Supplementary Information

Psoriasis-associated variant Act1 D10N with impaired regulation by Hsp90

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**Supplementary Figure 1.** Hsp90 inhibitor attenuates IL-17-mediated pulmonary inflammation. (a) Total BAL cell and differential cell counts from DMSO or PU-H71 pretreated mice (75mg/kg of inhibitor injected intraperitoneally, once daily for two days before IL-17 injection) left unchallenged (Control) or challenged for 24 h by intranasal injection of IL-17 (1µg). Data are graphed as mean fold induction over unchallenged  $\pm$  SEM. (b) Lung tissues isolated from mice treated as in (a) were subjected to RT-PCR analysis for *II6*, *CxcI1*, and *Csf2* induction. (c) ELISA assay for CXCL1 in BAL fluid from mice treated as in (a). \**p* < 0.05 (Student's *t*-test). Data are representative of two independent experiments with 6 mice per group.



**Supplementary Figure 2.** Act1 1-100 is sufficient to interact with Hsp90. (a) HEK293 cells were transiently transfected with Flag-Hsp90 plus HA-vector, HA-Act1 WT, HA-Act1 1-300, or HA-Act1 300-574. Cell lysates were immunoprecipitated with anti-HA, followed by immunoblot analysis for Flag and HA. Arrows indicate Act1. Asterisks indicate non-specific bands. (b) HEK293 cells were transiently transfected with Flag-Hsp90 plus HA-vector, HA-Act1 WT, HA-Act1 1-100, HA-Act1 101-200, or HA-Act1 201-300. Cell lysates were immunoprecipitated with anti-HA, followed by immunoblot analysis for Flag and HA. Arrows indicate Act1. Asterisks for Flag and HA. Arrows indicate Act1. Asterisks (\*) denote none-specific bands. Data are representative of two independent experiments.



**Supplementary Figure 3.** Loss of interaction between Act1 (D10N) and CD40. HEK293 cells were transiently transfected with Flag-CD40 plus HA-vector, HA-Act1 WT or HA-Act1 (D10N). Cell lysates were immunoprecipitated with anti-HA, followed by immunoblot analysis for Flag, TRAF3 and HA. Data are representative of two independent experiments.



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**Supplementary Figure 4.** Act1 (T6BM) is a loss-of-function mutant. (a) Schematics of Act1 (T6BM) mutation (PVEVDE  $\rightarrow$  PVAVAA). (b) *Act1<sup>-/-</sup>* MEFs transduced with Act1 (WT) or Act1 (T6BM) were treated with IL-17 (50 ng/ml) for the indicated times. Cell lysates were immunoprecipitated with anti-Act1, followed by immunoblot analysis for Hsp90, TRAF6, IKKi and Act1. (c) *Act1<sup>-/-</sup>* MEFs transduced vector, Act1 (WT) or Act1 (T6BM) were treated with IL-17 (50 ng/ml) for the indicated times, followed by immunoblot analysis. (d) *Act1<sup>-/-</sup>* MEFs transduced as in (c) were treated with IL-17 (50 ng/ml) for the indicated times followed by RT-PCR analysis for *II6, Cxcl1* and *Csf2* induction. Untreated (UT). \**p* < 0.05, \*\**p* < 0.01 (Student's *t*-test). Data are representative of three independent experiments.



**Supplementary Figure 5.** Loss of mRNA stability in Act1 (D10N). *Act1<sup>-/-</sup>* MEFs were transduced with vector, Act1 (WT), or Act1 (D10N) and pretreated with TNF (10ng/ml) for 1 hour. Cells were then treated with IL-17 (50ng/ml) and actinomycin D (ActD) (5µg/ml) for the indicated times followed by RT-PCR analysis for *II6*, *CxcI1*, and *Csf2* expression. Gene expression prior to IL-17 and ActD treatment was defined as 100%. mRNA stability is defined as the percentage of mRNA that remains after Act1D treatment. Data are one representative of three independent experiments.



**Supplemental Figure 6.** RT-PCR analysis of *ll17a, ll17f,* and *ll22* transcripts in the lymph nodes of 6 weeks old Act1 WT or Act1<sup>-/-</sup> mice (n= 5 per group). \*p < 0.05, \*\*p < 0.01 (Student's *t*-test). Data are from three independent experiments.



**Supplementary Figure 7.** RT-PCR analysis of *S100a8, S100a9, Defb1* and *II8* expression in human keratinocytes treated with IL-22 (50ng/ml) for three hours. Data are presented as fold induction over untreated. \*p < 0.05, \*\*p < 0.01 (Student's *t*-test). Data are representative of two experiments.







**Supplementary Figure 8.** Skin inflammation in *ll17rc<sup>-/-</sup>* mice. (a) Cells were isolated from the lymph nodes of WT or *ll17rc<sup>-/-</sup>* mice and stimulated with PMA and ionomycin for 5 hours followed by intracellular staining for IL-17 and IL-22. Flow plots are gated on CD4<sup>+</sup> T cells. Right graph indicates the percentage of IL-17<sup>+</sup> and IL-22<sup>+</sup> CD4<sup>+</sup> T cells. (b) Skin sections from WT or *ll17rc<sup>-/-</sup>* mice stained with H&E or with anti-CD3. Scale bar indicates 50µm. (c) RT-PCR analysis of gene transcripts the skin of WT or *ll17rc<sup>-/-</sup>* mice. Data are graphed as mean  $2^{-\Delta Ct} \pm SEM$ . (d) Cytokine production by skin infiltrates isolated from the skin of WT and *ll17rc<sup>-/-</sup>* mice. Cytokine production was normalized to tissue weight. (e) Skin sections from isotype or anti-IL-22-treated *ll17rc<sup>-/-</sup>* mice. \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.005, not significant (NS) (Student's *t*-test). Data are representative of two independent experiments.

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**Supplementary Figure 9.** Loss of IL-23 signaling attenuates the hyper T<sub>H</sub>17 response in *Act1<sup>-/-</sup>* mice. (a) RT-PCR analysis of *II17a*, *II17f*, and *II22* transcripts in the lymph nodes of 6 weeks old WT, *Act1<sup>-/-</sup>*, *II23<sup>r/-</sup>*, and *Act1<sup>-/-</sup> II23<sup>r/-</sup>* mice. (b) Serum IgE levels of WT,  $Act1^{-/-}$ ,  $II23r^{/-}$ , and  $Act1^{-/-}$   $II23r^{/-}$  mice at 6mos of age. \*p < 0.05, \*\*\*p < 0.005 (Student's *t*-test). Data are representative of two independent experiments.