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693 **Supplementary Materials:**

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695 Supplemental Table 1

Measure	HV (N = 26)	Atopic Derm. (N = 20)	AD-HIES (N = 31)
Colonization	8%	85%*	97%*
<i>S. aureus</i> abscess	0	10%	94%*
<i>S. aureus</i> pneumonia	0	0	42%*
Any pneumonia	0	0	81%*

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697 **Table S1: Clinical history of *S. aureus* complications.** Healthy volunteers (HV), patients with atopic dermatitis
698 without genetic mutations, and patients with hyper IgE syndrome (AD-HIES) underwent skin culture swabs as
699 previously described (Myles, et al. *BMC Microbiol* 2016;16(1):60). Rates of *S. aureus* positivity on culture are
700 shown. Chart review results from the same patients establishing history of *S. aureus* skin abscess, microbiologically
701 confirmed *S. aureus* pneumonia, or total pneumonia (with and without identified pathogen). * = statistically
702 significant versus HV by ANOVA with Bonferroni's adjustment.

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	17A (ng/mL)		17F (ng/mL)		TNF α (ng/mL)		
HV	Saline	MRSA	Saline	MRSA	Saline	MRSA	
1	0.30	0.52	1.43	0.67	0.69	0.07	
2	0.28	0.46	1.39	0.51	0.37	0.06	
3	0.40	1.10	1.92	2.50	0.08	0.27	
4	0.35	1.05	1.86	2.48	1.00	0.25	
5	0.31	0.33	0.59	0.32	0.03	0.10	
6	0.37	0.14	0.59	0.74	0.01	0.11	
7	0.12	0.50	0.27	0.11	0.55	0.21	
8	0.09	0.19	0.22	0.11	0.03	0.20	
9	0.20	0.16	0.21	0.12	0.36	0.30	
10	0.20	2.06	0.21	0.82	0.08	0.56	
11	0.13	0.73	0.26	1.75	0.16	0.13	
12	0.61	0.37	0.54	2.55	0.16	0.09	
	17A (ng/mL)		17F (ng/mL)		TNF α (ng/mL)		
AD-HIES	Saline	MRSA	Saline	MRSA	Saline	MRSA	Mutation
1	0.54	0.38	3.39	1.90	7.20	3.08	1144C>T; R382W
2	0.52	0.36	3.43	1.94	0.02	6.09	1145G>A; R382Q
3	0.82	0.50	1.97	1.08	0.41	0.89	1387delGTG; V463del
4	0.55	0.48	1.94	1.03	0.42	0.88	1909G>A; V637M
5	0.61	0.34	0.67	0.70	0.13	0.49	1861T>G; F621V
6	0.63	0.85	0.40	0.94	0.22	3.20	1970A>G; Y657C
7	0.30	0.49	0.60	0.62	0.31	1.29	c.1997T>C; p.L666R

8	0.72	0.62	2.83	3.51	6.87	8.59	1853G>A; G618D
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708 **Table S2:** STAT3 deficiency is associated with overproduction of TNF alpha. 12 healthy volunteers (HV) and eight
709 patients with AD-HIES underwent blister induction and challenge of the wounded area with lethally irradiated
710 bacteria or diluent control. IL-17A, IL-17F and TNF α concentrations in blister fluid after 20 hours of incubation
711 with saline or irradiated methicillin resistant *S. aureus* (MRSA) are shown.

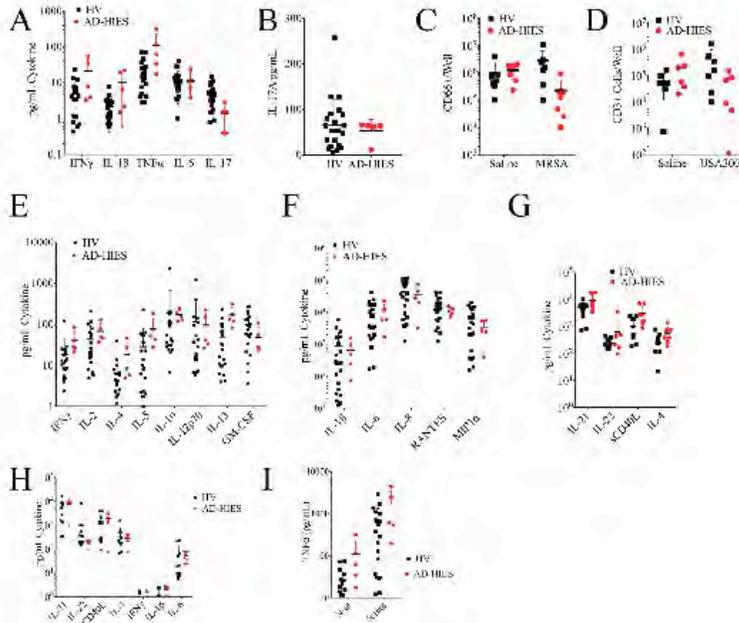
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715 **Supplemental Figure Legends:**

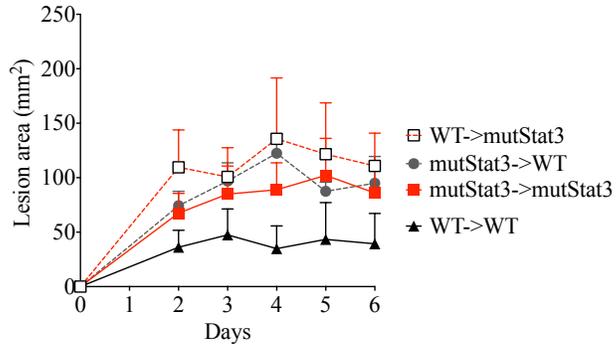
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717 **Supplemental Figure 1: Cytokine responses in blister challenge fluid.** 21 healthy volunteers (HV) and five
 718 patients with HIES underwent blister induction and challenge of the wound with 70% autologous serum. (A)
 719 cytokine levels from resultant neat fluid, as well as IL-17A from 16 hours (B) are shown. (C-D) Flow
 720 cytometry on blister exudate cells was performed on a separate cohort of seven HV and seven patients with
 721 AD-HIES 20 hours after blister challenge with either saline or irradiated MRSA. Cell count numbers for
 722 CD66+ (C) and CD3+ cells (D) are shown. (E-F) Cytokine levels from 16-hour inflammatory exudates in
 723 response to autologous serum (21 HV and five AD-HIES patients). (G-H) Cytokines levels at 20 hours from
 724 12 HV and 8 patients with HIES in response to saline (G) and irradiated MRSA (H). (I) TNF α concentrations
 725 in neat fluid and after 20 hours exposure to autologous serum for 21 HV and 5 patients with HIES. Data
 726 shown are a combination of experiments on individual patients and displayed as mean+sd. Significance
 727 determined by ANOVA with Bonferroni's correction versus HV.
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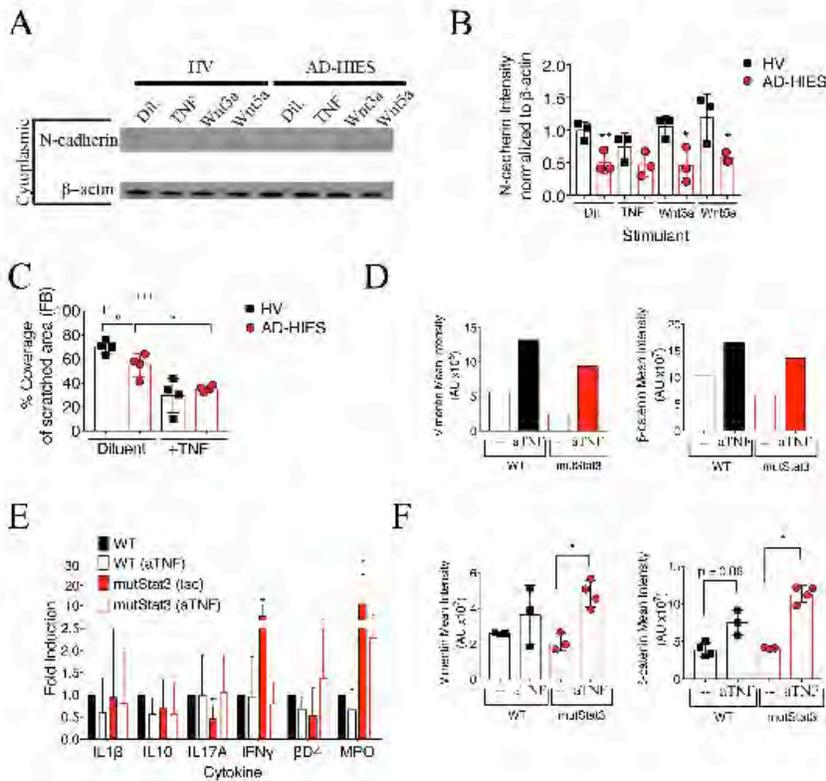
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Supplemental Figure 2: Bone marrow transplantation fails to rescue WT responses in mutStat3

mice. MutStat3 and WT mice underwent bone marrow transplantation. WT bone marrow was transplanted into WT mice (WT->WT) or mutStat3 mice (WT-> mutStat3); bone marrow from mutStat3 mice was transplanted in mutStat3 (mutStat3->mutStat3) or WT mice (mutStat3->WT). Infections performed as in Fig. 2A.



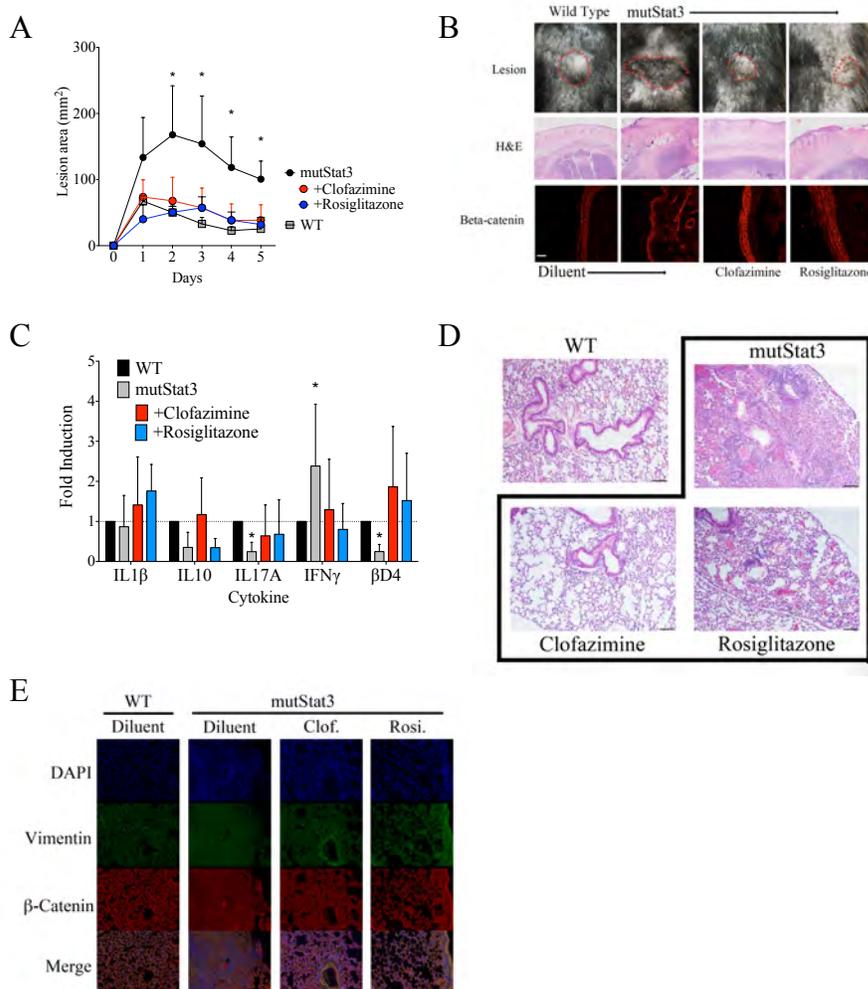
Supplemental Figure 3: Epithelial cells from patients with HIES have disordered outcomes in EMT models. (A-B) Fibroblasts from two HV and two patients with AD-HIES were stimulated with diluent (Dil.), TNF α (TNF), Wnt3a, or Wnt5a. Western blot for N-cadherin (A) and fluorescent quantification normalized to beta-actin (B) are shown for one patient and one HV within one experiment. Experiments were identical for both patient and HV samples and reproducible between experiments. Nuclear fraction western blot did not reveal any N-cadherin (not shown). (C) As in Figure 3G using fibroblasts (FB) incubated with TNF (see Video S3), data pooled from two experiments using two independent cell lines per group per experiment. (D) Quantification of fluorescent intensity for vimentin and beta-catenin from Figure 4D; average intensity taken across entire image. Images are representative of two experiments. (E) Littermate wild type (WT) and mutStat3 mice were treated with daily intraperitoneal injection of anti-TNF α (aTNF) or isotype control (Iso) for four days prior to infection with 10⁶ colony-forming units (CFU) of MRSA (N= 3-5 mice per group). Day 5 cytokine transcript levels relative to diluent treated (normalized to GAPDH). (F) Quantification of fluorescent intensity from Figure 4F as in panel D. Data shown are

761 representative of two or more independent experiments, unless indicated, and displayed as mean+sd.
762 Significance determined by ANOVA with Bonferroni's correction versus corresponding HV diluent or isotype
763 treated control. * = $p < 0.05$, ** = $p < 0.01$, ****= $p < 0.0001$.

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771 **Supplemental Figure 4: Drugs with reported EMT modulation improve outcomes in mutStat3 mice.**

772 (A-C) Wild type (WT) and mutStat3 mice were treated with daily gavage of clofazimine, rosiglitazone, or

773 diluent control for four days prior to subcutaneous infection with 10^6 colony-forming units (CFU) of

774 MRSA (N= 4-7 mice per group). Mean resultant lesion area over time (A), images of skin lesion, 20X

775 H&E histology, and 10X immunofluorescent (IF) images for beta-catenin in median mouse (B), and day 5

776 cytokine transcript levels relative to diluent treated (normalized to GAPDH; C) are shown. (D-E) WT and

777 mutStat3 mice were treated with clofazimine and rosiglitazone as above before nasal inoculation of 10^8

778 CFU of MRSA. Day 7 40X H&E (D) and 20X IF images (E) of lung are shown. Data shown are

779 representative of three independent experiments and displayed as mean+sd. Significance determined by
780 ANOVA with Bonferroni's correction versus WT diluent or isotype treated control. * = $p < 0.05$
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783 Video S1, S2, and S3 – uploaded separately

784 **Supplemental video 1:** Keratinocytes (KC) from healthy volunteers (HV) or patients with autosomal dominant
785 Hyper IgE Syndrome (AD-HIES) were placed into the scratch assay as described in the methods. Cells were pre-
786 incubated with diluent, clofazimine, or rosiglitazone prior to scratch assay. Video images run from time of scratch
787 to 22 hours later. Dotted lines added digitally to indicate starting point of KC at time zero. Images are
788 representative of three independent experiments.

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790 **Supplemental video 2:** Fibroblasts from healthy volunteers (HV) or patients with autosomal dominant Hyper IgE
791 Syndrome (AD-HIES) were placed into the scratch assay as described in the methods. Cells were pre-incubated
792 with diluent, clofazimine, or rosiglitazone prior to scratch assay. Video images run from time of scratch to 24 hours
793 later. Dotted lines added digitally to indicate starting point at time zero. Images are representative of three
794 independent experiments.

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796 **Supplemental video 3:** Fibroblasts from healthy volunteers (HV) or patients with autosomal dominant Hyper IgE
797 Syndrome (AD-HIES) were placed into the scratch assay as described in the methods. Cells were pre-incubated
798 with diluent or 2ng/mL of recombinant human TNF α . Video images run from time of scratch to 22 hours later.
799 Dotted lines added digitally to indicate starting point at time zero. Images are representative of three independent
800 experiments.

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