

Preliminary safety and efficacy of ELVN-001, a selective active site inhibitor of BCR::ABL1, in patients with CML driven by atypical fusion transcripts

Andreas Hochhaus¹, Susanne Saussele², Valerie Coiteux³, Dong-Wook Kim⁴, Yingsi Yang⁵, Qi Wang⁵, Brianna Hoffner⁵, M. Damiette Smit⁵, Fabian Lang⁶

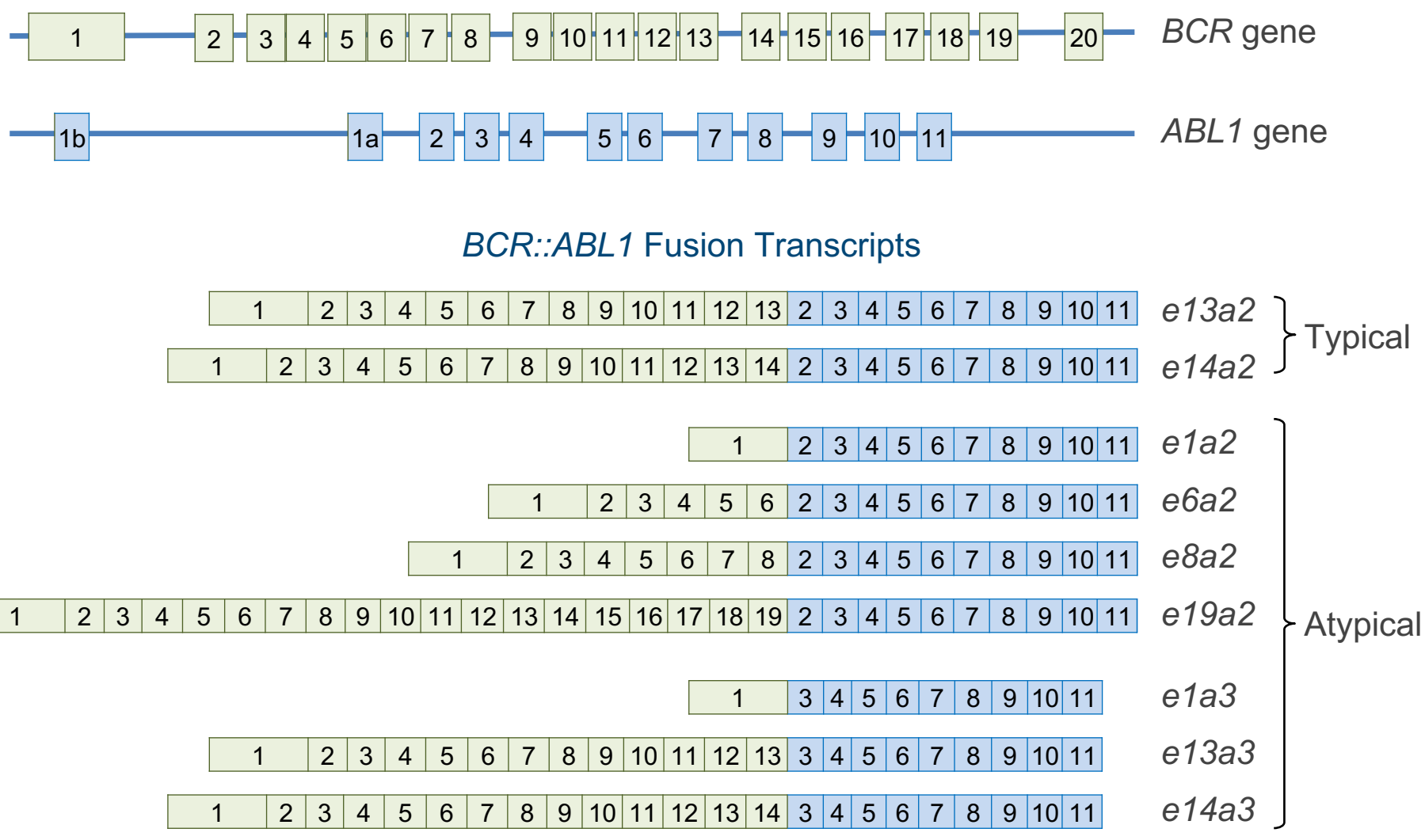
¹Universitätsklinikum Jena, Jena, Germany; ²Medizinische Fakultät Mannheim der Universität Heidelberg, Heidelberg, Germany; ³Service des Maladies du Sang Hôpital Huriez - CHRU de Lille, Lille, France; ⁴UiJeongbu Eulji Medical Center, Geumo-dong, Uijeongbu-si, South Korea; ⁵Enliven Therapeutics, Boulder, CO, USA; ⁶Department of Hematology and Oncology, Goethe University Hospital Frankfurt, Frankfurt, Germany



INTRODUCTION

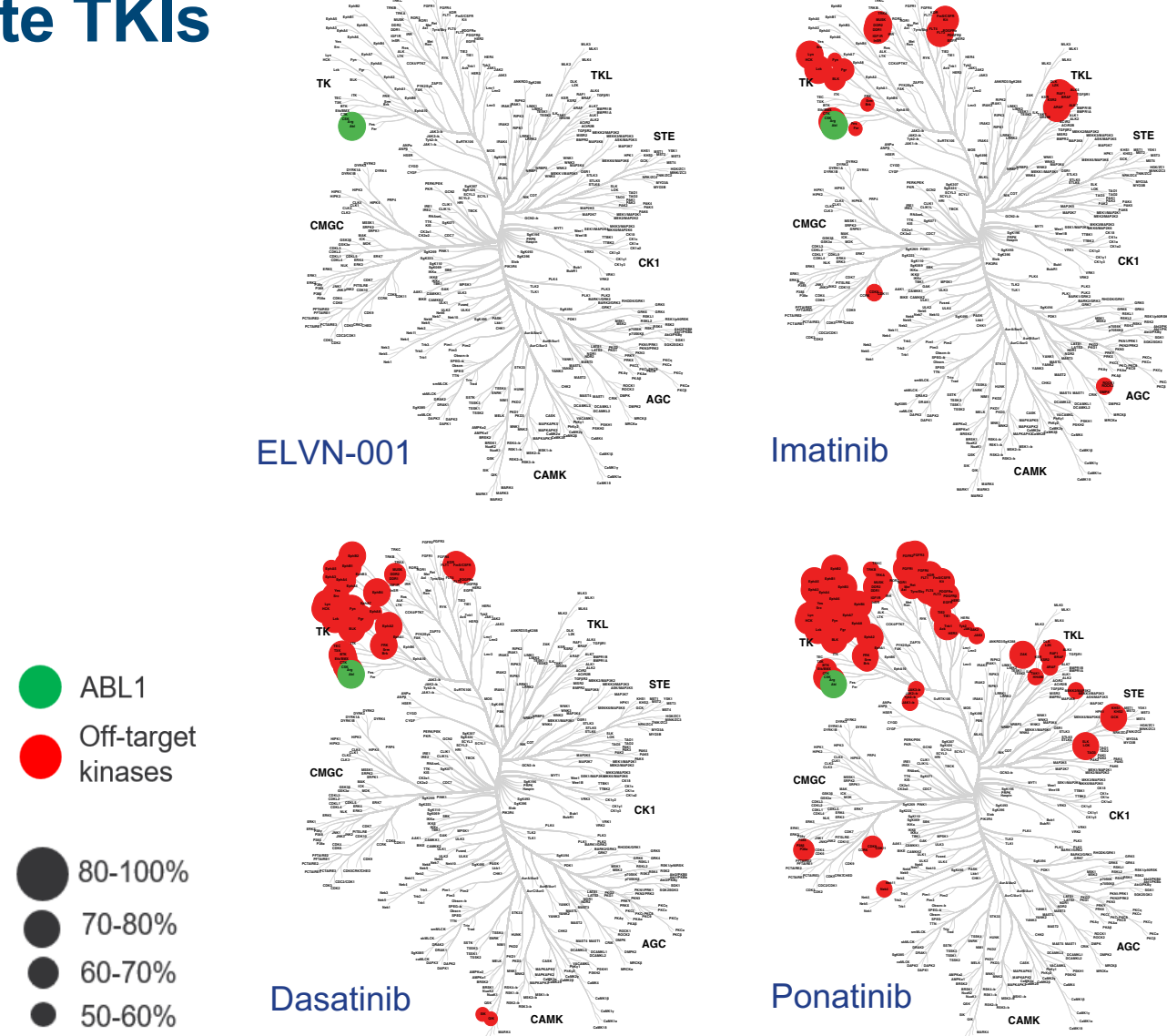
- CML is characterized by a reciprocal translocation of t(9;22)(q34;q11), leading to the *BCR::ABL1* fusion gene¹
 - Most patients have a typical fusion transcript: e13a2 and/or e14a2. However, approximately 2% have an atypical fusion transcript for which there is limited data available about prognosis and response to tyrosine kinase inhibitors (TKIs)
 - Emerging data show that atypical *BCR::ABL1* fusions lacking *ABL1* exon a2 (e.g., e13a3, e14a3, and e1a3) cannot be inhibited by allosteric TKIs targeting the myristoyl pocket (e.g., asciminib^{2,3})
 - In contrast, these translocations should be sensitive to active site TKIs
- ELVN-001: Highly Selective ATP-competitive Inhibitor of BCR::ABL1**
- ELVN-001 binds to a unique P-loop “folded-in” active conformation in the ATP-binding pocket, which:
 - Provides greater selectivity than available ATP inhibitors⁴, with potential for better tolerability
 - Creates a narrow tunnel allowing binding to T315I and other mutations
 - The ELVN-001 pharmacokinetic profile supports once daily dosing with or without food, which, in addition to low potential for DDIs, addresses key challenges with currently available TKIs
 - In patients with typical transcripts, ELVN-001 demonstrated favorable safety and tolerability across a wide range of doses, and encouraging preliminary efficacy was observed

Figure 1: *BCR::ABL1* Genes



NOTE: Figure is for illustrative purposes and not drawn to scale. This is not a full list of all possible atypical transcripts/variants. There are other rare transcripts/variants which are not listed. This figure is adapted from Soverini et al, licensed under CC BY 4.0 (<http://creativecommons.org/licenses/by/4.0/>).⁵

Figure 2: Kinome Selectivity of ELVN-001 vs Other Active-Site TKIs



Each of these active site TKIs was profiled at 30x their respective ABL1 IC₅₀ values against a panel of 377 protein kinases in biochemical assays with 100 mM ATP concentration in the screen (Reaction Biology).

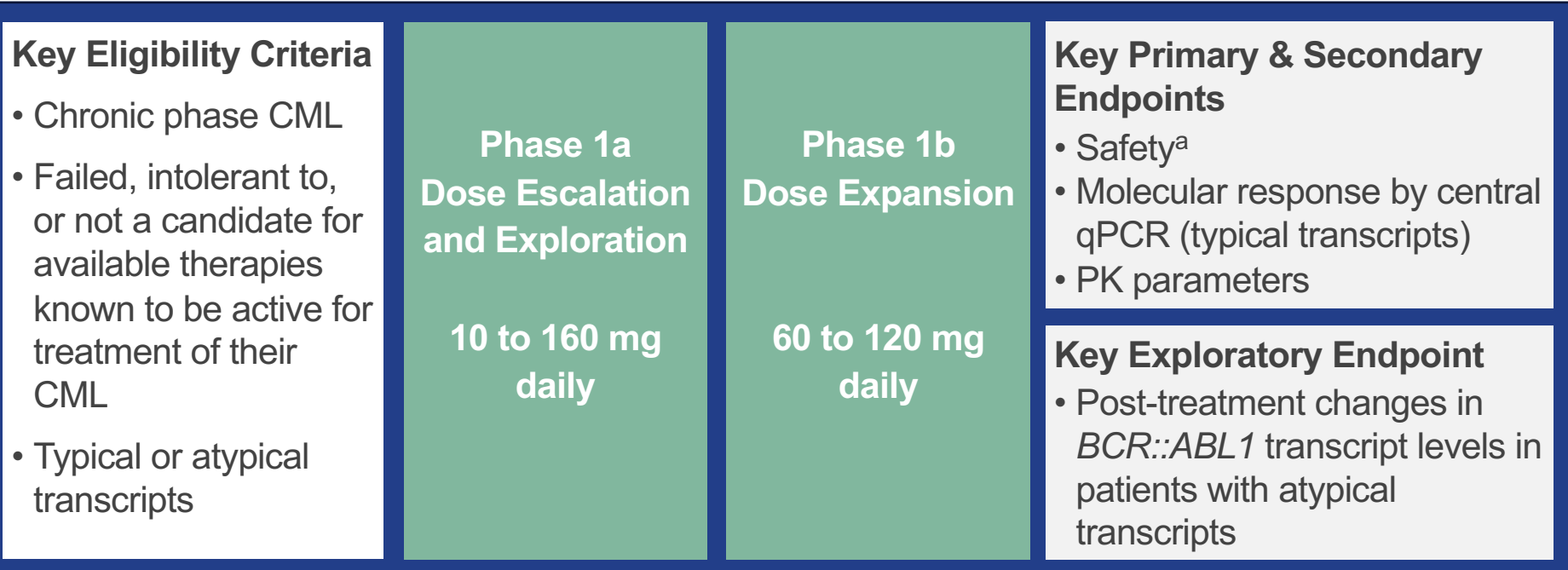
OBJECTIVE

- Here, we first report tolerability and anti-CML activity of ELVN-001 in patients with atypical transcripts enrolled in ENABLE (NCT05304377), a phase 1 study of ELVN-001 in patients with previously treated CML

METHODS

- Patients with atypical transcript received ELVN-001 orally at doses from 20 mg to 160 mg daily in the dose escalation phase of the ENABLE study; intrasubject dose escalation was allowed if specific criteria were met
- Testing for molecular response is non-standardized for atypical transcripts and was therefore assessed locally by individual molecular response⁶; testing for *BCR::ABL1* mutations was done centrally at baseline by NGS

Figure 3: ELVN-001: Phase 1 Trial Design



^aIncidence of DLTs, AEs, clinically significant laboratory and ECG abnormalities.

Abbreviations: AE, adverse event; ATP, adenosine triphosphate; BCR::ABL1, breakpoint cluster region-Abelson leukemia virus 1; BID, twice daily; CML, chronic myelogenous leukemia; DDI, drug-drug interaction; DLI, donor lymphocyte infusion; DLT, dose-limiting toxicity; ECG, electrocardiogram; GFR, glomerular filtration rate; GGT, gamma-glutamyl transferase; IC₅₀, half-maximal inhibitory concentration; MDS, myelodysplastic syndrome; NGS, next generation sequencing; NR, not related; PK, pharmacokinetics; QD, once daily; qPCR, quantitative reverse transcriptase polymerase chain; R, related; SCT, stem cell transplant; TEAE, treatment-emergent adverse event; TKI, tyrosine kinase inhibitor.

RESULTS

Anti-CML Activity in Patient with e13a3 Transcript

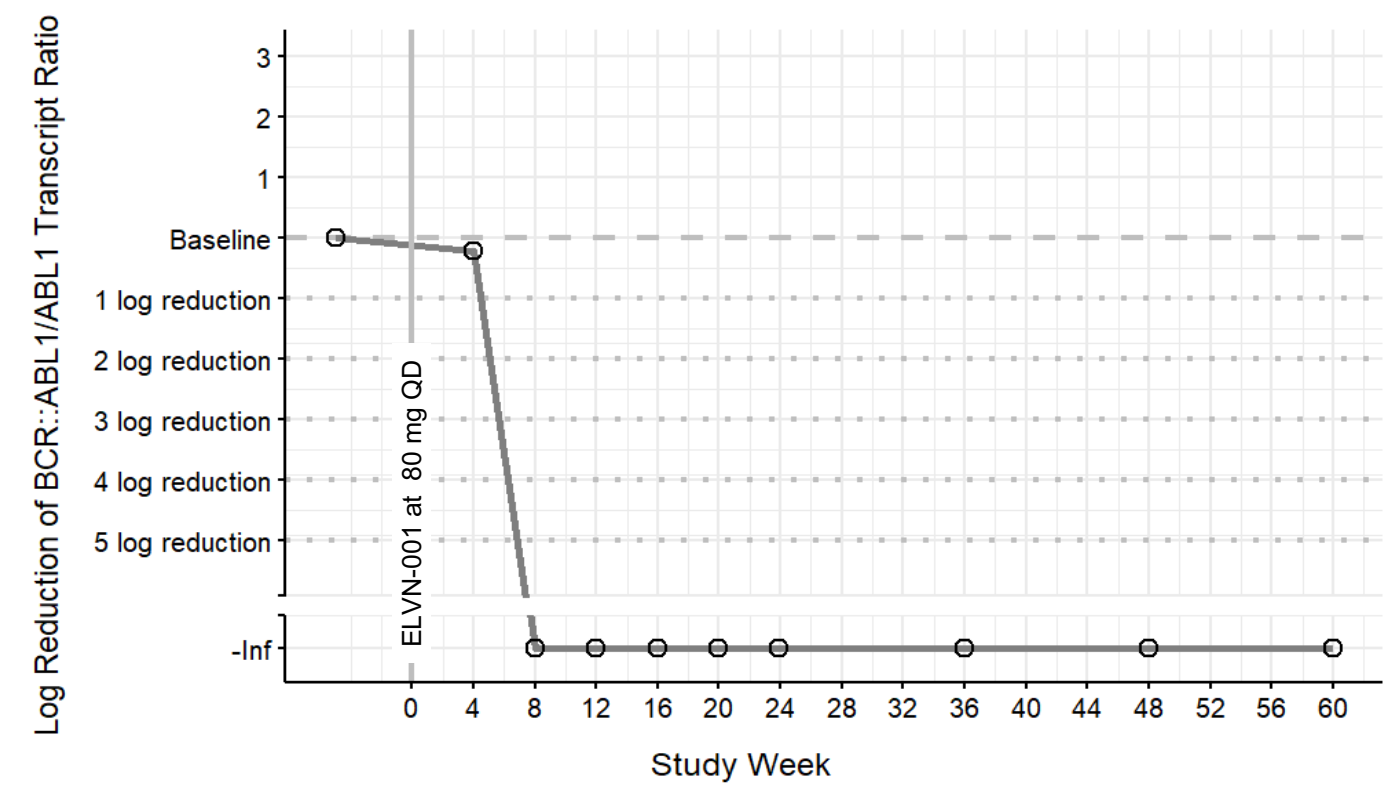
Patient Information

- 65-year-old female with primary diagnosis of CML in 2001
- Medical history: right bundle branch block, vulvar carcinoma, pulmonary emphysema, osteoporosis, MDS
- BCR::ABL1 mutations at baseline: none

TEAEs on ELVN-001

- Grade 1 increased cholesterol (NR)

>1 Log Reduction of *BCR::ABL1/ABL1* Transcript Ratio



NOTE: A transcript level of zero indicates the result is below the limit of quantification.

Anti-CML Activity in Patient with e19a2 Transcript

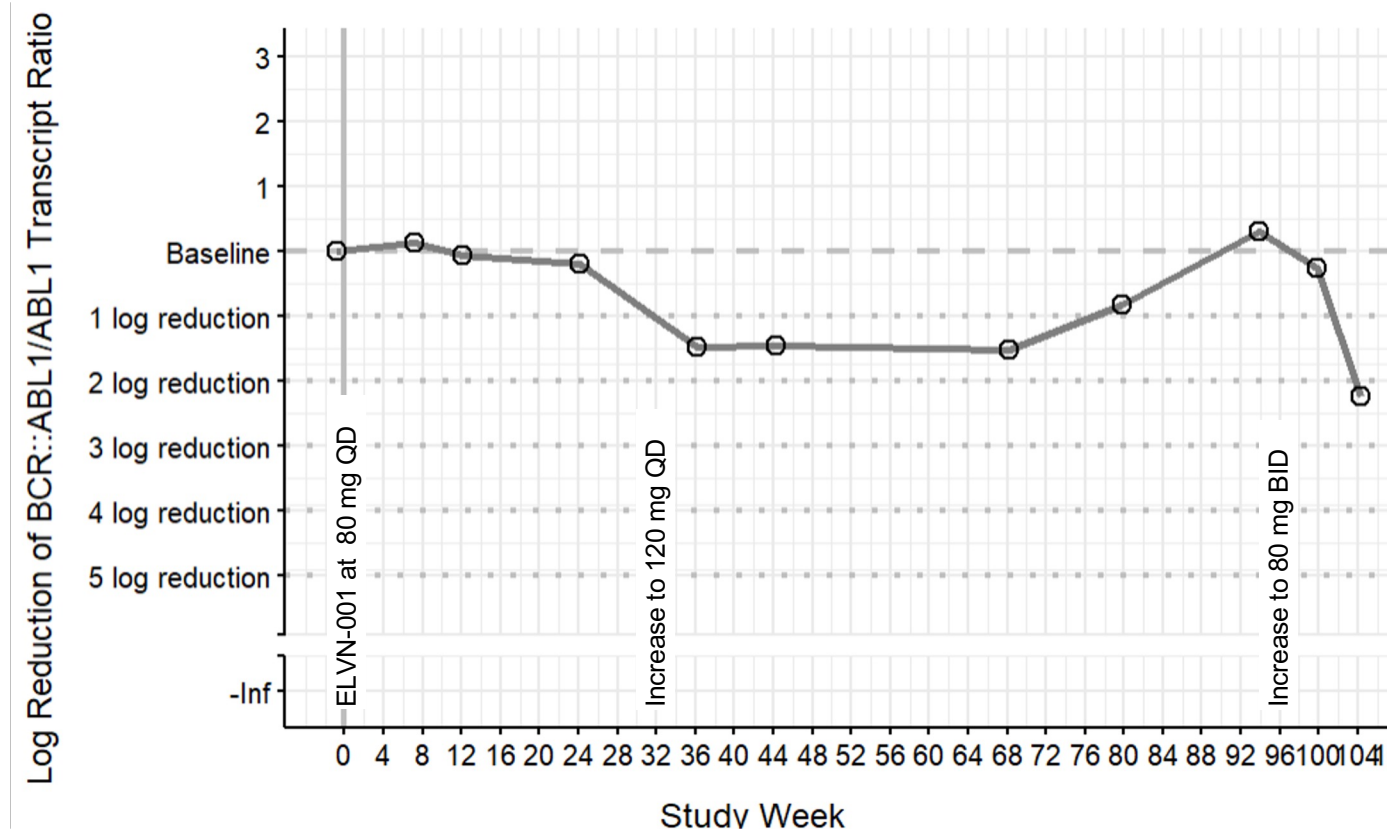
Patient Information

- 40-year-old male with primary diagnosis of CML in 2012
- Medical history: geographic tongue
- BCR::ABL1 mutation at baseline: T315I (100%)

TEAEs on ELVN-001

- Grade 1 dry skin (R), Grade 2 speech disorder, (NR) Grade 1 vomiting (NR), Grade 1 diarrhea (NR), Grade 2 back pain (NR)

BCR::ABL1/ABL1 Transcript Ratio in Patient with T315I Mutation



Anti-CML Activity in Patient with e1a2 Transcript

Patient Information

- 58-year-old female with primary diagnosis of CML in 2023
- Medical history: vitreous body detachment, hypertension, hypothyroidism, Hashimoto-thyroiditis, glaucoma, hypercholesterolemia
- BCR::ABL1 mutation at baseline: none

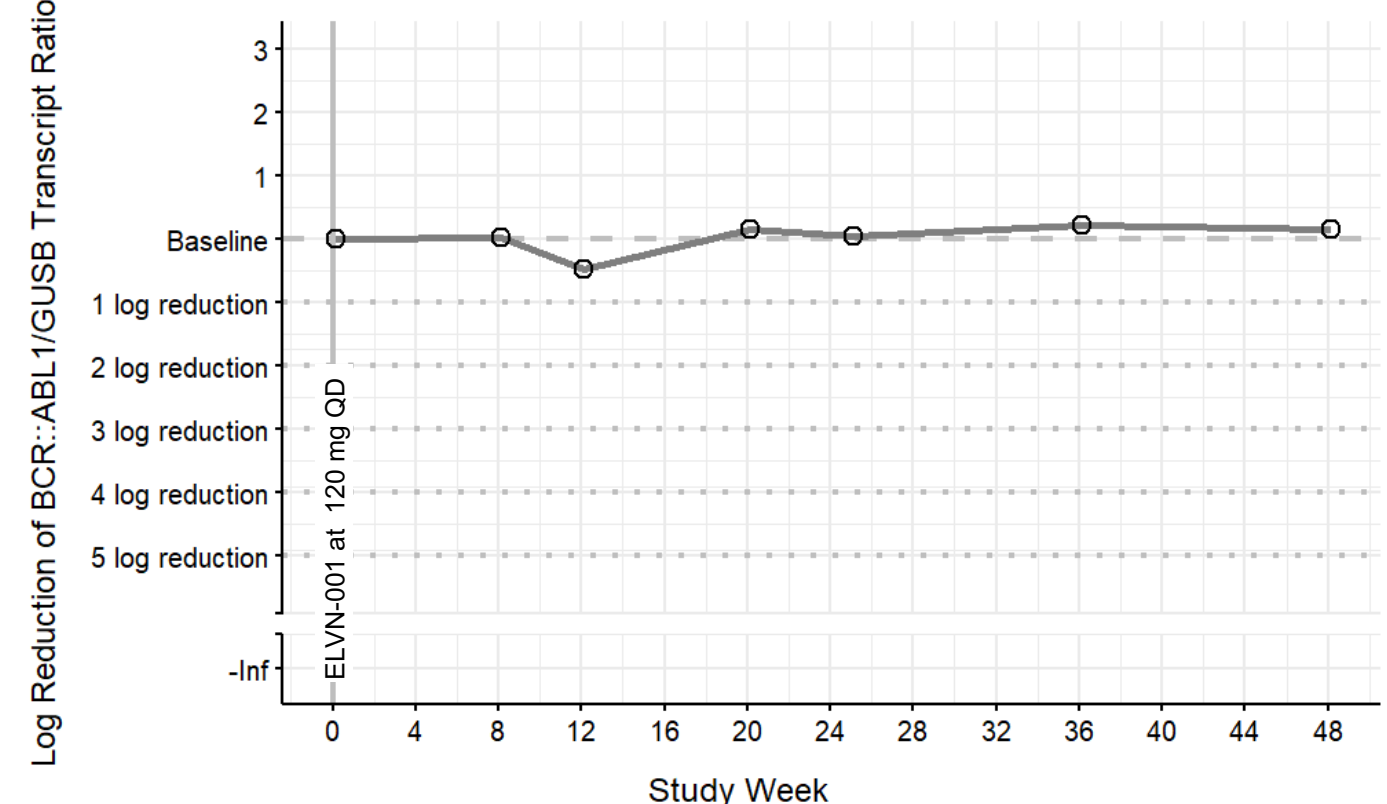
Prior Line of Therapy

Prior Line of Therapy	Best Transcript Category	Reason for Discontinuation
Bosutinib	>10%	Lack of efficacy
Dasatinib	>10%	Lack of efficacy

TEAEs on ELVN-001

- Grade 1 COVID-19 (NR), Grade 1 pruritus (R), Grade 1 GFR decreased (R), Grade 1 dry skin (R)

Stable *BCR::ABL1/GUSB* Transcript Ratio



Anti-CML Activity in Patient with e13a3 Transcript

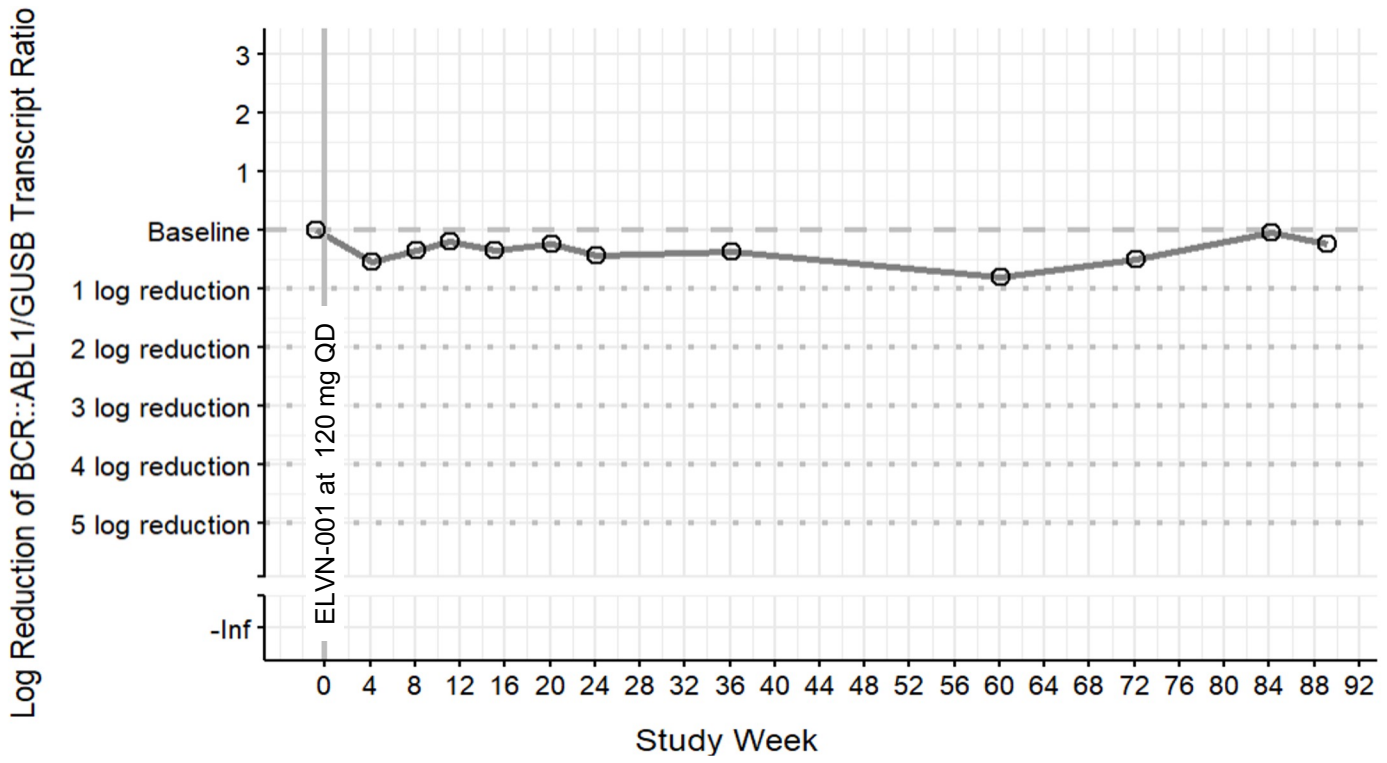
Patient Information

- 61-year-old female with primary diagnosis of CML in 2013
- Medical history: heart failure, stenosis of aortic valve, aortic insufficiency, hypertension, coronary 3-vessel disease, GGT increase, peripheral artery disease (Grade 1)
- BCR::ABL1 mutation at baseline: T315I (100%), S348L (34%)

TEAEs on ELVN-001

- Grade 1 muscle spasms (R), Grade 2 peripheral arterial occlusive disease (NR), Grade 2 lipase increase (NR), Grade 2 chronic kidney disease (NR), Grade 1 diarrhea (NR)

Stable *BCR::ABL1/GUSB* Transcript Ratio in Patient with T315I Mutation



Anti-CML Activity in Patient with e1a3 Transcript

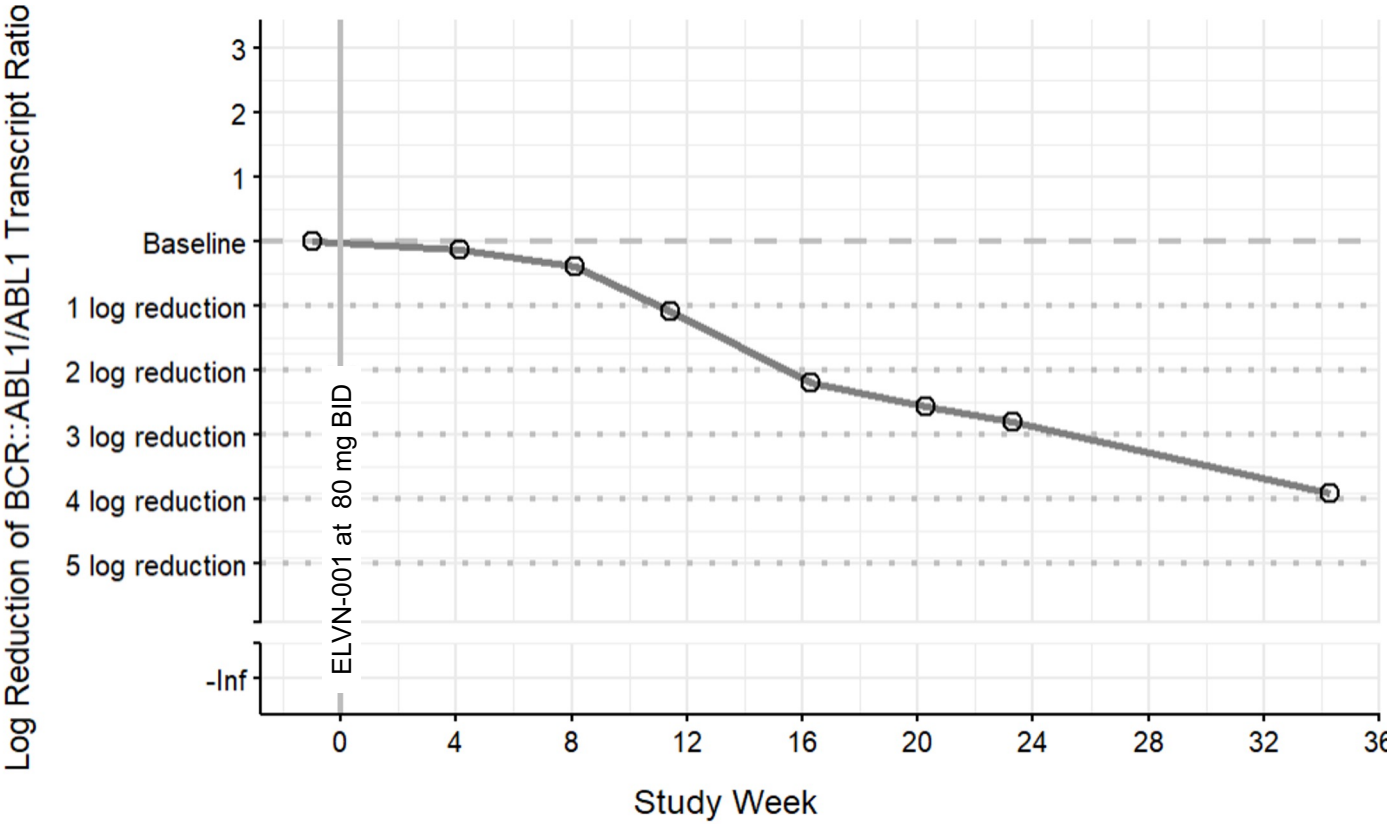
Patient Information

- 63-year-old female with primary diagnosis of CML in 2020
- Medical history: history of tobacco use
- BCR::ABL1 mutation at baseline: none

TEAEs on ELVN-001

- Grade 2 bronchiolitis (NR), Grade 1 thrombocytopenia (R), Grade 2 vomiting (NR), Grade 2 diarrhea (NR), Grade 1 asthenia (R), Grade 2 viral bronchitis (NR), Grade 2 pulmonary mass (NR)

>1 Log Reduction in *BCR::ABL1/ABL1* Transcript Ratio



Anti-CML Activity in Patient with e1a2 Transcript

Patient Information

- 60-year-old female with primary diagnosis of CML in 2021
- Medical history: laparoscopic partial hysterectomy, spinal stenosis surgery, articular inflammation both knees, onychomycosis, tinea pedis, thrombocytopenia, neutropenia, lipase elevation
- BCR::ABL1 mutation at baseline: none

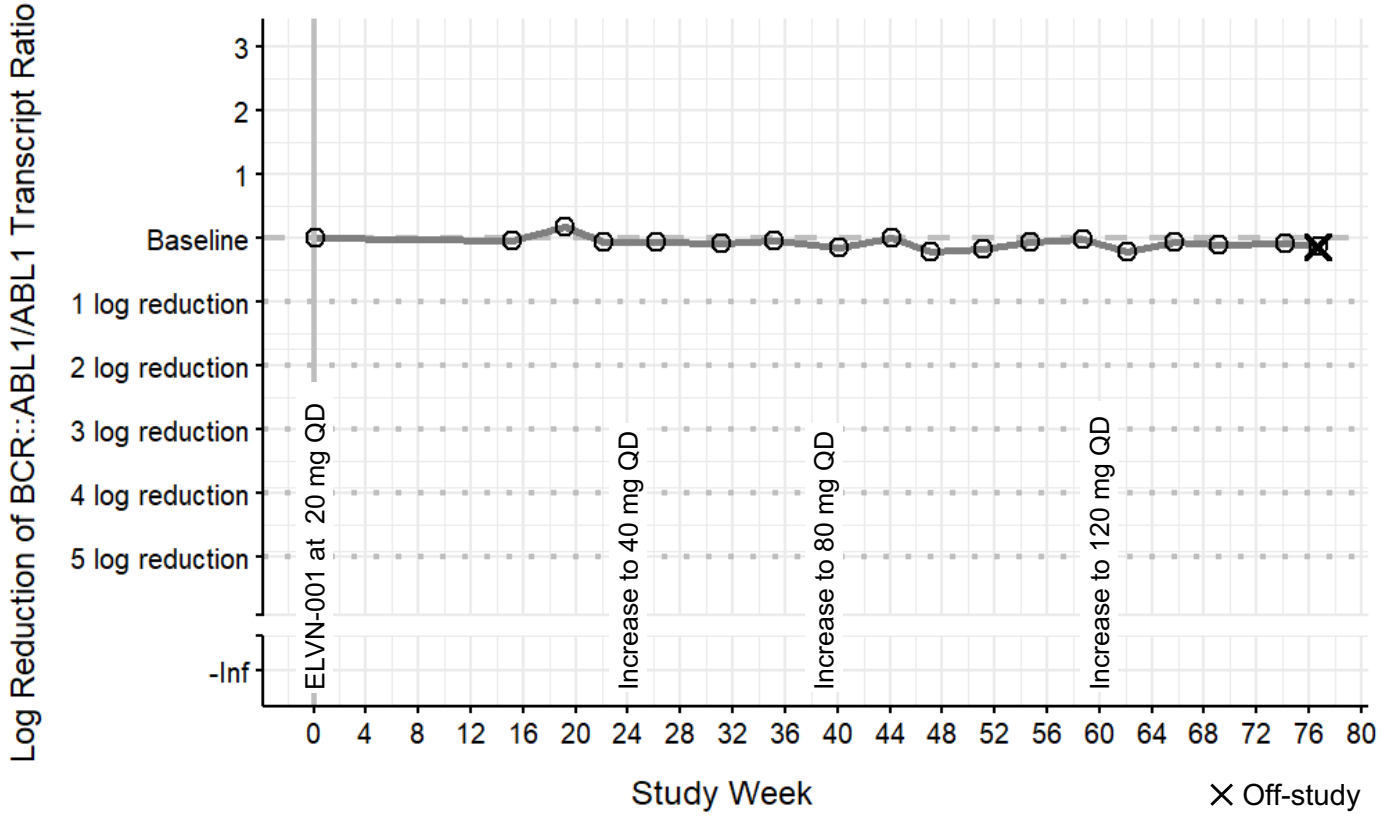
Prior Line of Therapy

Prior Line of Therapy	Best Transcript Category	Reason for Discontinuation
Imatinib	>10%	Lack of efficacy
Radotinib	>10%	Lack of efficacy
Dasatinib	>10%	Lack of efficacy

TEAEs on ELVN-001

- Grade 2 rash (R), Grade 1 pyrexia (NR)

Stable *BCR::ABL1/ABL1* Transcript Ratio



CONCLUSIONS

- The safety of ELVN-001, a novel selective active-site inhibitor of BCR::ABL1, in patients with atypical transcripts was favorable and consistent with the safety profile of ELVN-001 previously reported in patients with typical transcripts⁷
- ELVN-001 demonstrated encouraging anti-CML activity in heavily pre-treated patients with atypical *BCR::ABL1* transcripts; this includes patients with atypical fusions lacking *ABL1* exon a2 (e.g., e1a3, e13a3), which are resistant to TKIs targeting the myristoyl pocket
- Although this is a limited sample size from the Phase 1a of the ENABLE study, these data support further evaluation of ELVN-001 in patients with CML with atypical transcripts
- The phase 1 ENABLE study is active and recruiting (NCT05304377)

Acknowledgments: Thank you to all investigators and site staff, to the patients and their families for their participation in the study, to the Enliven medical team and to Ingrid Koo, PhD, who provided editorial support. This study was funded by Enliven Therapeutics, Inc.

References:

- Zhou X, et al. Oncol Rep. 2025.
- Leske JB, et al. Leukemia. 2024.
- Leyte-Vidal A, et al. Leukemia. 2024.
- Metz KS, et al. Cell Systems. 2018.
- Soverini et al. Mol Cancer. 2018.
- Schäfer et al. J Cancer Res Clin Oncol. 2021.
- Hochhaus A, et al. EHA 2025.