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Topical 11 β -hydroxysteroid dehydrogenase type 1 inhibition corrects cutaneous features of systemic glucocorticoid excess in female mice

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Abstract

Glucocorticoid (GC) excess drives multiple cutaneous adverse effects including skin thinning and poor wound healing. The ubiquitously-expressed enzyme 11 β -hydroxysteroid dehydrogenase type 1 (11 β -HSD1) activates mouse corticosterone from 11-dehydrocorticosterone (and human cortisol from cortisone). We previously demonstrated elevated 11 β -HSD1 activity during mouse wound healing but the interplay between cutaneous 11 β -HSD1 and systemic GC excess is unexplored. Here, we examined effects of 11 β -HSD1 inhibition by carbenoxolone (CBX) in mice treated with corticosterone (CORT) or vehicle for 6 weeks. Mice were treated bi-daily with topical CBX or vehicle 7 days before wounding and during wound healing. CORT mice displayed skin thinning and impaired wound healing but also increased epidermal integrity. 11 β -HSD1 activity was elevated in unwounded CORT skin and was inhibited by CBX. CORT mice treated with CBX displayed 51%, 59% and 100% normalization of wound healing, epidermal thickness and epidermal integrity, respectively. Gene expression studies revealed normalization of interleukin-6, keratinocyte growth factor, collagen-1, collagen-3, matrix metalloproteinase-9 and tissue inhibitor of matrix metalloproteinase-4 by CBX during wound healing. Importantly, pro-inflammatory cytokine expression and resolution of inflammation was unaffected by 11 β -HSD1 inhibition. CBX did not regulate skin function or wound healing in the absence of CORT. Our findings demonstrate that 11 β -HSD1 inhibition can **limit** the cutaneous effects of GC excess which may improve the safety profile of systemic steroids and the prognosis of chronic wounds.

44 **Précis**

45 Topical 11 β -hydroxysteroid dehydrogenase type 1 inhibition normalizes steroid-induced skin
46 thinning, epidermal integrity, poor healing and gene dysregulation despite ongoing systemic steroid
47 exposure.

Introduction

Systemic glucocorticoid (GC) therapy remains mainstream treatment for many inflammatory diseases e.g. lupus, asthma, chronic obstructive pulmonary disease, inflammatory bowel disease, multiple sclerosis, polymyalgia rheumatica and giant cell arteritis. Recent estimates indicate that 1.2% of the US population (>2.5 million people) were prescribed oral GC between 1999 and 2008, 28% of whom reported use of >5 years (1). Despite anti-inflammatory benefits, chronic GC excess drives adverse side-effects including weight gain, hypertension, hyperglycemia, osteoporosis, muscle weakness, glaucoma and depression (2). GC therapy also substantially increases healthcare costs. In patients with systemic lupus erythematosus, GC use was associated with a three-fold increase in annual expenditure (3). In a study of patients with severe asthma, high dose GC use was associated with annual increments of \$5479 relative to low GC exposure (4). Moreover, the incidence and impact of GC-related side-effects is under-reported (5) and further compounded by GC over-prescription (6).

In skin, GC excess causes acne, thinning, dryness, atrophic striae, telangiectasia, bruising, impaired wound healing (WH) and increased infection risk (2,7). Skin bruising and thinning have also been reported with low-dose (<7.5 mg/day) GC therapy (8). In a survey of asthma physicians, cutaneous manifestations were the second most frequently reported complication of inhaled GC (9). Despite recent advances in GC mimetics and GC-independent immune suppressants, the “holy grail” of dissociating GC anti-inflammatory benefits from side-effects remains elusive (10).

It is now established that human skin has corticosteroidogenic capability that is regulated in a complex way (11,12), although this function is lacking in mouse skin (13). Peripheral GC availability is also regulated by 11 beta-hydroxysteroid dehydrogenase type 1 (11 β -HSD1) which converts biologically inert cortisone and 11-dehydrocorticosterone to cortisol and corticosterone in humans and mice, respectively. In intact tissues and cells, 11 β -HSD1 functions predominantly as a reductase in an NADPH-dependent manner (14). We previously reported increased 11 β -HSD1 activity during

mouse skin WH (13), in aged mouse and human skin (15,16) and accelerated WH in aged 11 β -HSD1-null mice (16). However, the role of 11 β -HSD1 in impaired WH is unknown.

We propose that 11 β -HSD1 is a causative agent of cutaneous side-effects during systemic GC excess such as skin atrophy and impaired WH. Since human 11 β -HSD1 inhibitors are readily available, such a demonstration could lead to rapid progress in ameliorating this major clinical problem. Here, we provide the first empirical evaluation of this causal link. We investigate effects of topical 11 β -HSD1 inhibition by carbenoxolone (CBX) on skin function in mice treated with oral corticosterone and examine inflammatory, growth factor and extracellular matrix gene expression responses during WH. We demonstrate that 11 β -HSD1 inhibition accelerates WH and reduces skin damage in steroid-treated mice without compromising the inflammatory response.

Materials and Methods

Declarations

Studies presented in this paper were approved by the Institutional Animal Care and Use Committee and San Francisco Veterans Affairs Medical Center Veterinary Medical Unit. Materials were obtained from Sigma-Aldrich unless otherwise stated.

Animal Studies

Female SKH1 hairless mice (8–10 weeks old) were obtained from Charles River Labs (Wilmington, MA, USA) and acclimatized for 2 weeks. Mice were group housed, supplied with a basal chow diet *ad libitum* and exposed to a standard 12h light/dark cycle. Drinking water was supplemented with corticosterone (100 μ g/ml) or vehicle (0.66% ethanol) for 6 weeks (replaced twice weekly). 7 days prior to wounding, 100 μ g/cm² topical CBX (20 μ l) or vehicle (70% propylene glycol, 30% ethanol) was massaged into 2cm² skin until fully absorbed. Treatments were repeated bi-daily for the remainder of the experiment and replaced with 10 μ l per wound post-wounding.

24h prior to wounding, epidermal barrier function was measured using a Tewameter 300 (Courage + Khazaka, Cologne, Germany) to evaluate trans-epidermal water loss (TEWL). Epidermal barrier recovery was determined at 2 and 4h following disruption by sequential D-Squame tape stripping to a TEWL level of $>30\text{g/h/m}^2$. Epidermal integrity was defined as the number of tape strips required to achieve barrier disruption.

WH was conducted as previously described (13). Briefly, mice anesthetized under 2% isoflurane were wounded on both dorsal flanks by 5mm punch biopsy (Acuderm, Fort Lauderdale, FL, USA). Wounds received 20 μl bupivacaine 0.25% (analgesic) and monitored daily.

At 2, 4 and 9 days post-wounding, mice were culled by cervical dislocation, wounds were digitally imaged and excised for 11 β -HSD1 activity assay, RNA extraction or histology. Wound areas were determined by ImageJ (NIH, Bethesda, MD, USA).

Experiments were also replicated in control mice (untreated drinking water) treated with vehicle or CBX.

11 β -HSD1 activity assay

Radioactive conversion of tritiated 11 β -HSD1 substrate is the gold standard measure of 11 β -HSD1 activity as tissue steroid extraction efficiencies are variable and less sensitive.

Freshly isolated skin (20-40mg) was incubated immediately in 1ml high glucose and pyruvate DMEM with 100nM 11-dehydrocorticosterone (Steraloids, Newport, RI, USA) and ~1500cpm [3H] 11-dehydrocorticosterone, generated in-house as previously described (13). Samples were incubated at 37°C for 13h. Subsequently, tissues were weighed and steroids extracted and separated by TLC in 186ml chloroform and 14ml ethanol for 90min (co-migrated with 10mM 11-dehydrocorticosterone / corticosterone standards). Plate regions identified under UV were excised and percentage conversion of 11-dehydrocorticosterone to corticosterone was determined following liquid scintillation.

Quantitative PCR

Fresh skin tissue (20-40mg) was snap-frozen and stored at -80°C. Samples were homogenized in 1ml Trizol and RNA extracted using a PureLink RNA Mini Kit (Life Technologies, Grand Island, NY, USA). cDNA was generated from 1.2µg RNA using a Tetro cDNA Synthesis Kit (Bioline, Taunton, MA, USA). qPCR was conducted in 10µl reactions using a SensiFAST Probe Kit (Bioline) with 900nM TaqMan primers (with 250nM FAM probe) or 50nM 18S rRNA primers (with 200nM VIC probe) mix (Life Technologies) and 10ng cDNA. Duplicate PCRs were performed and analyzed, as previously described (13).

Serum corticosterone

For serum corticosterone, 400µl of blood was obtained by terminal cardiac puncture at 11 AM and incubated for 1h at 5°C before centrifuging at 1000 x RCF for 5min. Corticosterone levels were determined using a corticosterone EIA Kit (Cayman Chemical, Ann Arbor, MI, USA).

Histology

Freshly isolated samples were stored in Formalde-Fresh (Fisher Scientific, Pittsburgh, PA, USA) and processed into paraffin blocks. 5µm sections were de-waxed, rehydrated, and stained with hematoxylin and eosin (Leica, Buffalo Grove, IL, USA and Thermo Scientific, Kalamazoo, MI, USA). Sections were examined with a Zeiss microscope (Jena, Germany) and digital images captured with AxioVision software (Carl Zeiss Vision, Munich, Germany).

For epidermal thickness, the average of 4 measurements was taken per image and dermal area was quantified using ImageJ. For quantification of epidermal cellularity (>90% of which is composed of keratinocytes), the total number of epidermal nuclei were counted per field of view and normalized to epidermal length (to account for rete ridge undulation).

Proliferation was evaluated by incubating rehydrated sections with biotinylated primary antibody against Proliferating cell nuclear antigen (PCNA), CalTag Laboratories, Burlingame, CA, USA) overnight at 4°C. After 3 x TBS washes, staining was detected with ABC-peroxidase Kit (Vector Lab, Burlingame, CA, USA) and quantified using ImageJ.

Statistical Analysis

Following confirmation of data displaying a normal distribution, significance levels were determined by Student's t-test, one-way or two-way ANOVA using GraphPad Prism (La Jolla, CA, USA) on untransformed data. For two-way ANOVA, post hoc testing included analysis of differences between time-points in each treatment group and differences between treatment groups at each time-point with p-values adjusted for multiple testing as detailed in Figure legends. Variation is displayed as standard error based on at least three biological replicates.

Results

Cutaneous 11 β -HSD1 expression and activity increase during wound healing and systemic glucocorticoid excess

In agreement with known suppressive effects of GC treatment on hypothalamic-pituitary-adrenal axis signaling, systemic GC treatment (CORT) reduced circulating corticosterone levels by 70% compared to vehicle mice (Fig. 1A) with body weight unaffected between groups (data not shown).

The effect of GC excess on 11 β -HSD1 expression and activity during mouse skin WH were previously unexplored. At baseline, 11 β -HSD1 mRNA was 10-fold higher in CORT skin compared to vehicle (Fig. 1B). Hexose-6-phosphate dehydrogenase (H6PDH), which supplies 11 β -HSD1 co-factor, showed a similar increase in expression at baseline (Fig. 1C). Correspondingly, 11 β -HSD1 activity was also 42% greater in unwounded CORT skin compared to vehicle (Fig. 1D).

In vehicle-treated mice, 11 β -HSD1 mRNA and activity increased by 43-fold and 2.6-fold at day 2 and by 17-fold and 1.7-fold at day 4 post-wounding, respectively (Fig. 1B, 1D). H6PDH mRNA was also increased by 7-fold at days 2 and 4 post-wounding (Fig. 1C). CORT further increased 11 β -HSD1 mRNA by 11-fold and 20-fold (independently of CBX) at days 4 and 9 post-wounding, respectively (Fig. 1B). H6PDH mRNA was also further upregulated by 5-fold at day 4 and 9-fold (independently of CBX) at day 9 post-wounding in CORT mice (Fig. 1C). However, 11 β -HSD1 activity was not affected by CORT treatment during WH (Fig. 1D).

In CORT mice, CBX treatment inhibited 11 β -HSD1 activity by 61% in unwounded skin and by 91% and 55% at days 2 and 4 post-wounding, respectively (Fig. 1D). In control mice (untreated drinking water), CBX inhibited 11 β -HSD1 activity by 30% in unwounded skin and by 93%, 94% and 73% at days 2, 4 and 9 post-wounding (Supplemental Fig. 1).

These findings indicate that GC excess induces cutaneous 11 β -HSD1 expression and activity in unwounded skin which is inhibited effectively by CBX treatment.

11 β -HSD1 inhibition limits glucocorticoid-mediated skin thinning

GC cause skin thinning and dermal atrophy. However, the effect of 11 β -HSD1 inhibition on epidermal thinning and proliferation following GC exposure was previously unreported. Epidermal thickness was reduced by 36% in CORT mice compared to vehicle and was restored by 59% with CBX (Fig. 2A, 2B). A 27% CORT-mediated reduction in epidermal cellularity was also prevented by CBX (Fig. 2A, 2C) with a similar trend in PCNA staining (Supplemental Fig. 2). Dermal area decreased by 21% with CORT treatment compared to vehicle and this was also prevented by CBX (Fig. 2D). No difference in epidermal thickness or epidermal cellularity was observed in control mice treated with CBX (Supplemental Fig. 3).

These results suggest that 11 β -HSD1 inhibition can limit GC-mediated skin thinning despite sustained exposure to systemic GC excess.

Impact of glucocorticoid excess and 11 β -HSD1 inhibition on epidermal barrier function

A functional epidermal barrier is indispensable for protection against dehydration, exposure to irritants and invasion by microorganisms. The regulation of epidermal barrier function, integrity and recovery by corticosterone excess or 11 β -HSD1 inhibition has not been investigated.

Trans-epidermal water loss (TEWL) is the gold standard measure of epidermal barrier function. Baseline TEWL was largely unaffected by CORT or CBX (Fig. 3A). Interestingly, the number of tape strips required to disrupt the barrier (indicative of barrier integrity) was 50% greater in CORT mice and this was prevented by CBX treatment (Fig. 3B). We observed a modest increase in barrier recovery in the CORT/CBX group 2 hours after barrier disruption but no difference was observed between CORT and CORT/CBX or between any of the groups 4 hours after disruption (Fig. 3C). No statistically significant effect of CBX on barrier function (baseline TEWL), barrier integrity or barrier recovery was observed in control mice (Supplemental Fig. 4).

These findings suggest that 11 β -HSD1 mediates increased epidermal integrity during systemic GC excess.

11 β -HSD1 inhibition improves skin wound healing during glucocorticoid excess

GC excess drives impaired WH and chronic wound management remains an unmet clinical need in diabetic and elderly patients. However, the regulation of impaired WH during GC excess by 11 β -HSD1 is unknown. Systemic GC excess induced a striking WH delay which was improved by CBX treatment (Fig. 4A).

Initially, WH progressed at similar rates between all groups with a 36%, 39% and 33% reduction in wound area at day 4 compared to day 2 in vehicle, CORT and CORT/CBX mice, respectively (Fig. 4B). No change in wound area was observed between days 4 and 6 for either group. Although healing continued from day 6 for vehicle and CORT/CBX mice, no further healing was observed for CORT/VEH mice. By day 9, wound areas were 94% and 48% healed compared to day 4 in vehicle and CORT/CBX mice, respectively (Fig. 4B).

H+E staining of day 9 wounds revealed more advanced granulation tissue formation, improved structural organization and extracellular matrix deposition in CORT/CBX compared to CORT mice (Fig. 4C). CBX treatment did not affect WH in control mice (Supplemental Fig. 5).

In summary, 11 β -HSD1 inhibition improved mouse skin WH despite sustained systemic GC exposure.

Effect of systemic glucocorticoid excess and 11 β -HSD1 inhibition on cytokine, extracellular matrix and growth factor gene expression during wound healing

To further investigate the mechanisms underlying improved WH with 11 β -HSD1 inhibition, we analyzed changes in cytokine, extracellular matrix and growth factor gene expression.

221 Pro-inflammatory cytokine interleukin (IL)-1 β mRNA increased at days 2, 4 and 9 post-wounding by
222 750-fold, 587-fold and 90-fold respectively, independently of CORT or CBX treatment (Fig. 5A). A
223 similar expression profile was seen for tumor necrosis factor alpha (TNF α) mRNA (Fig. 5B).

224 The immunomodulatory cytokine IL-6 regulates a broad range of biological functions including
225 antibody-secreting B-cell differentiation and generation of Th17 cells. IL-6 mRNA expression was
226 undetectable in the majority of unwounded skin samples but increased at day 2 and day 4 before
227 declining at day 9 post-wounding in vehicle mice (Fig. 5C). IL-6 displayed a trend towards higher
228 expression at day 2 post-wounding in CORT compared to vehicle mice which was prevented by CBX
229 (Fig. 5C). IL-6 mRNA was also 14-fold greater in CORT mice at day 9 post-wounding compared to
230 vehicle mice (Fig. 5C).

231 Keratinocyte growth factor (KGF) is produced during WH and stimulates wound re-epithelialization.
232 KGF mRNA expression peaked at day 4 with a 9.9-fold and a 3.5-fold induction in vehicle and
233 CORT/CBX mice compared to unwounded skin, respectively (Fig. 5D). No induction in KGF was seen
234 at day 4 in CORT mice. At day 9 post-wounding, KGF was 17-fold higher in CORT compared to vehicle
235 mice and this was reversed by CBX (Fig. 5D).

236 Type I and III collagen constitute >70% of the dermis by dry weight. Collagen type I alpha 1 (COL-1 α 1)
237 and collagen type III alpha 1 (COL-3 α 1) mRNA expression increased 15-fold at day 4 post-wounding
238 in vehicle mice compared to unwounded skin (Fig. 5E, 5F). This was delayed in CORT mice, which
239 displayed a 3.5-fold and 14-fold induction of COL-1 α 1 and COL-3 α 1 at day 9 post-wounding
240 compared to vehicle controls both reversed by CBX (Fig. 5E, 5F).

241 Extracellular matrix turnover is regulated by collagen-degrading matrix metalloproteinases (MMP)
242 and their inhibitors, tissue inhibitor of matrix metalloproteinases (TIMP). MMP-9 mRNA in vehicle
243 mice increased 7.7-fold and 9.5-fold at day 2 and day 4 post-wounding, respectively, before
244 returning to baseline levels by day 9 (Fig. 5G). However, in CORT (but not CORT/CBX) mice, MMP-9

expression at day 9 remained elevated at 7.2-fold higher than vehicle mice (Fig. 5G). TIMP-1 displayed a similar mRNA profile, with 20-fold and 33-fold higher expression at day 2 and day 4 post-wounding respectively, in vehicle mice compared to unwounded skin (Fig. 5H). At day 9 post-wounding, MMP-9 mRNA was also 10-fold greater in CORT (but not CORT/CBX) compared to vehicle mice (Fig. 5H).

These findings oppose reports of increased inflammatory cytokine expression following 11 β -HSD1 blockade in other models. Improved WH appears to be mediated, in part, by restoration of KGF signaling and appropriate collagen, MMP and TIMP expression.

Discussion

Although GC have been used for over 65 years for their anti-inflammatory properties (17), adverse side-effects limit their use to low-dose, short-term intervention (2). GC-mediated skin atrophy, bruising, impaired WH and increased infection risk are of concern to both patients and physicians alike (18). In a survey of patients taking GC for >60 days, skin bruising or thinning was the second most prevalent self-reported side-effect (8). We have demonstrated for the first time that topical 11 β -HSD1 inhibition can **limit** these cutaneous side-effects. This key insight opens a wide range of avenues for 11 β -HSD1 inhibitors as adjunct therapies to safely increase steroid doses or treatment duration.

Morgan *et al.* recently demonstrated that global 11 β -HSD1-null mice were protected from the systemic side-effects of oral corticosterone (19). However, this study did not evaluate inflammation, which others report to be exacerbated by 11 β -HSD1 inhibition (20). Here, we evaluated the ability of 11 β -HSD1 inhibition to *reverse* cutaneous effects of GC excess.

Reduced morning serum corticosterone levels in CORT mice are consistent with the reported adrenal suppression at this sampling time (21) and total body weight was unaffected by GC treatment as previously described (19). The forward-feedback induction of 11 β -HSD1 mRNA and activity by GC in

269 unwounded skin is also in agreement with our previous observations in human skin (15) and may
270 exacerbate the 11 β -HSD1-mediated effects observed in unwounded CORT mouse skin.

271 Our results demonstrate that 11 β -HSD1 inhibition prevents GC-mediated reductions in epidermal
272 cellularity and dermal area, despite sustained exposure to systemic GC excess. Interestingly, we
273 observed that CBX also reversed a GC-mediated induction in epidermal integrity. This is supported
274 by studies demonstrating that exogenous GC treatment promote epidermal keratinocyte
275 differentiation and increase corneocyte numbers (22), reflecting a greater degree of disruption
276 required for epidermal barrier perturbation. Conversely, endogenous GC excess or topical GC
277 treatment cause GC-mediated decreases in epidermal integrity indicating that model-specific
278 differences should also be taken into consideration (23-26).

279 Baseline TEWL (epidermal barrier function) was unaffected by CORT treatment in agreement with
280 previous reports (23-26). In contrast to our findings, other studies reported impaired barrier
281 recovery following endogenous GC excess and GC treatment, although this was associated with
282 impaired barrier integrity (23,25,26). In our model, integrity and barrier recovery were both
283 stimulated by systemic GC excess. This may be due to differences in GC potency, formulation,
284 treatment duration, administration, strain, gender, age, diet or environment. Whilst integrity
285 appeared to be 11 β -HSD1-mediated, the modest induction in barrier recovery was not reversed by
286 CBX. This may be due to regulation by other hypothalamic-pituitary-adrenal axis pathways such as
287 β 2-adrenergic receptor signaling (27).

288 Between days 0 to 4, wound closure was similar between all groups, suggesting that the initial
289 contractile phase of WH (28) was unaffected. We observed termination of WH kinetics from day 4
290 post-wounding in CORT mice, supported by previous reports of GC-mediated impairment of
291 keratinocyte proliferation, migration and increased epidermal differentiation (29,30). Strikingly, our
292 results demonstrate that CBX treatment improved WH in CORT mice. The lack of complete WH
293 rescue by CBX may be due to exposure to serum corticosterone before cellular

compartmentalization is fully restored. Others have also demonstrated benefits of 11 β -HSD1 blockade on cutaneous WH in animal models of metabolic disease and endogenous GC excess (31,32), however, 11 β -HSD1 inhibition, inflammatory effects and mechanistic insights were lacking in these studies.

Our findings indicate that the cutaneous side-effects of GC excess are mediated through increased 11 β -HSD1 substrate availability as 11 β -HSD1 activity was not further increased by CORT treatment and CBX treatment did not affect skin thickness, epidermal integrity or WH in control mice not exposed to systemic GC excess. This suggests that 11-dehydrocorticosterone availability is insufficient in young, healthy mice to regulate skin function.

Recent reports have raised concerns over increased inflammation as consequence of 11 β -HSD1 blockade (33-36). We found no evidence of this our model; there was no exacerbation of IL-1 β , TNF- α or IL-6 mRNA expression during early WH and no delay in resolution of these pro-inflammatory cytokines in CBX mice. On the contrary, IL-6 expression increased during CORT mouse WH. This is supported by reports of pro-inflammatory GC actions (37,38), indicating greater regulatory complexity than previously thought (39-41).

KGF is a key factor that promotes wound re-epithelialization and is indispensable to the WH process (30,42,43). We found a lack of induction of KGF at day 4 post-wounding in CORT mice (coinciding with the cessation of wound closure) which was restored by CBX. The induction of collagen type I and III mRNA during the granulation phase of WH were attenuated in CORT mice, consistent with previous reports (30,44) and were also normalized by CBX, further supported by our histological findings.

MMP-9 and TIMP-1 mRNA expression were elevated at day 9 post-wounding in CORT mice and were normalized by CBX treatment. Although the underlying mechanisms remain poorly understood,

317 elevated MMP-9 and TIMP-1 mRNA are associated with slower healing in mouse wounds (45) and
318 MMP-9 is elevated in diabetic foot ulcers (46).

319 GC are able to activate the mineralocorticoid receptor (MR) with strong affinity and recent work by
320 Boix *et al.* elegantly demonstrated that MR epidermal knockout mice were resistant to GC-induced
321 epidermal thinning (47). Differentiating between glucocorticoid receptor and MR-mediated effects
322 was beyond the scope of the current study, but is an area of ongoing interest.

323 CBX has been used extensively in human and mammalian models. Administered systemically, it also
324 inhibits 11 β -HSD2 (which conducts the opposing reaction to 11 β -HSD1) resulting in renal
325 mineralocorticoid excess due to GC-mediated MR activation (48). However, 11 β -HSD2 is not
326 expressed in unwounded mouse skin or during WH (13) and topical administration limits the risk of
327 systemic exposure. CBX is also able to block gap junctions but at several orders of magnitude less
328 potently than 11 β -HSD1 (48). Further, we observed no difference between CBX and vehicle mice not
329 exposed to CORT, suggesting our findings are most likely due to cutaneous 11 β -HSD1 inhibition.

330 In summary, we report the ability of topical 11 β -HSD1 inhibition to **limit** cutaneous side-effects of
331 systemic GC excess including skin thinning and delayed WH. Our findings may improve the safety
332 profile of systemic steroids and the prognosis of chronic wounds, particularly those associated with
333 elevated circulating GC levels (49).

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336 the experiments, analyzed the data and prepared the manuscript. MH prepared tissue sections for
337 histology. TM and PME contributed essential equipment. All authors evaluated and approved the
338 final manuscript.

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494 Figure Legends

495 Figure 1 Cutaneous 11 β -HSD1 expression and activity increase during wound healing and 496 systemic glucocorticoid excess

497 A) Serum corticosterone levels were downregulated in mice treated with oral corticosterone (CORT)
498 compared to vehicle (VEH, n=3-6, Student's t-test), B) 11 β -HSD1 and C) H6PD mRNA expression were
499 upregulated by CORT in unwounded skin compared to vehicle mice (VEH). Increased expression
500 post-wounding was exacerbated by CORT independently of 11 β -HSD1 inhibitor (CBX) treatment
501 (n=3-6, two-way ANOVA), D) 11 β -HSD1 enzyme activity was elevated by CORT in unwounded skin
502 and increased during healing independently of CORT. Activity was inhibited by CBX in unwounded
503 skin and during healing (n=3-5, two-way ANOVA). Multiple comparisons for 1B-1D: * = vs. baseline
504 (day 0) in each treatment group (Dunnet's test), # = vs. VEH/VEH at each time-point and † = vs.
505 VEH/CORT at each time-point (Tukey's test). Significance: * = p<0.05, ** = p<0.01, *** = p<0.001

506 Figure 2 11 β -HSD1 inhibition limits glucocorticoid-mediated skin thinning

507 A) Representative H+E staining showing decreased epidermal thickness and epidermal cellularity in
508 CORT mice and improvement following 7 days CBX treatment (n=4, one-way ANOVA), B)
509 Quantification of epidermal thickness (n=4), C) Quantification of epidermal cellularity (n=4, one-way
510 ANOVA) and D) Quantification of dermal area (n=4, one-way ANOVA). Scale bar 50 μ m. Multiple
511 comparisons for 2B-2D: # = vs. VEH/VEH and † = vs. VEH/CORT (Tukey's test). Significance: * = p<0.05,
512 ** = p<0.01, *** = p<0.001

Figure 3 Impact of glucocorticoid excess and 11 β -HSD1 inhibition on epidermal barrier function

A) Baseline trans-epidermal water loss (TEWL) in CORT mice and following 7 days CBX treatment (n=8-10, one-way ANOVA), B) Increased barrier integrity (number of tape strips required to disrupt barrier) in CORT mice and reversal by CBX co-treatment (n=8-10, one-way ANOVA) and C) Barrier recovery relative to baseline TEWL (n=7-8, two-way ANOVA). Multiple comparisons for 3A and 3B: # = vs. VEH/VEH and † = vs. VEH/CORT (Tukey's test) and for 3C: * = vs. 2 hour time-point in each treatment group (Sidak's test) and # = vs. VEH/VEH at each time-point. Significance: * = p<0.05, *** = p<0.001

Figure 4 11 β -HSD1 inhibition improves skin wound healing during glucocorticoid excess

A) Representative wound appearance at day 9 (n=8), B) Quantification of wound areas showing cessation of healing from day 4 in CORT mice and improvement of healing by CBX treatment (n=8-24, two-way ANOVA), C) Representative day 9 H+E staining showing advanced granulation tissue (arrows) formation in mice co-treated with CBX compared to CORT alone (n=3). Scale bar A) 13mm, C) 50 μ m. *=vs. day 4. Multiple comparisons for 4B: * = vs. day 4 in each treatment group (Dunnet's test), # = vs. VEH/VEH at each time-point (Tukey's test). Significance: * = p<0.05, ** = p<0.01, *** = p<0.001

Figure 5 Effect of systemic glucocorticoid excess and 11 β -HSD1 inhibition on cytokine, extracellular matrix and growth factor gene expression during wound healing

A) IL-1 β and B) TNF- α mRNA increased during WH independently of CORT or CBX, C) IL-6 mRNA showed a trend towards induction by CORT at day 2 which was suppressed by CBX, D) KGF mRNA induction at day 4 was suppressed by CORT but not CORT/CBX and induced by CORT but not CORT/CBX at day 9. E) COL-1 α 1 and F) COL-3 α 1 mRNA was induced by CORT but not CORT/CBX at day 9. G) MMP-9 mRNA induction by CORT at day 9 and was reversed by CORT/CBX. H) TIMP-1 mRNA induction at day 4 was suppressed by CORT and reversed by CORT/CBX. (n=3-6, two-way

537 ANOVA). Multiple comparisons for 1B-1D: * = vs. baseline (day 0) in each treatment group (Dunnet's
538 test), # = vs. VEH/VEH at each time-point and † = vs. VEH/CORT at each time-point (Tukey's test).
539 Significance: * = $p < 0.05$, ** = $p < 0.01$, *** = $p < 0.001$

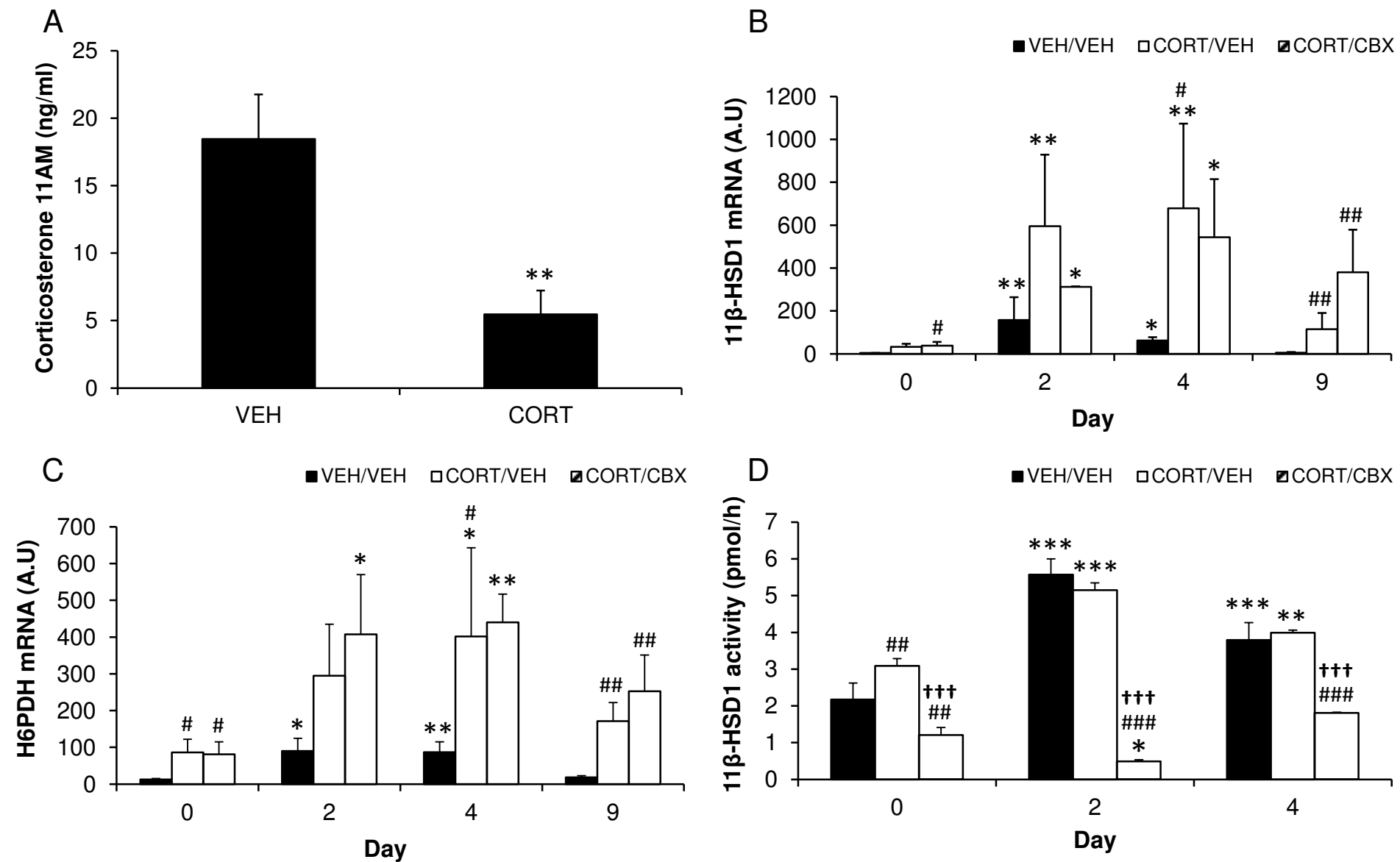
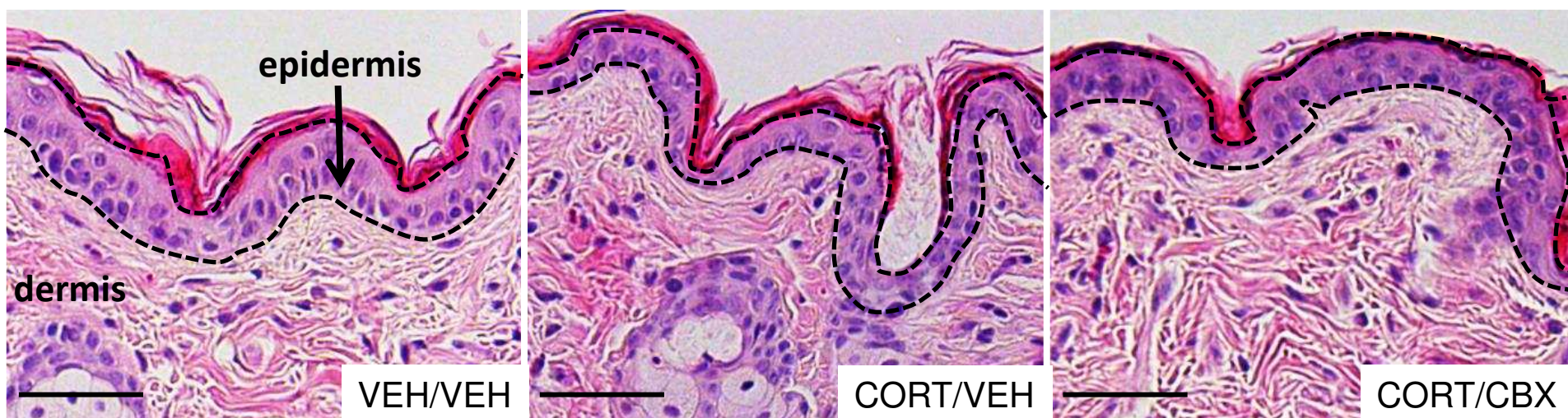
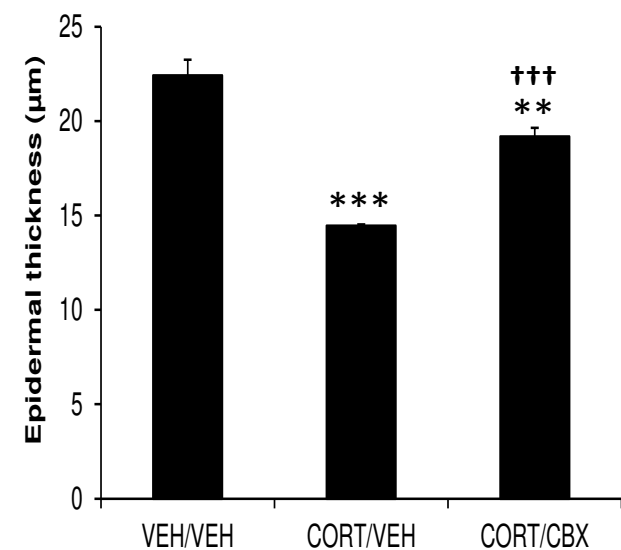


Figure 1

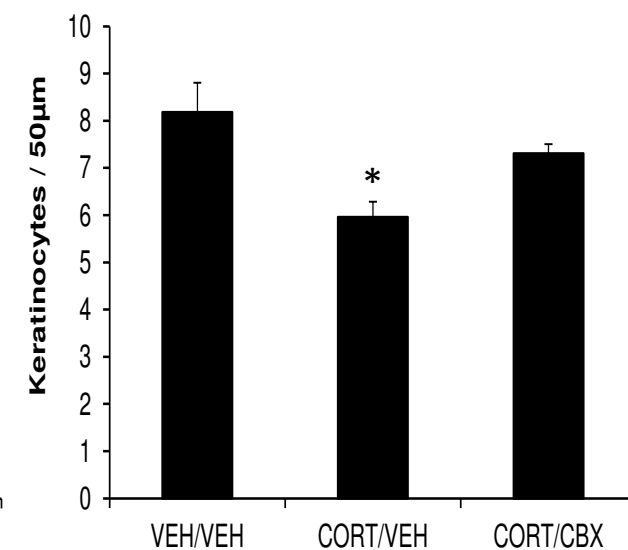
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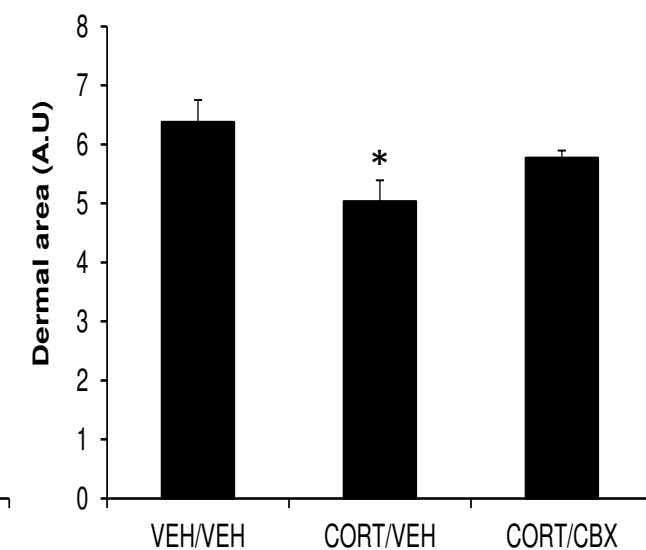


Figure 2

Figure 3

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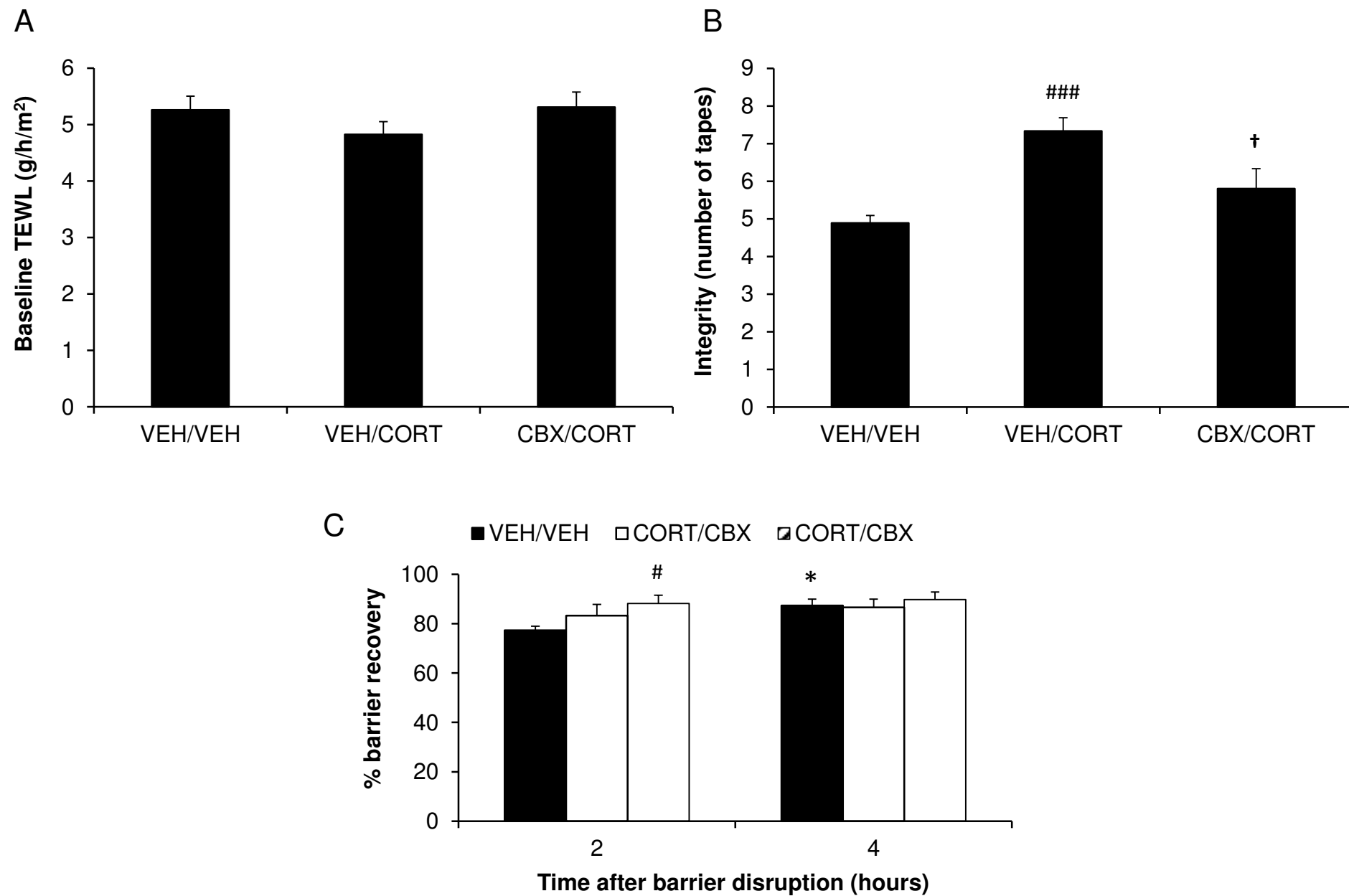


Figure 3

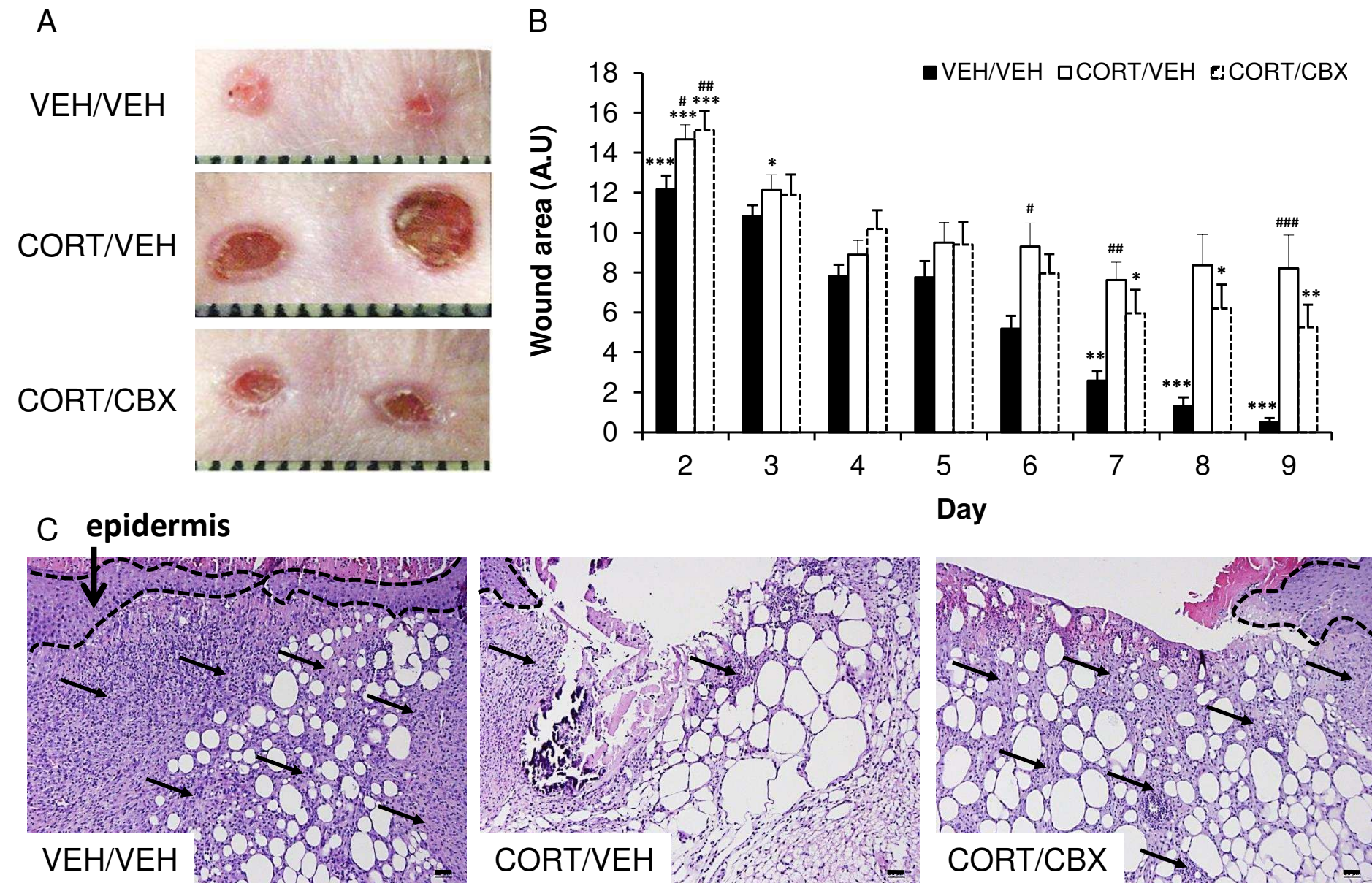


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Figure 5

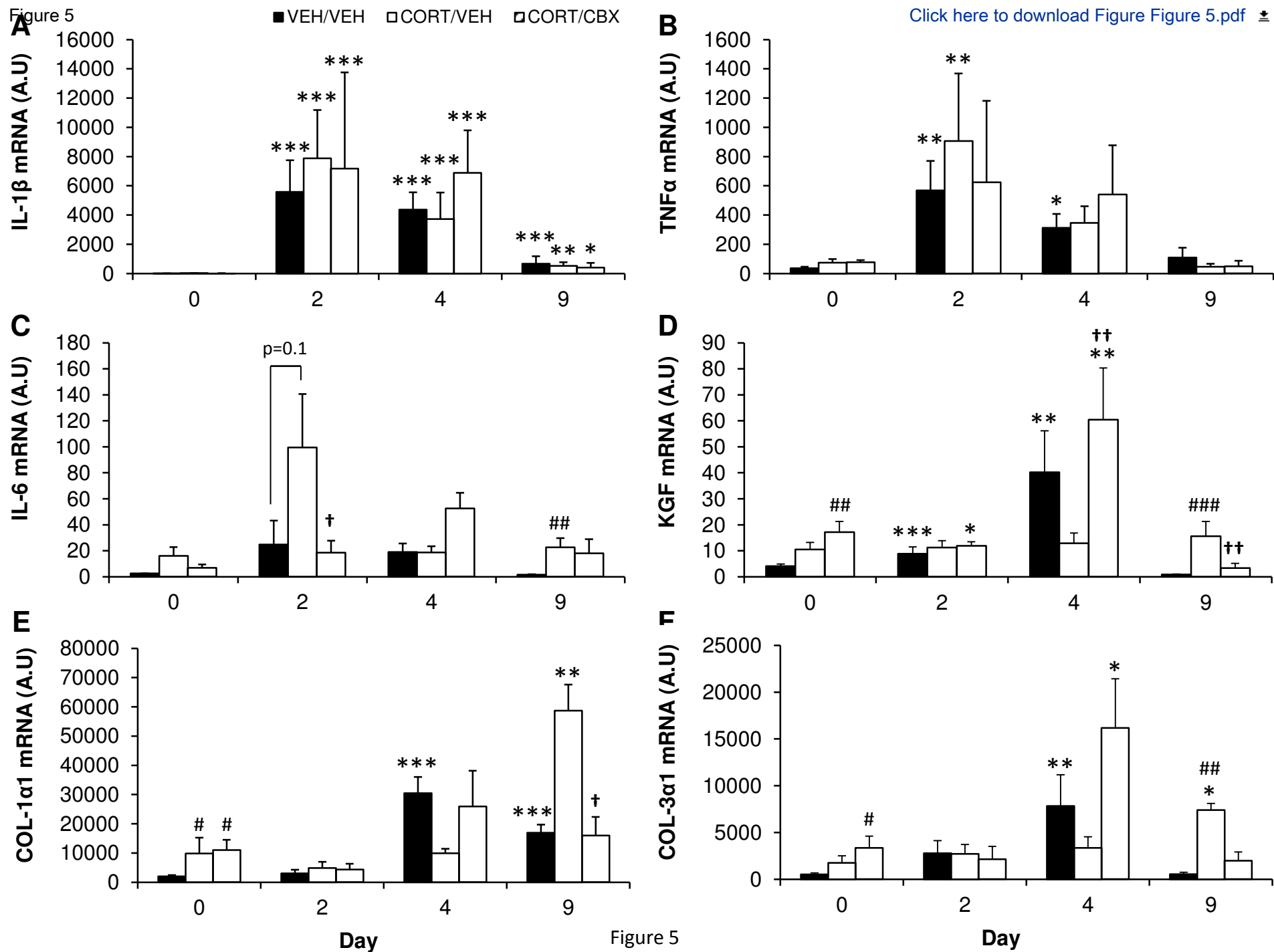


Figure 5

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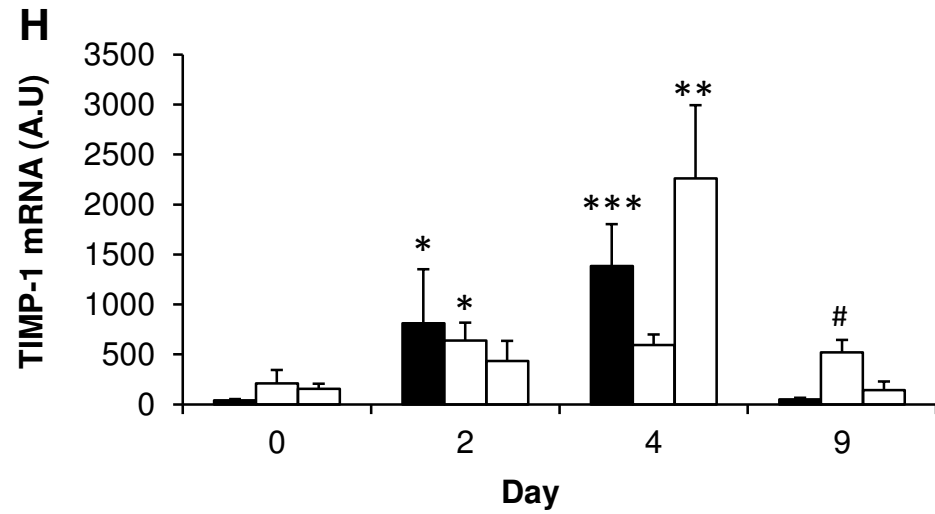
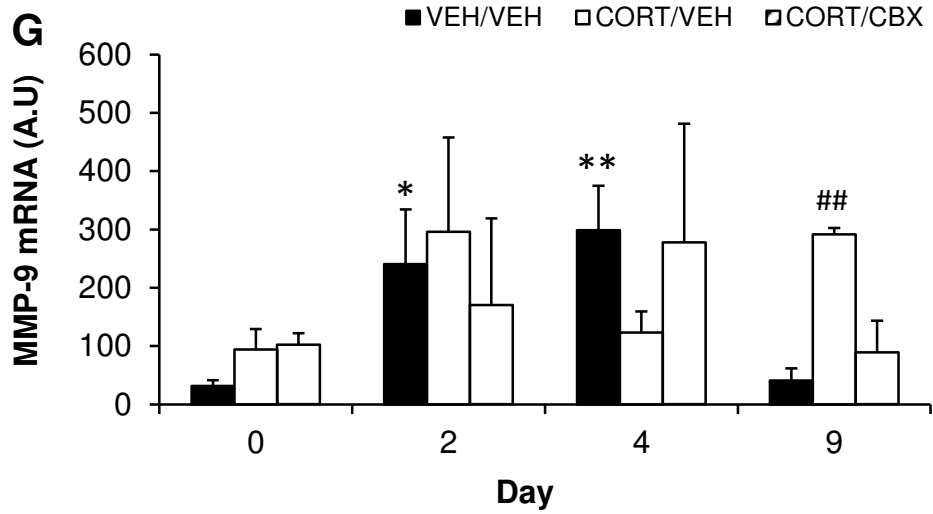


Figure 5