### PacBi

### **Accelerating innovation in genomics** 41<sup>st</sup> Annual J.P. Morgan Healthcare Conference

Christian Henry, President and CEO January 11, 2023

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## MISSION

Enabling the promise of genomics to better human health



We believe our new products will accelerate our growth

Revio, our transformative long-read system, exceeds our order expectations with 76 systems ordered in Q4

Onso, our highly accurate short-read system, is receiving excellent feedback in beta

Customers and ecosystem partners are increasingly working with PacBio across a range of applications to drive adoption

We have a strong balance sheet and intend to become cash flow positive during 2026



### Who is PacBio?



### 1,000+

Sequencers sold to date in 40 countries



#### ~9,000 Publications to date



### **400+** Exclusive issued US patents held<sup>1</sup>



### ~770 employees<sup>1</sup> ~230 commercial ~415 research + ops



#### **Instruments**



### Consumables



### Software/informatics

### Two differentiated technologies to sequence DNA in both long + short reads



6

### Using the **best-suited technology** in each application = **better results**

![](_page_6_Picture_1.jpeg)

#### **HiFi sequencing**

Delivers long reads with the highest accuracy<sup>1</sup> — even in hard-to-sequence regions

![](_page_6_Picture_4.jpeg)

#### **SBB** sequencing

Promises significant accuracy improvements over conventional NGS approaches

Complex disease research Plant + animal sciences Neuroscience Immunology

> Rare + inherited disease Gene editing confirmation Infectious disease/microbiology Targeted clinical panels

> > Therapy selection Noninvasive prenatal screening Early-stage cancer screening Cancer recurrence monitoring

![](_page_6_Picture_10.jpeg)

We serve a large market, ~\$7B today, growing to an estimated \$14B by 2026<sup>1</sup>

![](_page_7_Figure_1.jpeg)

Growth predicted across all segments, with highest growth in human germline + oncology

Increased investment in translational studies, including population-scale programs + expansion of sequencing into routine clinical testing

### In 2022, we had three strategic priorities

![](_page_8_Picture_1.jpeg)

![](_page_8_Picture_2.jpeg)

![](_page_8_Picture_3.jpeg)

### **Execution matters**

Leverage commercial investment to drive continued HiFi and Sequel II/IIe adoption

### **Delight our customers**

Deepen customer collaboration in clinical + a growing list of new applications

### **Progress product pipeline**

Demonstrate SBB as differentiated short-read offering + drive future HiFi system development **2022** preliminary financial performance

![](_page_9_Figure_1.jpeg)

### Continued to lay the foundation for HiFi with Sequel IIe expansion

512 374 Sequel I peak installed base 203 RS peak installed base 114 2019 2020 2021 2022

**Sequel II/IIe installed base** 

SMRT Cells shipped

![](_page_10_Figure_3.jpeg)

### Est. total petabase/year

![](_page_10_Figure_5.jpeg)

**37%** 2022 growth in Sequel II/IIe installed base 24%

2022 growth in Sequel II/IIe SMRT Cells

### **68**%

2022 growth in data generated from Sequel II/IIe

### Collaborations showing that HiFi uncovers variants missed by other sequencing technologies

![](_page_11_Picture_1.jpeg)

- Multiomic collab to include 5mC
- Early Revio adopter to add capacity; >1K rare disease genomes sequenced to date
- >4-fold rare coding SVs vs short-read genomes<sup>1</sup>

PacBi

![](_page_11_Picture_5.jpeg)

 Applying PacBio HiFi to unsolved short-read cases, which has led to improved diagnosis<sup>2</sup>

![](_page_11_Picture_7.jpeg)

- Aims to develop new methods to identify disease-causing mutations in Japanese population
- Intends to employ PacBio's tandem repeat genotyper (TRGT)

![](_page_11_Picture_10.jpeg)

- Plans to sequence over 300 rare disease genomes
- Increased diagnostic yield by using PacBio HiFi to identify variants missed by short-read sequencing<sup>3</sup>

1. https://doi.org/10.1016/j.gim.2022.02.007

2. https://www.ashg.org/wp-content/uploads/2022/09/ASHG2022-PosterAbstracts.pdf - pg 1450

3. https://www.ashg.org/wp-content/uploads/2022/09/ASHG2022-PosterAbstracts.pdf - pg 1442

### 2022 was a year of extraordinary innovation across the entire workflow

![](_page_12_Figure_1.jpeg)

Introducing the PacBio Compatible program, created to provide customers a curated portfolio across workflow in both long and short reads

![](_page_13_Figure_1.jpeg)

### Revio system designed from the ground up to deliver HiFi at scale

![](_page_14_Picture_1.jpeg)

![](_page_14_Picture_2.jpeg)

### Scale25M ZMW SMRT Cell

- 4 independent stages
- 24-hour cycle time

![](_page_14_Picture_6.jpeg)

### Ease of use

- 50% reduction in consumables
- Load-in-advance capability
- No N<sub>2</sub> requirement

![](_page_14_Picture_11.jpeg)

### **Compute power**

- 20× the relative compute power
- GPU-based architecture
- Google DeepConsensus on board

![](_page_14_Picture_16.jpeg)

### Affordability

- Sub \$1,000 human HiFi genome
- <1 minute to load instrument
- >50% decrease in file size

![](_page_14_Picture_21.jpeg)

### **Revio internal testing demonstrating robust performance**

![](_page_15_Picture_1.jpeg)

### In less than 3 months, Revio is off to best start in PacBio history

![](_page_16_Figure_1.jpeg)

Revio is designed to enable larger-scale genome projects and clinical researchers to sequence thousands of HiFi genomes

![](_page_17_Picture_1.jpeg)

Multi-unit Revio order to propel genomic medicine discovery in rare disease and cancer in Dubai

![](_page_17_Picture_3.jpeg)

### Radboudumc

Multi-unit Revio order to significantly ramp Darwin *Tree of Life* program and increase long reads in human applications such as single-cell transcriptomics + variant detection

The program has already solved several genetic mysteries using PacBio<sup>1</sup> and with Revio expects to ramp from 100s to 1,000s of genomes

### Onso designed to enable extraordinary accuracy for short-read sequencing

![](_page_18_Picture_1.jpeg)

![](_page_18_Picture_2.jpeg)

### Onso beta demonstrating ultra-high accuracy in customers' labs

![](_page_19_Picture_1.jpeg)

"We believe that more accurate reads will be transformative for many genomic applications, including oncology. We are excited to be evaluating the Onso platform with this in mind."

— Niall Lennon, PhD,
 Senior Director of Translational Genomics,
 Institute Scientist,
 Broad Institute

![](_page_19_Figure_4.jpeg)

![](_page_19_Picture_5.jpeg)

"We've been extremely impressed by Onso's levels of accuracy. This accuracy can open exciting new opportunities to transform agricultural biotechnology, specifically in areas like gene editing specificity analysis."

— Gina Zastrow-Hayes, Biotechnology and GT-Genomics Technology Manager, Corteva Agriscience

![](_page_19_Picture_8.jpeg)

Includes price of cluster generator; standalone instrument priced at \$225K
 Represents estimated list price of \$1,995 for the 2×150 kit and estimated yield
 Internal calculation sensitivity at 6,000x coverage with non-UMI SBB exceeds >24,000x coverage at 0.05% and 0.1% VAF

### PacBio entering next phase in strategic journey

![](_page_20_Figure_1.jpeg)

### **Our strategic priorities for 2023**

![](_page_21_Picture_1.jpeg)

![](_page_21_Figure_2.jpeg)

Drive rapid adoption of Revio by converting existing Sequel II/IIe customers + attracting new PacBio customers Demonstrate Onso's extraordinary level of accuracy in the field and show how it can transform research in needle-inhaystack applications

![](_page_21_Figure_5.jpeg)

Progress development of ultrahigh-throughput + benchtop long-read sequencers + next generation SBB sequencer

![](_page_21_Picture_7.jpeg)

Leverage current infrastructure to drive toward positive cash flow

Expand partnerships across ecosystem + workflow to drive customer adoption of SBB + HiFi

![](_page_21_Picture_11.jpeg)

### 2023 is first step on path to achieve >\$500M in revenue in 2026

![](_page_22_Figure_1.jpeg)

Revio planned to begin shipping in Q1 2023

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	1	

First Onso shipment anticipated in Q2 2023

![](_page_22_Figure_5.jpeg)

Enable new scaled projects as customers transition to Revio

![](_page_22_Figure_7.jpeg)

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