

New Selective Inhibitors of ERG Positive Prostate Cancer: ERGi-USU-6 Salt Derivatives

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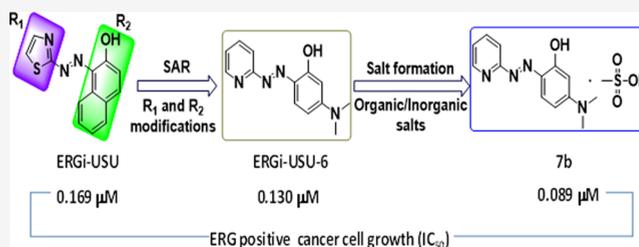
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ABSTRACT: Prostate cancer is among the leading causes of cancer related death of men in the United States. The *ERG* gene fusion leading to overexpression of near full-length *ERG* transcript and protein represents most prevalent (50–65%) prostate cancer driver gene alterations. The *ERG* oncoprotein overexpression persists in approximately 35% of metastatic castration resistant prostate cancers. Due to the emergence of eventual refractoriness to second- and third-generation androgen axis-based inhibitors, there remains a pressing need to develop drugs targeting other validated prostate cancer drivers such as *ERG*. Here we report the new and more potent *ERG* inhibitor ERGi-USU-6 developed by structure–activity studies from the parental ERGi-USU. We have developed an improved procedure for the synthesis of ERGi-USU-6 and identified a salt formulation that further improves its activity in biological assays for selective targeting of *ERG* harboring prostate cancer cells.

KEYWORDS: *ERG*, Oncogene, *TMPRSS2-ERG*, *RIOK2*, Inhibitor, Small molecule, Prostate cancer, Metastatic castration resistant prostate cancer, SAR, Precision medicine



The 2021 projection of prostate cancer (PCa) prevalence predicts alarming increases that have not been seen in the past two decades. PCa is one of the most prevalent noncutaneous malignancy and second leading cause of cancer deaths in men in the United States.¹ PCa is a global public health issue, exceeding one million people succumbing to the disease each year.² It is estimated that globally one in eight men will be diagnosed with PCa worldwide over his lifetime. Surgery, radiation therapy, and active surveillance are the few choices for the management of localized disease. The main line of treatment for advanced PCa is androgen deprivation therapy (ADT), which is widely used for treating PCa that has escaped the primary treatment regimens.³

It is remarkable that approximately half of primary PCas harbor gene fusions (prevalent fusion of the *TMPRSS2* gene promoter with the protein coding region of the *ETS* Related Gene [*ERG*]) which drive overexpression of the *ERG* fusion transcripts and *ERG* oncoprotein.^{4,5} In the gene fusion context *ERG* expression is driven by the androgen receptor (AR) due to the male hormone activated *TMPRSS2* promoter. The *ERG* was originally discovered as a retroviral oncogene.⁶ In castration resistant prostate cancer (CRPC) and metastatic CRPC the promiscuous activation of the AR can keep *ERG* levels high in a large number of cases. The global assessment of *ERG* in PCa even after accounting for racial/ethnic differences⁷ indicated that *ERG* is proposed as one of the most prevalent oncologic targets known in cancer. Oncogenic

activation of *ERG* is not restricted to PCa, aggressive forms of acute myeloid leukemia (AML) and Kaposi sarcoma-endotheliomas are also driven by this oncogene.⁸ In the context of malignancies, *ERG* activates cell invasion, abrogates normal differentiation, and facilitates a broad range of cancer cell survival mechanisms.

The realization of *ERG* fusion and overexpression in PCa has led to a significant effort in developing potential treatment by targeting the *ERG* oncoprotein.^{9–12} The *ETS* transcription factors including *ERG* are emerging drug targets in cancer.¹³ Initial small molecule screening for inhibition of *ERG* oncoprotein expression using mouse monoclonal anti-*ERG* antibody (9FY) developed in our lab^{14,15} resulted in the identification of ERGi-USU ((*E*)-1-[2-thiazolylo]-2-naphthol) that selectively inhibited the growth of *ERG* positive cancer cells without affecting normal cells *in vitro* and *in vivo*.¹² Following up on the intriguing hypothesis that an upstream regulator kinase is responsible for the cancer selective *ERG* inhibition, we subsequently identified an atypical kinase, *RIOK2*^{12–17} as the direct binding partner of ERGi-USU.

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This was accomplished by a kinome screen of a broad kinase panel in a substrate competition assay and by the tryptophan fluorescence quenching assay. The compound ERGi-USU (1-[2-thiazolylazo]-2-naphthol (NSC139021) has been tested for its chemopreventive properties in the context of cancer.^{18,19}

The objective of the current study was to develop new derivatives of ERGi-USU utilizing a structure–activity design approach, which show enhanced biological activity.^{20–24} Therefore, we designed new analogs around ERGi-USU and identified a more effective inhibitor, ERGi-USU-6. This led to development of salt formulations of the ERGi-USU-6. Approximately half of all therapeutic molecules are available in salt forms, as they provide benefits such as solubility, dissolution rate, permeability, and potency over the corresponding parent molecules.²⁵ Salt selection is a lucid and well-known process, where they are evaluated for their precise characteristics that consist of crystallinity, hygroscopicity, and polymorphism.^{26–28} First, we developed a new facile synthetic procedure that can be applicable for the synthesis of the ERGi-USU-6 in high yields. Subsequently, a set of pharmaceutically acceptable ERGi-USU-6 salts were prepared and evaluated for their efficacies for inhibition of growth of the ERG positive prostate cancer cell line, ERG protein, and RIOK2 protein read-out as an indicator associated with ERG inhibition. Herein, we report the results of our studies, describing the synthesis and pharmacological inhibition of ERG in prostate cancer experimental models.

We have previously identified (*E*)1-[2-thiazolylazo]-2-naphthol (ERGi-USU, Figure 1) as a selective inhibitor of

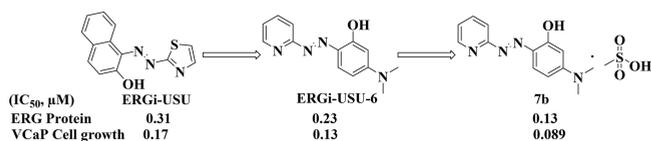


Figure 1. Identification of lead compound ERGi-USU-6 from ERGi-USU and its optimization to identify a salt derivative **7b**.

ERG positive cancer cells from a library of 2407 compounds from the National Cancer Institute diversity set.¹² The demonstration of the nontoxic effect of the parental compound ERGi-USU in primary normal endothelial HUVEC cells and in *in vivo* experiments led to further development of ERGi-USU derivative compounds. As described before, ERGi-USU binds to the RIOK2 atypical kinase and affects the ERG levels presumably through ERG upstream mechanisms selective for the context of cancer cells.¹²

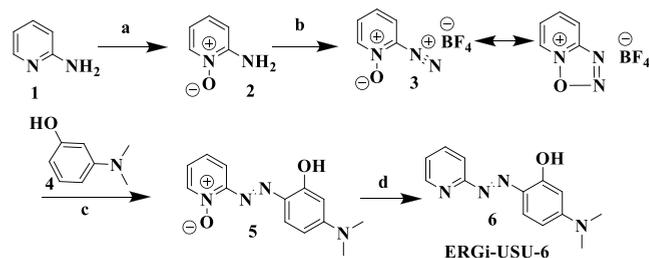
Since, RIOK2 protein levels are consistently decreased in response to ERGi-USU, we also measured the RIOK2 protein levels as indicators throughout our experiments in both ERG positive cancer cells and in ERG positive normal endothelial HUVEC cells. In order to improve the efficacy of ERGi-USU, structural activity relationship studies defined 134 compounds that were synthesized and tested for ERG protein inhibition levels and cell growth inhibition in the ERG positive prostate cancer cell line (VCaP), ERG negative prostate cancer cell line (LNCaP), and ERG positive normal primary endothelial cells (HUVEC). For high throughput screening, we utilized a highly specific anti-ERG mouse monoclonal antibody, 9FY, in an In-Cell Western assay developed by our group.¹²

Evaluation of 57 (see the supporting document for the structures) prioritized ERGi-USU analogs by In-Cell Western, Western blot, and cell growth assay resulted in the

identification of 6 compounds with similar or better activity in relation to the parental ERGi-USU. Among these, ERGi-USU-6 (Figures 1 and S1) was designated as the lead compound based on its favorable characteristics and biological activity. In the next step, we focused on preparing pharmaceutically acceptable ERGi-USU-6 salts to further improve efficacy. Though the ERGi-USU compounds show similarities to chemical structures of pan-assay interference compounds (PAINS)²⁹ complexing metal ions with pleiotropic activities, we observed that ERGi-USU-6 selectively inhibits the growth of ERG positive cancer cells with no detectable effect on ERG positive normal cells or ERG negative cancer cells, removing any concern of its pleiotropic biological activity. In the future, we also plan to investigate the possibility of the involvement of metal ion complexing mechanisms in the context of ERG positive cancer cells.

Acquiring ERGi-USU-6 in large quantities was found to be a major limitation for preparation of a variety of ERGi-USU-6 salts, as the reported procedures did not afford the desired compound in good yield and were inconsistent with previous literature reports.^{30–32} In addition, the method described by Grudpan and Taylor³³ required harsh reaction conditions and laborious workup procedure to obtain the desired products. Moreover, the yields obtained in this method were very low and may not be applicable to synthesize ERGi-USU-6 in large quantities. Therefore, we first modified the literature procedure and developed a facile route to synthesize ERGi-USU-6 in large quantities (Scheme 1), that was also used to synthesize similar analogs.³⁴

Scheme 1. Synthesis of ERGi-USU-6^a

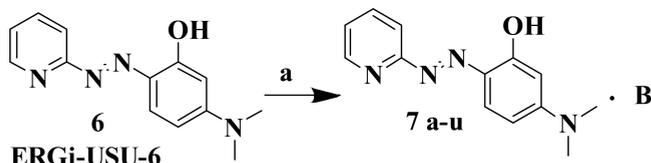


^aConditions (a) *m*-CPBA, acetone, room temperature; (b) NaNO₂, HBF₄, −10 to −5 °C; (c) 3-(dimethylamino)phenol, methanol, 10–15 °C; (d) MoO₂[(C₂H₅)₂NCS₂]₂, Ph₃P, acetone, room temperature.

The synthetic path utilizes 2-aminopyridine **1** as the starting material, which was then reacted with *m*-CPBA in acetone to afford 2-aminopyridine-1-oxide **2**. Treating **2** with sodium nitrite in 40% HBF₄ solution resulted the key intermediate, pyridin-2-diazonium 1-oxide tetrafluoroborate **3**, which usually exists in mesomeric form. Isolation of **3** is crucial in order to significantly improve the yields of the final compound. Since the compound **3** is hygroscopic, upon completion of the reaction, it was collected under a nitrogen atmosphere by filtration. Compound **5** was prepared by a diazotization reaction of **3** with 3-(dimethylamino)phenol **4**. Treating **5** with a freshly prepared Mo(VI) complex,³⁵ MoO₂[(C₂H₅)₂NCS₂]₂ and PPh₃ afforded the final compound. ¹H and ¹³C NMR and LC-MS data confirmed the identity of compound **6**. A highly deshielded ¹H NMR signal at δ_H 15.86 of **6** (Supporting Information) was observed, demonstrating that the H in phenolic OH may participate in an intramolecular hydrogen-bonding with the adjacent N in **6**.

A general scheme to synthesize salts of ERGi-USU-6 is outlined in Scheme 2. Reacting ERGi-USU-6 (free base) with

Scheme 2. Synthesis of ERGi-USU-6 Salts^a



^aConditions: (a) counter ion B, solvent, room temperature.

an appropriate solvent and the counterion precursor (Table 1) resulted in a variety of pharmaceutically acceptable organic and inorganic salts of the parent compound.

Table 1. Details of Counterions, Solvents, and Methods for the Preparation of ERGi-USU-6 Salts and Their Inhibition Concentrations

| Compound ID | B | Solvent | Synthetic Method | ERG Protein inhibition (IC ₅₀ , μM) | Cell inhibition (IC ₅₀ , μM) |
|----------------|--------------------------------|---------------|------------------|--|---|
| 6 (ERGi-USU-6) | -- | -- | -- | 0.23 | 0.13 |
| 7 a | HCl | 1,4-Dioxane | A | 0.22 | >0.5 |
| 7 b | | Ethanol | A | 0.13 | 0.089 |
| 7 c | | Isopropanol | A | 0.21 | 0.139 |
| 7 d | | Methanol | A | 0.24 | >0.5 |
| 7 e | | Acetonitrile | A | 0.26 | 0.136 |
| 7 f | HBr | Ethyl acetate | A | 0.24 | >0.5 |
| 7 g | H ₂ SO ₄ | Ethyl acetate | A | 0.26 | >0.5 |
| 7 h | Na ⁺ | Water | A | 0.11 | >0.5 |
| 7 i | HNO ₃ | Ethanol | A | 0.16 | >0.5 |
| 7 j | | Water | B | >0.5 | >0.5 |
| 7 k | | Acetone | B | 0.22 | >0.5 |
| 7 l | | Ethanol | A | 0.16 | >0.5 |
| 7 m | | Ethanol | A | 0.15 | >0.5 |
| 7 n | | Ethanol | B | >0.5 | >0.5 |
| 7 o | | Ethyl acetate | A | >0.5 | >0.5 |
| 7 p | | Ethanol | A | >0.5 | >0.5 |
| 7 q | | Ethanol | A | 0.24 | >0.5 |
| 7 r | | Ethanol | B | >0.5 | >0.5 |
| 7 s | | Ethanol | A | >0.5 | >0.5 |
| 7 t | | Ethanol | A | >0.5 | >0.5 |
| 7 u | | Acetonitrile | A | >0.5 | >0.5 |

In order to enhance the solubility while maintaining the selectivity, we synthesized formulations of 21 salt derivatives (Table 1). Among these, 7b was found to be the most effective salt derivative which exhibited increased solubility, inhibition of ERG and RIOK2 protein levels in ERG positive prostate cancer cells (VCaP) in three independent experiments (Figure 2).

We then assessed whether 7b has any adverse effect on ERG positive normal endothelial cells (HUVEC). To this end, HUVEC were treated with same dose concentration of 7b as were VCaP cells. It was observed that HUVEC were not affected by 7b salt derivatives at its highest doses (Figure 3).

We next examined the maximum tolerated dose (MTD) of ERGi-USU-6 *in vivo*. The MTD was estimated based on the

maximum at which all animals survived. Athymic nude mice were used to evaluate the MTD of ERGi-USU-6 in parallel comparison with ERGi-USU. Based on the dosages, the animals were divided into five groups. Group 1 designated as control/vehicle (90% PEG + 10% DMSO). Group 2 (50 mg/kg in control vehicle), group 3 (100 mg/kg in control vehicle), group 4 (150 mg/kg in control vehicle), and group 5 (200 mg/kg in control vehicle). Dosing and survival details are summarized in Figure 4. All mice receiving doses up to 100 mg/kg survived, but mortality occurred in 2 mice out of 5 at 150 mg/kg and 4 out of 5 at 200 mg/kg in the ERGi-USU-6 treatment group. To further define MTD, the overall toxicity as revealed by body weights was monitored throughout the study. The mice receiving the highest dose of 200 mg/kg showed significant weight loss (Figure 4C and D). Gross examination of major organs also revealed damage in tissues and vasculature as a result of the compound administration at the highest dose. In some events, localized inflammation at the site of the injection was observed at the highest dose raising solubility concerns. We therefore conclude that the MTD for ERGi-USU-6 is 100 mg/kg when compared with a 150 mg/kg dose of the parental compound ERGi-USU. During the observation period, no deterioration in health was observed in mice treated with the 100 mg/kg dose. These results point out that ERGi-USU-6 at a dose range of 100 mg/kg is safe and support the use of ERGi-USU-6 for further tumor xenograft experiments.

The TMPRSS2-ERG fusion driven activation of the ERG oncogene is one of the main cancer driver gene alterations that occurs in early stages of PCA and is retained in a subset of metastatic CRPCs. Early studies including reports from our laboratory have demonstrated that knockdown of ERG by siRNAs resulted in growth inhibition of TMPRSS2-ERG positive CaP cells in cell culture and xenograft models.¹² Along these lines emerging data from *in vitro* and *in vivo* models continue to underscore biological roles of ERG in PCA initiation and progression.¹³

These studies and the unprecedented prevalence and emerging PCA mortality reports provide a strong rationale for the development of ERG targeted therapies. Our goal has been to develop ERG inhibitors with tolerable systemic toxic effects. During this endeavor we consistently monitor the selectivity of ERG inhibitors for cancer cells harboring ERG overexpression and likely driven by oncogenic addiction and lack of inhibitory effect ERG inhibitors on normal endothelial cells, where endogenous regulated ERG expression plays essential function in cell renewal.³⁶ In other contexts, ERG is known to be expressed during development, restricted to precartilaginous and hematopoietic tissues including megakaryocytes.^{37,38}

Our previous and current *in vitro* and *in vivo* work and SAR studies on ERGi-USU led to the discovery of improved derivatives. Among these ERGi-USU-6 was found to be more effective and chosen for further studies. Drug molecules administered through salt form offer many potential benefits, as it can improve the solubility, dissolution rate, permeability, and efficacy of the drug. Salt selection is a rational and stepwise process in which salts are analyzed with regards to particular properties that include crystalline, hygroscopicity, and polymorphism assessment. Our primary objective was to identify an ERG inhibitor which can encounter pharmacological, technological conditions, possibilities and restrictions during development, and manufacturing and storage of the dosage forms. For this goal, selecting an optimal salt form of ERG

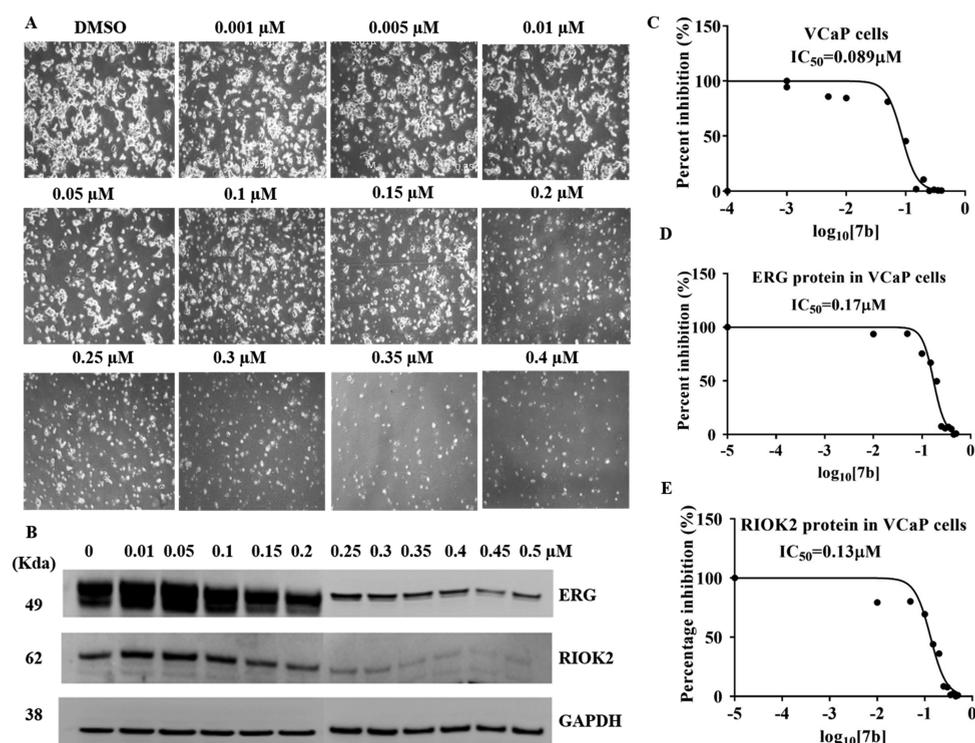


Figure 2. ERGi-USU-6 salt derivative **7b** showing improved activity. (A) **7b** inhibits the growth of ERG positive VCaP cells. (B) **7b** is a potent inhibitor of ERG and RIOK2 proteins. (C–E) **7b** showed improved cell growth, ERG protein, and RIOK2 protein inhibition with IC_{50} values of 0.089, 0.17, and 0.13 μM , respectively.

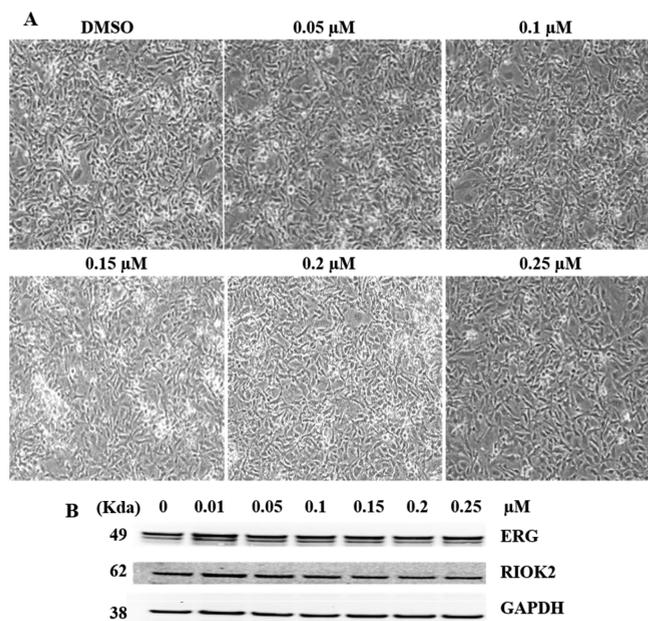


Figure 3. Cell growth, ERG protein, and RIOK2 protein levels are not affected in ERG positive normal endothelial HUVEC cells in response to ERGi-USU-6 salt formula **7b** treatment. (A) Cell growth inhibition response of HUVEC was tested in the 0–0.25 $\mu\text{mol/L}$ concentration range selected within the IC_{50} (0.089 μM) of VCaP prostate cancer cells. (B) No significant decrease of endogenous ERG protein and RIOK2 protein inhibition was observed in normal endothelial HUVEC cells.

USU-6 with an appropriate dosage form was crucial. From these studies we identified the most suitable new salt formula **7b** through SAR studies.

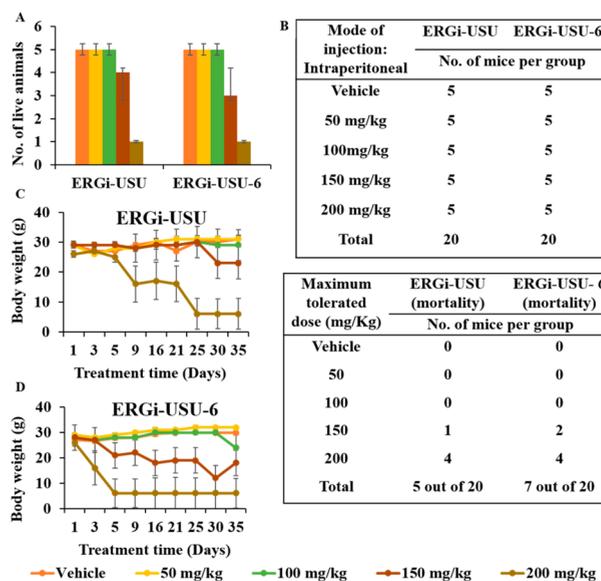


Figure 4. Maximum tolerated dose of ERGi-USU-6 is 100 mg/kg in athymic nude mice. (A, B) Assessment of maximum tolerated dose (MTD) of ERGi-USU and ERGi-USU-6. The table and bar diagram summarize the survival and mortality in each treatment group. (C and D) Body weight at different time points of the experiment. At the highest doses of ERGi-USU-6, body weight losses were observed when compared to the parental in ERGi-USU.

While there is no evidence for the direct binding of ERGi-USUs to ERG protein itself, there has been recurring themes on similar inhibitory profiles for various ERGi-USU derivatives for ERG and RIOK2 proteins in a given ERG positive cancer cell line. Although the mechanistic connection between RIOK2

and ERG is currently not known, based on our observations on direct binding of ERGi-USU to RIOK2, it is conceivable that a high-resolution crystal structure of ERGi-USU and RIOK2 will advance our knowledge by defining a new direct inhibitor of this atypical kinase. This is significant as pioneering research endeavors already point to the role of RIOK2 in key cancer driver pathways.^{16,17,39,40}

Previously, we reported the discovery of ERGi-USU as a selective inhibitor of ERG positive cancer cells.¹² Subsequent, development of its analogs and medicinal chemistry campaign has led to the development of ERGi-USU-6 through a new and more efficient chemical synthesis method for the large-scale production. Analysis of new ERGi-USU analogs and various salts resulted in a compound that inhibits the growth of ERG positive PCa cells at 0.089 μM IC₅₀. To the best of our knowledge, this is the best small molecule inhibitor of ERG protein and merits further development for therapeutic applications.¹³ Encouraged from this data, our future studies will include the pharmacokinetics of this lead compound determination of favorable administration route, evaluations of its effect in immune-competent mice

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsmmedchemlett.1c00308>.

Cell lines, reagents, general protocol for the protein inhibition studies, cell growth inhibition studies, maximum tolerated dose assay, statistical analysis, and synthesis of ERG-USU-6 and its derivatives (PDF)

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Notes

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The authors declare no competing financial interest.

■ ABBREVIATIONS

PCa, prostate cancer; CRPC, castration resistant prostate cancer; ERG, ETS related gene; ADT, androgen deprivation therapy; TMPRSS2, transmembrane protease, serine 2; AR, androgen receptor; AML, acute myeloid leukemia; RIOK2, RIO kinase 2; VCaP, vertebral cancer of the prostate; HUVEC, human umbilical vein endothelial cells; MTD, maximum tolerated dose; DMSO, dimethyl sulfoxide; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

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