

# Supplementary Materials for

## Modulating glycosphingolipid metabolism and autophagy improves outcomes in pre-clinical models of myeloma bone disease

Houfu Leng, Hanlin Zhang, Linsen Li, Shuhao Zhang, Yanping Wang, Selina J Chavda, Daria Galas-Filipowicz, Hantao Lou, Adel Ersek, Emma Morris, Erdinc Sezgin, Yi-Hsuan Lee, Yunsen Li, Ana Victoria Lechuga-Vieco, Mei Tian, Jianqing Mi, Kwee Yong, Qing Zhong, Claire M Edwards, Anna Katharina Simon\*, Nicole J. Horwood\*.

\* To whom correspondence should be addressed:

Prof.Katja Simon: [katja.simon@imm.ox.ac.uk](mailto:katja.simon@imm.ox.ac.uk); Prof.Nicole Horwood: [n.horwood@uea.ac.uk](mailto:n.horwood@uea.ac.uk)

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**SUPPLEMENTARY TABLE****Supplementary Table 1: Baseline MM patient characteristics**

<b>Patient Characteristics (n=9)</b>	
<b>Sex</b>	M 8 (89%) F 1 (11%)
<b>Median Age</b>	61 (41-62)
<b>ISS</b>	
<b>Stage 1</b>	1 (11%)
<b>Stage 2</b>	4 (44%)
<b>Stage 3</b>	1 (11%)
<b>Unknown</b>	3 (34%)
<b>Cytogenetics</b>	
<b>Standard Risk</b>	4 (44%)
<b>High Risk</b>	3 (34%)
<b>Unknown</b>	2 (22%)
<b>Disease response:</b>	
<b>CR/VGPR</b>	5 (56%)
<b>PR/MR</b>	2 (22%)
<b>PD</b>	2 (22%)
<b>Median lines of chemotherapy</b>	2 (22%)
<b>ASCT</b>	7 (78%)
<b>Bony disease at time of diagnosis</b>	8 (89%)
<b>Spinal disease</b>	7 (78%)
<b>Pelvic disease</b>	1 (11%)
<b>Bisphosphonate therapy (zoledronic acid)</b>	8 (89%)
<i>Abbreviations: ISS Stage 1: B2 microglobulin &lt; 3.5mg/L and Albumin&gt;35g/L, ISS Stage 3: B2 microglobulin&gt;5.5mg/L, ISS stage 2: patients not fulfilling criteria for Stage 1 or 3. Adverse cytogenetics defined as per IMWG criteria: [( t(4;14), t(14;16), t(14;20) 1q gain or del 17p]. Disease response defined as per IMWG criteria CR: complete response, VGPR: very good partial response, PR: partial response, MR: minimal response, PD: progressive disease. ASCT: autologous stem cell transplantation</i>	

**Supplementary Table 2: Reagents**

Reagent	Source	Catalog number	Note
RPMI-1640	Sigma-Aldrich	R8758-500ML	
Fetal Bovine Serum	Gibco	10500064	
	Sigma-Aldrich	F2442	
Penicillin-Streptomycin	Sigma-Aldrich	P0781	1:100 dilution
L-Glutamine	Sigma-Aldrich	G7513-100ML	1:100 dilution
Minimum Essential Medium Non-Essential Amino Acids	Gibco	11140050	1:100 dilution
Sodium Pyruvate	Gibco	11360-070	1:100 dilution
Dulbecco's Modified Eagle Medium (DMEM)	Sigma-Aldrich	D6429-500ML	
Alamar blue Cell Viability Reagent	Thermo Fisher	DAL1025	
RBC Lysis Buffer	Sigma	R7757-100ML	
NR12S dye	Bio-Techne Ltd.	7509	
Eliglustat	Selleckchem	S7852	stored at -20°C until diluted in DMSO immediately before use in culture medium (0.1–50 $\mu$ M)
	provided by Genzyme, a Sanofi Company	NA	
Murine sRANK Ligand (RANKL)	peprotech	315-11C-10	
Mouse M-CSF	Miltenyi Biotec	130-101-704	
Recombinant Human sRANK Ligand (E.coli derived)	peprotech	310-01	
Recombinant Human M-CSF Protein	R&D Systems	216-MC	
Minimum Essential Medium (MEM) Alpha	Gibco	12571-063	
Zoledronic acid	Sigma	1724827-150MG	
Iodomethane-d <sub>3</sub>	Merck	176036-5G	
C16 Glucosyl( $\beta$ ) Ceramide (d18:1/16:0); C16GlcCer	Avanti Polar Lipids	860539P-5MG	dissolved in 100% methanol at 37°C and then stored at -20°C until use for cell culture
C16 Lactosyl( $\beta$ ) Ceramide (d18:1/16:0); C16LacCer	Avanti Polar Lipids	860576P-5MG	
C24 Lactosyl( $\beta$ ) Ceramide (d18:1/24:0); C24LacCer	Avanti Polar Lipids	860577P-5MG	
TRAP kit	Sigma-Aldrich	386A	
Methyl Green	Bioenno Lifesciences	003027	
Osmium Tetroxide	Honeywell Fulka	251755-2ml	
D-threo-PDMP (D-PDMP)	Matreya LLC	1756	
	Santa Cruz	sc-280659	

Mouse IgG2b $\kappa$ ELISA Quantification Set	Bethyl Laboratories, Inc	E90-109	
Immunofluorescence staining mounting medium	FluorSave™ Reagent	345789-20ML	
Lysosome Isolation Kit	Sigma-Aldrich	LYSISO1	

**Supplementary Table 3: Antibodies**

Reagent	Source	Clone	Catalog number	Dilution	Validation
Antibodies used for flow cytometry					
CD16/CD32 Monoclonal Antibody (FcR Block)	Thermo Fisher	93	14-0161-85	200	C57BL/KaL wRijHsd and CD45.1 <sup>+</sup> B6.SJL splenocytes and bone marrow cells
Zombie Aqua™ Fixable Viability Kit	BioLegend	NA	423102	400	C57BL/KaL wRijHsd and CD45.1 <sup>+</sup> B6.SJL splenocytes and bone marrow cells
Brilliant Violet 605™ anti-mouse/human CD11b Antibody	BioLegend	M1/70	101237	200	C57BL/KaL wRijHsd bone marrow cells
APC anti-mouse CD115 (CSF-1R) Antibody	BioLegend	AFS98	135510	200	C57BL/KaL wRijHsd bone marrow cells
PE anti-mouse CD117 (c-Kit) Antibody	BioLegend	2B8	105808	200	C57BL/KaL wRijHsd bone marrow cells
Alexa Fluor® 700 anti-mouse/human CD45R/B220 Antibody	BioLegend	RA3-6B2	103232	200	C57BL/KaL wRijHsd bone marrow cells
Brilliant Violet 421™ anti-mouse CD3 Antibody	BioLegend	17A2	100227	200	C57BL/KaL wRijHsd bone marrow cells
Pacific Blue™ anti-mouse	BioLegend	A20	110722	100	CD45.1 <sup>+</sup> B6.SJL bone

CD45.1 Antibody					marrow cells
Alexa Fluor® 700 anti-mouse CD45.2 Antibody	BioLegend	104	109821	100	CD45.1 <sup>+</sup> B6.SJL bone marrow cells
CD11b Monoclonal Antibody, APC	eBioscience	M1/70	17-0112-82	200	CD45.1 <sup>+</sup> B6.SJL bone marrow cells
CD11c Monoclonal Antibody, FITC	eBioscience	N418	11-0114-82	200	CD45.1 <sup>+</sup> B6.SJL bone marrow cells
Antibodies used for Western blot					
LC3	Sigma-Aldrich	Polyclonal	L8918	1000	RAW264.7 cell
TRAF3	Cell Signaling	Polyclonal	4729	1000	C57BL/6J bone marrow cells and RAW264.7 cell
P62 (SQSTM1)	MBL	Polyclonal	PM045	1000	RAW264.7 cell
TRAF6	abcam	Monoclonal	ab33915	1000	C57BL/6J bone marrow cells
IκBα	Cell Signaling	44D4	4812S	1000	C57BL/6J bone marrow cells
LAMP1	abcam	Polyclonal	ab24170	1000	RAW264.7 cell
βActin	Cell Signaling	8H10H10	3700	5000	C57BL/6J bone marrow cells and RAW264.7 cell
IRDye 800CW Donkey Anti-Rabbit IgG Secondary Antibody	LI-COR		926-32213 RRID AB_621848	10,000	validated by manufacturer
IRDye 680LT Donkey Anti-Mouse IgG Secondary Antibody	LI-COR		926-68022 RRID AB_10715072	20,000	validated by manufacturer
Antibodies used for confocal microscopy					
LAMP2	Santa Cruz Biotechnology	H4B4	sc18822	200 (Sup. Fig. 6a)	U2OS cell
Anti-LAMP2 antibody	abcam	GL2A7	ab13524	200 (Figure	RAW264.7 cell

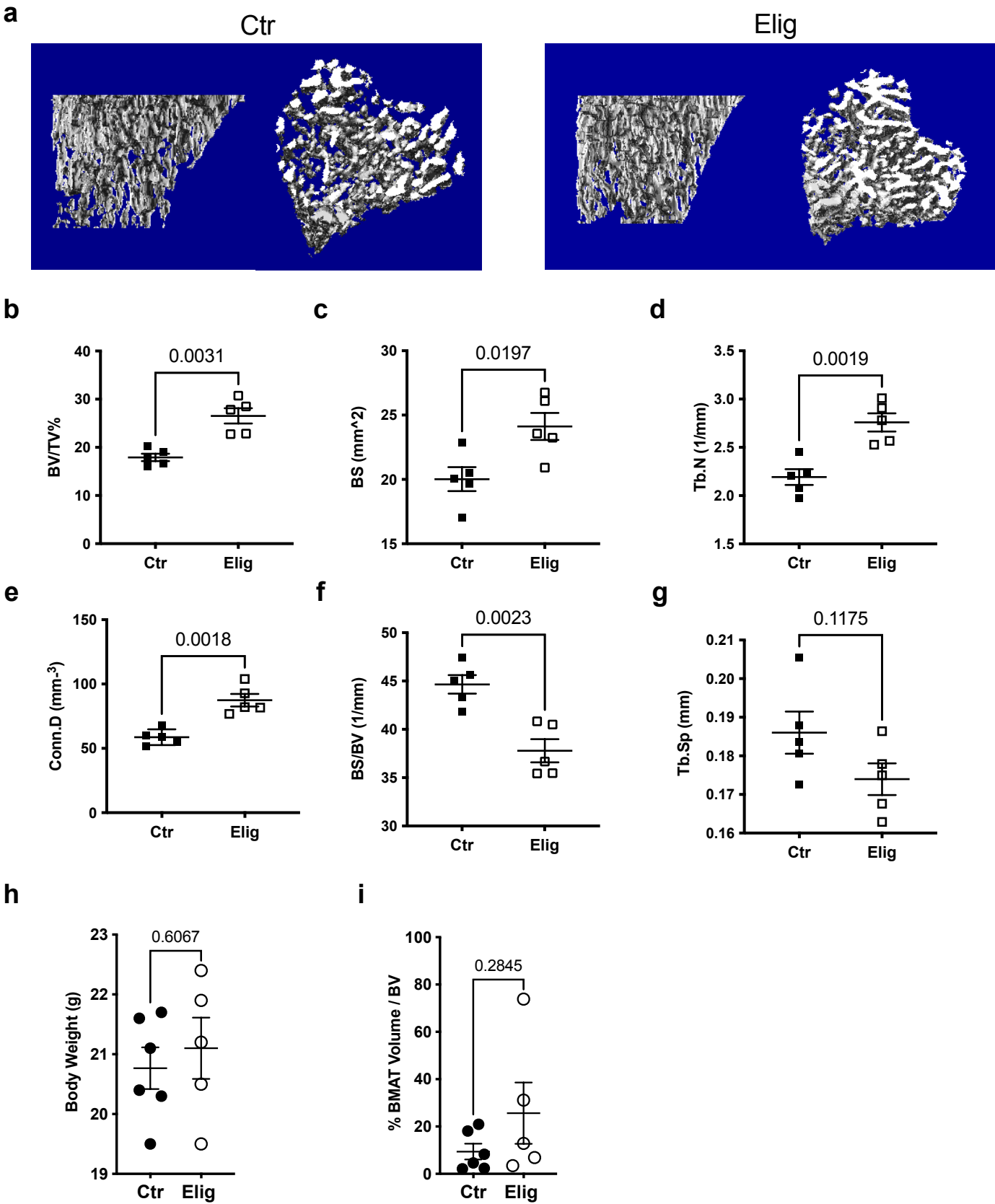
				60 and Sup. Fig. 6c)	
Cy3-AffiniPure Goat Anti-Mouse IgG (H+L), Secondary	Jackson ImmunoRes earch	Polyclonal	115-165-003	500 (Sup. Fig. 6a)	U2OS cell
CoraLite®488- conjugated TRAF3 Monoclonal antibody	Proteintech	1E3F4	CL488-66310	150 (Figure 6o and Sup. Fig. 6c)	RAW264.7 cell
Alexa Fluor™ 647 chicken anti- rat IgG (H+L)	Invitrogen™ M	Polyclonal	A21472	500 (Figure 6o)	RAW264.7 cell
DAPI	Thermo Scientific		62248	2µg/ml	U2OS cell and RAW264.7 cell

**Supplementary Table 4:** TaqMan® probes

Gene	Source	Identifier
<i>traf3</i>	Thermo Fisher	Mm00495752_m1
<i>gapdh</i>	Thermo Fisher	Mm99999915_g1

## **SUPPLEMENTARY FIGURES**

Supplemental Figure 1



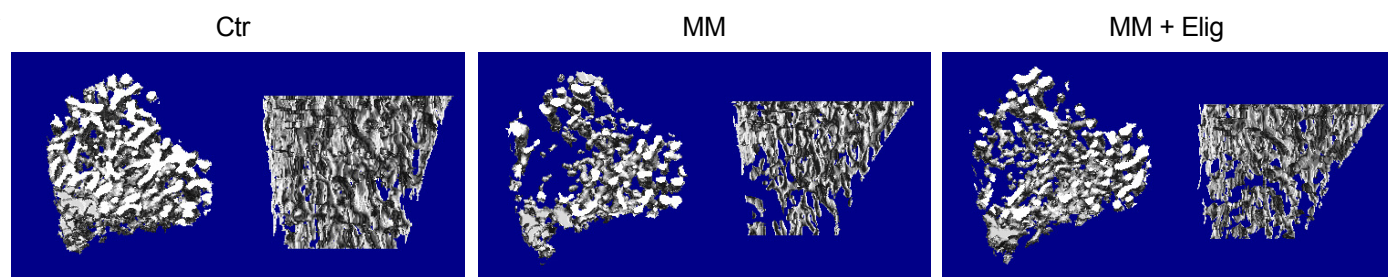


**Supplementary Figure 1.** Eliglustat increases trabecular bone in healthy male mice by inhibiting OC *in vivo*.

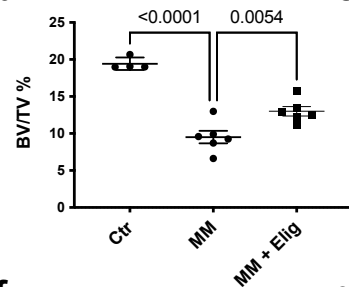
(a) Representative micro-CT reconstruction images of tibiae from 8-week-old male C57BL/6J mice treated with normal chow (Ctr, n=5 biologically independent animals) or eliglustat (Elig, n=5 biologically independent animals) chow for 19 days. (b-g) Micro-CT analysis of tibiae: BV/TV, BS, Tb.N, Conn.D, BS/BV and Tb.Sp (Ctr, n=5 and Elig, n=5 biologically independent animals). (h, i) The body weight (h) and BMAT volume (i) of the mice from respective group (Ctr, n=5 and Elig, n=5 biologically independent animals). Data are presented as mean values  $\pm$  SEM. Exact *p* values are depicted in the figure. Statistical analysis was performed using unpaired two-tailed Student's *t* test. Source data are provided as a Source Data file.

# Supplemental Figure 2

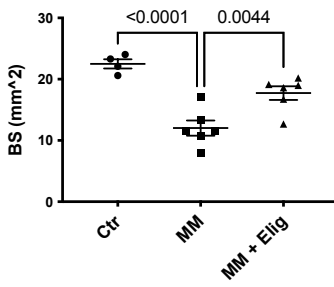
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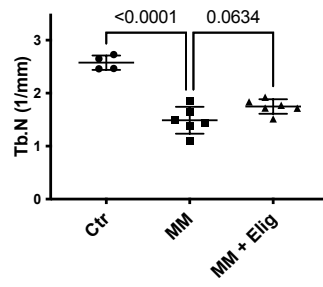
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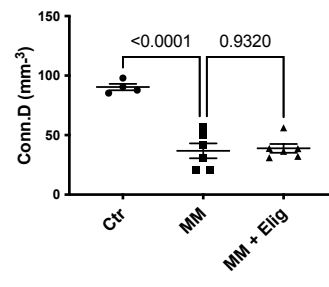
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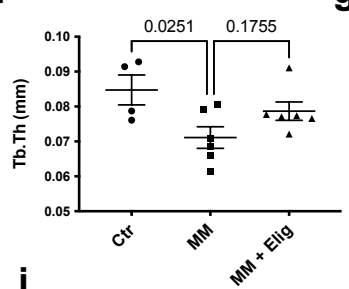
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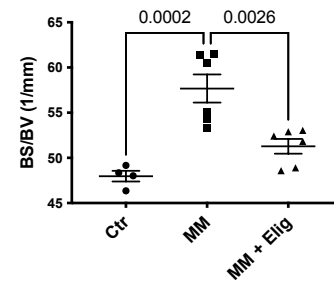
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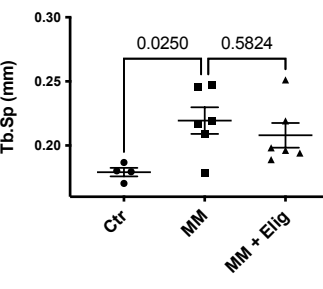
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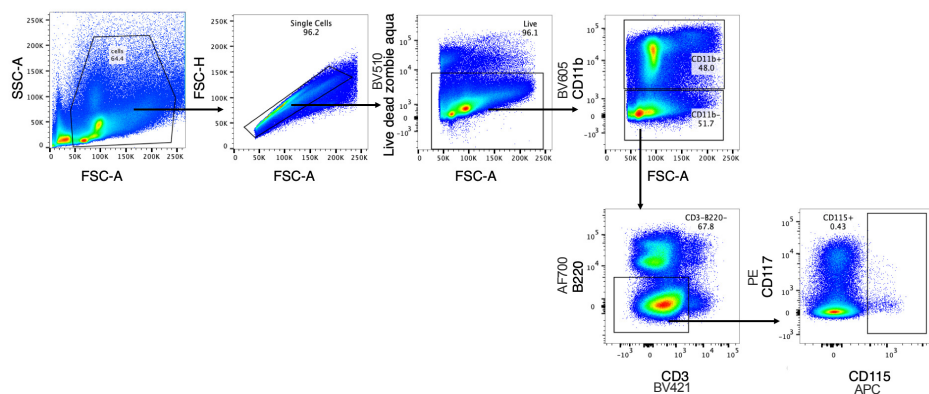
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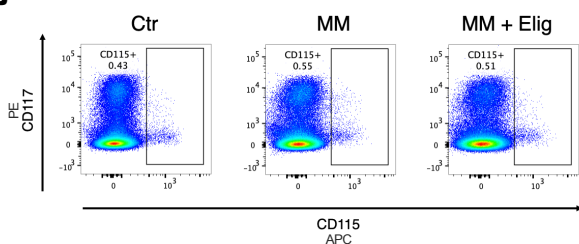
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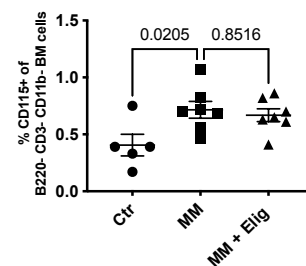
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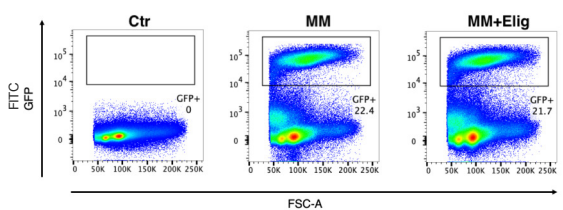
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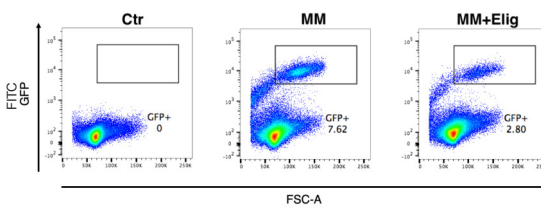
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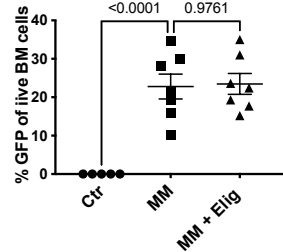
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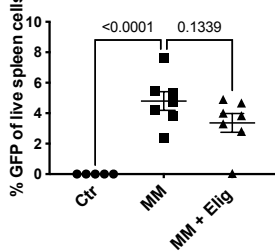
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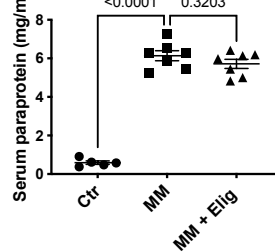
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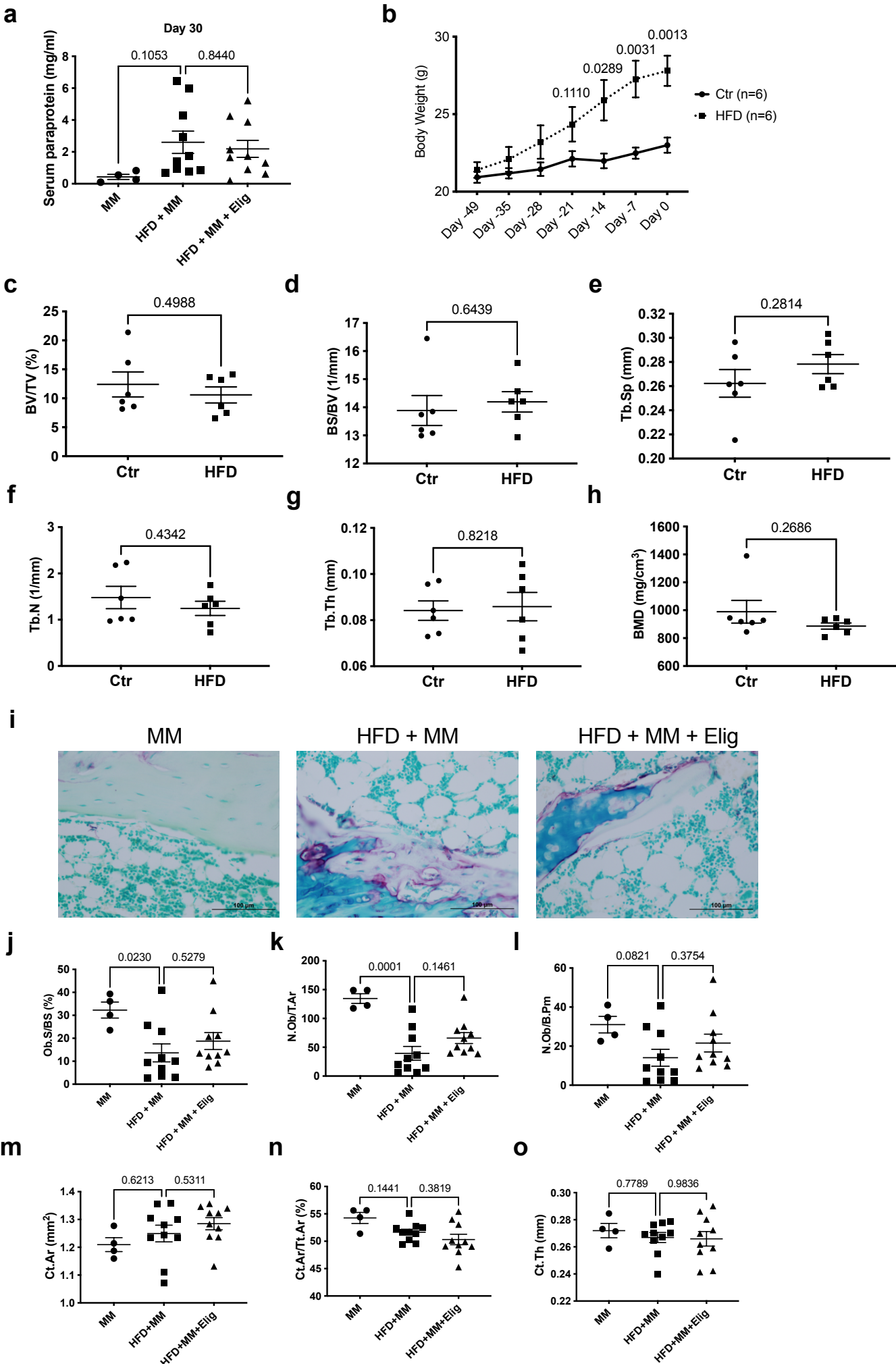
**Supplementary Figure 2.** Eliglustat ameliorates 5TGM1-GFP MM cell-induced bone disease in male mice but has no effect on OC progenitors or MM tumour burden.

5TGM1-GFP MM cells were injected to 8-week-old male C57BL/KaLwRijHsd mice to generate the MM model. Eliglustat chow (150 mg/kg/day) was administered from day 4 post tumour injection (until day 23). **(a)** Representative micro-CT reconstruction images of tibiae trabecular bones from naive control (Ctr, *n*=4 biologically independent animals), MM mice with normal chow (MM, *n*=6 biologically independent animals), or MM mice with eliglustat chow (MM+Elig, *n*=6 biologically independent animals). **(b-h)** Tibiae trabecular bone parameters were assessed: BV/TV, BS, Tb.N, Conn.D, Tb.Th, BS/BV and Tb.Sp (Ctr, *n*=4; MM, *n*=6 and MM+Elig, *n*=6 biologically independent animals).

**(i-k)** Flow cytometry gating strategy of the OC progenitors (CD11b<sup>-</sup> B220<sup>-</sup> CD3<sup>-</sup> CD115<sup>+</sup>) in BM cells **(i)** from female C57BL/KaLwRijHsd mice, either untreated controls (Ctr), MM mice with normal chow (MM), or MM mice with eliglustat chow (MM+Elig). Representative plots of OC progenitors (CD11b<sup>-</sup> B220<sup>-</sup> CD3<sup>-</sup> CD115<sup>+</sup>) from the indicated female groups **(j)** and the quantified percentage **(k)** (Ctr, *n*=5; MM, *n*=7 and MM+Elig, *n*=7 biologically independent animals). **(l-o)** Female mice tumour burden in BM **(l)** and spleen **(m)** was assessed by quantifying GFP<sup>+</sup> cells using flow cytometry **(n-o)** (Ctr, *n*=5; MM, *n*=7 and MM+Elig, *n*=7 biologically independent animals). **(p)** Serum paraprotein IgG2b $\kappa$  secreted by 5TGM1-GFP cells was quantified using ELISA (Ctr, *n*=5; MM, *n*=7 and MM+Elig, *n*=7 biologically independent animals).

Data are presented as mean values  $\pm$  SEM. Exact *p* values are depicted in the figure. Statistical analysis was performed using One-way ANOVA. Source data are provided as a Source Data file.

Supplemental Figure 3



**Supplementary Figure 3.** Eliglustat did not alter OB and cortical bone in HFD induced MGUS condition.

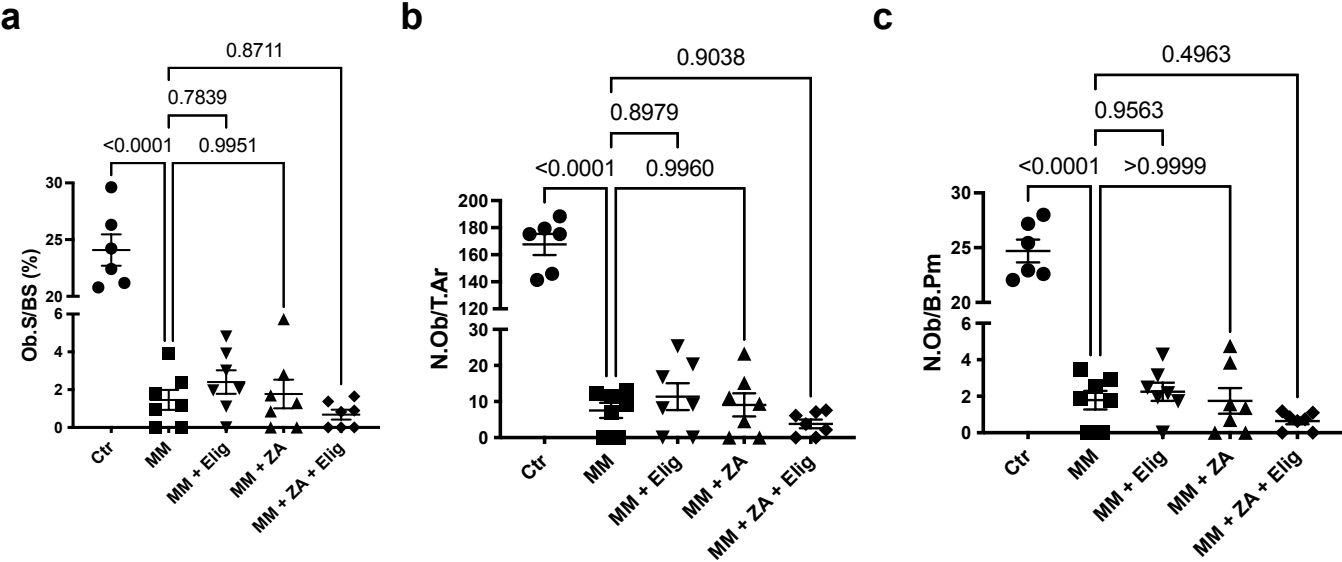
(a) Eliglustat does not alter serum paraprotein levels in MM, HFD+MM and HFD+MM+Elig groups measured at day 30 post MM cell injection (MM,  $n=4$ ; HFD+MM,  $n=10$  and HFD+MM+Elig,  $n=10$  biologically independent animals). Statistical analysis was performed using One-way ANOVA for **a**.

(b) C57BL6J mice were divided into normal diet group (Ctr,  $n=6$  biologically independent animals) or HFD group (HFD,  $n=6$  biologically independent animals), body weight changes were observed over 49 days. (c-h) Micro-CT analysis parameters of tibiae: BV/TV, BS/BV, Tb.Sp, Tb.N, Tb.Th and BMD for mice on Ctr diet and HFD (Ctr,  $n=6$  and HFD,  $n=6$  biologically independent animals). Statistical analysis was performed using unpaired two-tailed Student's *t* test for **b-h**.

(i-l) Histology images and bone histomorphometry analysis for OB including Ob.S/BS, N.Ob/T.Ar and N.Ob/B.Pm in MM, HFD+MM and HFD+MM+Elig groups (MM,  $n=4$ ; HFD+MM,  $n=10$  and HFD+MM+Elig,  $n=10$  biologically independent animals). The result is representative of two independent experiments. (m-o) Cortical bone parameters including Ct.Ar (cortical area), Ct.Ar/Tt.Ar (cortical area over total area) and Ct.Th (cortical thickness) for MM, HFD+MM, HFD+MM+Elig groups (MM,  $n=4$ ; HFD+MM,  $n=10$  and HFD+MM+Elig,  $n=10$  biologically independent animals). Statistical analysis was performed using One-way ANOVA for **j-o**.

Data are presented as mean values  $\pm$  SEM. Exact *p* values are depicted in the figure. Source data are provided as a Source Data file.

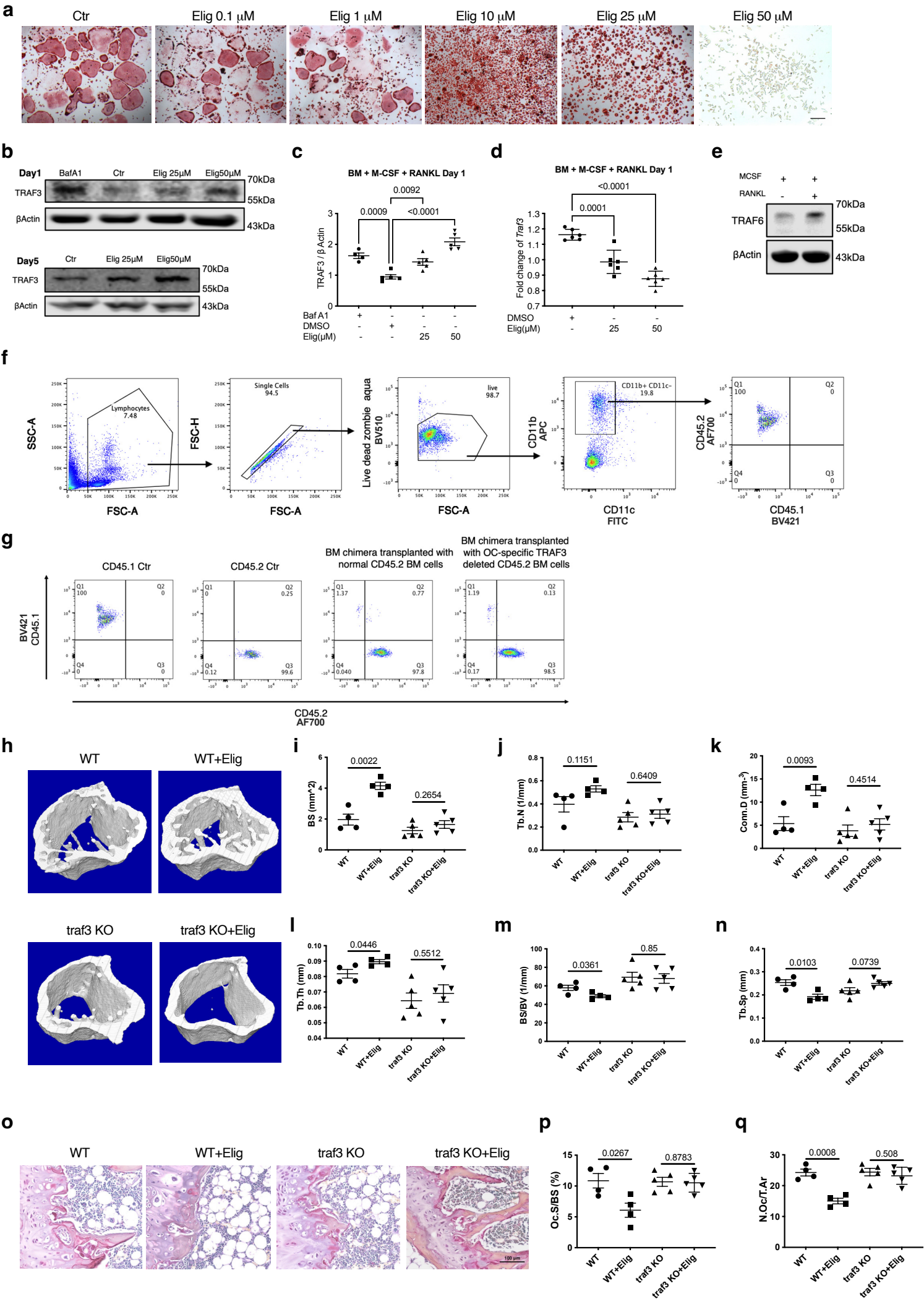
Supplemental Figure 4



**Supplementary Figure 4.** Eliglustat combined with ZA does not enhance OB number or area.

(a-c) Bone histomorphometric analysis for OB parameters including Ob.S/BS, N.Ob/T.Ar and N.Ob/B.Pm were based on morphology in paraffin sections stained with TRAP and methyl green and quantified using *Osteomeasure software* (Ctr,  $n=6$ ; MM,  $n=7$ ; MM+Elig,  $n=7$ ; MM+ZA,  $n=7$ ; MM+ZA+Elig,  $n=7$  biologically independent animals). Data are presented as mean values  $\pm$  SEM. Statistical analysis was performed using One-way ANOVA. Exact  $p$  values are depicted in the figure. Source data are provided as a Source Data file.

Supplemental Figure 5





**Supplementary Figure 5.** Eliglustat inhibits OC formation via TRAF3.

(a) 8-week-old C57BL/6J mouse BM cells were treated with M-CSF and RANKL to form OCs in 96-well plates. Different doses of Eliglustat (0.1, 1, 10, 25, 50  $\mu$ M) were added and OCs were identified by TRAP staining on day 6. Scale bar represents 200 $\mu$ m.

(b-c) TRAF3 protein levels were quantified by Western blot on day 1 of OC differentiation (similar pattern was observed throughout the differentiation period including on day 5). BafA1 (10nM) was added 2 hours before protein harvest; BafA1, n=4; DMSO control, n=5; 25 $\mu$ M Elig, n=5; 50 $\mu$ M Elig, n=5 biologically independent samples. The result is representative of four independent experiments. Statistical analysis was performed using One-way ANOVA for c.

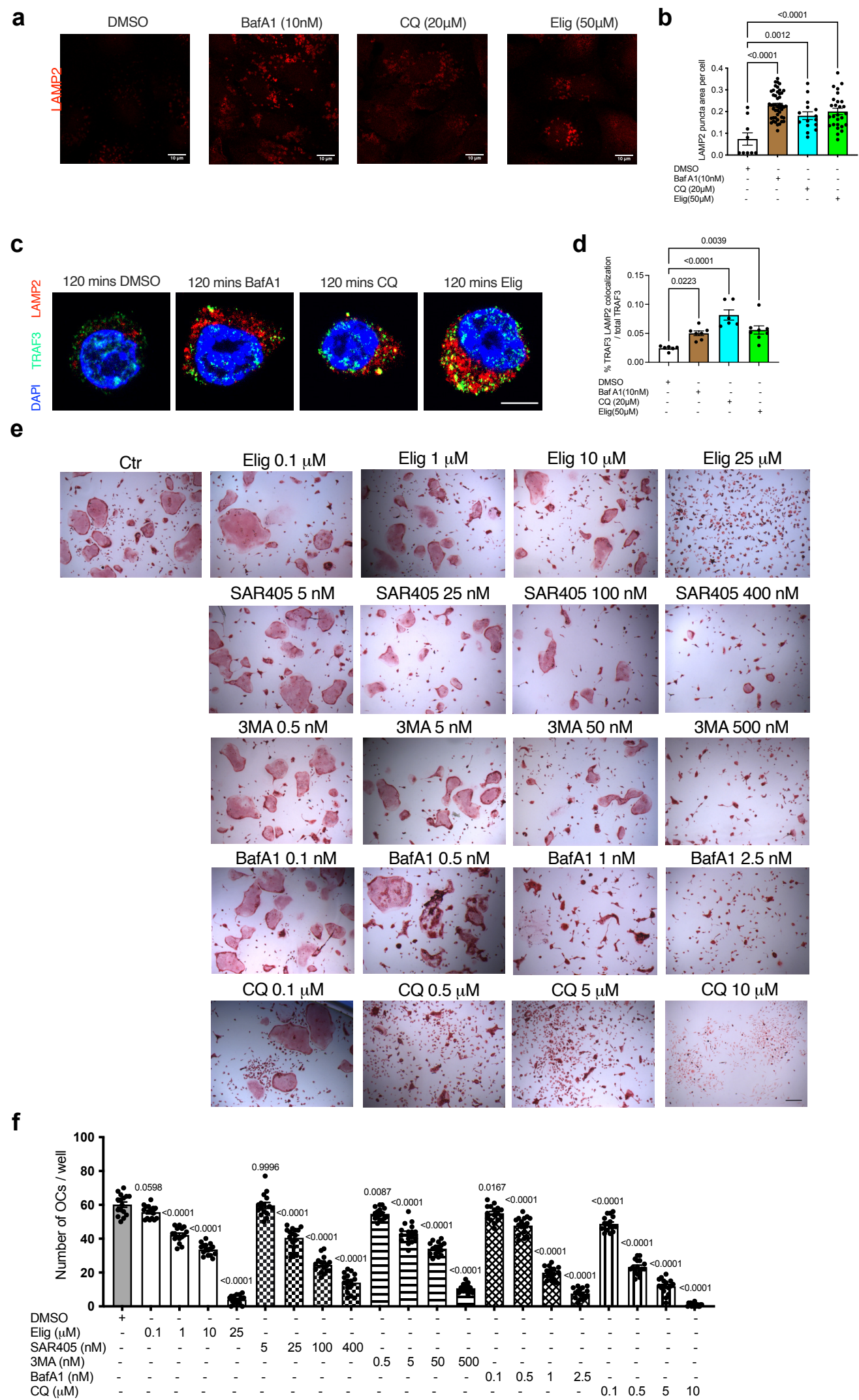
(d) *traf3* mRNA levels on day 1 were quantified by qRT-PCR; DMSO control, n=6; 25 $\mu$ M Elig, n=6; 50 $\mu$ M Elig, n=6 biologically independent samples. Statistical analysis was performed using One-way ANOVA for d.

(e) 50ng/ml RANKL increased TRAF6 protein level in primary BM cells during OC formation after 2 hours treatment. The result is representative of three independent experiments.

(f, g) CD45.1<sup>+</sup> recipient mice were lethally irradiated and reconstituted with CD45.2<sup>+</sup> wildtype BM cells or CD45.2<sup>+</sup> OC-specific TRAF3-deleted BM cells. Gating strategy for checking the reconstitution efficacy of the chimeric mice by evaluating the peripheral blood CD45.2<sup>+</sup> myeloid population (f). Representative flow cytometry plots showing the distribution of CD45.1<sup>+</sup>/CD45.2<sup>+</sup> of myeloid cells in peripheral blood from an unirradiated CD45.1 mouse, unirradiated CD45.2 mouse, irradiated CD45.1<sup>+</sup> mouse transplanted with wildtype CD45.2<sup>+</sup> BM cells and CD45.1<sup>+</sup> mouse transplanted with CD45.2<sup>+</sup> OC-specific TRAF3-deleted BM cells (g). (h) Micro-CT reconstruction images of WT, WT+Elig, *traf3* KO and *traf3* KO+Elig mice. (i-n) Micro-CT analysis parameters of tibiae including: BS, Tb.N, Conn.D, Tb.Th, BS/BV and Tb.Sp (WT, n=4; WT+Elig, n=4; *traf3* KO, n=5; *traf3* KO + Elig, n=5 biologically independent animals). (o) Representative TRAP/0.2% methyl green stained tibial sections showing red OCs on the

endocortical bone surface from each group. The result is representative of two independent experiments. (**p, q**) Bone histomorphometry parameters including Oc.S/BS and N.Oc/T.Ar (WT,  $n=4$ ; WT+Elig,  $n=4$ ; traf3 KO,  $n=5$ ; traf3 KO + Elig,  $n=5$  biologically independent animals). Statistical analysis was performed using unpaired two-tailed Student's *t* test for **i-n** and **p, q**. Data are presented as mean values  $\pm$  SEM. Exact *p* values are depicted in the figure. Source data are provided as a Source Data file.

Supplemental Figure 6



**Supplementary Figure 6.** Eliglustat accumulates LAMP2 and TRAF3 in a cell, in addition, autophagy inhibitors including SAR405, 3MA, BafA1 and CQ prevent OC formation.

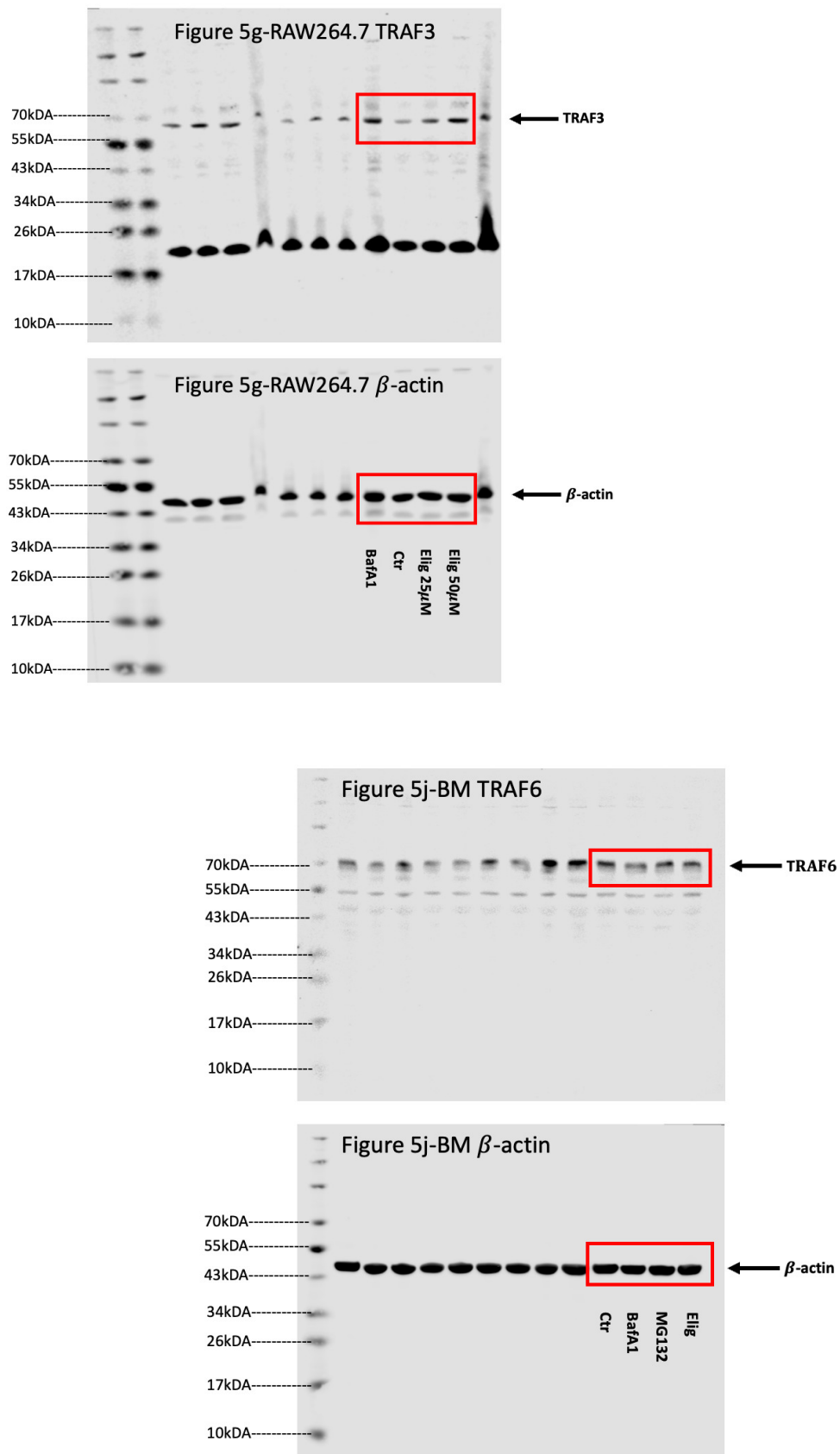
**(a, b)** Confocal images **(a)** and quantification **(b)** of the LAMP2 puncta area per eliglustat-treated U2OS cell (arbitrary units). CQ and BafA1 served as the positive control.  $n=9$  cells for DMSO control,  $n=47$  cells for BafA1,  $n=15$  cells for CQ and  $n=27$  cells for Elig group. The result is representative of three independent experiments.

**(c, d)** Confocal images of TRAF3 and LAMP2 in Pre-OCs derived from RAW264.7 cells were treated with BafA1, CQ and eliglustat for 120 minutes, Scale bar represents  $10\mu\text{m}$ ,  $n\geq 4$  **(c)**; the percentage of TRAF3 and LAMP2 colocalization is quantified **(d)**.  $n=6$  cells for DMSO control,  $n=7$  cells for BafA1,  $n=6$  cells for CQ and  $n=8$  cells for Elig group. The result is representative of three independent experiments.

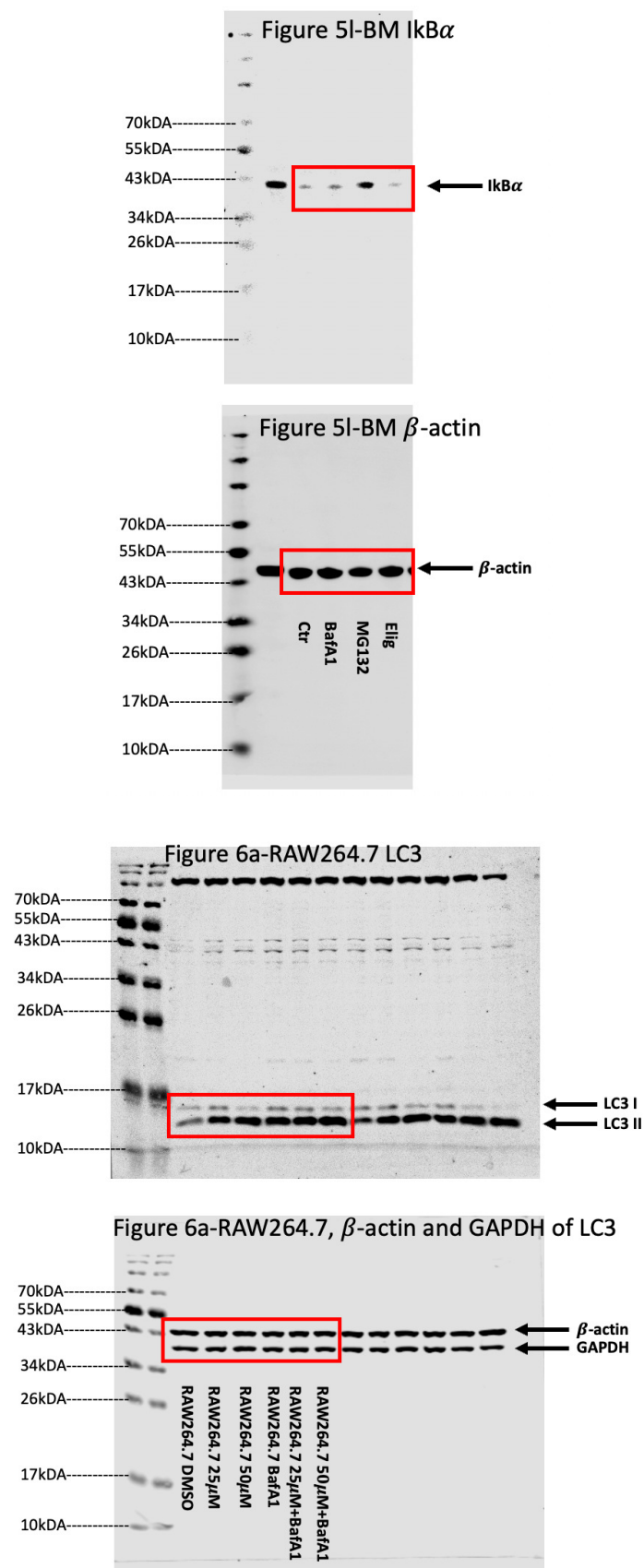
**(e)** BM cells were differentiated into OCs with M-CSF and RANKL. Different doses of eliglustat ( $0.1, 1, 10, 25\ \mu\text{M}$ ), SAR405 ( $5, 25, 100, 400\ \text{nM}$ ), 3MA ( $0.5, 5, 50, 500\ \text{nM}$ ), BafA1 ( $0.1, 0.5, 1, 2.5\ \text{nM}$ ) and CQ ( $0.1, 0.5, 5, 10\ \mu\text{M}$ ) were present throughout the culture period and OCs were identified by TRAP staining on day 5. Scale bar represents  $200\mu\text{m}$ . **(f)** Number of OCs per well was quantified.  $n=16$  biologically independent samples examined over 4 independent experiments.

Data are presented as mean values  $\pm$  SEM. Exact  $p$  values are depicted in the figure. Statistical analysis was performed using One-way ANOVA. Source data are provided as a Source Data file.

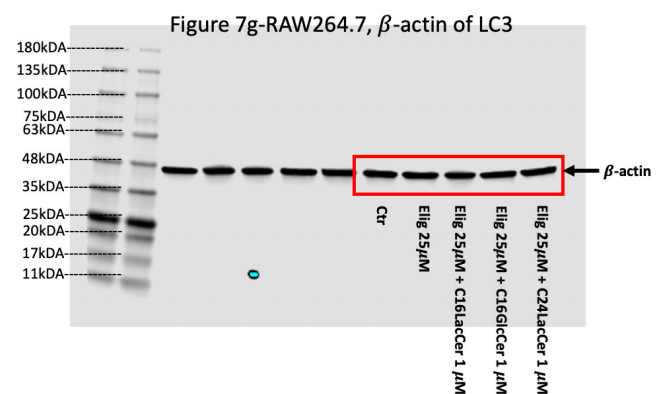
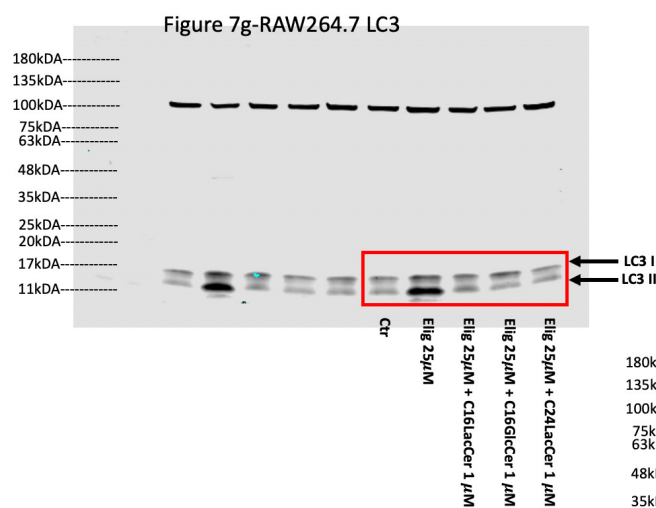
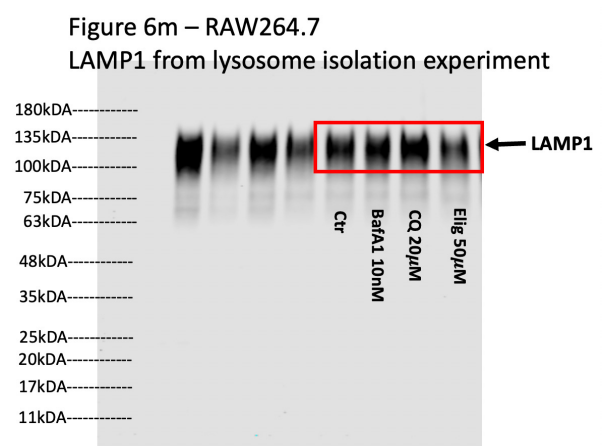
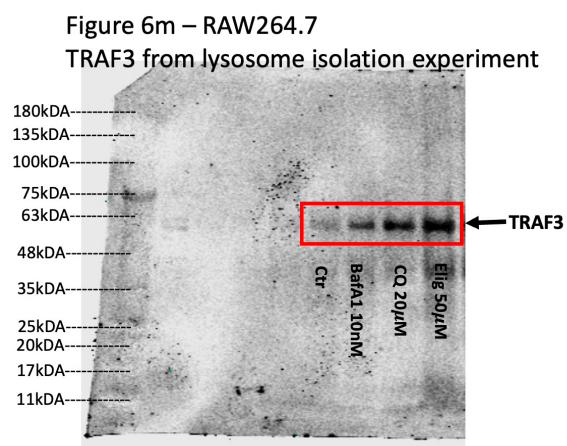
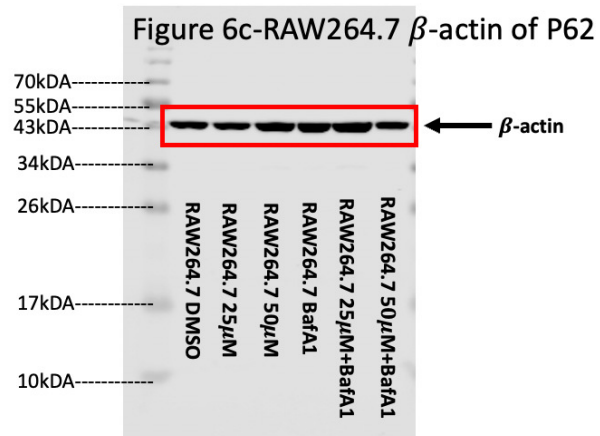
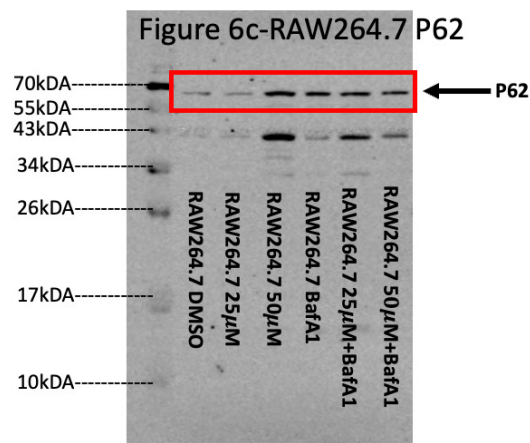
**Supplementary Figure 7.** Western blot full scans.



**Supplementary Figure 7.** Western blot full scans (continued).

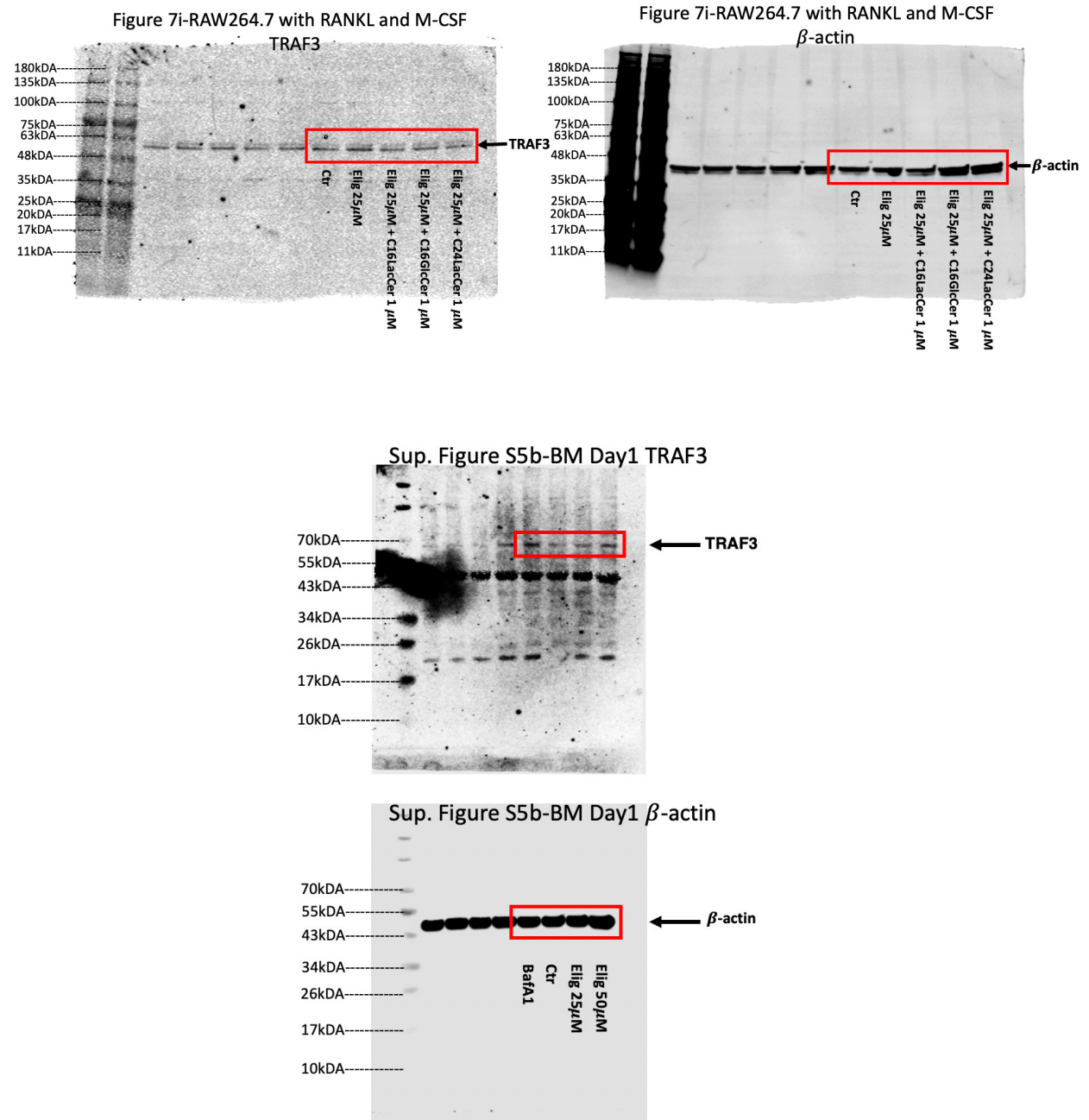


**Supplementary Figure 7.** Western blot full scans (continued).





**Supplementary Figure 7.** Western blot full scans (continued).





**Supplementary Figure 7.** Western blot full scans (continued).

