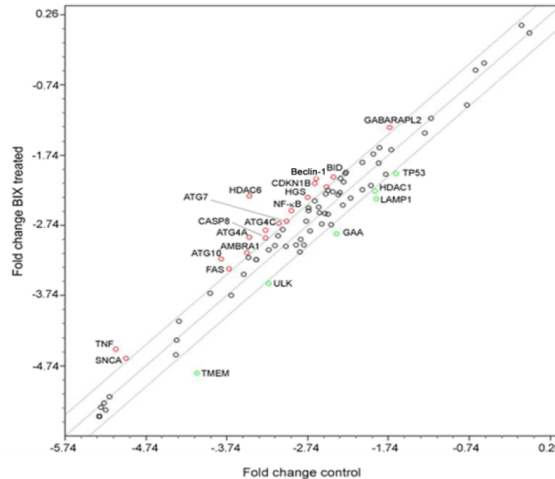


SUPPLEMENTARY FIGURES

**a** Genes up- and down-regulated in BIX 4h treated



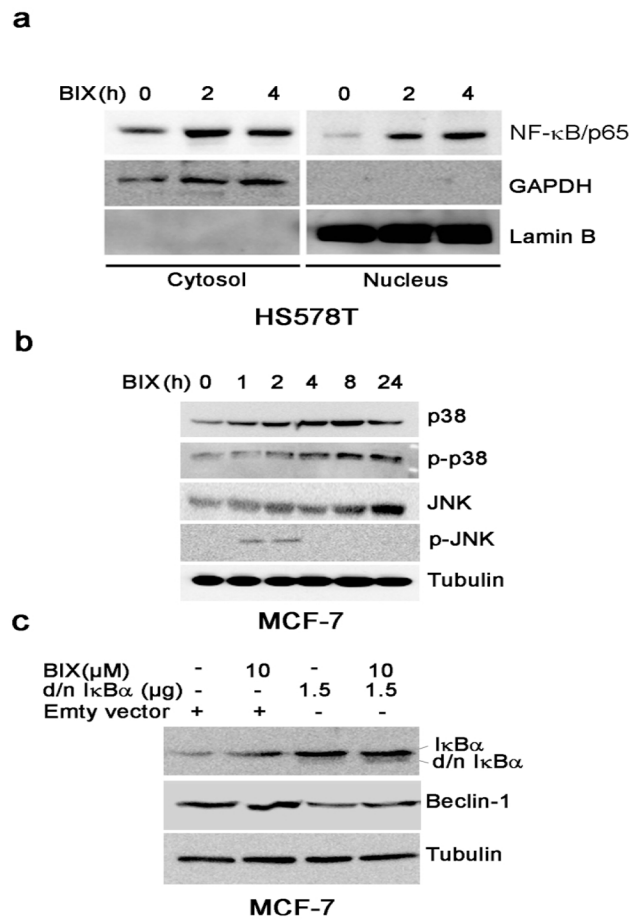
**b** Genes up-regulated or down-regulated in BIX 4h treated

Gene Symbol	Fold Regulation	Gene Symbol	Fold Regulation
HDAC6	13.6737	TMEM74	-5.3889
TNF	4.0652	GAA	-3.0667
ATG10	3.8727	LAMP1	-2.9417
Beclin-1	3.6469	TP53	-2.2815
ATG4A	3.5884	HDAC1	-2.2346
CDKN1B	3.2415	ULK2	-2.1936
ATG4C	2.7895		
NFKB1	2.5432		
HGS	2.4909		
ATG7	2.4116		
GABARAPL2	2.395		
BID	2.308		
AMBRA1	2.308		
SNCA	2.2553		
FAS	2.2501		
CASP8	2.2038		
ATG12	2.0658		
BAD	2.0658		

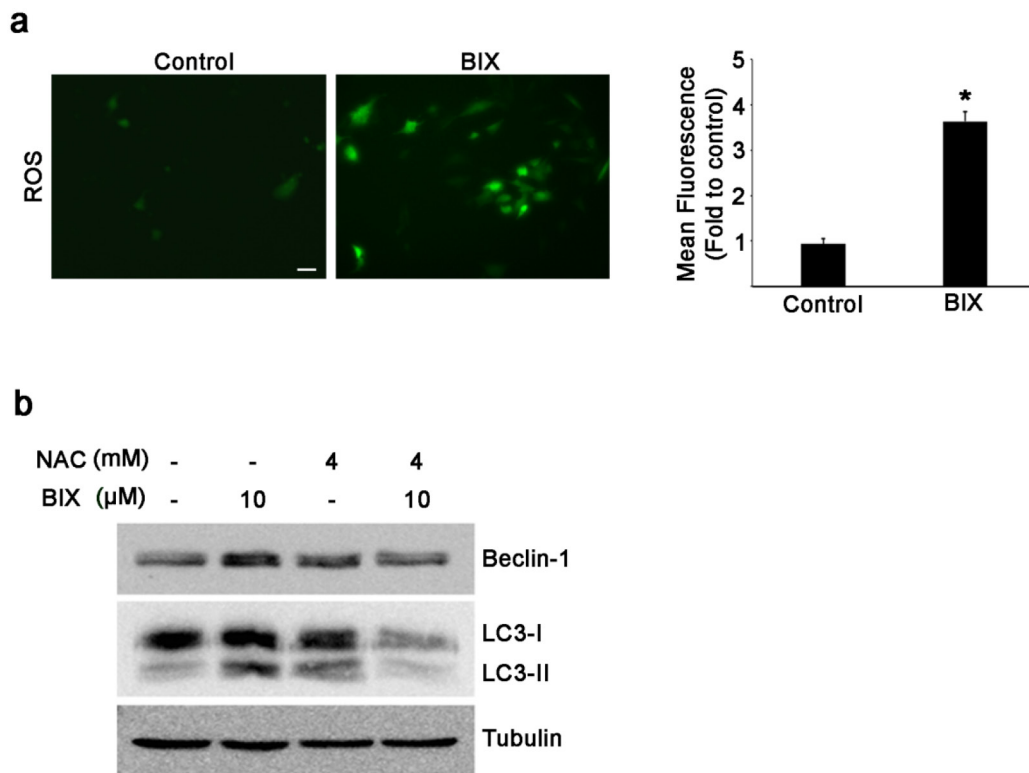
Genes up-regulated in BIX 24h treated

Gene Symbol	Fold Regulation
TNF	65.5426
ATG4B	11.1659
WIP1	3.2967
Beclin-1	3.2363
GABARLAPL1	3.0831
MAP1LC3A	3.0688
MAP1LC3B	2.8832
BNIP3	2.8436
NFKB1	2.684
SNCA	2.2569
ATG4A	2.0294

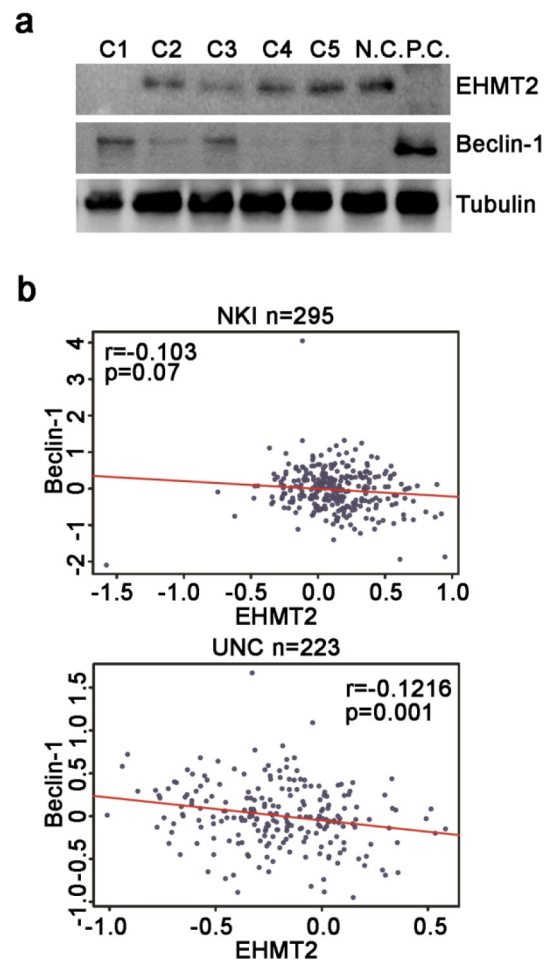
**Supplementary Figure S1: RT2 Profiler PCR array analysis for autophagy related gene expression in BIX-treated MCF7 cells. a. and b.** The expression of 84 key genes involved in autophagy were compared between control and cells without or with 10  $\mu$ M BIX for 4, 24 h. Graph representing fold change in the level of mRNA in cells treated with BIX against untreated cells. Each circle represents an autophagy gene; those with largest fold-changes are indicated in red. The central line represents no changes in expression; above the central line, genes whose expression is increased: below, those with reduced levels. Grey lines indicate 2-fold increase or decrease.



**Supplementary Figure S2: a.** BIX-induced NF-κB nuclear translocation. HS-578T cells were treated with 10 μM BIX-01294 (BIX) for indicated times. The nuclear and cytosolic fractions were separately isolated from each group and blotted with antibodies to NF-κB/65, GAPDH (cytoplasmic marker) and Lamin B (nuclear marker), respectively. **b.** Western blot analysis was performed with the antibodies specified (associated with activation of NF-κB) using the lysates from MCF-7 cells exposed to 10 μM BIX for indicated times. **c.** Inhibition of NF-κB activation and NF-κB-mediated Beclin-1 expression in MCF-7 cells by transfected d/n IκBα (IκBα dominant/negative vector). MCF-7 cells transfected with an empty vector (lane 1,2), and MCF-7 stably transfected with d/n IκBα (lanes 3,4).



**Supplementary Figure S3: BIX induces intracellular ROS-mediated autophagy in HS578T cells.** **a.** HS578T cells were treated with 10  $\mu$ M BIX for 48 h, and stained with 1  $\mu$ M CM-H2DCF-DA for 30 min. The level of ROS was examined using fluorescence microscopy. Scale bar: 50  $\mu$ m. Bar graph represents fold increase of ROS of three independent experiments expressed as means  $\pm$  SEM \*P < 0.05 by one-way ANOVA. **b.** HS578T cells were treated with 10  $\mu$ M BIX for 4 h in the presence or absence of 4 mM NAC. Western blot analysis was performed using the specified antibodies.



**Supplementary Figure S4: Comparison of EHMT2 expression in normal and tumor tissues.** **a.** Comparing the expression of EHMT2 and Beclin-1 in the lysates from tumors of 5 colon cancer patients (C 1–5), and normal colon tissue (N.C.) using Western blot. EHMT2 lysate was used as a EHMT2 positive control (P.C.) **b.** Correlation of Beclin-1 and EHMT2 expression in Netherlands Cancer Institute, NKI (upper box) and University of North Carolina, UNC (lower box) cohort breast cancer patients was estimated by Pearson's correlation test.