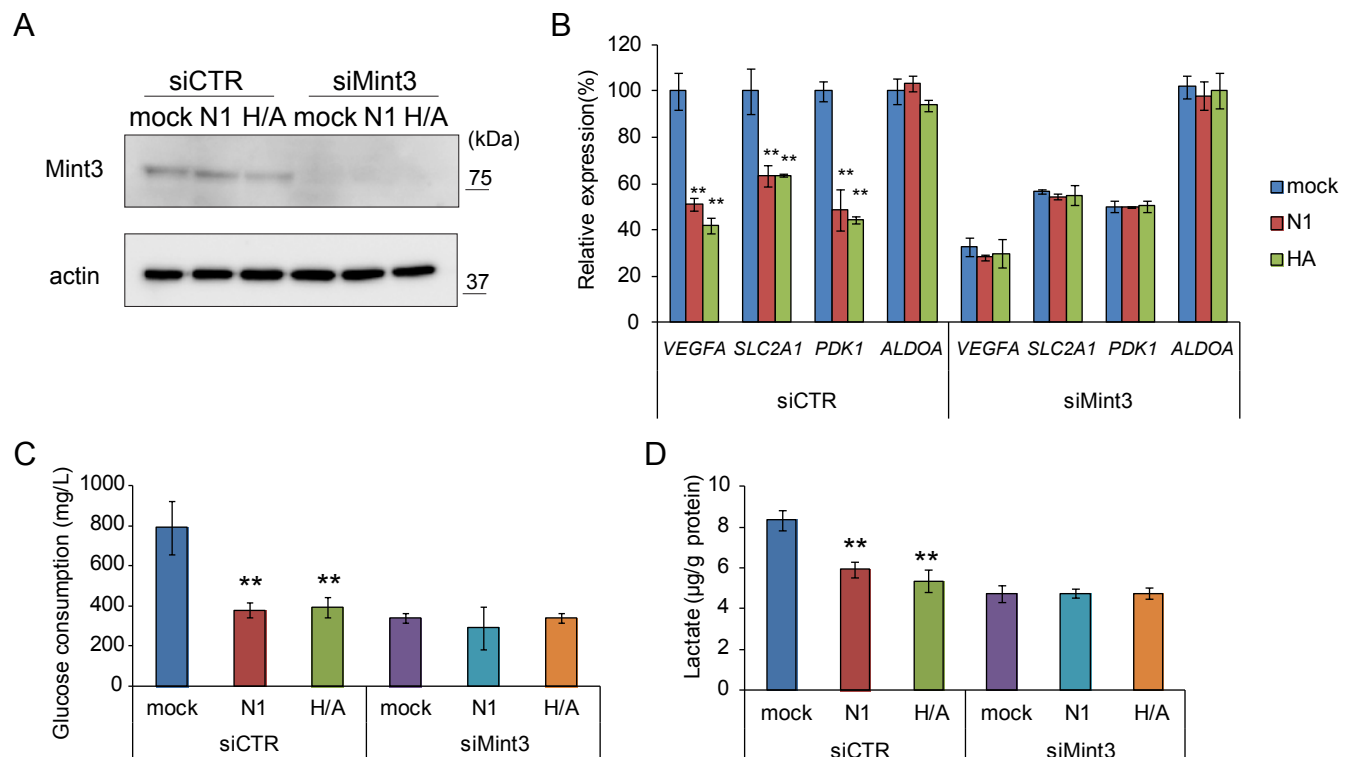
**Figure S2. Immunostaining of Mint3, calnexin, and EEA1 in NECAB3-depleted HT1080 cells.**



**Figure S3. Wild-type and C2 mutant of NECAB3 restores glycolysis during normoxia in NECAB3-depleted HT1080 cells.**

(A) Expression of V5-tagged NECAB3 and its mutants in NECAB3-depleted HT1080 cells.

(B, C) Glucose consumption (B) and lactate production (C) in NECAB3-depleted HT1080 cells expressing NECAB3 derivatives. In B and C, error bars indicate s.d. (n = 3), and data were analysed by t-test. \*, p < 0.05 and \*\*, p < 0.01.

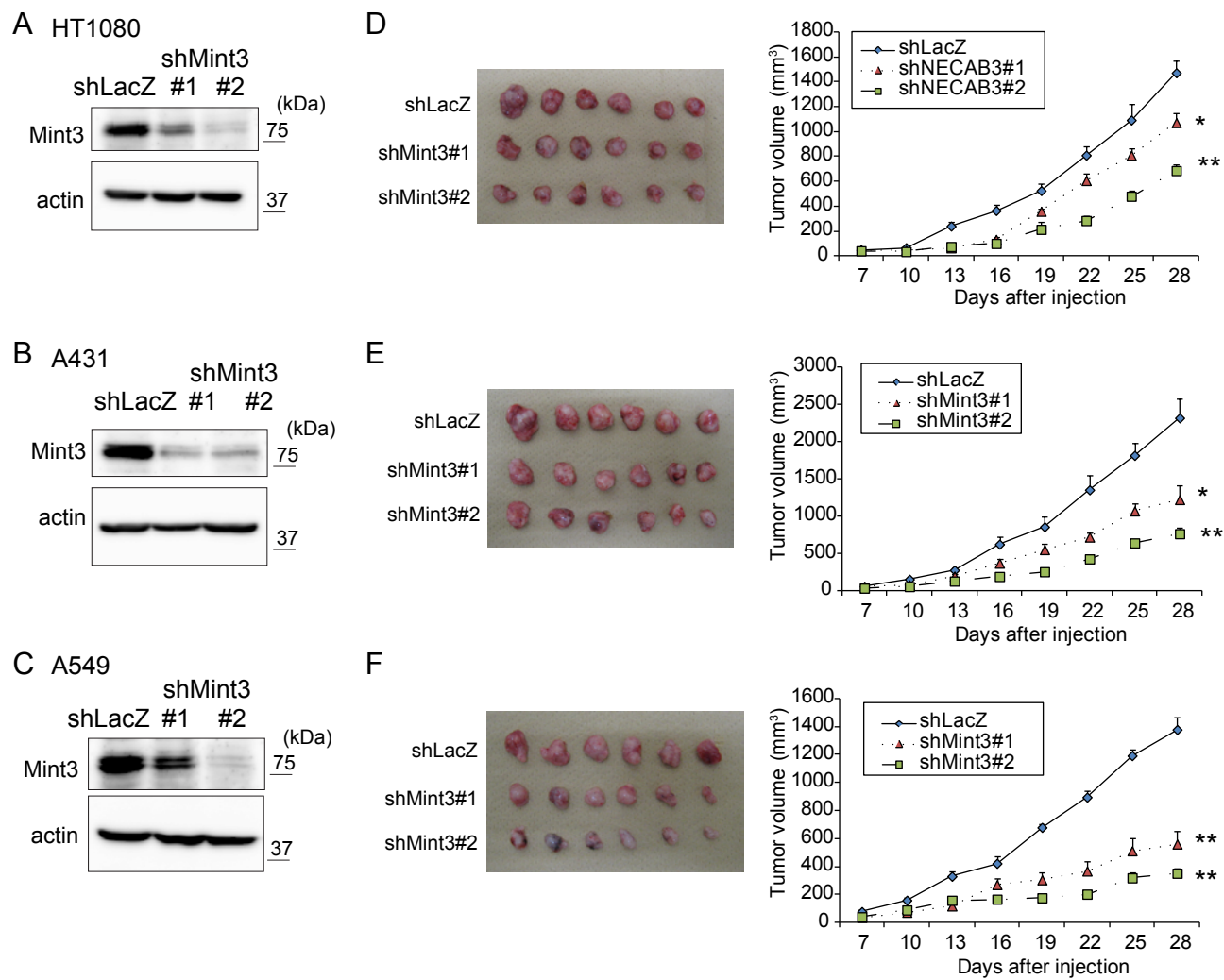


**Figure S4. Dominant-negative effects of NECAB3 mutants require Mint3.**

(A) Immunoblotting of Mint3 and actin in whole lysates from HT1080 cells expressing mock, NECAB3 N1, or H/A mutants and transfected with control or Mint3 siRNA.

(B-D) Gene expression (B), glucose consumption (C), and lactate production (D) in HT1080 cells expressing mock or NECAB3 dominant negative mutants and transfected with control or Min3 siRNA.

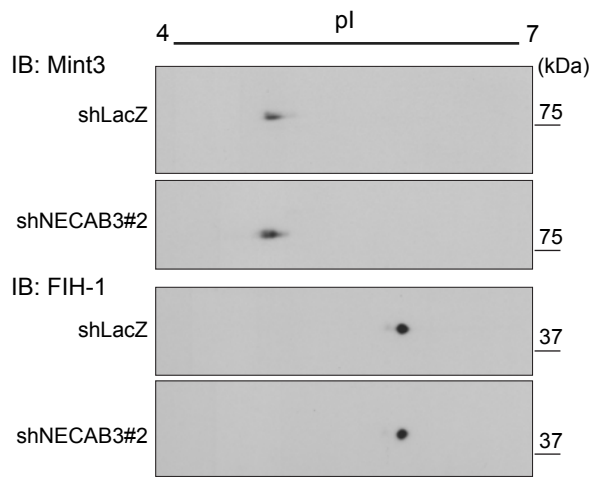
In B-D, error bars represent s.d. (n = 3), and data were compared pairwise by t-test. \*\*, p < 0.01.



**Figure S5. Mint3 depletion attenuates growth of cancer cells implanted in vivo.**

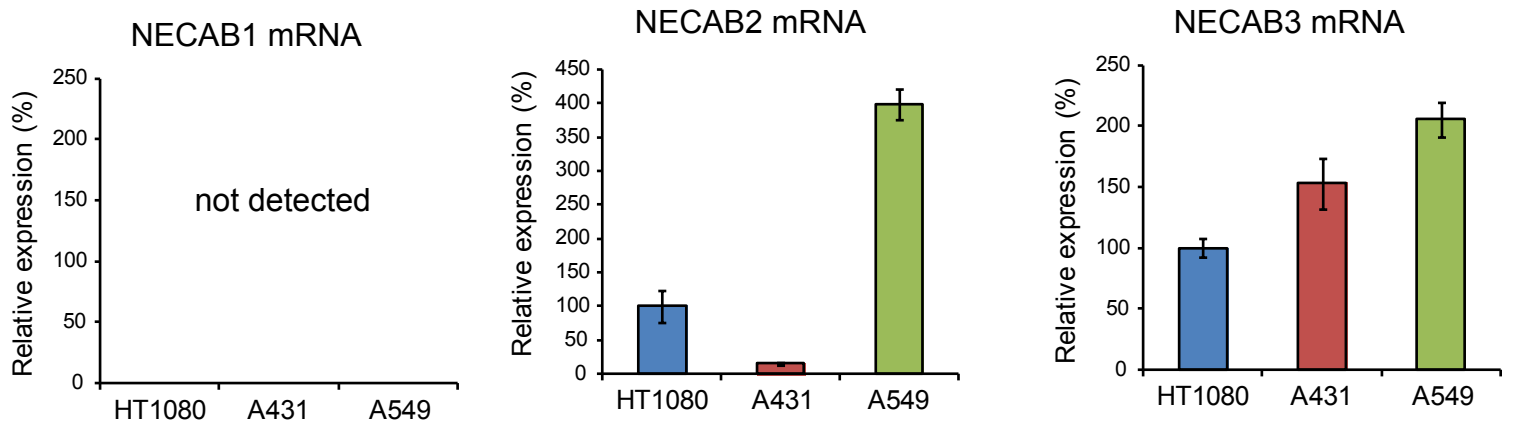
(A-C) Immunoblotting of Mint3 and actin in whole cell lysates from control (shLacZ) or Mint3-depleted (shMint3) HT1080 (A), A431 (B), and A549 cells (C).

(D-F) Growth of Mint3-depleted HT1080 (D), A431 (E), and A549 cells (F) implanted into immunodeficient mice. Photos show tumours at day 28 after injection. Error bars indicate s.e.m. (n = 6), and data were analysed by Mann-Whitney U-test. \*, p < 0.05 and \*\*, p < 0.01.



**Figure S6. NECAB3 depletion does not affect phosphorylation of Mint3.**

The acidity of Mint3 and FIH-1 in control (shLacZ) or NECAB3-depleted HT1080 cells was analysed by 2D electrophoresis and immunoblotting.



**Figure S7. Expression of NECAB genes in cancer cells.**

Expression of NECAB1, NECAB2, and NECAB3 was analysed by real-time PCR in HT1080, A431, and A549 cells. NECAB1 expression was not detectable in all cells.

gene name	reference sequence	start	stop
NECAB3	NM_031231.2	336	972
		546	1019
APLP2	NM_001642.1	2124	2581
		2127	3191
		2139	3190
		2142	3248
		2160	3084
		2166	2579
		2184	3238
		2184	3188
		2184	3252
		2184	3238
		2208	3558
		2235	3080
BTBD15	NM_014155.3	51	1602
		360	1599
		687	1576
		1032	1485
ESE-1	U73844.1	645	1229
HSD17B1	NM_000413.1	-385	694
		-376	318
		-373	699
SOX13	NM_005686.2	903	1594
SOX7	NM_031439.2	-70	860
		-58	871
ZNF143	NM_003442.4	330	1448

**Table S1. Prey fragments identified from yeast two-hybrid screens against Mint3.**