

Tob negatively regulates NF- κ B activation in breast cancer through its association with the TNF receptor complex.

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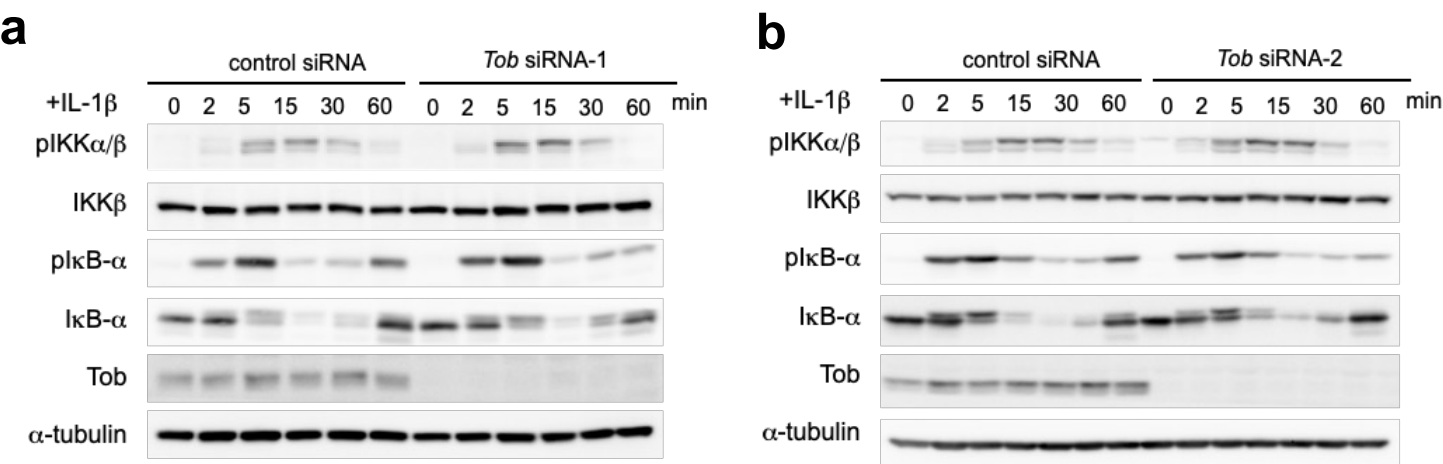
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Competing Interests

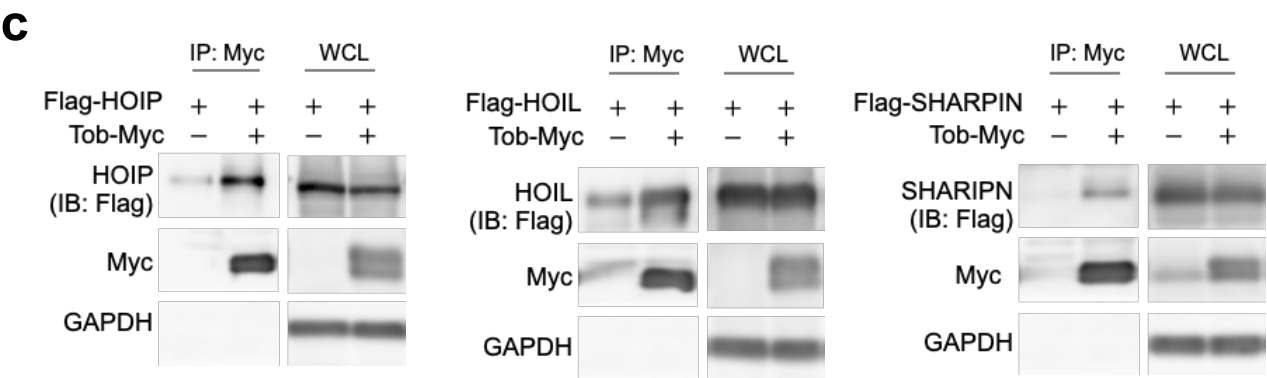
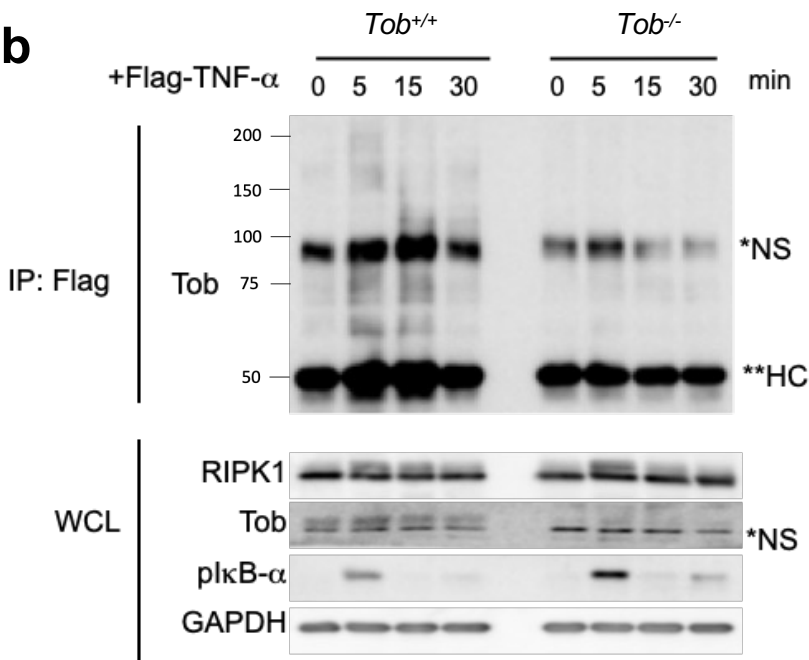
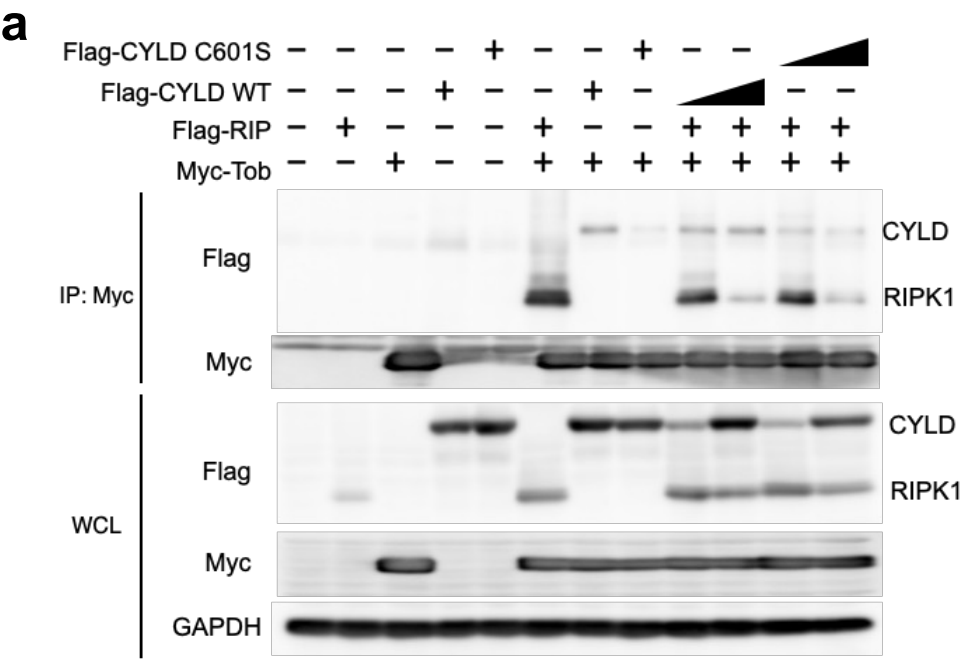
We have nothing to disclose.

Supplemental Figure 1



Supplemental Figure 1: *Tob* doesn't affect IL-1 β induced NF- κ B activation in MCF-7 cells. (a, b) MCF-7 cells were transiently transfected with a control, *Tob* siRNA-1 (a) or siRNA-2 (b). After 72 h, cells were treated with 10 ng/mL IL-1 β .

Supplemental Figure 2



Supplemental Figure 2: High expression of CYLD prevents Tob and RIPK1 interaction independently of de-ubiquitylation activity. (a) HEK293T cells were transiently transfected with C-terminal Myc-tagged Tob (1 μ g), N-terminal Flag-tagged RIPK1 (0.2 μ g), and N-terminal Flag-tagged CYLD WT (0.3 μ g or 3 μ g) or CYLD^{C601S} mutant (0.3 μ g or 3 μ g). After 24 h, cells were harvested and subjected to immunoprecipitation with an anti-Myc tag antibody and protein G-conjugated beads. WCLs were also prepared from the same cell lysates before immunoprecipitation. (b) *Tob*^{+/+} or *Tob*^{-/-} MEFs were stimulated with Flag-tagged TNF- α (2 mg/mL). Cells were harvested at the indicated times and were subjected to immunoprecipitation with anti-Flag tag antibody-conjugated beads (refer to Figure 5c). IP samples were subjected to western blotting with anti-Tob antibody. (c) HEK293T cells were transiently transfected with C-terminal Myc-tagged pME18s-Tob and N-terminal Flag-tagged HOIP, HOIL, or SHARIPN. After 24 h, cells were harvested and subjected to immunoprecipitation with an anti-Myc tag antibody and protein G-conjugated beads. WCLs were also prepared from the same cell lysates before immunoprecipitation.

Supplemental Table 1

Cell line	subtype	ER	PR	ErbB2
CRL1500	luminal	+	+ /—	—
UACC-812	luminal	+	+ /—	+
HCC1419	luminal	—	—	+
T47D	luminal	+	+	—
YMB-1-E	ND			
MCF-7	luminal	+	+	—
MDA-MB-453	luminal	—	—	+
MDA-MB-361	luminal	+	+	+
HCC2157	basal	—	—	—
MDA-MB-175VII	luminal	+	—	—
HCC1500	luminal	+	+	—
HCC2218	luminal	—	—	+
UACC-893	luminal	—	—	+
MDA-MB-415	luminal	+	+ /—	—
BT-483	luminal	+	+ /—	—
HCC70	basal	—	—	—
BT-474	luminal	+	+	+
MDA-MB-134VI	luminal	+	—	—
ZR-75-30	luminal	+	—	+
DU4475	basal	—	—	—
MDA-MB-157	basal	—	—	—
Hs 578T	basal	—	—	—
CAMA-1	luminal	+	+ /—	—
HCC202	luminal	—	—	+
BT-20	basal	—	—	—
HCC1954	basal	—	—	+
SK-BR-3	luminal	—	—	+
MDA-MB-231	basal	—	—	—
HCC1937	basal	—	—	—
MDA-MB-468	basal	—	—	—
BT-549	basal	—	—	—
HCC38	basal	—	—	—
HCC1143	basal	—	—	—
MDA-MB-436	basal	—	—	—
HCC1395	basal	—	—	—

Supplemental Table 1: Subtypes of breast cancer cell lines, presented in references 9, 26, 27.

Supplemental Table 2

Gene Symbol	Entrez Gene ID	TOB1		TOB2	
		correlation (r)	p value	correlation (r)	p value
RELB	5971	-0.4184	8.45E-53	-0.1396	0.000001001
NFKBIE	4794	-0.367	3.86E-40	-0.2836	5.72E-24
NFKB2	4791	-0.3379	6.50E-34	-0.1407	8.29E-07
TNIP1	10318	-0.3309	1.63E-32	0.1069	0.0001859
CXCL1	2919	-0.3172	7.20E-30	0.02792	0.3302
CCL20	6364	-0.2771	6.52E-23	-0.1148	0.00005905
SDC4	6385	0.2734	2.55E-22	-0.05254	0.0668
CXCL3	2921	-0.2593	3.70E-20	0.08411	0.003307
CD83	9308	-0.241	1.47E-17	0.01743	0.5434
GFPT2	9945	-0.2382	3.62E-17	0.1316	0.000004057
TRAF1	7185	-0.2308	3.48E-16	-0.02989	0.2973
IL8	3576	-0.2154	2.96E-14	0.08135	0.004497
REL	5966	0.2096	1.48E-13	0.2623	1.32E-20
TNFAIP2	7127	-0.2034	7.66E-13	0.06684	0.01966
PTGS2	5743	-0.1595	2.18E-08	0.2203	7.44E-15
TNFAIP3	7128	-0.1544	6.16E-08	0.1506	1.28E-07
IRF1	3659	-0.1539	6.72E-08	-0.05156	0.07207
CXCL2	2920	-0.1482	2.05E-07	0.1492	1.70E-07
TNFRSF9	3604	-0.1369	0.00000161	-0.06676	0.0198
IL6	3569	-0.1244	0.00001334	0.1736	1.08E-09
PLK2	10769	0.1086	0.0001464	0.1227	0.00001767
FST	10468	-0.08853	0.001985	0.09241	0.001244
NFKB1	4790	-0.08431	0.003234	0.2505	6.94E-19
NFKBIA	4792	-0.06722	0.01896	0.02084	0.4673
GCH1	2643	0.06611	0.02104	-0.133	0.000003212
CTGF	1490	-0.05939	0.03822	0.1334	0.000002981
ZFP36	7538	0.04923	0.08588	0.2804	1.93E-23
IL7R	3575	-0.01387	0.6287	0.1452	3.61E-07

Supplemental Table 2: Gene set from GESA TIAN_TNF_SIGNALING_VIA_NFKB. Genes in red are those shown in Figure 2b and c, and the three with the smallest *p*-values are shown.