

### Supplemental material

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Figure S1. Targeting of IL1RAP with antibodies reduces growth of AML cells by inducing differentiation and apoptosis. (A) IL1RAP expression by flow cytometry in THP-1 cells using a fluorescently conjugated IL1RAP antibody. One representative experiment is shown. (B) Cell proliferation of THP-1 cells treated with different doses of two IL1RAP monoclonal antibodies. Data represent the mean ± SD of two independent experiments. P-values were calculated using unpaired two-tailed t tests and multiple comparisons were corrected for using the Holm-Sidak method. (C) Morphology of THP-1 cells after treatment with 150 µg/ml IL1RAP mAbs or isotype control mAbs for 24 hrs. 100× magnification, 1.25 NA. (D) Photos of THP-1 cells in liquid culture after treatment with IL1RAP antibodies. 10×, 0.25 NA. (E) Expression of macrophage differentiation markers in THP-1 cells by flow cytometry 24 h after addition of 150 µg/ml IL-1RAP mAb 2. Representative histograms are shown (top). Bar graphs (bottom) represent the mean ± SD of four independent experiments. P-values were calculated using ratio-paired two-tailed t tests comparing raw MFI values of isotype control mAb 2 treated cells versus IL1RAP mAb 2 treated cells. (F) Percentage of apoptotic (annexinV<sup>+</sup>) THP-1 cells after treatment with 150 µg/ml IL1RAP mAb 1 or mAb 2 for 72 hrs. Data represent the mean ± SD of four independent experiments. P-values were calculated using unpaired two-tailed t tests. (G) Percentage of apoptotic (annexinV<sup>+</sup>) THP-1 cells after treatment with 600 nM IL1RAP pAb, 600 nM IL-1Ra, or 6  $\mu$ M IL-1Ra for 72 h. Data represent the mean ± SD of three independent experiments. P-values were calculated using unpaired two-tailed *t* tests. (H) CD13 expression by flow cytometry in THP-1 cells using a fluorescently conjugated CD13 antibody. One representative experiment is shown (left). Cell proliferation of THP-1 cells treated with a CD13 pAb. Data represent the mean ± SD of two technical replicates (right). (I) IL1RAP expression by flow cytometry in KG-1a cells using a fluorescently conjugated IL1RAP antibody. One representative experiment is shown (left). Cell proliferation of KG-1a cells treated with 150 μg/ml IL1RAP pAb. Data represent the mean ± SD of two independent experiments (right). \*, P < 0.05; \*\*, P < 0.01; \*\*\* P < 0.001.





Figure S2. **Targeting of ILIRAP reduces growth of primary human AML cells without affecting healthy hematopoietic cells. (A)** ILIRAP shRNA transduction efficiencies of 2/5 primary AML samples used in Fig. 1 F. **(B)** Photos of individual colonies from primary AML colony assays in Fig. 1 F. 10×, 0.25 NA. **(C)** ILIRAP expression by flow cytometry in CD34<sup>+</sup>CD38<sup>+</sup> and CD34<sup>+</sup>CD38<sup>+</sup> cells from a cord blood specimen using a fluorescently conjugated ILIRAP antibody. One representative experiment is shown. **(D)** PCR amplification of the *FLT3* gene on genomic DNA isolated from THP-1 cells transduced with MSCV-GFP empty control or MSCV-FLT3-ITD-GFP. The WT allele gives a 329-bp fragment, while the transgene (which includes the *FLT3*-ITD coding sequence only) gives a 239-bp sequence (indicated by blue arrow). **(E)** PB counts of *wt* and *Il1rap<sup>-/-</sup>* mice every 2 mo until 14 mo of age. WBC, white blood cells. Dotted lines indicate normal range. **(F)** Flow cytometry analysis of PB of *wt* and *Il1rap<sup>-/-</sup>* mice at 7.5 mo of age. B lymphocytes were identified as CD11b<sup>-</sup>Gr1<sup>-</sup>B220<sup>+</sup>; T lymphocytes, CD11b<sup>-</sup>Gr1<sup>-</sup>CD3<sup>+</sup>; granulocytes, CD11b<sup>+</sup>Gr1<sup>+</sup>; monocytes, CD11b<sup>+</sup>Gr1<sup>-</sup>. Data represent the mean ± SD of six mice per group. **(G)** Percentage of c-Kit<sup>+</sup>Sca1<sup>+</sup> and c-Kit<sup>+</sup>Sca1<sup>-</sup> cells in the Lineage<sup>-</sup> population of BM of *wt* and *Il1rap<sup>-/-</sup>* mice by flow cytometry. Data represent the mean ± SD of three mice per group. **(H)** Percentage of donor cells in the PB of mice transplanted *wt* or *Il1rap<sup>-/-</sup>* cells and *wt* competitor cells. Data represent the mean ± SD of 13 mice transplanted with *wt* versus *wt* cells and 15 mice transplanted with *Il1rap<sup>-/-</sup>* versus *wt* cells.

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Figure S3. **Down-regulation, antibody targeting, or genetic deletion of ILIRAP inhibits AML pathogenesis in vivo. (A)** ILIRAP expression by flow cytometry in HEL cells using a fluorescently conjugated ILIRAP antibody. One representative experiment is shown. **(B)** Flow cytometry histograms showing ILIRAP protein knockdown achieved by ILIRAP shRNAs in cell lines used for xenografts in Fig. 2 B. **(C)** ILIRAP expression was measured in ILIRAP shR-NA-transduced HEL cells isolated from BM of xenografted mice (Fig. 2 B) at day 30. HEL cells were identified by lack of mouse CD45.1 expression and presence of GFP (contained in the shRNA vectors). **(D)** Total body weights of antibody-treated THP-1 xenograft mice over the time course of the experiment in Fig. 2 C. tx, treatment. **(E)** Representative images of livers from antibody-treated THP-1 cell infiltration (white arrowheads) and residual normal mouse liver (black arrowheads) are indicated. 4×, 0.13 NA. **(F)** ILIRAP expression in BM aspirates from MLL-AF9 leukemic mice by flow cytometry. MLL-AF9-transduced BM cells (GFP<sup>+</sup>, green histograms) and residual nonleukemic BM (GFP<sup>-</sup>, blue histograms) are shown. **(G)** Leukemic cell chimerism in BM of MLL-AF9 primary recipient mice 4 wk after transplant (experimental scheme is shown in Fig. 2 F). Data represent the mean ± SD of nine mice per group. **(H)** Kaplan-Meier plot showing survival of primary recipient mice transplanted with *wt or Il1rap<sup>-/-</sup>* MLL-AF9 cells. P-value was calculated using log-rank (Mantel-Cox) test. **(I)** Immunophenotypes of *wt* and *Il1rap<sup>-/-</sup>* MLL-AF9 donor cells used for secondary transplant in Fig. 2 (G and H). **(J)** Left, spleen weights of *wt*, homozygous *FLT3*-ITD knock-in mice (*Il1rap<sup>+/+</sup>FLT3<sup>ITD/ITD</sup>*) and *Il1rap<sup>-/-</sup>FLT3<sup>ITD/ITD</sup>* mice at 12 mo of age. Right, flow cytometric analysis of spleen cell populations in *wt*, *Il1rap<sup>+/+</sup>FLT3<sup>ITD/ITD</sup>*, and *Il1rap<sup>-/-</sup>FLT3<sup>ITD/ITD</sup>* mice at 12 mo of age. P-values were calculated using unpaired two-tailed t-tests. \*, P < 0.05; \*\*\*,

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Figure S4. Down-regulation, antibody targeting, or genetic deletion of IL1RAP inhibits FLT3 ligand and SCF-mediated signaling and growth, and ILIRAP interacts with their respective receptors. (A) Top, surface expression of ILIRAP, IL-1RI, FLT3, and c-KIT by flow cytometry in THP-1 cells. One representative experiment is shown. Bottom, flow cytometry histograms showing IL1RAP protein knockdown achieved by IL1RAP shRNA#2 in THP-1 cells used for western blots in Fig. 3 A. (B) Top, surface expression of IL1RAP, IL-1RI, FLT3, and c-KIT by flow cytometry in TF-1 cells. One representative experiment is shown. Bottom, flow cytometry histograms showing IL1RAP protein knockdown achieved by IL1RAP shRNA#1 in TF-1 cells used for western blot in Fig. 3 F. (C) Surface expression of IL1RAP, FLT3, and c-KIT by flow cytometry in HEL cells. One representative experiment is shown. (D) Left, quantification of p-FLT3 (Y591) from Western blot in Fig. 3 A. Right, quantification of p-c-KIT (Y719) from western blot in Fig. 3 F. Dotted line indicates unstimulated, control shR-NA-transduced cells. Protein levels relative to actin are shown. (E) THP-1 cells were treated with IL1RAP mAb, IL1RAP pAb, or isotype controls (100 µg/ml) for 30 minutes. Protein lysates were examined for p-FLT3 (Y591) and total FLT3 by immunoblotting. (F) Quantification of p-FLT3 (Y591) from Western blots in Fig. 3 (B and C). Dotted line indicates unstimulated, isotype control antibody treated cells. Protein levels relative to actin are shown. Error bars represent the mean ± SD of independent experiments. (G) 293T cells were transfected with HA-tagged IL1RAP and examined for IL1RAP and p-FLT3 (Y591) 18 h later by Western blot (top). Surface overexpression of IL1RAP in 293T cells by flow cytometry (bottom). (H) Quantification of p-c-KIT (Y719) from Western blots in Fig. 3 (G and H). Dotted line indicates unstimulated, isotype control antibody treated cells. Protein levels relative to actin are shown. Error bars represent the mean ± SD of independent experiments. (I) Quantification of p-FLT3 (Y591) from western blots in Fig. 3 D. Dotted line indicates unstimulated, isotype control antibody treated cells. Protein levels relative to ACTIN are shown. (J) Quantification of Western blot in Fig. 3 I. Protein levels relative to ACTIN are shown. Data represent the mean ± SD of two independent experiments. P-values were calculated using unpaired two-tailed t tests. (K) Surface expression of IL1RAP, IL-1RI, FLT3, and c-KIT by flow cytometry in AML cell lines used in Fig. 4 (A-G). "Positive" gates were set using a fluorescent isotype control antibody. Data represent the mean ± SD of two or more independent experiments. (L) Coimmunoprecipitation of endogenous c-KIT in protein lysates from THP-1 cells using an independent c-KIT antibody. Immunoblot was performed for c-KIT and IL1RAP. \*, P < 0.05; \*\*, P < 0.01.

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#### Table S1. Absolute colony number output for IL1RAP shRNA-transduced primary AML samples used in Fig. 1 F

	Control shRNA	IL1RAP shRNA#1	IL1RAP shRNA#2
MDS01	48	14	13
MDS02	20	8	0
AML03	182	124	57
AML07	5538	3268	2320
AML13	10	7	5
AML15	46		26

Table S2. Flow cytometer configuration used for determination of FRET in Fig. 4 (C and G)

	Photomultiplier tube	Dichroic mirror	Band pass filter	Channel
Blue laser, 488 nm	A	735LP	780/60	PE-Cy7
	В	635LP	660/20	FRET
	C	595LP	610/20	PE-Texas red
	D	556/LP	575/26	PE
	E	502LP	530/30	FITC/GFP/Alexa488
	F		488/10	SSC
Red laser, 633 nm	A	755LP	780/60	APC-Cy7
	В	690LP	730/45	Alexa700
	С		660/20	APC/Alexa647
Violet laser, 407 nm	A	505LP	530/30	AmCyan/Pacific orange
	В		450/50	Pacific blue/eFluro450/BV421/CFP/Dapi