1	Supplementary Materials for
2 3 4	Targeting rapid TKI-induced AXL upregulation overcomes adaptive ERK reactivation and exerts antileukemic effects in <i>FLT3</i> /ITD acute myeloid leukemia
5 6	Tessa S. Seale, Li Li, J. Kyle Bruner, Melody Chou, Bao Nguyen, Jaesung Seo, Ruiqi Zhu, Mark J. Levis, Christine A. Pratilas, and Donald Small
7	Correspondence to: donsmall@jhmi.edu
8	This PDF file includes:
9	Supplementary Figure Legends
10	Figures. S1 to S8
11	Tables S1 to S2
12	
13	
14	Supplementary Figure Legends
15	Supplementary Table 1. Combination Index (CI) values for FLT3 TKI in combination with
16	AXL inhibitors. Data was collected 48 hours after treatment, with the exception of MV4;11
17	gilteritinib and TP-0903 Annexin V combination, which was 72 hours after treatment. CI values
18	were calculated at ED50 using CompuSyn software.
19	Supplementary Table 2. AML Patient Characteristics
20	Supplementary Figure 1. FLT3 TKI treatment increases AXL activation and lowers GAS6
21	levels in FLT3/ITD cell lines. MV4;11 cells were treated with 25 nM of sorafenib for 48 hours.
22	Protein lysates were collected at 0 hour and 48 hours and assayed with PathScan® RTK
23	Antibody Array panels (A). (B) The signal intensities of untreated and 48-hour duplicates were
24	quantified and adjusted against the local background using Image Lab software. Next, the
25	adjusted signal intensity was first normalized against the positive control intensities on each
26	array, and then the 48 hour was normalized against the untreated. Error bars indicate average

27 signal intensity ± SD. (C) Molm14 cells were treated with 25 nM of sorafenib for 24 hours. 28 Protein lysates were subjected to immunoblot analysis against the indicated antibodies. (D) MV4:11 and Molm14 cells were treated with 25 nM of sorafenib for 24 hours and GAS6 mRNA 29 30 expression was measured at the indicate time points using quantitative RT-PCR as normalized 31 to GAPDH. Error bars indicate average mRNA expression ± SEM. (E) The indicated cell lines 32 are treated with 100 nM of gilteritinib or 10 nM of guizaritinib for 24 hours. Protein lysates are collected at the indicated time points and subjected to immunoblot analysis. CK means 33 cytokines, and + Ctrl means positive control, which is Molm14 cells are treated with 100 nM of 34 35 gilteritinib for 24 hours. Supplementary Figure 2. AXL overexpression increases ERK phosphorylation and 36 37 proliferation (A) Molm14 cells transduced with either a control vector (Control) or a constitutive 38 AXL overexpression vector (AXL+) were treated with the indicated drugs for 4 hours. Protein 39 lysates were collected at the indicated time points and subjected to immunoblot analysis. The 40 black lines intersecting the blots indicate that the bands on either side of the line were not originally next to each other on the original immunoblot. (B) Molm14 cells transduced with 41 either a doxycycline inducible non-specific control (indControl) or an inducible AXL 42 43 overexpression construct (indAXL+) were treated with sorafenib and 0.5 µg/mL of doxycycline for 8 hours. The protein lysates were collected at the indicated time points and subjected to 44 45 immunoblot analysis. The black lines intersecting the blots indicate that the bands on either side of the line were not originally next to each other on the original immunoblot. (C) ) Molm14 and 46 47 MV4;11 cells transduced with either a control vector (Ctrl) or a constitutive AXL overexpression 48 vector (AXL+) were treated with the indicated concentrations of gilteritinib for 48 hours and subjected to MTT analysis. OD stands for optical density and error bars are average OD ± SD. 49 50 Supplementary Figure 3. AXL levels are unaffected by MEK inhibition but are reduced by 51 PI3K and YAP inhibition. (A) Molm14 and MV4;11 cells were treated with 25 nM sorafenib and either 10 µM of U0126 or DMSO for 24 hours. Protein lysates were collected at the indicated time points and subjected to immunoblot analysis against the indicated antibodies. The black lines intersecting the blots indicate that the bands on either side of the line were not originally next to each other on the original immunoblot. (B) Molm14 and MV4;11 cells were treated with 100 nM of gilteritinib, 3 nM of trametinib, or the combination for 24 hours and AXL mRNA expression was measured using qRT-PCR normalized to untreated control. Error bars indicated average expression ± SEM. (C) Molm14 and MV4;11 cells were treated with the indicated drug concentrations for 24 hours. Protein lysates were collected at the indicated time points and subjected to immunoblot analysis against the indicated antibodies. The line bisecting the blot indicates that the lanes were not next to each other originally and were cropped as a result. (D) MV4:11 cells transfected with either a control siRNA (ctrl) or siRNA against YAP (siYAP) were treated with 100 nM of gilteritinib for 24 hours. Protein lysates were collected at the indicated time points and subjected to immunoblot analysis against the indicated antibodies. (E, F) Molm14 and MV4;11 cells were treated with gilteritinib alone or with verteporfin (E) or VT107 (F) for 24 hours. Protein lysates were collected at the indicated time points and subjected to immunoblot analysis against the indicated antibodies. The line bisecting the blot indicates that the lanes were not next to each other originally and were cropped as a result. Supplementary Figure 4. Combination FLT3 TKI and AXL inhibitors decrease proliferation and increase apoptosis. (A) MV411 and Molm14 cells were treated with the indicated drug treatments for 48 hours. Proliferation was measured by MTT assay. Error bars indicate mean percent OD ± SD. (B) MV4;11 cells were treated the indicated drugs for 72 hours. Apoptosis was assayed with Annexin V staining. Error bars indicate average fold change of % Annexin V vs. untreated ± SD. Asterisks directly above the error bar indicate P-value compared to untreated (\*P <0.05, \*\*P < 0.01, \*\*\*P <0.001). (C) Molm14 (left) and MV4;11 (right) cells were assayed with the indicated MTT or Annexin V assays with the indicated drug combinations. The

52

53

54

55

56

57

58

59

60

61

62

63

64

65

66

67

68

69

70

71

72

73

74

75

76

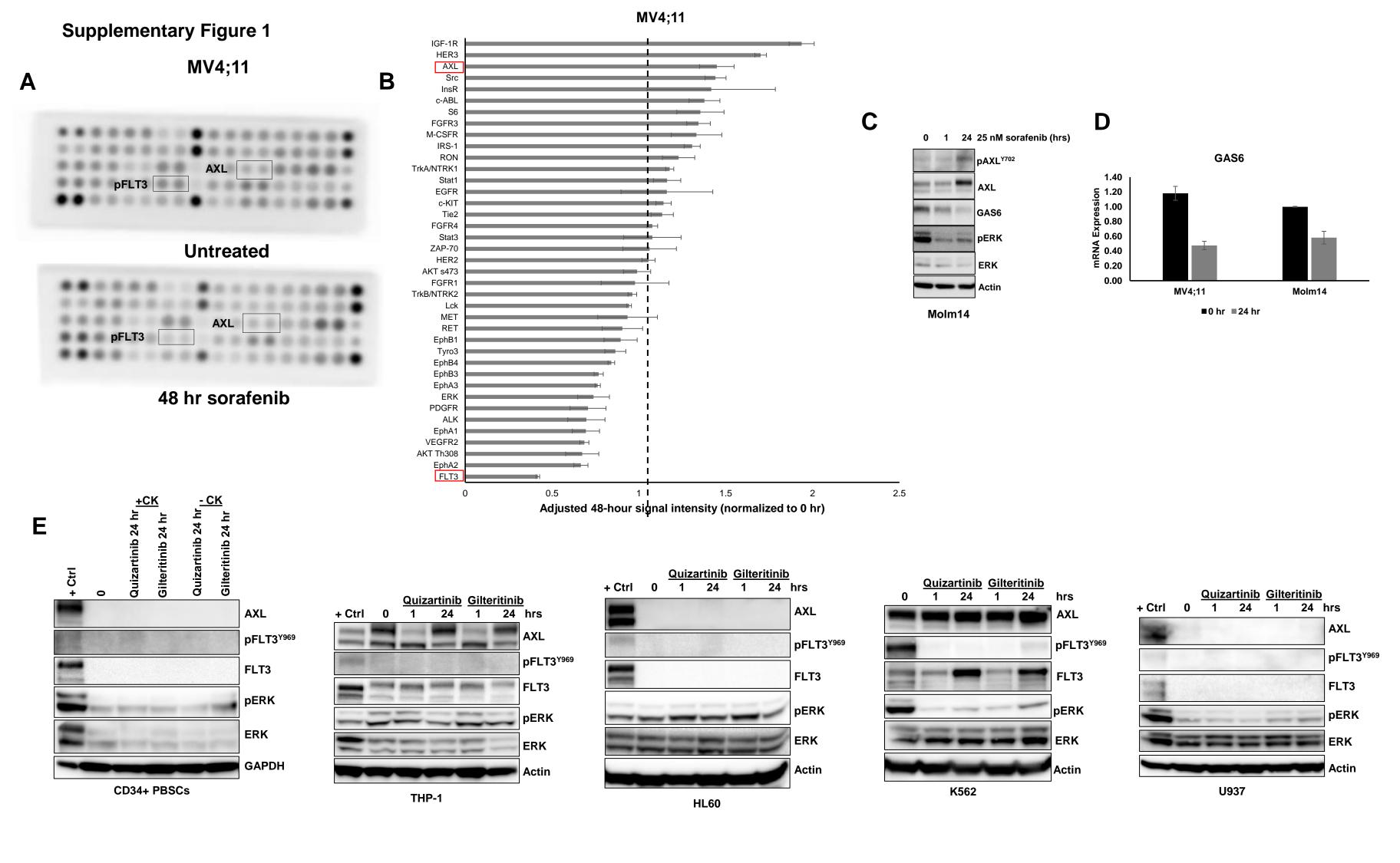
- dose reduction index (DRI) was calculated for sorafenib in combination versus sorafenib alone.
- 78 The dotted lines indicate the normalized sorafenib alone value.
- 79 Supplementary Figure 5. **AXL inhibition through genetic knockdown and ligand trapping**
- affects pERK rebound. (A) Molm14 cells stably transfected with a shRNA construct against
- AXL (shAXL-2) were treated either with or without 0.5 µg/mL of doxycycline for 24 hours.
- 82 Protein lysates were subjected to immunoblot analysis. (B) Molm14 cells stably transfected with
- either a non-silencing control vector (control) or with an shRNA construct targeting AXL (shAXL-
- 84 2) were treated with 25 nM of sorafenib and 0.5 μg/mL of doxycycline for 24 hours. Protein
- lysates were subjected to immunoblot analysis against the indicated antibodies. The blots are
- representative of at least two independent experiments. (C) Molm14 (left) and MV4;11 (right)
- cells are treated with 25 nM of sorafenib and either 3 µg/mL of Ctrl-Fc or AXL-Fc for 24 hours.
- 88 Protein lysates were collected at the indicated time points and subjected to immunoblot assay.
- 89 (D) MV4;11 cells were treated with the indicated drug combinations (sorafenib concentration: 25
- 90 nM, Ctrl-Fc and AXL-Fc concentrations: 1 µg/mL) for a total of three days. Cells were counted at
- 91 the indicated time points and percent cell viability was assessed using trypan blue exclusion
- 92 assay. The experiments were done in triplicate and the error bars show the average percent
- 93 viability  $\pm$  SD (\*\*P < 0.01).
- 94 Supplementary Figure 6. **AXL and FLT3 inhibition decrease FLT3/ITD primary cell viability**.
- The indicated primary cell samples are treated with 80 nM of gilteritinib, 80 nM of TP-0903, 1 M
- 96 of R428, or the combination for 24 hours. Cell viability was measured using Trypan blue
- 97 exclusion assay.
- 98 Supplementary Figure 7. TKI-induced AXL perturbations are observed in other RTK-driven
- 99 cancers (A) HER2-amplified SKBR3 cells were treated with 75 nM of lapatinib, 100 nM of TP-
- 100 0903 and the combination for 24 hours. Protein lysates were analyzed with immunoblot
- analysis. (B) SKBR3 cells were treated with 75 nM of lapatinib. mRNA was subjected to

quantitative RT-PCR for AXL and GAS6. Expression values were measured in triplicate relative to GAPDH levels. Error bars indicate average expression ± SEM (C) EGFR-mutated HCC827 cells were treated with 100 nM of erlotinib for 24 hours, and then the cells were washed and replated and treated with either 100 nM of erlotinib, 1000 nM of R428, and the combination for an additional 1 hour. The vertical line indicates point of initial drug removal. Protein lysates were subjected to immunoblot analysis. (D) HCC827 cells are treated with 750 nM of lapatinib. mRNA was subjected to quantitative RT-PCR for AXL and GAS6 relative to GAPDH. Expression values were measured in triplicate. Error bars indicate average fold change ± SEM.

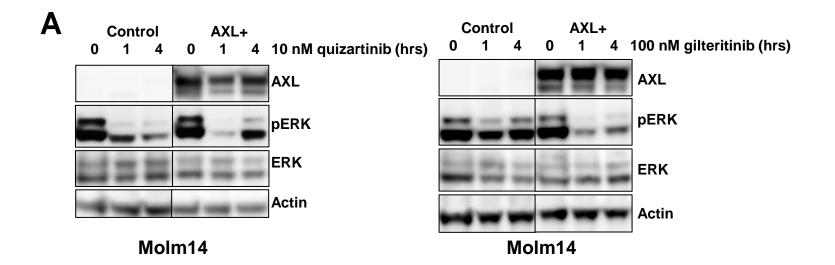
Supplementary Fig 8. Recovery leukemia burden of shAXL knockdown mice. Inducible shAXL-2 and control Molm14 cells were injected into NSG mice. 72 hours after injection, the mice were randomized into groups and treated with either 15 mg/kg, 30 mg/kg of gilteritinib, or vehicle once daily (excluding weekends). 500 µg/mL of doxycycline was provided daily via acidified drinking water from the animal facility. Recipients were treated for 20 days. 21 days post the end of treatment, bone marrow was collected via aspiration from surviving mice of the indicated groups and evaluated for percent of human CD45 positive leukemia cells. Error bars indicate the average % hCD45 ± SD. (B) Human CD34+ PBSCs from healthy donors are treated with gilteritinib, R428, TP-0903 or the combination and plated in methylcellulose for a colony formation assay with three biological replicates. Error bars represent average number of colonies ± SD.

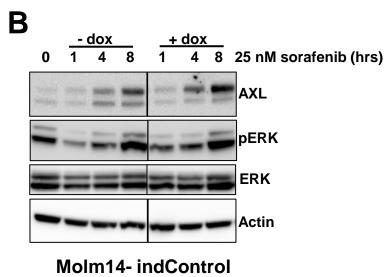
Drug Combination	Molm14	MV4;11
Sorafenib + R428: MTT	0.49	0.077
Gilteritinib + R428: MTT	0.85	0.29
Gilteritinib + R428: Annexin V	0.66	0.23
Gilteritinib + TP-0903: Annexin V	0.61	0.34

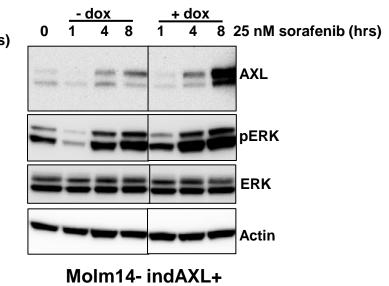
Sample	Age/Gender	FLT3 mutation Status	WBC (cells/µL)	Cytogenetics	Newly- diagnosed/Relapse
863	82/F	FLT3/ITD, 96 bp insertion, VAF 46%	116,000	Normal karyotype	Newly-diagnosed
092	59/F	FLT3/ITD, 54 bp insertion, VAF 73.7%	17,600	del16q	Relapsed
148	76/F	FLT3/ITD, 60 bp insertion, VAF, 29.8%	135,600	Normal karyotype	Newly-diagnosed
842	70/M	FLT3/ITD 352 bp insertion, VAF 181%	93.750	t(4;12)(q12;p13),del(7)( q31q35)	Relapsed
310	62/F	FLT3/ITD, 364 bp insertion, VAF 55%; 349 bp insertion, VAF <5%	318,160	+8	Newly-diagnosed
431	44/F	FLT3/ITD	N/A	Normal karyotype	Newly-diagnosed
336	28/F	FLT3/ITD	N/A	Normal karyotype	Newly-diagnosed



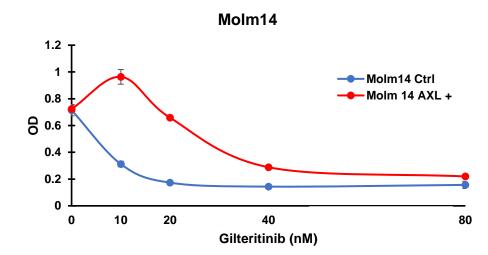
## **Supplementary Figure 2**

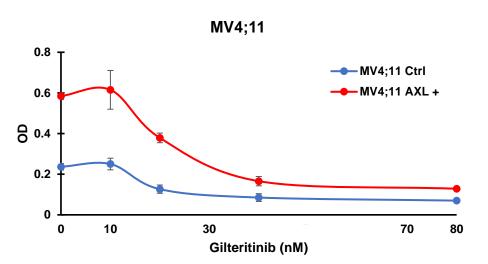


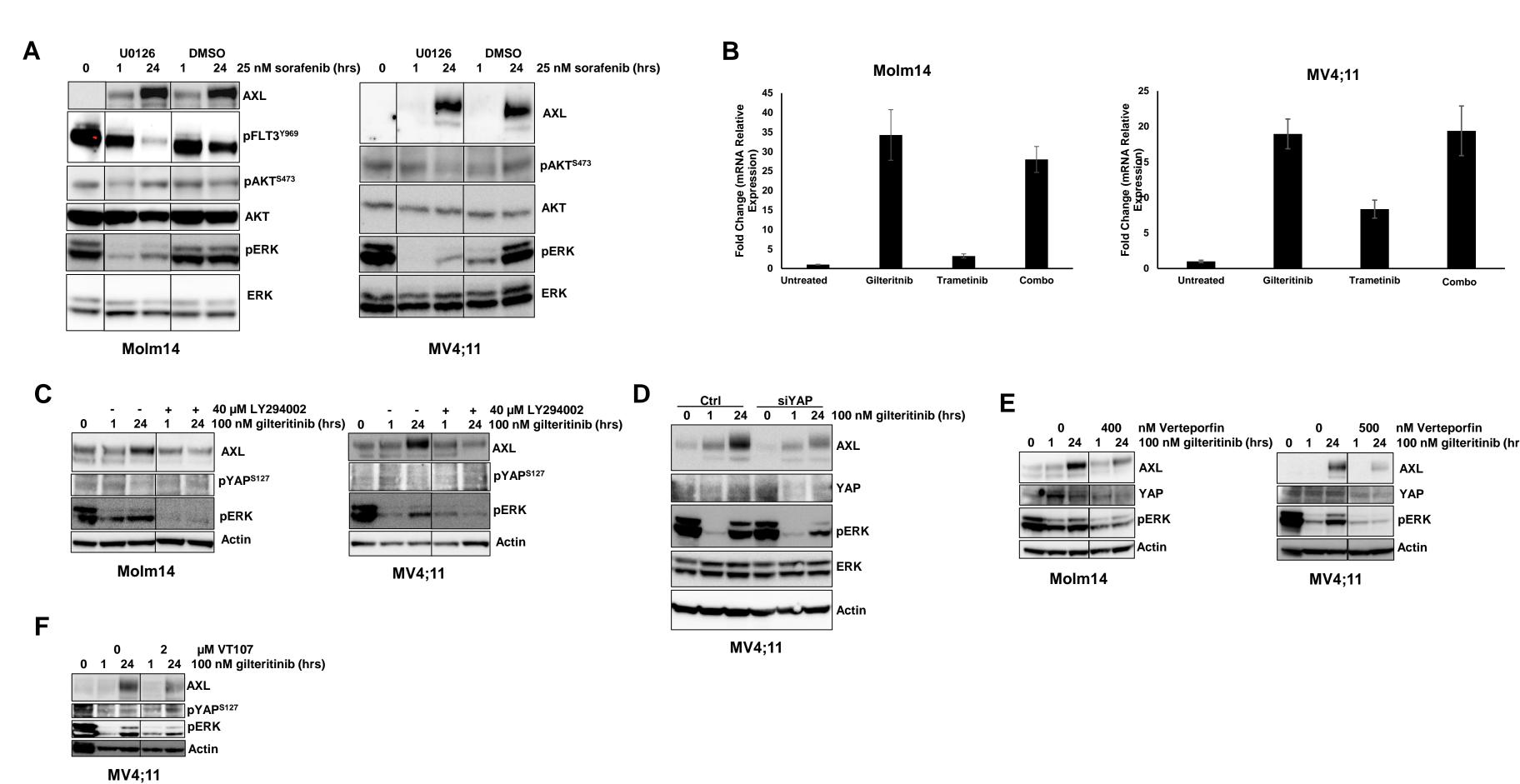




C



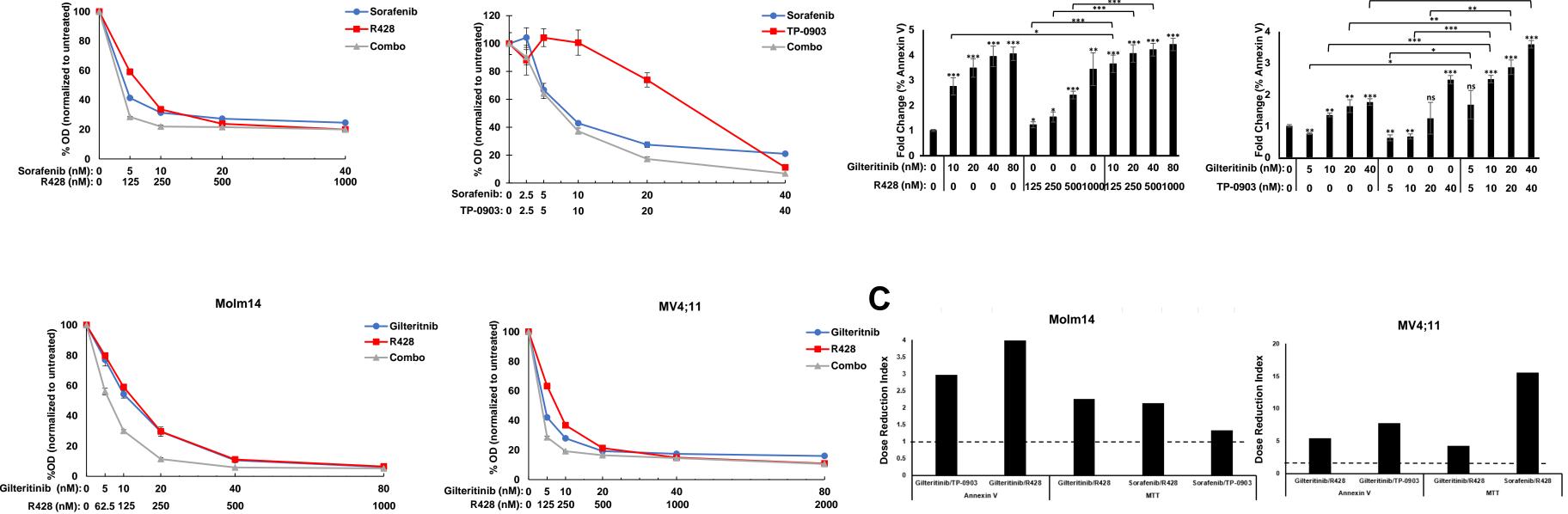




## **Supplementary Figure 4**

MV4;11

A

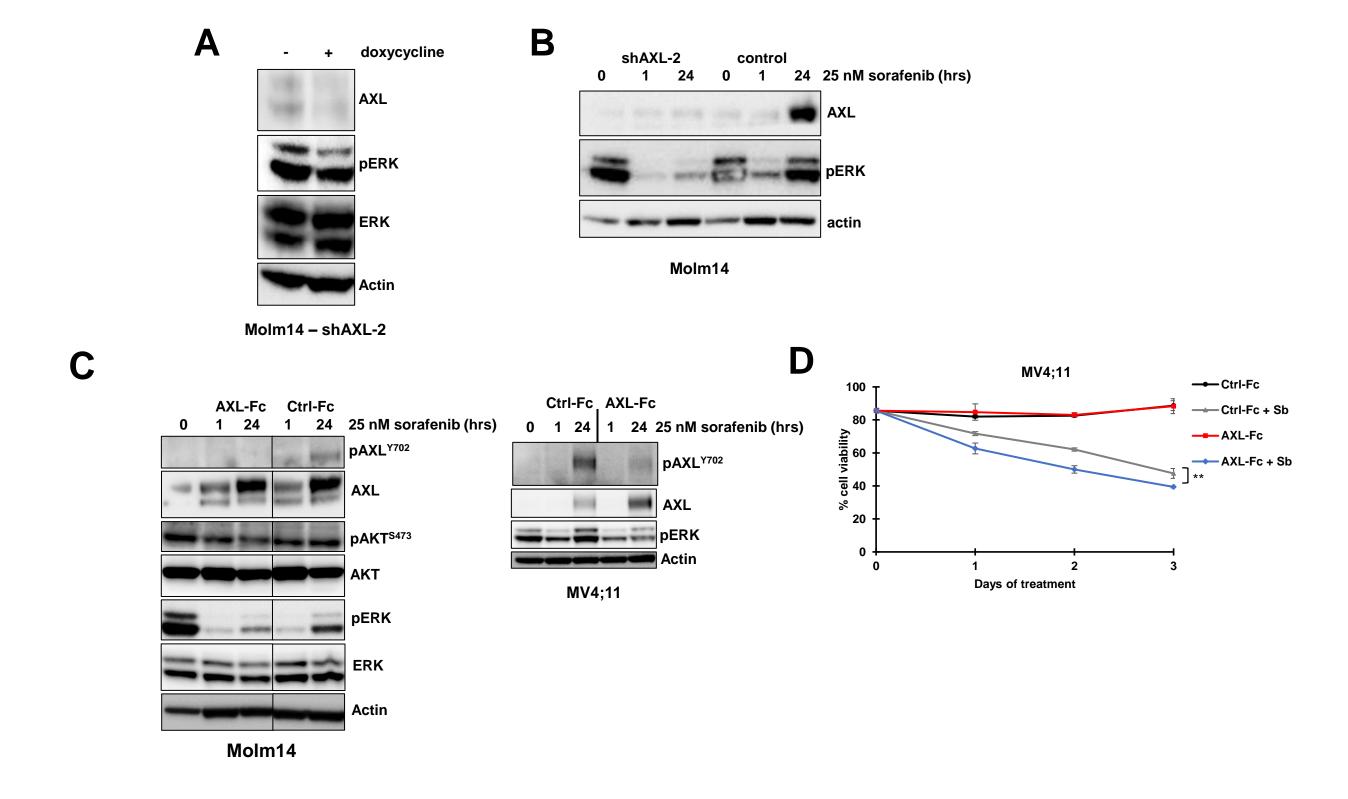


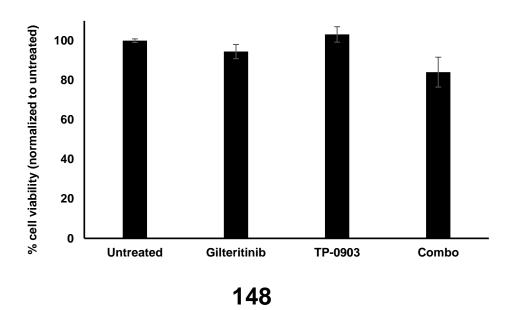
MV4;11

B

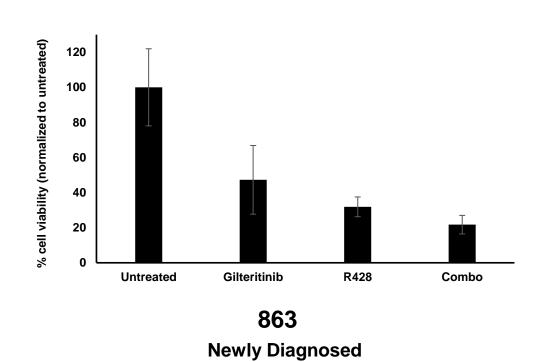
MV4;11

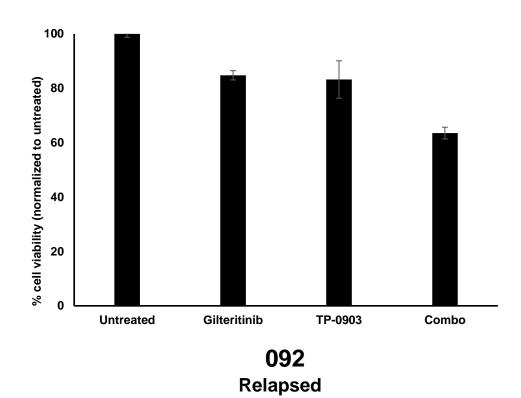
MV4;11



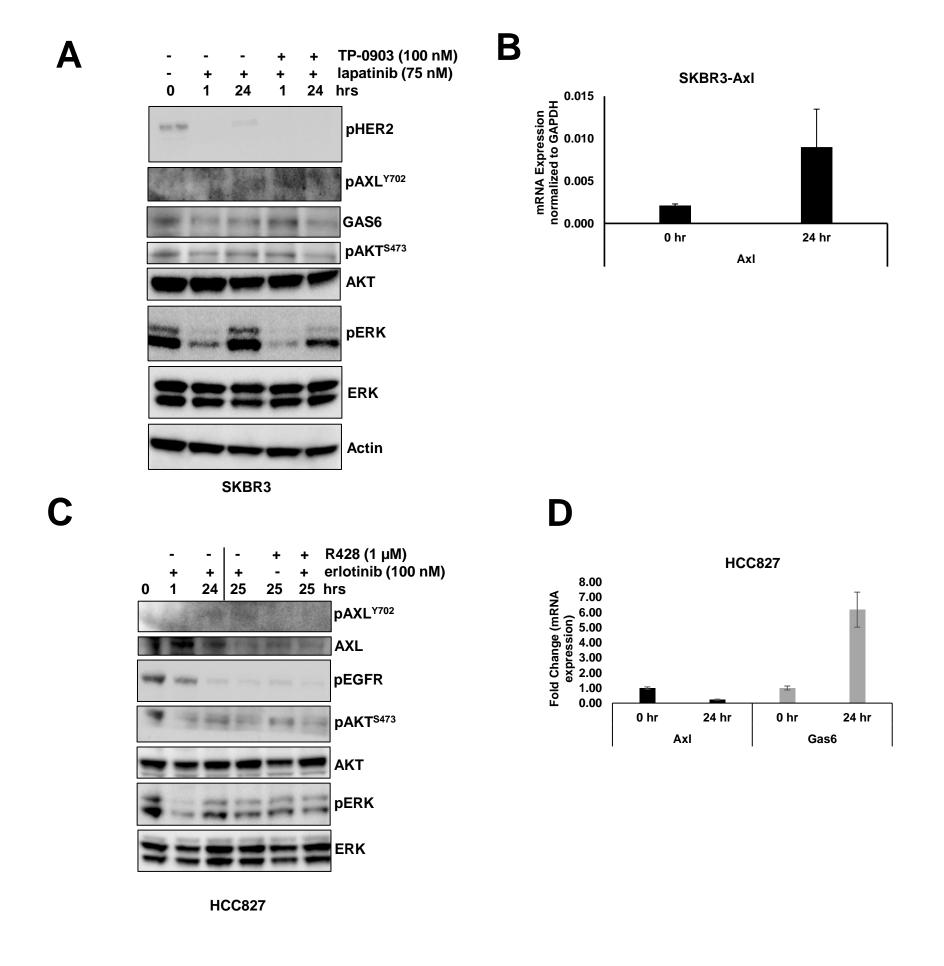


**Newly Diagnosed** 





## **Supplementary Figure 7**



SKBR3-Gas6

Gas6

24 hr

0 hr

3.00

