In vivo stabilization of a less toxic asparaginase variant leads to a durable antitumor response in acute leukemia

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Abstract

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Asparagine is a non-essential amino acid since it can either be taken up via the diet or synthesized by asparagine synthetase. Acute lymphoblastic leukemia (ALL) cells do not express asparagine synthetase or express it only minimally, which makes them completely dependent on extracellular asparagine for their growth and survival. This dependency makes ALL cells vulnerable to treatment with L-asparaginase, an enzyme that hydrolyzes asparagine. To date, all clinically approved L-asparaginases have significant L-glutaminase co-activity, associated with non-immune related toxic side effects observed during therapy. Therefore, reduction of L-glutaminase co-activity with concomitant maintenance of its anticancer L-asparaginase effect may effectively improve the tolerability of this unique drug. Previously, we designed a new alternative variant of Erwinia chrysanthemi (ErA; Erwinaze) with decreased L-glutaminase co-activity, while maintaining its L-asparaginase activity, by the introduction of three key mutations around the active site (ErA-TM). However, Erwinaze and our ErA-TM variant have very short half-lives in vivo. Here, we show that the fusion of ErA-TM with an albumin binding domain (ABD)-tag significantly increases its in vivo persistence. In addition, we evaluated the in vivo therapeutic efficacy of ABD-ErA-TM in a B-ALL xenograft model of SUP-B15. Our results show a comparable long-lasting durable antileukemic effect between the standard-of-care pegylated-asparaginase and ABD-ErA-TM L-asparaginase, but with fewer co-glutaminase-related acute side effects. Since the toxic side effects of current L-asparaginases often result in treatment discontinuation in ALL patients, this novel ErA-TM variant with ultra-low L-glutaminase co-activity and long in vivo persistence may have great clinical potential.

Introduction

L-asparaginases (L-ASNases) are cornerstone drugs in the treatment of acute lymphoblastic leukemia (ALL) and have greatly improved patients' outcomes over the last 40 years.^{1,2} Since the introduction of L-ASNases, mortality rates in pediatric ALL patients have drastically decreased, currently reaching overall 5-year survival rates of almost 90%.³ While the depletion of blood asparagine (Asn) represents a critical component of the multiple-drug ALL treatment regimen,³⁻⁵ the use of L-ASNase is also associated with a variety of toxic side effects.⁶⁻⁹ A large part of these un-

wanted effects can be attributed to the secondary L-glutaminase co-activity displayed by all clinically approved bacterial L-ASNases.¹⁰ These L-ASNases not only convert Asn into L-aspartate (Asp) and ammonia, but also hydrolyse glutamine (Gln) into L-glutamate (Glu) and ammonia.¹¹⁻¹³ For all clinically approved L-ASNases, this L-glutaminase coactivity ranges from 2-10% of their primary L-ASNase activity depending on their microbiological origin.^{10,14,15} Importantly, Gln is much more abundant than Asn in the blood, with levels that range between 500-800 μ M compared to ~50 μ M for Asn.^{16,17} Because Gln levels are much higher, extensive amounts of ammonia are produced which might contribute to L-ASNase-induced neurotoxicity.¹⁸⁻²⁰ The L-glutaminase co-activity also has an impact on protein synthesis in the liver and pancreas resulting in hepatotoxicity, pancreatitis, hyperglycemia, leukopenia and thrombosis.^{6,14,21}

Over the past years, several attempts have been made to develop less toxic, low-glutaminase variants from various biological sources with better enzymatic parameters.^{10,21,22} Important criteria to keep in mind while designing alternative L-ASNases are efficacy, toxicity, kinetic properties, pharmacokinetics (i.e. in vivo half-life) and immunogenicity.²²⁻²⁴ The active degradation of native L-ASNase enzymes and their short half-life represents one of the biggest problems observed in the L-ASNase research field.²⁵ Currently, the first-line L-ASNase used in the clinic is polyethylene glycol (PEG)-asparaginase, a L-ASNase derived from Escherichia coli (EcA) which is conjugated to a PEG-tag that not only improves the half-life of the enzyme, but also the solubility and masks the bacterial enzyme from the immune system.²⁵⁻³¹ However, studies have shown that PEG itself can be immunogenic and that this anti-PEG immune response facilitates clearance in vivo.^{10,32,33} Considering the bacterial origin of clinically used L-ASNases, immune responses against the enzyme itself are also commonly observed during therapy, including bronchospasms, anaphylactic shock, and hypersensitivity reactions.^{21,34,35} Currently, debate is ongoing whether the L-glutaminase coactivity of clinically approved L-ASNases is essential, contributory or fully dispensable for their anticancer effect based on contradictory results.^{10,36-39} Several studies have shown that the cancer cell killing potential of L-glutaminase co-activity has been linked to asparagine synthetase (ASNS) expression levels.¹⁰ Chan et al. found that L-glutaminase co-activity is not required for the cell killing potential of L-ASNase against ASNS-negative ALL cancer cells in vitro using a newly designed L-glutaminase-deficient EcA Q59L mutant (EcA^{Q59L}),⁴⁰ while other groups reported that Gln depletion contributed to the overall anti-leukemic effect.^{38,40,41} In addition, Panosyan et al. showed that deamination of Gln may enhance the anti-leukemic effect of L-ASNase.⁴² Recently, Nguyen et al. engineered a variant of Erwinia chrysanthemi (ErA) with three mutations, ErA-TM, that acquired ultra-low L-glutaminase activity while maintaining its L-ASNase activity.¹⁵ ErA has a very short in vivo half-life, but the authors reported that the addition of a His-SUMO tag N-terminally to *ErA*-TM enhanced the drug's *in vivo* stability.⁴³ Using short-term His-SUMO-*ErA*-TM treatment, in vivo antileukemic efficacy was successfully demonstrated in a B-ALL xenograft model of SUP-B15 cells and a patient-derived T-ALL xenograft model,⁴³ confirming the previous *in vitro* results from Chan *et al.*⁴⁰ However, it was recently shown that the EcA^{Q59L} of Chan et al. did not induce a durable in vivo anticancer effect against xenografts of the ASNS-negative SUP-B15 leukemia cell line in

NOD/SCID gamma mice (NSG) compared to the L-ASNase wild-type (L-ASNase^{WT}),⁴⁴ therefore again contradicting previous reports.⁴⁰

To clarify whether L-glutaminase co-activity is required for a durable long-term anti-leukemic response, we took advantage of the superior tolerability of our low-glutaminase ErA-TM variant. To further improve the in vivo persistence (even longer than that afforded by the His-SUMO tag⁴³), we introduced an albumin-binding domain (ABD) N-terminal to ErA-TM, which we refer to as ABD-ErA-TM. We show that the introduction of an ABD-tag stabilizes ErA-TM and dramatically enhances its half-life. In addition, we evaluated the in vivo efficacy of this low-glutaminase variant in a SUP-B15 ALL xenograft model. Our results show a comparable long-lasting durable antileukemic effect between the mice treated with the standard-of-care PEG-asparaginase and ABD-ErA-TM, but with fewer co-glutaminase related nonimmunological toxicities. In the future, the implementation of this new L-ASNase variant in the clinic could improve ALL patients' outcomes and quality of life.

Methods

Expression and purification of ABD-*ErA***-TM and ABD-***ErA***-WT, crystal structure of ABD-***ErA***-TM, surface plasmon resonance and isothermal titration calorimetry** For detailed methodology, see the *Online Supplementary Appendix*.

Enzyme activity assay

The catalytic activities of the L-ASNase variants were determined using the aspartic acid β -hydroxamate (AHA) method as previously described.⁴⁵⁻⁴⁷ A detailed protocol can be found in the Online Supplementary Materials and Methods.

In vivo ABD-*ErA*-TM and PEG-asparaginase pharmacokinetics and pharmacodynamics study

The pilot pharmacodynamic study was performed at The University of Texas MD Anderson Cancer Center under protocol ACUF #00001658-RN01 following published methods.^{44,48} For the other pharmacodynamic and pharmacokinetic study, healthy wild-type C57BL/6J were randomly divided into vehicle phosphate-buffered saline (PBS) (n=5), PEG-asparaginase (n=15) and ABD-*ErA*-TM (n=15) treatment groups. One single administration/dose of 2,500 IU/kg PEG-asparaginase or ABD-*ErA*-TM was administered via intraperitoneal injection. For the activity determinations, 50 μ L of peripheral blood was collected and samples were processed according to the protocol described by Nguyen *et al.*⁴³ For the amino acid level determinations, 120 μ L of peripheral blood was collected and samples were measured as previously described.⁴³ For a more detailed

description of the methods used, see the Online Supplementary Appendix.

Single-dose acute toxicity study

Healthy wild-type C57BL/6J (Charles River) mice (9 weeks old; n=18) received one single administration/dose (2,500 IU/kg) of PEG-asparaginase or ABD-*ErA*-TM via intraperitoneal injection. After 96 hours, blood was collected and hematologic (Vetscan[®] HM5 Hematology Analyzer; Abaxis) and biochemical analyses (VetScan[®] VS2 Chemistry Analyzer; Abaxis) were performed. For detailed methodology, see the *Online Supplementary Appendix*.

In vivo treatment of a SUP-B15 cell line xenograft model with L-ASNases

NSG (#005557; The Jackson Laboratory) female mice (n=15) were intravenously injected at 6 weeks of age with 150 μ L PBS containing 5 × 10⁶ luciferase-positive SUP-B15 cells. At regular time points, the bioluminescence was measured using the IVIS Lumina II imaging system (PerkinElmer). After evidence of leukemic cell engraftment in the spleen and/or bone marrow, the mice were randomly divided into vehicle PBS (n=5), PEG-asparaginase (n=5) and ABD-*ErA*-TM (n=5) treatment groups. PEG-asparaginase was administered every 10 days via intraperitoneal injection at a dose of 2,500 IU/kg (total of 5 administration/doses), while ABD-*ErA*-TM was given at the same dose every 7 days (total of 7 administration/doses). During the experiment, leukemic burden was evaluated via bioluminescent imaging at regular time points.

The ethical committee on animal welfare at Ghent University Hospital approved all the *in vivo* mice experiments.

Statistical analysis

Statistical power was calculated before every experiment (80% statistical power, α =0.05). The Shapiro-Wilk test was used to check for normality of the data. The amino acid level data were not normally distributed and therefore non-parametric Mann-Whitney U tests were performed. A two-sided *P* value <0.05 was considered statistically significant. For the toxicity data, Mann-Whitney U tests, followed by Bonferroni correction for multiple comparisons, were performed. Results are expressed as mean ± standard deviation where appropriate. GraphPad Prism 9 (GraphPad Software Inc., La Jolly, CA, USA) was used to analyze the data.

Results

Introduction of an ABD-tag extends the *in vivo* persistence of *ErA*-TM L-ASNase variants

Dennis *et al.*⁴⁹ used phage display to select peptides that bind to serum albumins. The 18-residue peptide SA21 with

the highest binding affinity to the human serum albumin (467 nM) contains two cysteines speculated to form an intramolecular disulfide bond that would stabilize the binding to albumin. We fused this SA21 sequence, from now on referred to as ABD, to our low-glutaminase ErA-TM variant (Figure 1A). The challenge in the purification was the undesired buildup of inter-ABD disulfide bonds instead of the desired intra-ABD disulfide bonds. The lack of cysteine residues in ErA/ErA-TM suggested that any undesired intermolecular disulfide bridges are between the cysteine of an ABD in a protomer and a cysteine of an ABD in another protomer. In fact, using gel filtration we could see that ~25% of the protein eluted as an octamer instead of the physiological tetramer, presumably due to this incorrect disulfide bond formation between ABD and cysteine residues located in different tetramers (Online Supplementary Figure S1). For the experiments described, only the physiological tetrameric fraction was used.

We first verified if the k_{cat} and K_m values of ABD-*ErA*-TM were comparable to the ones from untagged *ErA*-TM and our previously reported His-SUMO-*ErA*-TM variant⁴³ (Table 1). The K_m property is key for obtaining complete Asn depletion *in vivo*. The introduction of the triple mutations to *ErA* increased the K_m from ~50 μ M to ~100 μ M, and the fusion of *ErA*-TM to ABD further increased the K_m to ~150 μ M.

To verify that ABD-ErA-TM can efficiently deplete Asn levels in vivo, we conducted a pilot pharmacodynamic properties study that showed a significant decrease of Asn levels upon ABD-ErA-TM administration (single administration/dose 2,500 IU/kg) (Online Supplementary Figure S2). At the same time, we were encouraged to see a significant increase in half-life compared to that of the His-SUMO-*ErA*-TM protein⁴³ (38.3 vs. 15.7 hours) (Table 2). Presumably, this increase in half-life is due to the interaction between the ABD moiety and the mouse serum albumin, and that this interaction slows down the degradation of the enzyme. To verify that our molecule binds serum albumins, we used surface plasmon resonance and isothermal titration calorimetry to determine the in vitro interaction with mouse and human serum albumins. Whereas ErA-TM shows no binding of either mouse or human serum albumin, ABD-ErA-TM binds with a low micromolar affinity to both (Table 3; Online Supplementary Figures S3 and S4). To test the prediction that a disulfide bridge between the two ABD cysteines is required for the interaction with albumins, we generated a variant in which these cysteines were mutated to serines $(ABD(C \rightarrow S)-ErA-$ TM). As expected, this variant exhibited no affinity to either mouse or human serum albumin. Indeed, the in vivo halflife of the ABD($C \rightarrow S$)-ErA-TM variant was essentially identical to that of ErA-TM (2.5 vs. 2.3 hours) (Table 2).

Finally, to obtain a molecular understanding of the ABD, we solved the crystal structure of ABD- $\it ErA$ -TM at 1.75 Å



Figure 1. Visualization of ABD-ErA-TM. (A) ABD-ErA-TM is expressed as a His-SUMO fusion protein. Cleavage with SUMO protease releases the molecule used in these studies. Two cysteine residues in the ABD region build the disulfide bond required for albumin binding. (B) The tetrameric ABD-ErA-TM as observed in the crystal structure solved at 1.75 Å resolution. A disulfide bridge is formed between two cysteines of ABD that acts to rigidify the loop structure. Thus, for the first time, we have a view of the ABD-conformation that is required for the interaction with serum albumins.

Table 1. L-ASNase kinetic parameters for ABD-ErA wild type and select mutants. This table shows the K_m and k_{cat} values of our low-glutaminase alternative ABD-ErA-TM variant compared to the ones from untagged ErA-TM and our previously reported His-SUMO-ErA-TM variant.43

| Enzyme | k _{cat} s⁻¹±SD | κ _m μM±SD | k _{cat} / K _m s ⁻¹ μΜ ⁻¹ | Specific activityª IU/mg±SD |
|-----------------------------|----------------------------|-------------------------|---|--------------------------------|
| ABD- <i>ErA</i> -WT | 311.3±5.5 | 39.3±3.0 | 7.92 | 504.9±30.1 |
| ABD- <i>ErA</i> -TM | 289.2±2.6 | 151.8±4.8 | 1.91 | 453.1±6.2 |
| ABD-ErA-TM C7S/C13S | 272.8±3.9 | 138.9±6.2 | 1.96 | 417.7±13.6 |
| <i>ErA</i> -WT ^ь | 207.5±3.6 | 47.5±3.5 | 4.37 | 353.2±6.1 |
| His-SUMO- <i>ErA</i> -TM⁵ | 261.2±2.8 | 95.0±3.5 | 2.75 | 443.5 ± 4.8 |

^a1 IU of enzyme is defined as the amount of enzyme used to produce 1 μmol of aspartate in 1 minute at 37°C. ^bKinetic parameters reported by Nguyen et al.¹⁵ The introduction of an albumin binding domain (ABD) increases the K_m value, a parameter for asparagine specificity, from ~40 µM (ABD-ErA-WT) to ~150 µM. The specific activity (IU/mg of protein) is a commonly used parameter to express specific L-ASNase activity. SD: standard deviation.

resolution. Data collection and refinement statistics are formation that is required for the interaction with serum shown in Online Supplementary Table S1. In the tetrameric molecule, one protomer had clear electron density for the ABD. Presumably, the ABD of the other three pro- In vivo pharmacokinetic and pharmacodynamic profiling tomers were not seen due to being highly flexible. As of ABD-ErA-TM in comparison to PEG-asparaginase shown in Figure 1B, the two cysteines of the ABD form a To further document the pharmacokinetic profile and disulfide bridge that acts to rigidify the loop structure. pharmacodynamic properties of our ABD-stabilized L-Thus, for the first time, we have a view of the ABD-con- ASNase variant in comparison to those of the standard-

albumins.

of-care, healthy C57BL/6J mice were injected intraperitoneally with 2,500 IU/kg of PEG-asparaginase or ABD-ErA-TM. The L-ASNase activity was measured at days 1, 3, 5, 7, 10 and 14. Although a similar pharmacokinetic profile was seen with the PEG-asparaginase, the activity of ABD-ErA-TM decreased more rapidly compared to that of the PEGasparaginase (Figure 2A). In addition, we determined the Asn, Asp, Gln, and Glu levels in blood plasma prior to the treatment and at days 1, 3, 5, 7, 10 and 14 after L-ASNase administration. Both enzymes efficiently depleted the blood Asn in a comparable manner (P=0.0079; Mann-Whitney U test), with levels slowly recovering after day 5 for ABD-ErA-TM and after day 7 for PEG-asparaginase (Figure 2B). In line with these results, plasma Asp levels increased significantly at days 1 and 3 (P=0.0079; Mann-Whitney U test) compared to baseline levels and then decreased to the pretreatment level by day 14 for both treatment groups (Online Supplementary Figure S5A). For Gln, only at 24 hours after administration was a minimal but significant reduction in Gln plasma levels seen for PEG-asparaginase, while no decrease of Gln levels was observed in ABD-ErA-TM mice (Online Supplementary Figure S5B). On the other hand, a consistent differential effect was demonstrated on the plasma Glu levels, with a significant increase in the PEG-asparaginase mice up to day 7 after administration, followed by a decrease towards baseline levels again. These increased Glu levels upon PEG-asparaginase treatment point out that long-term L-glutaminase co-activity is maintained. Notably, a minimal increase was observed in the ABD-ErA-TM group but not to the extent as in the PEG-asparaginase mice, as to be expected from this lowglutaminase variant (Online Supplementary Figure S5C).

Toxicity profiling of ABD-ErA-TM

Next, we evaluated the acute toxicity of ABD-*ErA*-TM *versus* PEG-asparaginase in healthy wild-type C57BL/6J female mice at a dose of 2,500 IU/kg. We compared the impact of both enzymes on the animals' body weight, one of the main parameters to evaluate L-ASNase toxicity. At

the end of the experiment, PEG-asparaginase mice lost a greater percentage of their starting body weight (*P*=0.0022; Mann-Whitney U test), while the ABD-ErA-TM mice showed no significant decrease in body weight (Figure 2C). In addition, we also analyzed hematologic parameters, including white blood cell count, red blood cell count and blood platelet levels. While no effect was observed on red blood cell counts, significant differences were seen in white blood cell and blood platelet levels (Online Supplementary Figure S6A-C). As previously mentioned, side effects of L-ASNase treatment, in particular liver and pancreas toxicity, are often observed. Blood chemistry analysis showed a significant decrease in globulin and albumin levels, both markers of hepatic dysfunction, in the PEG-asparaginase group compared to the vehicle group (P=0.0366 and P=0.0084, respectively) (Online Supplementary Figure S6D-E). Other blood parameters, including alanine aminotransferase (Online Supplementary Figure S6F), total bilirubin (Online Supplementary Figure S6G), blood urea nitrogen (Online Supplementary Figure S6H) and amylase (Online Supplementary Figure S6I) did not show any significant changes between groups.

In vivo therapeutic evaluation of ABD-*ErA*-TM in a B-cell acute lymphoblastic leukemia cell line xenograft model We evaluated the *in vivo* efficacy of long-term Asn deprivation using ABD-*ErA*-TM in a luciferase positive SUP-B15

vation using ABD-*ErA*-TM in a luciferase positive SUP-B15 xenograft mouse model. To examine the need for co-glu-

Table 2. Half-life of *ErA* variants.

| Protein | Half-life, (hours) |
|--------------------------|--------------------|
| <i>ErA</i> -TM | 2.3 |
| His-SUMO- <i>ErA</i> -TM | 15.7 |
| ABD- <i>ErA</i> -TM | 38.3 |
| ABD(C→S)- <i>ErA</i> -TM | 2.5 |

ABD-*ErA*-TM shows a significant increase in half-life compared to the His-SUMO-ErA-TM protein⁴³ (38.3 vs. 15.7 hours). The *in vivo* half-life of the ABD(C \rightarrow S)-*ErA*-TM variant was essentially identical to that of *ErA*-TM (2.5 vs. 2.3 hours), demonstrating the need for a disulfide bridge between the two ABD cysteines.

Table 3. Binding affinities determined by surface plasmon resonance and isothermal titration calorimetry.

| | Human serum albumin | | | Mouse serum albumin | | | | |
|-----------------------------|--|---|------------------------|------------------------|--|---|------------------------|------------------------|
| | SPR | | ITC | SPR | | | ІТС | |
| | k _a (x 10 ³ M ⁻¹ s ⁻¹) | k _d (x 10 ⁻² s ⁻¹) | κ _⊳ (μM) | κ _d (μΜ) | k _a (x 10 ³ M ⁻¹ s ⁻¹) | k _d (x 10 ⁻² s ⁻¹) | κ _⊳ (μM) | κ _d (μΜ) |
| <i>ErA</i> -TM | NB | NB | NB | NT | NB | NB | NB | NT |
| ABD- <i>ErA</i> -TM | 1.78±0.55 | 1.41±0.30 | 8.11±0.80 | 4.37 | 1.71±0.64 | 0.88±0.30 | 5.20±0.41 | 4.95±0.49 |
| ABD(C→S)- <i>ErA</i> -TM | NB | NB | NB | NT | NB | NB | NB | NT |

Surface plasmon resonance (SPR) and isothermal titration calorimetry (ITC) were used to determine the *in vitro* interaction with mouse serum albumin (MSA) and human serum albumin (HAS). While *ErA*-TM shows no binding of either MSA or HSA, ABD-*ErA*-TM binds with a low micromolar affinity to both MSA and HSA. Values are means ± standard deviations from four measurements each for SPR, two measurements for ITC with MSA, and a single measurement for ITC with HSA. NB: no binding; NT: not tested.





Figure 2. ABD-ErA-TM shows reduced toxicity in mice compared to **PEG-asparaginase but displays a similar pharmacokinetic profile.** (A) L-ASNase activity measurements in blood plasma samples of mice obtained at days 1, 3, 5, 7, 10 and 14 after 2,500 IU/kg of L-ASNase show a comparable pharmacokinetic profile for both enzymes, although the activity of ABD-ErA-TM decreased more rapidly compared to that of the PEG-asparaginase. (B) Measurement of asparagine levels in blood plasma samples of mice obtained at days 1, 3, 5, 7, 10 and 14 after 2,500 IU/kg of PEG-asparaginase or ABD-ErA-TM. A significant depletion of asparagine is observed for both enzymes (P=0.0079; Mann-Whitney U-test, with levels slowly recovering after day 5 for ABD-ErA-TM and day 7 for PEGasparaginase. (C) After 96 hours, PEG-asparaginase mice lost a greater percentage of their starting body weight (P=0.0022; Mann-Whitney U test), while the ABD-ErA-TM mice showed no significant decrease in body weight after L-ASNase administration. For each figure, the mean and standard deviation are shown.

taminase activity to kill ALL cells, we compared our stabilized low-glutaminase variant ABD-ErA-TM with PEG-asparaginase. Based on the results of our pharmacokinetic and pharmacodynamic studies, a dose of ABD-ErA-TM (2,500 IU/kg) was administered every 7 days, while PEGasparaginase (2,500 IU/kg) was given every 10 days guaranteeing continuous Asn deprivation in the blood with minimal residual L-ASNase activity building up before each new dose. In total, mice received either five doses of PEGasparaginase or seven doses of ABD-ErA-TM (Figure 3A). Results not only show that both ABD-ErA-TM and PEG-asparaginase were equally capable of rapidly lowering the leukemic burden, but also that a comparable long-term survival could be accomplished (Figure 3B). During the treatment time course, both PEG-asparaginase and ABD-*ErA*-TM treated mice displayed lower bioluminescent signals, below the background detection levels at the end of the treatment, while PBS control mice clearly showed an increase in leukemic burden (Figure 3C). At day 55, control mice had to be euthanized due to ethical humane endpoints, while both PEG-asparaginase and ABD-ErA-TM

mice showed no sign of ALL burden up to day 204, with the exception of one PEG-asparaginase treated mouse that relapsed at day 180 and clearly showed clinical signs of tumor formation.

The L-ASNase activity in blood plasma samples was measured 24 hours after administration of ABD-*ErA*-TM (at days 1, 8, 15, 22, 36 and 43) and PEG-asparaginase (at days 1, 11, 21 and 41). Results indicate that acceptable stable enzyme activity levels were maintained during the whole experiment (threshold for adequate activity >100 IU/L) without any excessive build-up of L-ASNase activity (Figure 3D). In addition, we also determined the amino acid levels in treated mice to confirm efficacy of Asn depletion during L-ASNase treatment. Both enzymes depleted the blood Asn, which was below detection at every time point (Figure 3E).

Besides the therapeutic effect, the impact and tolerability of both drugs was evaluated on the body weight of the animals. At the end of treatment, we observed a mean weight loss of 8.8% in the ABD-*ErA*-TM treatment group, while the PEG-asparaginase mice had a mean weight loss



Figure 3. The low L-glutaminase ABD-ErA-TM eliminates SUP-B15 cells as effectively as the high L-glutaminase PEGasparaginase, but with reduced toxicity. (A) Mice were injected with luciferase-expressing SUP-B15 cells and monitored for engraftment via bioluminescent imaging (BLI). Once engraftment was observed, mice were treated intraperitoneally (i.p.) every 10 days with PEG-asparaginase (n=5), every 7 days with ABD-*ErA*-TM (n=5) (drug dose for both enzymes 2,500 IU/kg) or vehicle (phosphate-buffered saline, PBS) (n=5). At regular time points during treatment, blood plasma samples were collected for amino acid determinations or activity measurements. (B) For each group, the BLI at regular time points is shown. Before the start of treatment, mice clearly show engraftment of SUP-B15 cells in the bone marrow and spleen area. At days 13 and 31, vehicle PBStreated mice show a very strong BLI signal consistent with a high disease burden, while ABD-ErA-TM and PEG-asparaginase mice show no sign of leukemia. After treatment termination, mice were followed up for survival up to day 204 after initiation of treatment. Only one PEG-asparaginase-treated mouse relapsed and had to be euthanized due to ethical reasons. (C) For both L-ASNase treatment groups, the radiance dramatically decreased returning to background levels by day 31, with the exception of one PEG-asparaginase-treated mouse that relapsed during the survival monitoring. (D) The L-ASNase activity in blood plasma samples was measured 24 hours after each L-ASNase administration. Results clearly show that L-ASNase activity was maintained during the treatment time course. For each time point, the mean and SD are shown. (E) Asparagine levels were measured prior to L-ASNase administration at days 6 and 27 for ABD-ErA-TM and at days 9 and 29 for PEG-asparaginase. Our data show that asparagine levels significantly decreased during L-ASNase therapy in the same manner for both treatment groups compared to day 0 (P=0.0079; Mann-Whitney U test). For each time point, the mean and SD are shown. (F) At the end of treatment, a mean weight loss of 8.8% in ABD-ErA-TM-treated mice was observed, while the PEG-asparaginase-treated mice had a mean weight loss of 13.9% (ABD-ErA-TM vs. PEG-asparaginase P=0.0011; Mann-Whitney U test), demonstrating a better tolerability of ABD-ErA-TM in these mice. For each time point, the mean and SD are shown.

of 13.9% (Figure 3F). At the end of this survival experiment (day 204), mice were sacrificed and analyzed for signs of leukemia development. No significant difference between spleen/body weight in both treated groups was observed (*Online Supplementary Figure S7A*). In addition, analysis of human CD45⁺ cells in peripheral blood, spleen and bone marrow showed undetectable levels of leukemic cells in both PEG-asparaginase- and ABD-*ErA*-TM-treated mice, with exception of the PEG-asparaginase mouse that relapsed during survival monitoring (*Online Supplementary Figure S7B*).

Together, these results convincingly show that the ABDtag stabilizes this *ErA*-TM variant *in vivo* and ABD-*ErA*-TM can lower ALL leukemic disease burden in a manner comparable to that of the high-glutaminase standard-of-care PEG-asparaginase, but with a better tolerability.

Discussion

Although L-ASNase is a key chemotherapeutic agent in childhood ALL, particularly during the induction phase, its administration is associated with multiple and severe toxicities. Unfortunately, these side effects prevent many patients from receiving their prescribed course of treatment, resulting in a detrimental impact on their outcome.^{50,51} Several studies showed that the L-glutaminase co-activity of clinically used L-ASNases contributes partially to these unwanted side effects.²¹ Therefore, lowering L-glutaminase co-activity while maintaining its anticancer effect might provide the required safety profile that reduces treatment interruptions.

Immune reactions during L-ASNase therapy occur frequently together with the development of anti-L-ASNase antibodies.⁵² Patients suffering from hypersensitivity reactions or silent inactivation to *EcA* are switched to native

ErA.^{50,52-54} Although the latter is a useful second-line therapy in ALL, some disadvantages are associated with this drug, including a much shorter half-life compared to that of PEG-asparaginase.^{6,55} As patients can also develop antibodies against the PEG-tag itself, thereby limiting tolerance and efficacy,^{33,56} additional research on more immunotolerant tags such as XTEN⁵⁷ and PASylation^{58,59} is currently ongoing. In our previous study,43 we presented an ErA-TM variant with significantly lower L-glutaminase but comparable L-ASNase activity relative to ErA-WT. We also achieved an extended circulation time by the fusion of a SUMO tag to these molecules. With the goal of further increasing blood circulation time, we analyzed several other tags, and discovered that the fusion of an ABD to ErA (WT or TM variant) resulted in respective molecules with much longer in vivo half-lives. In this study, we present ABD-ErA-TM, a novel ErA derived variant that is superior to our previous ones⁴³ by combining both low L-glutaminase co-activity and an exceptionally extended in vivo persistence.

Acute toxicological evaluation of ABD-*ErA*-TM revealed reduced non-immunological toxicity compared to the current first-in-line EcA. Our toxicity data show a clear correlation between reduced L-glutaminase co-activity and reduced drug toxicity, especially on the average percentage of body weight loss. We note that other biochemical parameters of liver and pancreas toxicity including alanine aminotransferase, total bilirubin, blood urea nitrogen and amylase levels did not show any changes. This may not be surprising since L-ASNase toxicity in patients is often only observed after multiple L-ASNase administration/doses.

The clinical relevance of L-glutaminase co-activity, present in all clinically Food and Drug Administration (FDA)approved bacterial L-ASNases, in the context of ALL remains largely unclear. First, it was shown that this L- glutaminase co-activity was not required to induce an in vivo L-ASNase effect in the leukemic SUP-B15 cell line.43 However, Chan et al.44 showed that a durable and longlasting antileukemic effect in the same SUP-B15 xenograft could only be obtained using a L-ASNase variant that retained its L-glutaminase co-activity.⁴⁴ Our results clearly contradict these findings and demonstrate that SUP-B15 leukemic cells can be durably killed with a L-ASNase variant with ultra-low glutaminase co-activity. We speculate that the lack of durable response noted by Chan et al.44 was due to insufficient long-term Asn depletion that resulted from using a non-stabilized L-ASNase variant, which means that it needs to be injected more frequently to observe a therapeutic effect. Our results clearly show a similar long-lasting durable anti-leukemic effect between PEG-asparaginase and ABD-ErA-TM.

Industrial production problems recently resulted in a shortage of ErA and reduced treatment options for patients who developed hypersensitivity or silent inactivation against PEG-asparaginase. To overcome these manufacturing issues, a recombinant version of ErA, JZP-458, was recently developed. While this JZP-458 has been approved by the FDA as part of a treatment regimen for children and adults with ALL or lymphoblastic lymphoma,⁶⁰ our ABD-*ErA*-TM has multiple benefits over the JZP-458 preparation. One is avoiding the toxicities that arise due to the glutaminase co-activity. Secondly, the in vivo persistence of JZP-458 is short whereas the albumin binding, provided by the ABD domain, increases the halflife >10-fold. This will provide a dramatic clinical advantage, allowing patients to be dosed once a week instead of three times a week as currently required for Erwinaze or JZP-458. In addition, our new Erwinia L-ASNase preparation requires no additional post-translational modifications resulting in high protein yields. As our ABD-ErA-TM would only need to be administered once a week, smaller amounts will be necessary to treat the same number of patients compared to current standard-of-care L-ASNases.

Until now, all immune-related side-effects were ignored in our experimental setup, as the *in vivo* efficacy study was done with human ALL cells xenotransplanted in immunodeficient NSG mice. Unfortunately, good preclinical models to evaluate the immunogenicity of L-ASNases are not available and this is one of the biggest challenges for future L-ASNase research.

In conclusion, this study convincingly shows that high Lglutaminase co-activity present in standard-of-care PEGasparaginase is not essential for durable anti-leukemic properties *in vivo*, therefore contradicting previous reports. Furthermore, our findings also suggest a decline in *in vivo* toxicity when L-glutaminase co-activity is reduced. In addition, the fusion of an ABD-tag to our alternative *ErA*-TM variant increases its half-life and *in vivo* stability. Identifying a L-ASNase variant with fewer side effects will not only improve the quality of life of ALL patients directly, but will also allow treatment protocols to be maintained or prolonged, which will translate into improved outcomes, especially for adult ALL patients. In addition, a few aggressive solid cancer subtypes with poor prognosis and a clinical unmet need have been identified as sensitive to Asn deprivation, including ovarian and metastatic breast cancer. However, because of the adverse side effects in adults, clinical trials have been discontinued so far. By identifying a L-ASNase variant with fewer unwanted toxicities, we hope to revive the clinical trials and provide extra therapeutic opportunities for these patients.

Disclosures

AMS, AL and HAN declare competing financial interests by being founders with equity stake in Enzyme by Design Inc.: a start-up developing new L-asparaginases. AL and HAN are inventors of the Int. Patent. Appl. PCT/US2017/020090 (WO2017151707A1) that describes the ABD-ErA-TM variant. None of the other authors declare any competing financial interests.

Contributions

MVT, AMS, HAN, TL, SG, PVV and AL designed the study; MVT, AMS, HAN, HL and WC performed experiments; LR and KV provided excellent technical assistance; TL, SG, PVV and AL supervised this study; MVT and AL drafted the manuscript; MVT, AL, HUN and YDV edited the figures; MVT, AMS, YDV, HAN, LR, KV, EP, YS, HL, PLL, WC, VM, BDM, TL, SG, PVV and AL proofread the original draft and made valuable adjustments; AMS, PLL, TL, SG, PVV, and AL obtained funding for this study. All authors have read and agreed to the published version of the manuscript.

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Data-sharing statement

The authors confirm that the data supporting the findings of this study are available within the article and its supplementary materials. More detailed data are available from the co-senior authors PVV and AL upon reasonable request.

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