

1 **Interleukin-1/inhibitory kappa B kinase epsilon-mediated increases in glycolysis**
2 **augment epithelial effector function and promote house dust mite-induced allergic**
3 **airways disease**

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38 **List of abbreviations:**

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40 NECs: Nasal epithelial cells

41 MTE cells: Mouse tracheal epithelial cells

42 WT: Wild-type

43 Rn: Newtonian resistance

44 G: Tissue resistance

45 H: Elastance

46 BAL: Bronchoalveolar lavage

47 ELISA: Enzyme-linked immunosorbent assay

48 qRT-PCR: Quantitative reverse transcription-polymerase chain reaction

49 ECAR: Extracellular acidification rate

50 OCR: Oxygen consumption rate

51 BSA: Bovine serum albumin

52 MAC: Macrophages

53 NEU: Neutrophils

54 EOS: Eosinophils

55 LYM: Lymphocytes

56 KC: Chemokine (C-X-C motif) ligand 1

57 CCL20: Chemokine (C-C motif) ligand 20;

58 TSLP: Thymic stromal lymphopoietin

59 GM-CSF: Granulocyte-macrophage colony-stimulating factor

60 Mct4: Monocarboxylate transporter 4

61 Pfk1: Phosphofructokinase, liver type

62 Hk2: Hexokinase 2

63 Glut: Glucose transporter

64 Pkm2: Pyruvate kinase isoenzyme type M2

65 Ldha: Lactate dehydrogenase A

66 Pgm1: Phosphoglucomutase 1

67 Pdk1: Pyruvate dehydrogenase kinase 1

68 Tpi1: Triosephosphate isomerase 1

69 Eno1: Enolase 1

70 Pgk1: Phosphoglycerate kinase 1

71 Gpi: Glucose-6-phosphate isomerase
72 Pfkfb3: 6-Phosphofructo-2-Kinase/Fructose-2,6-Biphosphatase 3
73 Mct1: Monocarboxylate transporter 1
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123 **Supplemental Materials and Methods**

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125 **Sputum induction**

126 Sputum was induced and processed, as described previously (1-3). Prior to sputum
127 induction, subjects inhaled 400 µg salbutamol using a metered-dose inhaler (+spacer).
128 Sputum was induced using an ultrasonic nebulizer (ultra-Neb 2000, Devilbiss; output set at
129 0.9 ml/min). Subjects inhaled hypertonic saline (NaCl 5%) when FEV1 post salbutamol was
130 $\geq 65\%$ predicted and isotonic saline (NaCl 0.9%) when FEV1 was $<65\%$ predicted. The
131 aerosol was inhaled for three consecutive periods of 5 min. FEV1 was monitored every 5
132 minutes for safety reasons, and when FEV1 dropped to 80% of the post-bronchodilator
133 values, the induction procedure was stopped. The whole sputum was weighted and three
134 volumes of PBS were added. After homogenizing by manual agitation for 30 sec and
135 centrifugation (800 g) for 10 min at 4°C, the cell pellet and supernatant were separated.
136 Cells were treated with Sputolysin® 0.1% (Calbiochem, Germany), washed with PBS and
137 resuspended in 1 ml. Total cell counts, % squamous cells and cell viability (trypan blue
138 staining) were determined with a manual hemocytometer. Sputum cell differentials were
139 determined by counting 500 cells non squamous cells on Cytospin samples that were
140 stained with RAPI-DIFF II stain (Atom Scientific, Manchester, United Kingdom).

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142 **Cell culture and treatments**

143 Primary human nasal epithelial cells (NECs) were isolated from 6 healthy volunteers
144 and 6 patients with allergic rhinitis and asthma by gentle stroking of the inferior turbinate
145 surface with a Rhino-Probe curette and cultured as recently described (4) in bronchial
146 epithelial cell growth medium (Lonza). Atopy was confirmed by positive skin tests and
147 elevated serum IgE (>100 IU/ml), and asthma was diagnosed by physicians, confirmed by
148 positive response to bronchodilator (≥ 200 cc and 12% improvement in FEV1 and/or FVC) or
149 a positive methacholine challenge test (PC20 < 8 mg/ml), and had rhinitis with a sinonasal
150 questionnaire (SNQ) score (5) of greater than 1. Healthy volunteers had no history of rhinitis

151 or asthma, negative skin tests, negative methacholine challenge tests, and a SNQ score of
152 less than 1. For experiments, NECs were plated on collagen-coated 12-well plates at a
153 density of 2×10^5 cells/well and cultured in a 1:1 mixture of bronchial epithelial cell basic
154 medium and DMEM-H with SingleQuot supplements (Cambrex), bovine pituitary extract (13
155 mg/ml), bovine serum albumin (1.5 μ g/ml), and nystatin (20 units). Following 2 h starvation in
156 basal medium, NECs were treated with 50ug/ml of HDM D. pteronyssinus; GREER, Lenoir,
157 NC; 144.9 endotoxin units/mg protein, lot 290903) for 24 h. Primary mouse tracheal
158 epithelial (MTE) cells were isolated from wild-type (WT) C57BL/6 mice or C57BL/6 mice
159 lacking the Inhibitor of κ B kinase ϵ gene (referred to herein as *Ikkbe*^{-/-}) and cultured as
160 previously described (6, 7). After reaching confluence, MTE cells were incubated for 16 h in
161 serum-free medium. Cells were stimulated with IL-1 α (10 ng/mL), IL-1 β (10 ng/mL), IL-6 (10
162 ng/mL), IL-13 (5 ng/mL), IL-33 (5 ng/mL), TGF- β 1 (5 ng/mL), TNF α (5 ng/mL), IL-17 (20
163 ng/mL), lipopolysaccharide (LPS, 1 μ g/mL), or HDM (10 μ g/mL or 50 μ g/mL as indicated in
164 the Figure Legend) for 2 or 24 hr.

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166 To address the importance of glycolysis or IKK ϵ /TBK1 in IL-1 β induced pro-inflammatory
167 responses in MTE cells, cells were pre-treated with the hexokinase inhibitor, 2-
168 Deoxyglucose (2-DG, 10 mM) for 1 h prior to exposure to IL-1 β for 24 h. Alternatively, cells
169 were incubated with the lactate dehydrogenase A inhibitor, oxamate (10 mM), or the
170 IKK ϵ /TBK1 inhibitor, Amlexanox (10-100 μ M, Tocris) overnight, followed by the stimulation
171 IL-1 β (10 ng/mL) for 24 h. To determine whether IL-1 β -induced glycolysis or IKK ϵ /TBK1
172 augmented the subsequent response to HDM, in select experiments, cells were washed post
173 inhibitor/IL-1 β treatment, incubated with DMEM/F12 medium for 2 h before stimulation with
174 HDM for an additional 2 hours according the schematic illustrations shown in the relevant
175 figures. To address the role of glycolysis in IL-1 α -dependent pro-inflammatory responses,
176 MTE cells were pre-incubated with 10 mM 2-DG for 1 h or 10 mM oxamate for 16 h prior to
177 exposure to IL-1 α for 24 h and subsequent assessment of pro-inflammatory mediators in

178 supernatants. Note that absolute values of KC vary between studies due to freezing of some
179 supernatants.

180 **Mouse studies**

181 Age-matched, 8- to 12-week-old mice were used (The Jackson Laboratory, Bar
182 Harbor, ME) for all experiments. Wild-type (WT, C57BL6/NJ), *Rag*^{-/-} (C57BL6/J), or *Ikkbe*^{-/-}
183 (C57BL6/J) mice along with their strain-matched controls were sensitized (Days 1 and 8),
184 challenged (Days 15-19), and rechallenged (Days 29, 32, 36, and 39) with HDM extract
185 containing 10 µg protein (GREER, Lenoir, NC; 144.9 endotoxin units/mg protein, lot 290903)
186 as shown in Figure 1A. Mice were euthanized and analyzed at different end points at Days 1
187 (2 h post HDM sensitization), 2 (24 h post HDM sensitization), 20 (24 h post the last
188 challenge of 5 consecutive HDM challenge), 29 (2 h post the first HDM re-challenge), 30 (24
189 h post the first HDM rechallenge), and/or 40 (24 h post the last HDM rechallenge). The
190 control group was subjected to saline as a vehicle control. In the *Ldha* siRNA knockdown
191 studies, WT C57BL6/NJ mice were anesthetized with isoflurane and subjected to 10 mg/kg
192 of siRNA targeting *Ldha* or scrambled small interfering siRNA oropharyngeally on days 26,
193 30, 33, and 37 post-initiation of the HDM exposure regimen, and mice were harvested in day
194 40 (24h post the last HDM re-challenge). In the IL-1 neutralization experiments, WT
195 C57BL6/J mice were challenged with HDM on days 15, 16, 17, and 18, and analyzed on day
196 20. Mice received 5 mg/kg of IL-1 Trap (Regeneron Pharmaceuticals, Tarrytown, NY) on
197 Day 14 and Day 17 by i.p. injections, based upon a previous study demonstrating that this
198 dosing regimen of IL-1 Trap attenuated cardiac remodeling after experimental acute
199 myocardial infarction in mice (8). In select experiments, 1 µg of IL-1β (R&D Systems,
200 resuspended in 0.1% BSA in PBS) was directly administered intranasally.

201 **Assessment of airway hyperresponsiveness**

202 Following completion of the HDM protocol, mice were anesthetized with
203 intraperitoneal pentobarbital sodium (90 mg/kg), tracheotomized, and mechanically
204 ventilated at 200 breaths/min. Mice were subjected to increasing doses of methacholine (0,
205 12.5, 25, 50, and 100 mg/mL) administered via ultrasonic nebulization, and respiratory

206 mechanics were assessed using a forced oscillation technique on a computer-controlled
207 small animal ventilator (SCIREQ, QC, Canada), as previously described (9, 10). Parameters
208 of Newtonian resistance (R_n), tissue resistance (G) and elastance (H) were calculated and
209 quantified by averaging the three highest measurements obtained at each incremental
210 methacholine dose for each mouse (9, 10).

211 **Assessment of mucus metaplasia:** Airway mucus was stained via Periodic acid Schiff
212 (PAS) and the staining intensity was evaluated by scoring of slides by two independent
213 blinded investigators(11). Levels of MUC5AC were evaluated in lung tissue or BAL via
214 ELISA (My Biosource).

215 **Bronchoalveolar lavage fluid processing**

216 After mice were euthanized, bronchoalveolar lavage (BAL) was performed using 1 ml
217 PBS. BAL was collected and total cell counts were determined using an Advia 120
218 Automated Hematology Analyzer. BAL was spun down at 1200xg for 5 min. Cells were
219 transferred to slides using a cytopsin, fixed in methanol and stained using the Hema3 kit
220 (Fisher Scientific, Kalamazoo, MI) and analyzed by counting a minimum of 300 cells per
221 mouse, as described elsewhere (11). Supernatants were flash frozen in liquid nitrogen and
222 stored at -80 °C until analysis.

223 **Enzyme-Linked Immunosorbent Assay (ELISA)**

224 IL-1 α , IL-1 β , IL-6, TNF α , IL-17, GM-CSF, and CCL20 were detected by ELISA in lung
225 homogenates (normalized for protein) or supernatants from cell culture, according to the
226 manufacturer's instructions (R&D Systems, Minneapolis, MN).

227 **Quantitative Reverse Transcription-Polymerase Chain Reaction (qRT-PCR)**

228 RNA was extracted using miRNeasy columns (Qiagen, Valencia, CA) as directed by
229 the manufacturer. One μ g of RNA was reverse transcribed to cDNA for gene analysis using
230 SYBR Green (Bio-Rad; Hercules, CA, USA) to assess expression of *Mct4*, monocarboxylate
231 transporter 4; *Pfkl*, phosphofructokinase, liver type; *Hk2*, Hexokinase 2; *Glut1*, glucose
232 transporter 1; *Glut3*, glucose transporter 3; *Pkm2*, pyruvate kinase isoenzyme type M2;
233 *Ldha*, lactate dehydrogenase A; *Pgm1*, phosphoglucomutase 1; *Pdk1*, pyruvate

234 dehydrogenase kinase 1; *Tpi1*, triosephosphate isomerase 1; *Eno1*, enolase 1; *Pgk1*,
235 phosphoglycerate kinase 1; *Gpi*, glucose-6-phosphate isomerase; *Pfkfb3*, 6-Phosphofructo-
236 2-Kinase/Fructose-2,6-Biphosphatase 3; *Mct1*, monocarboxylate transporter 1; *Glut2*,
237 glucose transporter 2; and *Ikbke*, inhibitory kappa B kinase ϵ . Expression values were
238 normalized to the house keeping gene cyclophilin. Detailed primer sequences are provided
239 in the online supplement, Table E3.

240 **Bioenergetics**

241 The extracellular acidification rate (ECAR) was measured using the Seahorse
242 Extracellular Flux (XF24) Analyzer (Agilent Technologies). MTE cells were seeded onto 24-
243 well seahorse plate at a density of 50,000 cells per well and cultured with or without 10
244 ng/mL IL-1 β for 24 h. Cells were then washed 3 times with Seahorse stress test glycolysis
245 assay media (DMEM without glucose, L-glutamine, phenol red, sodium pyruvate, and
246 sodium bicarbonate [Sigma-Aldrich] supplemented with 1.85 g/l sodium chloride, 2mM L-
247 glutamine, and 3 mg/l phenol red [GlycoStress Assay], pH 7.35). The plate was incubated in
248 a 37°C non-CO₂ incubator for 1 h. The plate was then transferred to the Seahorse XF24
249 Analyzer for analysis and subjected to ECAR measurements followed by successive
250 treatments with glucose (10 mM), oligomycin (0.25 μ M), and 2-deoxyglucose (100 mM).

251 **Glucose measurements**

252 Glucose consumption and uptake in MTE cells were measured 24 h post stimulation with IL-
253 1 β , by measuring glucose concentration in the cell culture supernatants (Eton Bioscience)
254 and cellular incorporation of fluorescent glucose analog [2-NBDG, 2-(N-(7-Nitrobenz-2-oxa-
255 1,3-diazol-4-yl)Amino)-2-Deoxyglucose, Life technology] using a plate reader (Biotek,
256 Winooski, VT).

257 **Immunohistochemistry**

258 Fixed sections were prepared for immunostaining by deparaffinizing with xylene and
259 rehydrating through a series of ethanols. For antigen retrieval, slides were heated for 20 min
260 in 95°C citrate buffer (pH 6.0), then rinsed in dis tilled water. Sections were then blocked for 1
261 h in blocking serum as per manufacturer's instructions (Vectastain Alkaline Phosphatase

262 Universal, Vector). Slides were then washed in TBS with 0.1% TWEEN-20 3x5 min, followed
263 by incubation with primary antibody for lactate dehydrogenase A overnight at 4°C. Sections
264 were washed again and incubated with a biotinylated universal secondary antibody
265 (Vectastain Alkaline Phosphatase Universal, Vector) for 30 min at room temperature. Slides
266 were washed and incubated with the Vectastain ABC-AP reagent (prepared as per
267 manufacturer's instructions) for 30 min at room temperature. Sections were then incubated
268 with Vector Red/Vector Blue Alkaline Phosphatase Substrate Kit I (Vector) for 10 min at
269 room temperature, rinsed with tap water, and counterstained with Mayer's Hemotoxylin.

270 **Immunofluorescence**

271 Following euthanization, left lobes were fixed with 4% paraformaldehyde, stored at
272 4°C overnight for fixation of the tissue, mounted in paraffin, and 5 µm sections were affixed
273 to glass microscope slides for histopathology as previously described (12). For antigen
274 retrieval, slides were heated for 20 min in 95°C citrate buffer (pH 6.0) with 0.05% TWEEN-20
275 then rinsed in distilled water. Sections were then blocked for 1 h in 1% bovine serum
276 albumin (BSA) in PBS, followed by incubation with primary antibody for IKBKE (Cell
277 Signaling Technology, Danvers, MA) at 1:100, overnight at 4°C. Slides were then washed
278 3x5min in PBS, incubated with Alexafluor 647, and counterstained with DAPI in PBS for
279 nuclear localization. Sections were imaged using a Zeiss 510-META confocal laser scanning
280 microscope.

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282 **Lactate assay**

283 The concentration of lactate in the medium, BAL, and lung homogenates was
284 assessed with a Lactate Assay Kit (Eton Bioscience) according to each manufacturer's
285 recommendations.

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302 **Figure legend**

303 **Figure E1:** Airway inflammation in mice exposed to house dust mite (HDM). **(A)** Total and
304 **(B)** differential cell counts in bronchoalveolar lavage (BAL) in response to saline or HDM,
305 (see schematic in **Figure 1A**). Data are expressed as means (\pm SEM) ($n = 5$ mice per
306 group). * $P < 0.05$ (Student's t test) compared with saline controls.

307 **Figure E2:** Assessment of levels of IL-1 α and IL-1 β in lung tissues from mice subjected to
308 vehicle control or IL1 Trap. Mice were challenged with HDM on days 15, 16, 17, and 18, and
309 received 5 mg/kg of IL-1 Trap or vehicle on days 14 and 17 intraperitoneally. Levels of IL-1 α
310 and IL-1 β in lung tissues homogenates were evaluated on day 20 via ELISA. * $P < 0.05$
311 compared to saline control, † $P < 0.05$ compared to HDM/Veh group (ANOVA).

312 **Figure E3:** Evaluation of pro-inflammatory mediators and airway inflammation in mice
313 exposed to interleukin (IL)-1 β . **A:** Lung tissue levels of KC, TSLP, CCL20 and GM-CSF
314 following intranasal administration of IL-1 β or vehicle. Total **(B)** and differential cell counts
315 **(C)** in BAL from the mice at multiple time points post intranasal administration of vehicle or
316 IL-1 β . Data are expressed as means (\pm SEM) ($n = 5$ mice per group). * $P < 0.05$ (ANOVA)
317 compared with Vehicle controls.

318 **Figure E4: A:** Assessment of viability following exposure to MTE cells to oxamate or 2-
319 deoxyglucose. Epithelial cells were exposed to IL-1 β in the presence or absence of
320 inhibitors. Cell survival was evaluated via crystal violet staining of cells. Results were
321 expressed as % survival compared to untreated control cultures. **B:** Impact of 2-DG or
322 oxamate on IL1 α -mediated increases in lactate and the indicated pro-inflammatory
323 mediators measured 24 hr post exposure to IL-1 α . * $P < 0.05$ (ANOVA) compared to the
324 sham group. † $P < 0.05$ compared to the IL-1 β treated Vehicle group (ANOVA).

325 **Figure E5: A:** Quantification of Western blots shown in Figure 8A. Data reflects $n=3$ healthy
326 subjects and $n=3$ asthmatics. Data were normalized to β -actin and are expressed as fold
327 change from healthy sham controls * $P < 0.05$ compared to the sham healthy group. † $P <$
328 0.05 compared to HDM healthy group (ANOVA). **B:** Sub-analysis of sputum lactate and IL-
329 1 β in asthmatics with normal neutrophils (cut off $\leq 61\%$) or high neutrophils (cut off $> 61\%$),

330 controlled (ACQ \leq 1.5) or uncontrolled asthma (ACQ $>$ 1.5). p-values (ANOVA or Wilcoxon
331 rank sum test) are provided in each of the figures.

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335 **References**

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Table E1. Demographic, functional, and inflammatory characteristics of the study cohort

	Healthy subjects	Asthmatic patients	<i>P</i> value
n	20	94	
Age (yr)	53 ± 3	52 ± 2	0.86
BMI (kg/m ²)	24.30 ± 1.05	27.28 ± 0.48	0.01
Age asthma onset (range)	-	41 (12-54)	-
Gender (M/F)	10/10	43/51	0.3
Atopy*, no. (%)	4 (20)	42 (45)	0.03
Positive Dpt*, no. (% atopy)	3 (75)	25 (60)	
FEV1 % predicted	101 ± 4.62	81.56 ± 2.07	0.0001
FEV1/FVC ratio	77.68 ± 2.51	72.41 ± 1.13	0.0291
FENO (ppb), median (IQR)	20 (17.3-26.8)	17.5 (10-36.25)	0.13
Eosinophils (%), median (IQR)	0.2 (0-0.9)	0.9 (0.2-5.45)	0.03
Medication use			
Not treated	-	5 (5)	-
ICS/LABA, no. (%)	-	58 (62)	-
eq Beclomethasone µg/ml	-	1000 (400-2000)	
ICS alone, no. (%)	-	2 (2)	-
OCS, no. (%)	-	3 (3)	-
SABA, no. (%)	-	64 (68)	-
SABA only, no. (%)	-	19 (20)	-
LTRA, no. (%)	-	24 (26)	-

Data are expressed as means ± SDs, or medians with interquartile range (IQR). *P* values are based on the student's *t* test (mean ± SD), the χ^2 test for proportions (sex), the Wilcoxon rank sum test (median [range]), or Poisson regression (atopy). Atopy is defined as positive test results for at least 1 specific IgE to common aeroallergens. FENO: Fraction of exhaled nitric oxide. * Dpt.: Dermatophagoid pteronyssinus. ICS: inhaled corticosteroid. LABA: long acting beta agonist. OCS: oral corticosteroids. SABA: Short acting beta agonist. LTRA: leukotriene receptor agonists. Ethnicity: All subjects are Caucasian except for 2 African subjects.

Table E2. Demographic, functional, and inflammatory characteristics of the study cohort enrolled at the University of Vermont Medical Center

	Healthy subjects	Asthmatic patients
n	7	6
Age (range)	23 (19-27)	23 (19-45)
Gender (M/F)	0/6	3/4
Age asthma onset (range)	-	4 (1-18)
BMI	23.2 (19.8-26.3)	28.6 (22.0-35.0)
Atopy*, no. (%)	-	7 (100)
Medication use		
ICS_LABA	-	3
ICS	-	3
SABA only	-	1
SNQ	0.4 (0-0.8)	1.4 (1.2-1.6)

Data are expressed as median (range). ICS_LABA: inhaled corticosteroid_long acting beta agonist. ICS: inhaled corticosteroid. SABA: short acting beta agonist. SNQ: sinonasal questionnaire. *All participants were allergic to house dust mite as determined either by positive skin prick test to *D. pteronyssinus* or positive serum IgE to *D. pteronyssinus*. All subject are Caucasian.

Table E3. The primers used in this study

Genes	Forward	Reverse
<i>Mct4</i>	5'-ATCGTGGGCACTCAGAAGTT-3'	5'-CGCCAGGATGAACACATACTT-3'
<i>Pfkl</i>	5'-CATATATGTGGGGGCCAAAG-3'	5'-GACACACAGGTTGGTGATGC-3'
<i>Hk2</i>	5'-GGGACGACGGTACACTCAAT-3'	5'-GCCAGTGGTAAGGAGCTCTG-3'
<i>Glut1</i>	5'-TCTCTGTCCGGCCTCTTTGTT-3'	5'-CCAGTTTGGAGAAGCCATA-3'
<i>Glut2</i>	5'-GCCTGTGTATGCAACCATTG-3'	5'-GAAGATGGCAGTCATGCTCA-3'
<i>Glut3</i>	5'-TGTCACAGGAGAAGCAGGTG-3'	5'-GCTCCAATCGTGGCATAGAT-3'
<i>Pkm2</i>	5'-CTGCAGGTGAAGGAGAAAGG-3'	AGATGCAAACACCATGTCCA-3'
<i>Ldha</i>	5'-GGAAGGAGGTTCAAGCAG-3'	5'-ACCCGCCTAAGGTTCTTCAT-3'
<i>Pgm1</i>	5'-TCAGGCCATTGAGGAAAATC-3'	5'-CGAACTTACCTTGCTCTCC-3'
<i>Pdk1</i>	5'-GGCGGCTTTGTGATTTGTAT-3'	5'-ACCTGAATCGGGGGATAAAC-3'
<i>Tpi1</i>	5'-CCTGGCCTATGAACCTGTGT-3'	5'-CAGGTTGCTCCAGTCACAGA-3'
<i>Eno1</i>	5'-CTGCCTCCGAGTTCTACAGG-3'	5'-CGCTTAGGGTTGGTCACTGT-3'
<i>Pgk1</i>	5'-CAAGGCTTTGGAGAGTCCAG-3'	5'-TGTGCCAATCTCCATGTTGT-3'
<i>Gpi</i>	5'-GTGGTCAGCCATTGGACTTT-3'	5'-CTGGAAATAGGCAGCAAAGC-3'
<i>Pfkfb3</i>	5'-CAGCTACCAGCCTCTTGACC-3'	5'-AACTTCTGCCTCTGCTGGA-3'
<i>Mct1</i>	5'-TCCAGTAATGATCGCTGGTG-3'	5'-AGTTGAAAGCAAGCCCAAGA-3'
<i>Ikake</i>	5'-CTGGATGTCCCAAAGTTCGT-3'	5'-AGGCTGCTGCTGAGGTAGAG-3'

Figure E1

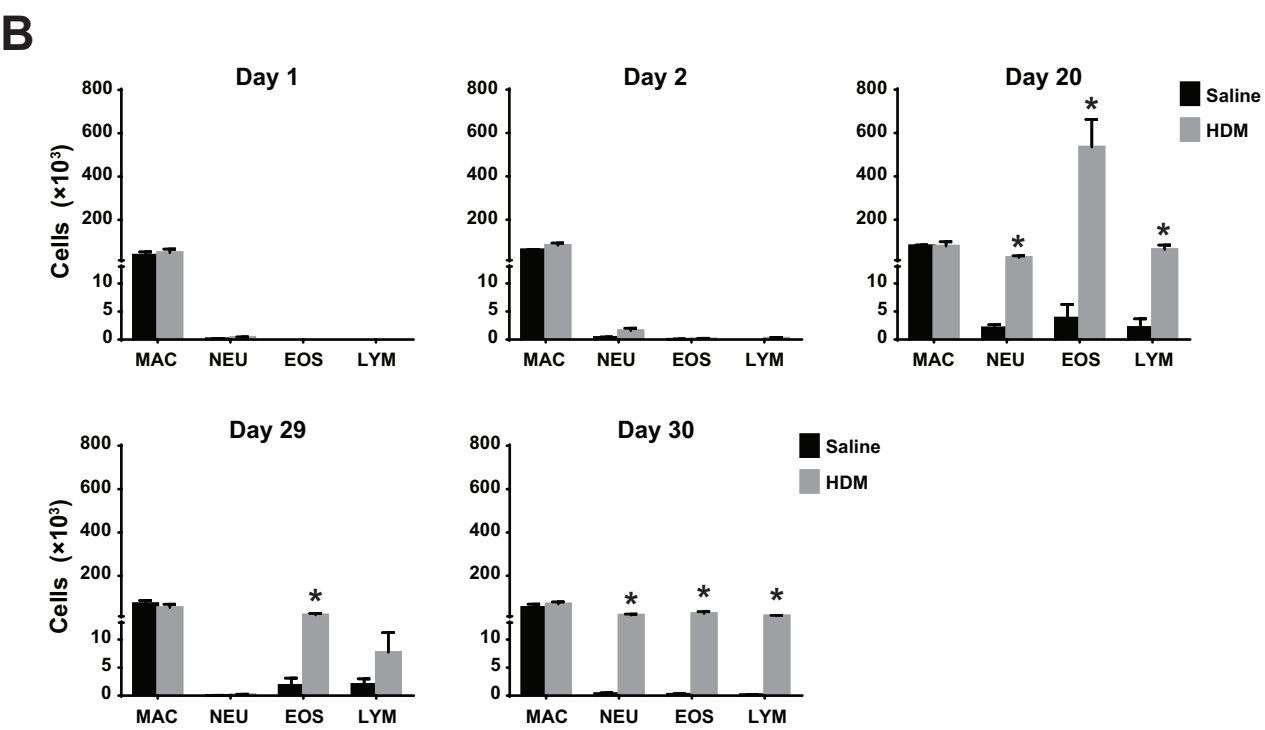
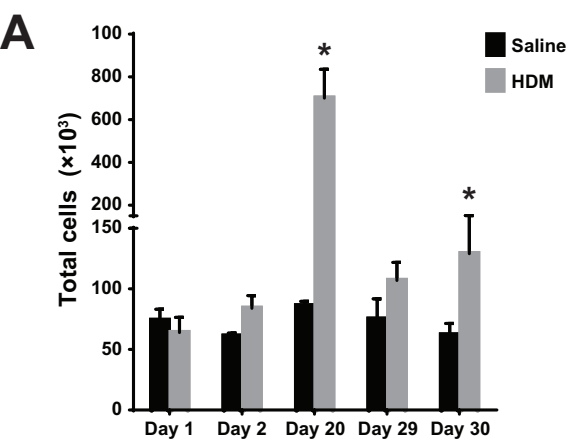
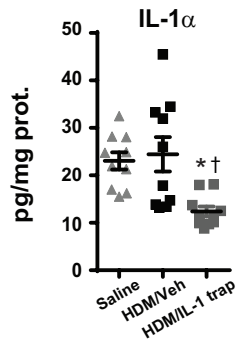
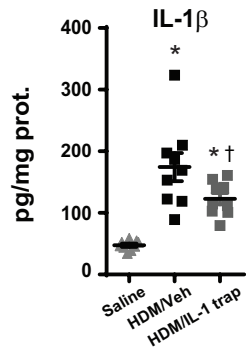


Figure E2

A



B



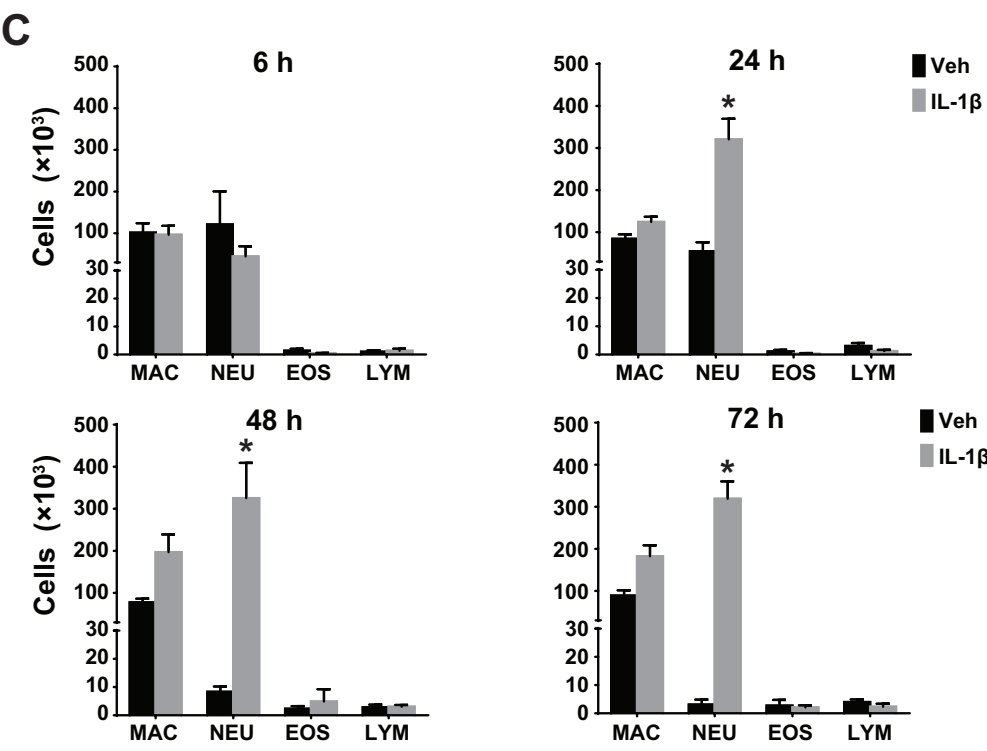
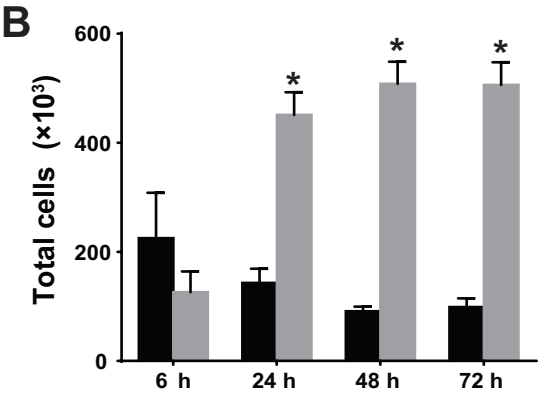
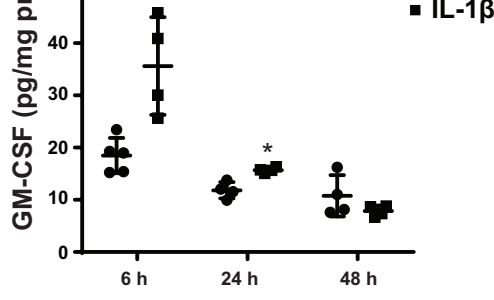
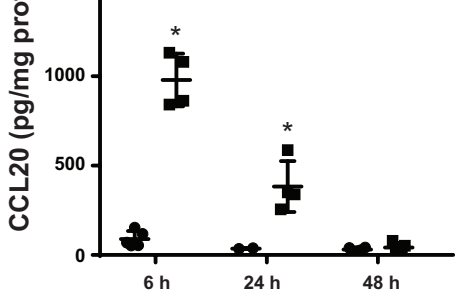
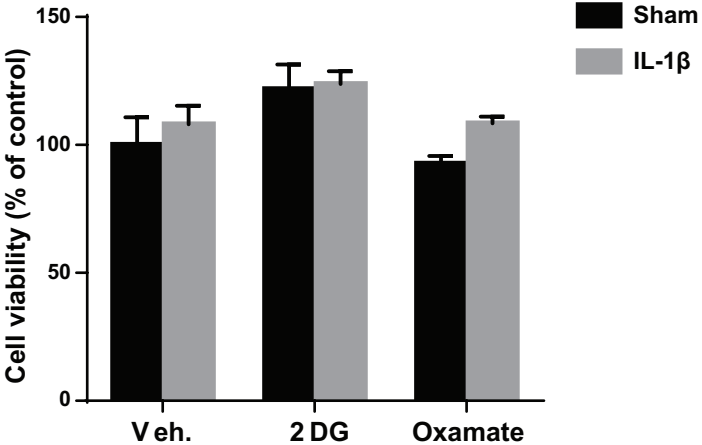


Figure E4

A



B

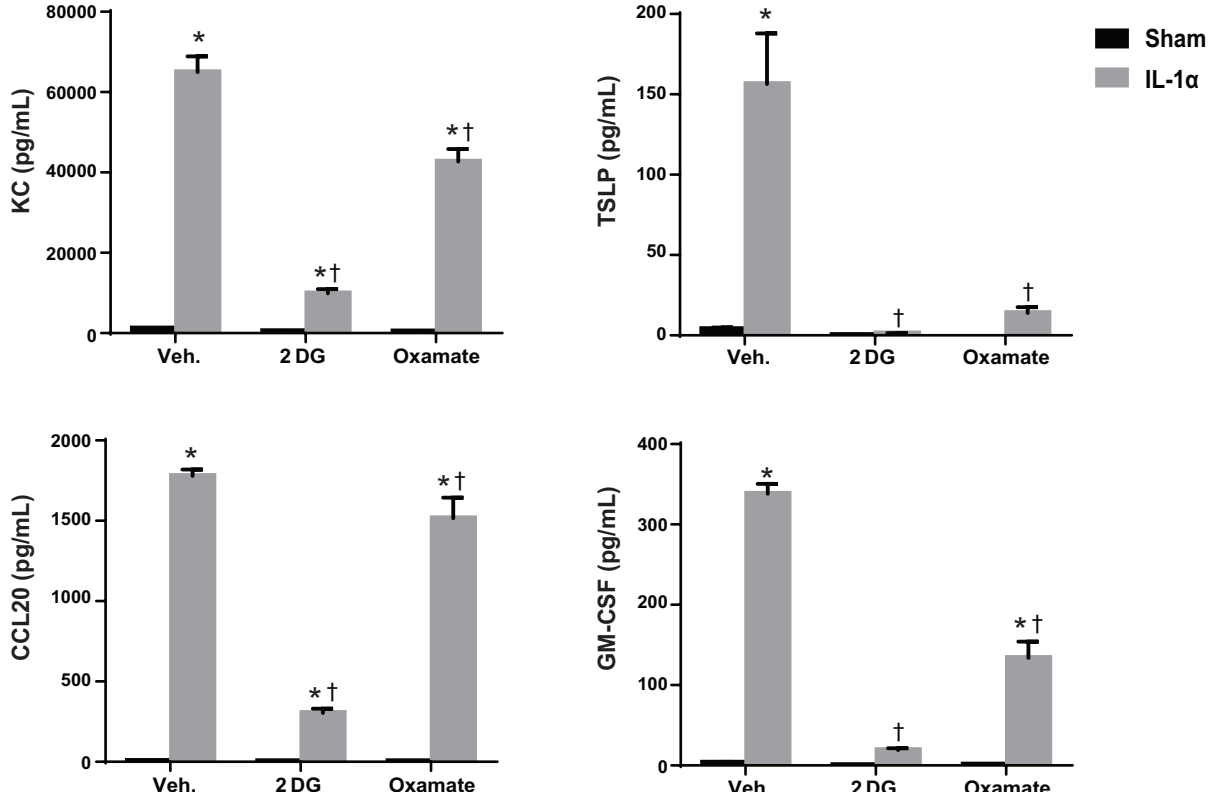
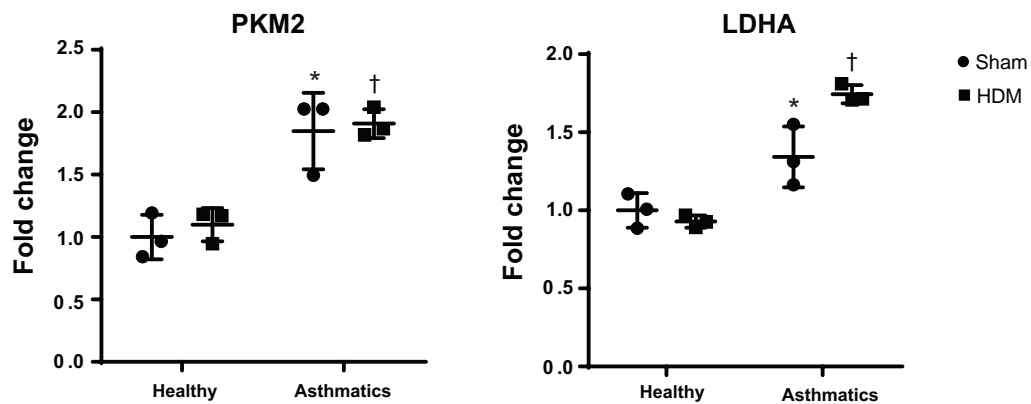


Figure E5

A



B

