

SM09419, a Novel, Small-Molecule CDC-like Kinase (CLK) Inhibitor, Demonstrates Strong Inhibition of the Wnt Signaling Pathway and Antitumor Effects in Tumor Protein p53 (TP53)-Mutant Acute Myeloid Leukemia Models

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Poster #3913

Background

- Acute myeloid leukemia (AML) with TP53 mutation (TP53mut) accounts for ~13% of AML cases and is an aggressive, treatment-resistant subtype of cancer with dismal prognosis and limited therapeutic options¹
- Aberrant activation of the Wnt signaling pathway is associated with AML initiation and progression; it is also required for the self-renewal and survival of leukemic stem cells, making Wnt signaling inhibition a potential therapeutic modality for adverse AML²⁻⁴
- CLKs regulate the activity of serine/arginine-rich splicing factors (SRSFs) that modulate spliceosome assembly, mRNA splicing, and gene expression^{5,6}
- SM09419 is a novel, oral, small-molecule pan-CLK inhibitor that potently inhibits the Wnt pathway⁷
- These studies examined the antitumor activity of SM09419 in preclinical models of TP53mut AML

Conclusions

- SM09419 demonstrated potent *in vitro* antileukemic effects and inhibited SRSF activation and Wnt pathway signaling in TP53mut AML cells
- SM09419 had strong *in vivo* antileukemic activity as a single agent and in combination with VEN and/or AZA; SM09419 was well tolerated, except in the triplet combination
- These results suggest that SM09419 is a potential treatment for hard-to-treat subtypes such as TP53mut AML
- A Phase 1 study assessing the safety, tolerability, and pharmacokinetics of SM09419 in subjects with advanced hematologic malignancies is being initiated

Results

Figure 1. SM09419 inhibited proliferation in AML cell lines and primary human AML cells regardless of TP53 mutation status, cytogenetics, or diagnosis

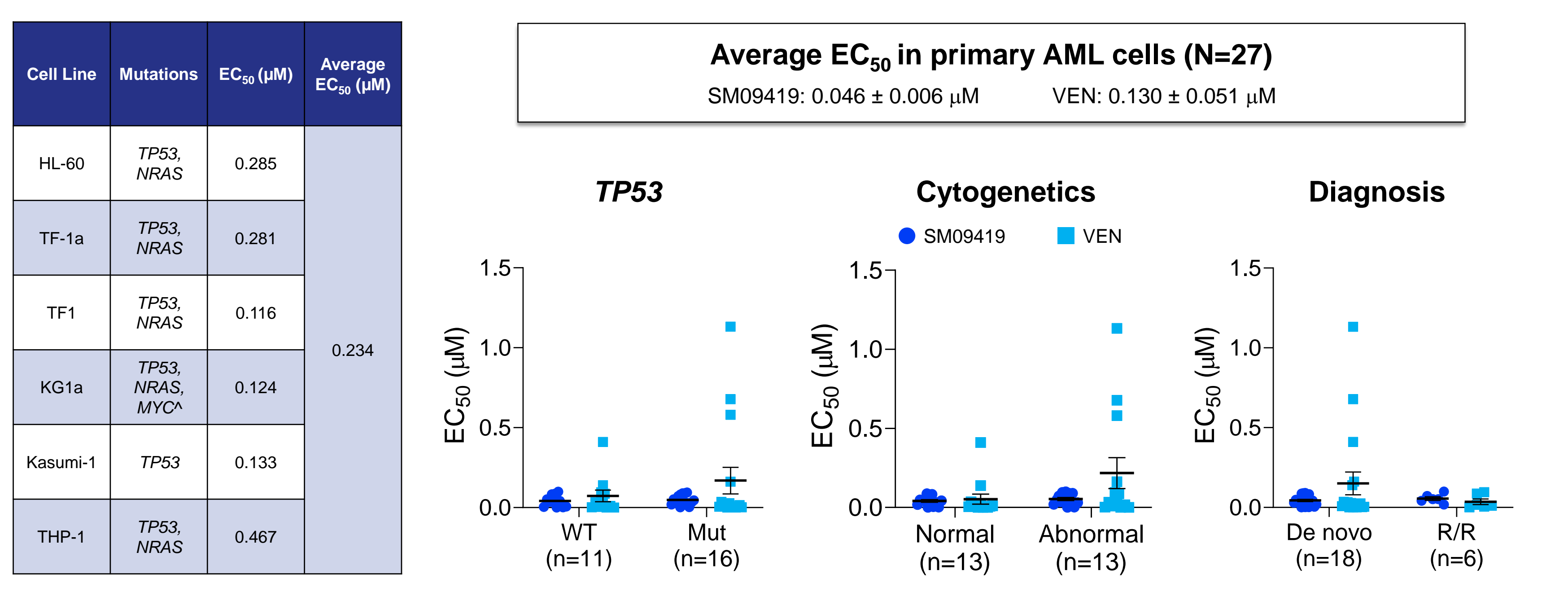


Figure 2. SM09419 induced apoptosis in TP53mut AML cells

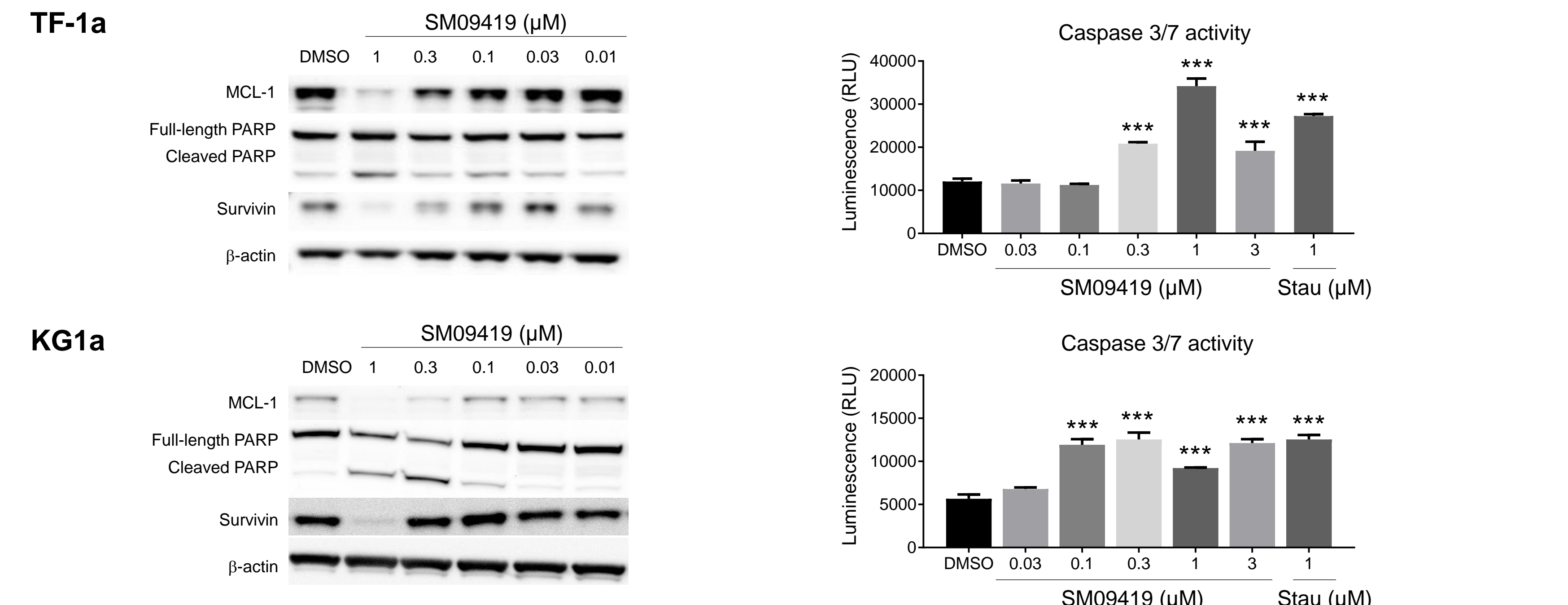


Figure 3. SM09419 dose-dependently inhibited SRSF6 phosphorylation and Wnt pathway-related gene and protein expression in TP53mut AML cells

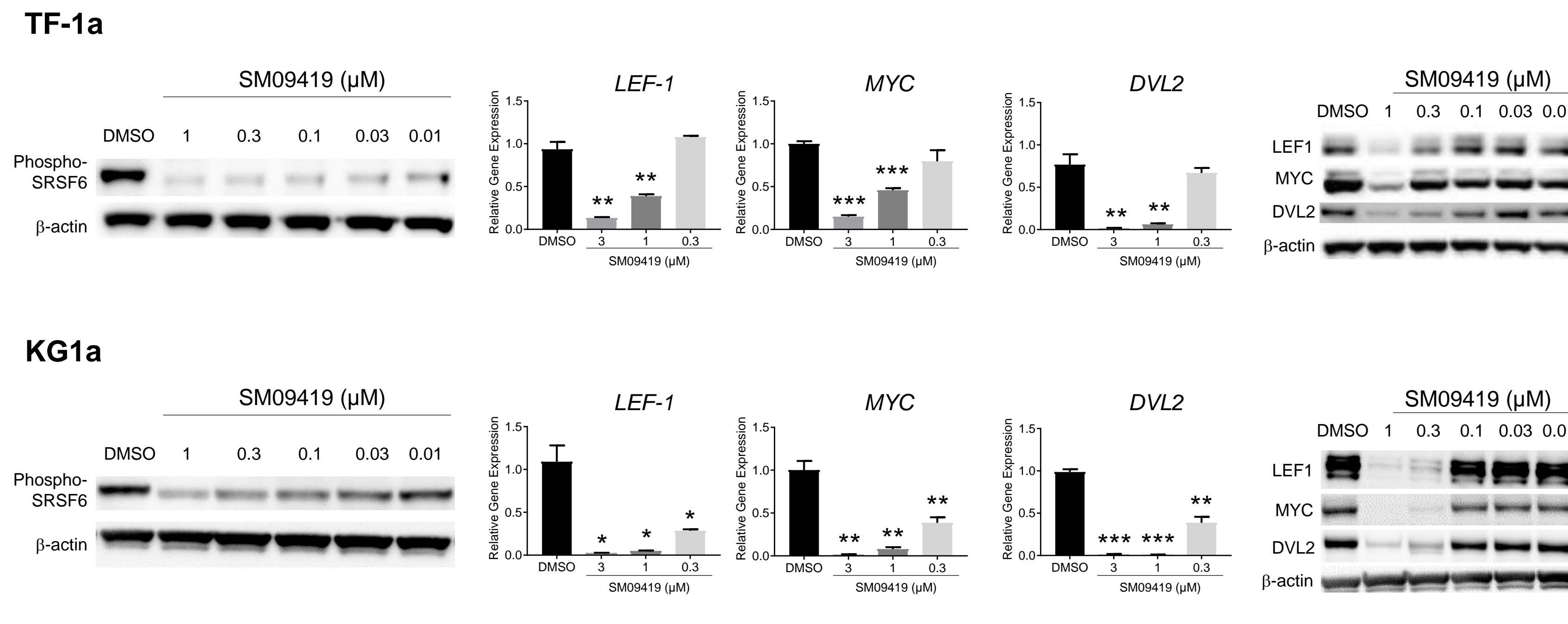


Figure 4. SM09419 demonstrated strong antileukemic effects as a single agent and in combination with Venetoclax, Azacitidine, or Cytarabine in AML xenografts

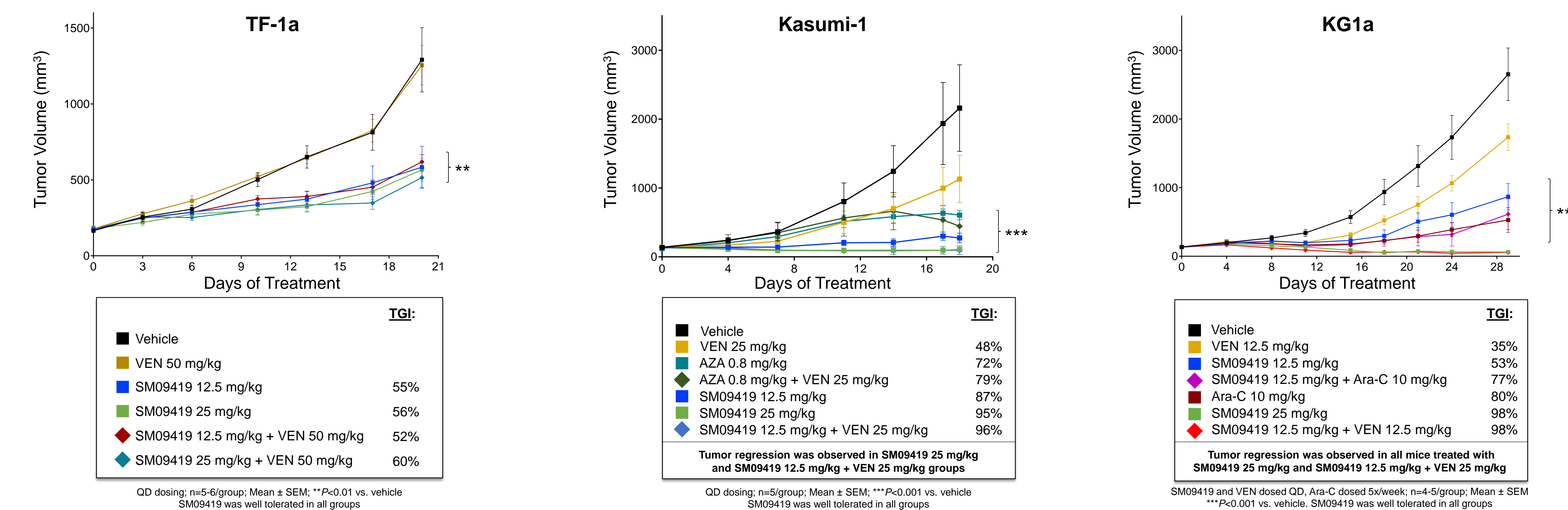
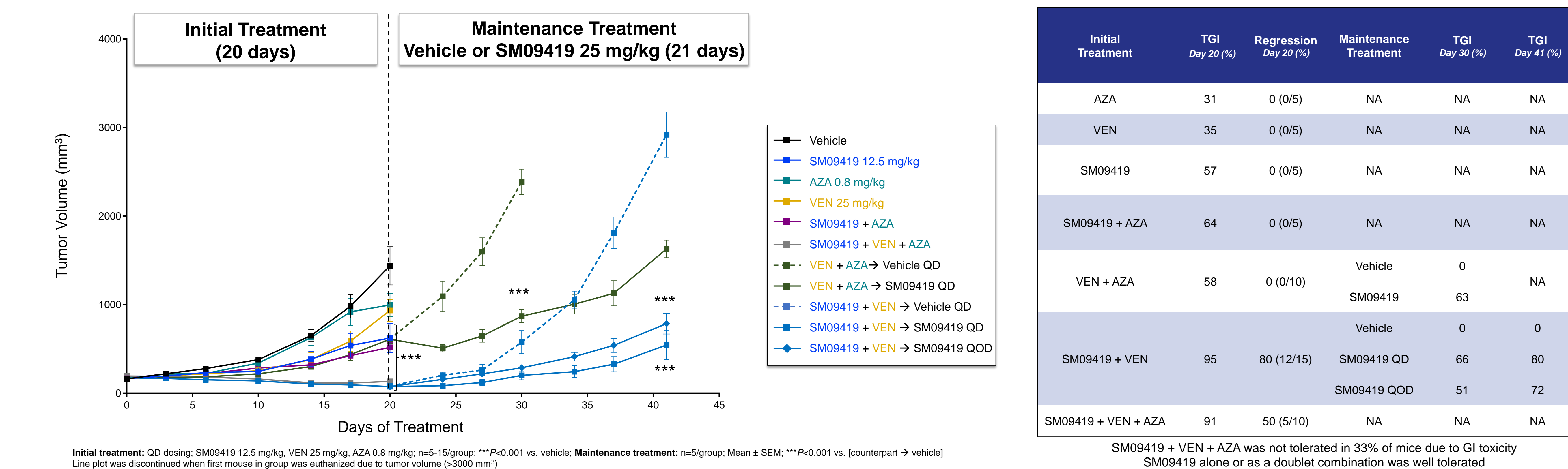


Figure 5. SM09419 + Venetoclax as initial treatment induced tumor regression and SM09419 as maintenance treatment slowed tumor regrowth in KG1a xenografts



Methods

In vitro assays:

- Cell proliferation in 6 TP53mut AML cell lines and 27 primary leukapheresis-derived human AML cell samples (Champions Oncology) treated with vehicle, SM09419, or Venetoclax was assessed using the CellTiter-Blue[®] assay and CellTiter-Glo[®] assay in 4 replicates, respectively. Differences in TP53 mutation status, cytogenetics, or AML diagnosis (de novo or relapsed/refractory [R/R]) were directly compared in primary human AML cells (Fig. 1)
- Apoptosis in AML cells treated with vehicle, SM09419, or staurosporine (Stau) for 48 hours was assessed by Western blot (PARP cleavage and expression of apoptosis regulators) and the Caspase-Glo[®] 3/7 assay kit (Fig. 2)
- Effects of SM09419 on SRSF phosphorylation and Wnt pathway-related protein expression in AML cell lines were measured by Western blot after 48 hours of treatment (Fig. 3)
- Gene expression in AML cell lines after 24 hours of exposure to vehicle or SM09419 was measured by qRT-PCR using TaqMan[®] primers and normalized to GAPDH expression (Fig. 3)

In vivo assays:

- Cell line-derived xenografts: SCID mice were implanted with TF-1a, Kasumi-1, or KG1a cells in the right flank and randomized into treatment groups when tumors reached ~100-200 mm³. Mice were orally treated with vehicle, SM09419, Venetoclax (VEN), Azacitidine (AZA), Cytarabine (Ara-C), or combinations of these drugs for indicated doses and times (Fig. 4 and Fig. 5)
- Tumor growth inhibition (TGI) was calculated relative to vehicle
- Tumor regressions were assessed according to the Response Evaluation Criteria in Solid Tumors (RECIST) guidelines: 30%-100% reduction in tumor volume relative to the start of the study
- Tolerability was determined by average bodyweight change from baseline (<15% loss considered well tolerated)

References

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