

BOUNDLESS BIO

Unlocking a New Paradigm in Cancer Treatment via ecDNA-Directed Therapies (ecDTx)

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X @BoundlessBio

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Boundless Bio (BOLD): clinical-stage public company establishing a new category in oncology that addresses oncogene amplified cancers via targeting extrachromosomal DNA (ecDNA)



Oncogene amplified cancer:

- Generally unresponsive to targeted therapy and immunotherapy
- Significant unmet medical need (worse survival)
- ~1.3M new patients per year in major markets¹

ecDNA:

- Cancer-specific circular DNA—a root cause of oncogene amplification
- Transformative emerging area of cancer biology
- Spyglass drug discovery platform identifies ecDNA synthetic lethalities

ecDNA-directed therapies (ecDTx):

- BBI-355: oral CHK1 inhibitor, Phase 1/2 initial clinical POC data expected in 2H 2025
- BBI-825: oral RNR inhibitor, Phase 1/2 initial clinical POC data expected in 2H 2025
- ecDTx 3: targets novel kinesin, advancing toward development candidate
- ECHO diagnostic identifies ecDNA+ cancers to enable patient selection

Experienced team:

- Track record of precision oncology drug and diagnostic approvals, multi-\$B M&A
- Leading ecDNA scientific founders, board, advisors
- Cash runway into Q4 2026; expected to fund BBI-355 and BBI-825 clinical programs through initial POC

ecDNA are cancer-specific, circular units of DNA that are a frequent driver of oncogene amplified cancer



Next-generation precision oncology pipeline targets ecDNA to address high unmet need cancer patients





CHK1: checkpoint kinase 1 ecDTx: ecDNA-directed therapy RNR: ribonucleotide reductase

ECHO: <u>ec</u>DNA <u>H</u>arboring <u>O</u>ncogenes; BBI's proprietary ecDNA diagnostic IRB: institutional review board

Accomplished leadership team has proven experience delivering value for patients and shareholders





Significant unmet need in oncogene amplified cancers



Cancers with oncogene amplifications: more aggressive, difficult to treat, and worse prognosis

Oncogene amplified cancers

- Oncogene amplification is a type of oncogenic alteration where extra copies (>2) of an oncogene (e.g., EGFR) drive tumor growth or resistance
- Patients with oncogene amplifications have worse survival than other cancer patients
- Unlike other alterations, oncogene amplified tumors are generally unresponsive to targeted therapies and immunotherapies



Patients with primary or metastatic cancers with amplifications, point mutations, skipping deletions or fusions of these genes: AR, ALK, ARAF, BRAF, CCND1, CDK4, CDK6, EGFR, ERBB2, FGFR1, FGFR2, FGFR3, FGFR4, HRAS, IDH1, IDH2, KIT, KRAS, MAP2K1, MAP2K2, MAP2K4, MDM2, MET, MYC, NF1, NRAS, NTRK1, NTRK2, NTRK3, PDGFB, PDGFRA, PDGFRB, PIK3CA, RAF1, RET, ROS1

1. cBioPortal analysis using MSK-IMPACT (N=14,674 patients) and MSK-MET (N=1,115 patients) data; p-value = < 0.0001 PFS: progression free survival Despite advancements in precision medicine, cancers with <u>gene amplifications</u> generally do not respond to targeted therapies



A new approach is needed to treat cancers driven by oncogene amplifications



Across most oncogenes, patients with gene amplified tumors derive little benefit from targeted therapies



Targeted therapies have not been approved for,

nor demonstrated robust clinical activity in, most <u>oncogene amplified</u> cancers*



ORR: overall response rate NSCLC: non-small cell lung cancer



ecDNA: a key driver of oncogene amplifications



ecDNA are a primary driver of oncogene amplified cancers and enable resistance to targeted therapies

ecDNA are circles of DNA, **Circular shape of ecDNA enhances** ecDNA asymmetrically segregate during mitosis, distinct from chromosomes. enabling exponential copy number increase or transcriptional activity, leading to that amplify full-length genes high oncogene expression decrease during cellular division and regulatory elements Non-Mendelian Inheritance ecDNA lack Unique gene sequence centromeres (DNASeq) Potential exponential 00 copy number 02 increase after each cell division 549: mother cell **Open chromatin** (ATACSeq) 5558 VOPP1 daughter cells **High transcriptional** activity (RNASeq) **Protein products of genes** amplified on ecDNA can provide a fitness advantage - driving cancer growth and resistance Active Gene Expression Large size: 2 – 5 Mbp MYCN amplifications on ecDNA in pediatric neuroblastoma

ecDNA are detected broadly across different cancer types, but not in normal tissue or blood



BOUNDLESS BIO Kim, Nguyen, Turner, et al. 2020 Nature Genetics

PCAWG: Pan-Cancer Analysis of Whole Genomes TCGA: The Cancer Genome Atlas WGS: Whole Genome Sequencing Cancer's most common high copy number oncogene amplifications frequently occur on ecDNA; when they do, it is associated with worse survival

~54% of high-copy number oncogene amplifications are detected on ecDNA

Most frequently amplified oncogenes,

segmented by amplification type

Patients with oncogene amplification on ecDNA have worse survival

> Survival of cancer patients, segmented by gene amplification status



ecDNA enable cancer cells to resist therapies by rapidly adapting oncogene dependency



In this model, ecDNA enable gastric cancer cells to rapidly switch oncogene dependency from FGFR2 to EGFR under therapeutic pressure

Tumors employ ecDNA to rapidly enable new oncogenic drivers, even under combination target inhibition



Only targeting oncogenes amplified on ecDNA is a futile therapeutic approach due to ecDNA-enabled process of continuous and rapid oncogene dependency switching

Driving a new treatment paradigm by targeting ecDNA pathways that enable tumor evolution and resistance

Traditional Targeted Therapy:

Develop targeted inhibitors against activated oncodriver targets, but cells typically develop **resistance**



Applying traditional targeted therapy approach to ecDNA enabled cancers is clinical 'Whac-a-Mole'

Next Generation Precision Oncology:

Exploit underlying vulnerabilities in **ecDNA-driven cells** to drug targets essential for ecDNA functionality in cancer





Spyglass reveals cancer targets, both novel and validated, that intersect distinct nodes of ecDNA lifecycle

CHK1

BBI-355: PHASE 1/2 ENROLLING

Novel, oral, selective inhibitor of CHK1 BBI-098: 2nd generation CNS-penetrant CHK1i CHK1 is master regulator of ecDNA-induced replication stress

Novel kinesin

ecDTx 3: LEAD OPTIMIZATION

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Kinesin required for proper segregation of ecDNA during cell division



RNR

BBI-825: PHASE 1/2 ENROLLING

Novel, oral, selective inhibitor of RNR

RNR is a rate-limiting enzyme for assembly and repair of ecDNA ECHO: novel investigational diagnostic test designed to detect ecDNA using routine clinical NGS data Non-significant risk ("NSR") determination granted by FDA for use in Phase 1/2 POTENTIATE trial of BBI-355





BBI-355: potentially best-in-class, oral, selective CHK1 inhibitor in Phase 1/2 POTENTIATE trial

First ecDTx; targets ecDNA-induced replication stress



ecDNA+ oncogene amplified cancer cells have significantly elevated replication stress (RS)





Inhibition of checkpoint kinase 1 (CHK1) is synthetic lethal in ecDNA+ cancer cells CHK1 is a master regulator of the RS response



BBI-355: novel, oral, selective CHK1 inhibitor designed to disrupt ecDNA and overcome limitations of prior and existing CHK1 inhibitors





BBI-355 demonstrated single agent activity across a wide variety of oncogene amplified tumor models



BBI-355 demonstrated synergistic combination activity in preclinical models of cancer indications in which single agent targeted therapies have not proven effective in the clinic



• FGFR2 inhibition with infigratinib resulted in minimal, transient anti-tumor activity, consistent with clinical experience

When combined with BBI-355, extended synergistic tumor regression observed

BBI-355 demonstrated *in vivo* proof of concept in multiple additional oncogene addicted xenograft models Oncogene amplified sarcoma and gastric cancer; synergistic activity in combination with targeted therapy



Combination of BBI-355 with targeted therapy in vivo resulted in:

- Deeper tumor regressions
- Longer duration of response

Phase 1/2 study of BBI-355 designed to drive to clinical proof of concept in multiple solid tumor settings



"POTENTIATE" Study: Precision Oncology Trial Evaluating Novel Therapeutic Interrupting Amplifications Tied to ecDNA

BOIN: Bayesian optimal interval design MTD: Maximum Tolerated Dose; RP2D: Recommended Phase 2 Dose HGSOC: High-grade serous ovarian carcinoma *Amplification of wildtype driver oncogene, patients with known pathogenic driver mutations or fusions of ANY oncogene are excluded

Part 3 Simon's optimal two-stage design (Goal 1^{st} stage RR $\geq 2/23$)

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Preliminary human pharmacokinetic (PK) data of BBI-355 showed dose-proportionality and achieved exposures in the predicted therapeutically active range at 60 mg PO Q2D, which is a tolerated dose level





- BBI-355 demonstrated good oral bioavailability in human subjects
- Average C_{max} and AUC showed dose proportionality from 20 to 80 mg Q2D
- Average T_{1/2}: ~40h, leading to drug accumulation of ~2 to 3-fold
- Moderate inter-subject variability observed

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*Dose limiting toxicities (DLTs) in 2 of 4 subjects (i.e., neutrophil count decreased, platelet count decreased) **DLTs in 3 of 3 subjects (i.e., neutrophil count decreased, platelet count decreased) *DLT in 1 of 5 subjects (i.e., neutrophil count decreased)

AUC: area under the curve PO: oral

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Evidence of BBI-355 pharmacodynamic activity observed in clinical samples across dose levels pCHK1-S345 induction in skin and tumor biopsies





BBI-355 summary: the first ecDTx in clinical development for oncogene amplified cancer

CHK1: checkpoint kinase 1

BBI-355: PHASE 1/2

- BBI-355 is a potentially best-in-class, oral, selective CHK1 inhibitor in development to address the unmet medical needs of patients with oncogene amplified cancer
- **Currently no cancer therapy** has been approved for patients with *EGFR*, *FGFR*, or *CDK4/6* amplifications, a large segment of cancer patients
- The **POTENTIATE** trial's modular design (<u>NCT05827614</u>) enables multiple avenues for expansion opportunities across diverse oncogene amplifications and tumor types
- Initial human PK data shows dose-proportionality with exposures in the predicted therapeutically active range
- Preliminary clinical data of BBI-355 as a single agent and in combination with EGFR or FGFR inhibitors in 2H 2025





BBI-825: first-in-class, oral, selective RNR inhibitor in Phase 1/2 STARMAP trial

Second ecDTx; targets ecDNA assembly & repair



Oncogene amplifications, often on ecDNA, are a frequent mechanism of clinical resistance to multiple therapeutic modalities

Tissue images from clinical specimens suggest resistance amplifications are frequently ecDNA-mediated -



MET (on ecDNA)

Ribonucleotide reductase (RNR) is the rate-limiting enzyme in the *de novo* synthesis of dNTPs, which are essential for the assembly and repair of ecDNA



BBI-825 is a novel, oral, selective RNR inhibitor designed to disrupt the assembly and repair of ecDNA





BBI-825 resulted in dNTP depletion, reduced ecDNA, and cytotoxicity in ecDNA amplified cancer cells



In *MYC* and *FGFR2* amplified ecDNA+ GI cancer cells, treatment with BBI-825 resulted in depletion of dNTPs and reduced ecDNA levels, leading to tumor cell death



BBI-825 demonstrated synthetic lethality in combination with KRAS^{G12C} inhibition in *KRAS^{G12C}*-addicted syngeneic colorectal cancer xenograft, both preventing and treating resistance post-emergence

Combination of BBI-825 with adagrasib prevented resistance to KRAS^{G12C} inhibition, resulting in durable tumor regressions



Single-agent BBI-825 treated resistance post-KRAS^{G12C} inhibition, resulting in significant anti-tumor activity



resistant to adagrasib via amplifications on ecDNA

BBI-825 overcame amplification-based resistance to BRAFi + EGFRi treatment of BRAF^{V600E} CRC cells in vitro



- BRAF^{V600E} mutant CRC cell lines developed rapid resistance to encorafenib + cetuximab via oncogene amplification
- Combination with BBI-825 prevented resistance and led to cancer cell death
- Additional BRAF^{V600E} melanoma and endometrial cancer models also demonstrated BBI-825 synergy with standard of care



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E: encorafenib C: cetuximab CRC: colorectal cancer

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Phase 1/2 study of BBI-825 evaluates prevention and treatment of amplification-mediated resistance to RTK/MAPK pathway inhibitors



"STARMAP": Study Treating Acquired Resistance; MAPK Amplifications



CRC: Colorectal cancer MAPK: Mitogen-activated protein kinases RP2D: Recommended Phase 2 Dose RTK: Receptor tyrosine kinase

MAPK-pathway amplification-mediated acquired resistance Initial clinical proof of concept (cPOC) and total addressable market (TAM)



While we seek to demonstrate initial POC in KRAS^{G12C} and BRAF^{V600E} CRC, BBI-825's TAM may be pan-tumor and pan-RAS



BBI-825 summary: the second ecDTx in clinical development for cancer with resistance gene amplifications



RNR: ribonucleotide reductase

BBI-825: PHASE 1/2

- BBI-825 is a **first-in-class**, **oral**, **selective** RNR inhibitor in development to address **patient populations** with MAPK-pathway activated cancer
- **Rapid resistance in multiple tumor types** is a limitation of current targeted therapies, presumably due to **amplification** of resistance genes
- Preclinically, BBI-825 demonstrated significant tumor growth inhibition, including regressions, in MAPK pathway-activated tumor models
- STARMAP trial modular design (<u>NCT06299761</u>) enables multiple avenues for expansion opportunities across pan-tumor and pan-RAS and activated cancers
- Preliminary clinical data of BBI-825 in combination with BRAF & EGFR or KRAS & EGFR inhibitors in *BRAF^{V600E}* and *KRAS^{G12C}* mutated colorectal cancer with resistance gene amplification in 2H 2025



Boundless Bio: leading a new area of cancer biology and targeting a large unmet need



Seeking to address oncogene amplification market by targeting oncogene agnostic nodes of ecDNA biology



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1: # of new patients per year in US/EU/Japan with EGFR, FGFR, CDK4/6 amplifications on ecDNA are estimated to be 15K, 50K, 15K, respectively

2: # of new CRC patients per year in US/EU/Japan with $KRAS^{G12C}$ and $BRAF^{V600E}$ are estimated to be 5K and 13K, respectively

Boundless Bio is leading a compelling and differentiated approach to address oncogene amplified cancers

Dedicated to Oncogene Amplified Cancers by targeting a unique cancer biology	 Oncogene amplifications: one of cancer's highest unmet medical needs, represents expansive addressable market ecDNA: a root cause of amplification; Boundless Bio is the leading ecDNA company Spyglass: ecDNA-focused discovery engine ecDTx: multiple clinical-stage programs with robust preclinical data ECHO: diagnostic designed to identify ecDNA+ cancers using routine NGS assays 					
Fortress Position, Track Record of Success, Well- Funded	 Founded by world's leading ecDNA experts; led by experienced management team with a track record of precision oncology drug approvals and multi-\$B M&A All ecDTx internally discovered and wholly-owned; IP life through at least 2041-2044 Approximately \$179M in cash and equivalents*, provides expected runway into Q4 2026 					
	ecDTx	Target	ecDNA Node	Anticipated Milestones		
	PDI 255		Poplication Stross	24 2025, Initial clinical POC from Phase 1/2 POTENTIATE trial		

Multiple Value Drivers	BBI-355	CHK1	Replication Stress	2H 2025: Initial clinical POC from Phase 1/2 POTENTIATE trial
	BBI-825	RNR	Assembly & Repair	2H 2025: Initial clinical POC from Phase 1/2 STARMAP trial
	ecDTx 3	Kinesin	Segregation	1H 2026: Submit IND

Boundless Bio: a breakthrough biotech company leading the next wave of innovation in cancer treatment





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Unbound by convention, bound to save lives

www.boundlessbio.com

Bibliography of recent reviews and publications covering ecDNA—active hyperlinks

2024	Paul Mischel, Howard Chang	Nature Reviews Cancer: Extrachromosomal DNA in cancer
2023	Paul Mischel, Vineet Bafna, Howard Chang, Roel Verhaak	Nature: Extrachromosomal DNA in the cancerous transformation of Barrett's oesophagus
2022	Anton Henssen (Charité Berlin), Roel Verhaak	Nature Reviews: Extrachromosomal DNA amplifications in cancer
2022	Vineet Bafna, Paul Mischel	Annual Reviews: Extrachromosomal DNA in Cancer
2022	Paul Mischel, Howard Chang	Nature Structural and Molecular Biology: Gene regulation on extrachromosomal DNA
2022	Samuel Bakhoum (MSKCC)	Cancer Genomics: Chromosomal instability as a source of genomic plasticity
2022	Rene Medema (Netherlands Cancer Inst.)	Chromosoma: Life of double minutes: generation, maintenance, and elimination
2022	Vineet Bafna, Howard Chang, Paul Mischel	Annual Reviews: Extrachromosomal DNA: An Emerging Hallmark in Human Cancer
2020	Anton Henssen, Howard Chang, Paul Mischel, Vineet Bafna, Roel Verhaak	Nature Genetics: Extrachromosomal DNA is associated with oncogene amplification and poor outcome across multiple cancers
2020	Paul Mischel, Charles Swanton (Crick Inst.)	Annals of Oncology: Extrachromosomal DNA—relieving heredity constraints, accelerating tumour evolution
2020	Christopher Ott (Mass Gen)	Cancer Cell: Circles with a Point: New Insights into Oncogenic Extrachromosomal DNA
2019	Roel Verhaak, Vineet Bafna, Paul Mischel	Nature Reviews: Extrachromosomal oncogene amplification in tumour pathogenesis and evolution

