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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 lethal targets

ecDNA-directed therapies (ecDTx):

- BBI-355: oral CHK1 inhibitor, POTENTIATE Phase 1/2 trial: initial clinical POC data expected in 2H 2025
- BBI-825: oral RNR inhibitor, STARMAP Phase 1/2 trial: not advancing at this time
- ecDTx 3: targets novel kinesin, advancing toward development candidate by mid-2025
- ECHO diagnostic identifies ecDNA+ amplified cancers: in use in POTENTIATE study

Experienced team:

- Track record of precision oncology drug and diagnostic approvals, multi-\$B M&A
- Leading ecDNA scientific founders, board, advisors
- Cash runway into 2027; funding BBI-355 through initial POC data and key milestones for ecDTx 3

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

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

ECHO: ecDNA Harboring Oncogenes; BBI's proprietary ecDNA diagnostic

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



Extended management team brings robust strategic and operational oncology experience



Tony Pinkerton, PhD	Peter Krein, PhD	Sara Weymer	Amy Berkley, PhD	Meredith Wesley
SVP, Drug Discovery	SVP, Precision Medicine	SVP, Clinical Operations	SVP, Program Team Leadership	SVP, Talent and Culture
Sanford Burnham Prebys	UNCOLOGY Ster	THERAPEUTICS	IONIS	ıgnyta
	QIAGEN	Z Biotech	Halozyme	ERASCA
Kalypsys.	Roche	Biogen	Lpath	SANDARUS, nc.º
	Selpercatinib capsules		Herceptin HYLECTA** trastuzumab and hyaluronidase-oysk Nufection FOR SUBCUTANEOUS USE 1. 600 mp.10.000 white	



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

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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, transcriptional activity, leading to enabling exponential copy number increase or 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 5493 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

Multiple clinical datasets support that ecDNA are detected broadly across major cancer types, but not in normal tissue or blood



Analysis of whole genome sequencing data from >3,000 tumor and matched normal samples from donors to TCGA and PCAWG; ~15,000 tumor samples from donors to GEL

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Kim, Nguyen, Turner, et al. 2020 Nature Genetics Bailey, Mischel, Swanton, et al. 2024 Nature 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



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

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



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: unique platform for interrogating ecDNA-driven tumors Proprietary target and drug candidate discovery engine



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

Kinesin required for proper segregation of ecDNA during cell division



RNR

RNR is a rate-limiting enzyme for assembly and repair of ecDNA

ECHO: investigational diagnostic test in clinical use 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

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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 PK data of single agent 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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Data cutoff of February 21, 2024

*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; Q2D: every two days 2ON/5OFF: two days on drug and 5 days off drug

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
- **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





ecDTx 3: Novel Kinesin Degrader

Targets ecDNA segregation



Targeting a novel Kinesin that regulates ecDNA segregation and viability of ecDNA bearing cells



during mitosis



Oobatake and Shimizu, Genes Chrom Canc 2019

- Chromosome segregation is primarily dependent on interactions between the mitotic spindle microtubules and the centromere, making some proteins non-essential in non-cancer cells
- Spyglass has revealed Kinesin, which is non-essential for chromosome segregation, but essential for proper ecDNA segregation
- No inhibitors of Kinesin have been disclosed ٠
- Genetic knockdown of Kinesin disrupts normal ecDNA segregation, reduces ecDNA, and shows synthetic lethality and anti-tumor activity in multiple ecDNA+ models
- Boundless has developed potent, selective, orally bioavailable ٠ degraders with anti-tumor activity in vivo in models insensitive or resistant to targeted therapy

Currently developing a potent, selective, oral Kinesin degrader as a potential first-in-class ecDTx

Selective Kinesin degrader causes ecDNA mis-segregation in FGFR2/MYC amplified tumor cells



- Kinesin targeted degrader (BBI-7769, 100 nM) induces misaligned ecDNA in mitosis
- Degrader-induced segregation defects phenocopy genetic knock-down

In vivo tumor growth inhibition demonstrated using Kinesin degrader tool compound



- Kinesin degrader demonstrated significant monotherapy TGI and blocks *MYCN* amplification; ~90% degradation of Kinesin *in vivo*
- Overcame resistance to HER2i in ERBB2/MYC amplified tumor model



ecDTx 3 summary: novel Kinesin degrader advancing towards development candidate

Novel Kinesin

ecDTx 3: Lead Optimization

- ecDTx 3 is a potentially **first-in-class**, **oral**, **selective** Kinesin degrader
- Kinesin is a novel cancer target that is **essential for ecDNA segregation** but **non-essential** for normal chromosome segregation
- In vitro cytotoxicity and *in vivo* proof of concept established in multiple oncogene amplified models
- Initial biomarker/patient selection strategy for monotherapy identified
- Oral Kinesin degrader on path towards Development Candidate in mid-2025 with plans for **IND Submission in 1H 2026**





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





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 patients per year in US/EU/Japan with *AR*, *MDM2*, *MET* amplifications on ecDNA are estimated to be 30K, 25K, 5K, 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 programs; first in the clinic ECHO: diagnostic designed to identify ecDNA+ amplified 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 \$167M in cash and equivalents*, provides expected runway into 2027

	ecDTx	Target	ecDNA Node	Anticipated Milestones
Multiple Value Drivers	BBI-355	CHK1	Replication Stress	2H 2025: Initial clinical POC from Phase 1/2 POTENTIATE trial
	ecDTx 3	Kinesin	Segregation	Mid-2025: Development Candidate; 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: Enhancing transcription-replication conflict targets ecDNA-positive cancers	
2024	Howard Chang, Paul Mischel, Charles Swanton	Nature: Origins and impact of extrachromosomal DNA	
2024	Ben Cravatt, Paul Mischel, Howard Chang	Nature: Coordinated inheritance of extrachromosomal DNAs in cancer cells	
2024	Vineet Bafna, Roel Verhaak	Nature Genetics: Mapping extrachromosomal DNA amplifications during cancer progression	
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	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	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)	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	
	Publed	ecDNA revolution	

