Transforming PacBio 40th Annual J.P. Morgan Healthcare

Christian Henry | President and Chief Executive Officer

Safe harbor statement

All statements in this presentation (and any accompanying oral presentation) that are not historical are forward-looking statements within the meaning of Section 27A of the Securities Act of 1933, as amended, Section 21E of the Securities Exchange Act of 1934, as amended, and the U.S. Private Securities Litigation Reform Act of 1995, including statements that generally relate to future events or our future financial or operating performance, future applications and availability, release dates, uses, accuracy, advantages, or quality or performance of, or benefits or expected benefits of using, our products or technologies, and the suitability or utility of our products or technologies for particular applications or projects. Such statements are based on management's current beliefs, estimates, assumptions and projections and on information available to management as of the date of this presentation. Forward-looking statements include, among other things, statements regarding our preliminary financial and operational results for the fourth quarter and full year ended December 31, 2021, including our expected revenue, cash balance, number of installed instruments, growth in SMRT Cells, growth relative to previous periods, data generated from Sequel II/IIe usage; rates and amounts of potential commercial opportunities and growth, including estimated sizes and penetration of certain markets; ability of HiFi Viral products to sequence viral genomes, including all circulating variants of SARS-CoV2; sequencing completeness and accuracy of our products and technology, including our HiFi and SBB technologies; estimates regarding our being the only company with both leading long read and short read technologies; expected purchases by certain customers upon future product release; attributes and sequencing advantages of, and potential upgrades and improvements to, our products and technology, including potential research and diagnostic benefits; strategic plans and priorities; research and development plans, including expected improvements and updates; commercial collaborations and partnerships; and other future events. Accordingly, you should not place undue reliance on forwardlooking statements because they involve known and unknown risks, uncertainties, changes in circumstances and other factors that are, in some cases, beyond the Company's control and could cause actual results to differ materially from the information expressed or implied by forward-looking statements made in the presentation. Factors that could materially affect actual results can be found in our filings with the Securities and Exchange Commission, including our most recent reports on Forms 8-K, 10-K and 10-Q, and include those listed under the caption "Risk Factors." The Company undertakes no obligation to revise or update information in this presentation to reflect events or circumstances in the future, even if new information becomes available.

The COVID-19 pandemic and efforts to control its spread have affected our financial results for the three and nine months ended September 30, 2021 and will likely continue to adversely impact our revenues for the fourth quarter of 2021. Due to the uncertain scope and duration of the pandemic, we cannot reasonably estimate the future impact to our operations and financial results. Even after the COVID-19 pandemic has subsided, we may continue to experience an adverse impact to our business as a result of its global economic impact, including any recession that has occurred or may occur in the future.



ENABLING THE PROMISE OF GENOMICS TO BETTER HUMAN HEALTH

We create some of the world's most advanced sequencing technologies.

Only company with both leading long-read and highly accurate short-read tech1

Three highly differentiated core technologies focused on accuracy, quality + completeness: HiFi, Nanobind, and SBB

50+ Products spanning entire genomics workflow

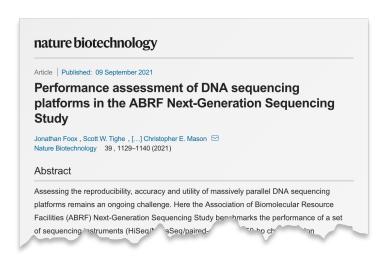
700+ Global employees

\$40B+ Total addressable market²

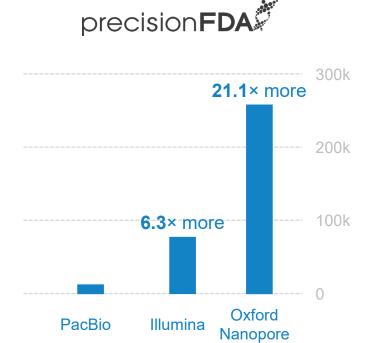


PacBio HiFi technology recognized as the most accurate + complete

Association of Biomolecular Resource Facilities



"PacBio [HiFi] provides the lowest error rate out of all technologies."



Total errors (SNV + indel + SV)²

Telomere-to-telomere Consortium

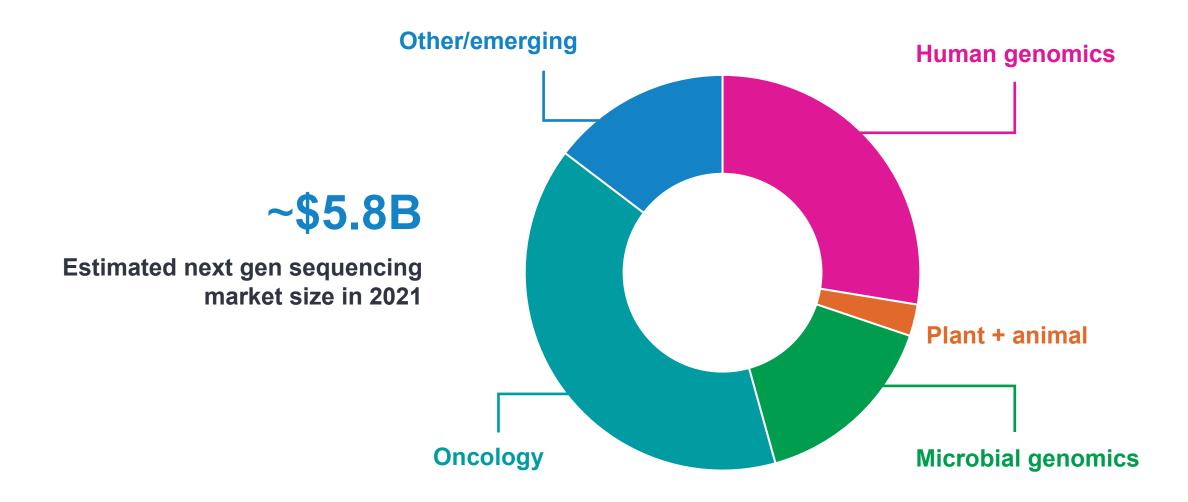
The complete sequence of a human genome

Sergey Nurk1., Sergey Koren1., Arang Rhie1., Mikko Rautiainen1., Andrey V. Bzikadze2, Alla Mikheenko3, Mitchell R. Vollger4, Nicolas Altemose5, Lev Uralsky6.7, Ariel Gershman8, Sergey Aganezov9, Savannah J. Hoyt¹⁰, Mark Diekhans¹¹, Glennis A. Logsdon⁴, Michael Alonge⁹, Stylianos E. Antonarakis¹², Matthew Borchers¹³, Gerard G. Bouffard¹⁴, Shelise Y. Brooks¹⁴, Gina V. Caldas¹⁵, Haoyu Cheng^{16,17}, Chen-Shan Chin¹⁸, William Chow¹⁹, Leonardo G, de Lima¹³, Philip C, Dishuck⁴, Richard Durbin²¹, Tatiana Dvorkina³, Ian T. Fiddes²², Giulio Formenti^{23,24}, Robert S. Fulton²⁵, Arkarachai Fungtammasan¹⁸, Erik Garrison^{11,26} Patrick G.S. Grady¹⁰, Tina A. Graves-Lindsay²⁷, Ira M. Hall²⁸, Nancy F. Hansen²⁹, Gabrielle A. Hartley¹⁰ Marina Haukness¹¹, Kerstin Howe¹⁹, Michael W. Hunkapiller³⁰, Chirag Jain^{1,31}, Miten Jain¹¹, Erich D. Jarvis^{23,24}. Peter Kerpedijev³². Melanje Kirsche⁹. Mikhail Kolmogorov³³. Jonas Korlach³⁰. Milinn Kremitzki²⁷ Heng Li^{16,17}, Valerie V. Maduro³⁴, Tobias Marschall³⁵, Ann M. McCartney¹, Jennifer McDaniel³⁶, Danny E. Miller^{4,37}, James C, Mullikin^{14,29}, Eugene W, Myers³⁸, Nathan D, Olson³⁶, Benedict Paten¹¹, Paul Peluso³⁰, Pavel A. Pevzner³³, David Porubsky⁴, Tamara Potapova¹³, Evgeny I. Rogaev^{6,7,39,40}, Jeffrey A. Rosenfeld⁴¹ Steven L. Salzberg^{9,42}, Valerie A. Schneider⁴³, Fritz J. Sedlazeck⁴⁴, Kishwar Shafin¹¹, Colin J. Shew²⁰, Alaina Shumate⁴², Yumi Sims¹⁹, Arian F. A. Smit⁴⁵, Daniela C. Soto²⁰, Ivan Sović^{30,48}, Jessica M. Storer⁴⁵ Aaron Streets^{5,47}, Beth A. Sullivan⁴⁸, Françoise Thibaud-Nissen⁴³, James Torrance¹⁹, Justin Wagner³⁶, Brian P. Walenz¹, Aaron Wenger³⁰, Jonathan M. D. Wood¹⁹, Chunlin Xiao⁴³, Stephanie M. Yan⁴⁹, Alice C. Young¹⁴, Samantha Zarate⁹, Urvashi Surti⁵⁰, Rajiv C. McCoy⁴⁹, Megan Y. Dennis²⁰, Ivan A. Alexandrov^{3,7,51} Jennifer L. Gerton¹³, Rachel J. O'Neill¹⁰, Winston Timp^{8,42}, Justin M. Zook³⁶, Michael C. Schatz^{9,49}, Evan E.

"PacBio's recent HiFi circular consensus sequencing...has resulted in unprecedented assembly accuracy..." 3

- 1. https://doi.org/10.1038/s41587-021-01049-5
- 2. https://doi.org/10.1101/2020.11.13.380741
- 3. https://doi.org/10.1101/2021.05.26.445798

Sequencing is multi-billion-dollar opportunity across diverse set of markets



2021 was focused on building the foundation for growth

1. EXPAND COMMERCIAL REACH



2. DRIVE PRODUCT DEVELOPMENT PIPELINE



3. MARKET LEADERSHIP IN WHOLE GENOME CLINICAL SEQUENCING



Expanded leadership team focused on driving growth



Christian Henry
PRESIDENT + CHIEF EXECUTIVE OFFICER
25+ years | Prior CCO at Illumina



Susan Kim
CHIEF FINANCIAL OFFICER
20+ years | Prior Tech CFO



Mark Van Oene
CHIEF OPERATING OFFICER
20+ years | Prior CCO at Illumina



Peter Fromen

CHIEF COMMERCIAL OFFICER

20+ years | Prior head of PopGen at Illumina



Catherine Ball
SVP, RESEARCH
25+ years | Prior CSO at AncestryDNA



Richard Shen

SVP, RESEARCH & DEVELOPMENT

25+ years | Prior President at Omniome



Kelvin LiuVP, TECHNOLOGY DEVELOPMENT

15+ years | Prior founder/CEO of Circulomics



Neil Ward

GM, EUROPE, MIDDLE EAST, AND AFRICA

20+ years | Prior Europe Sales Director at Illumina



Jason Kang
GM, ASIA PACIFIC
20+ years | Prior sales leadership at Illumina



Lara Toerien

GM, AMERICAS

20+ years | Prior sales leadership at Illumina

Significant commercial investment enables us to reach more customers

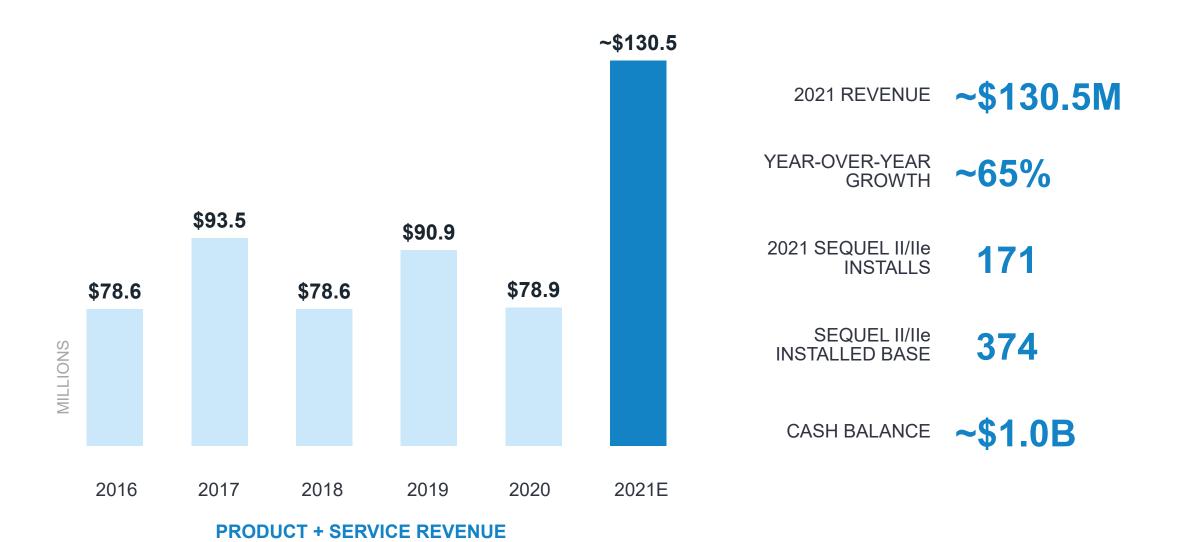


6
Global locations

>200
Commercial employees
66 service + support staff
48 quota-carrying reps

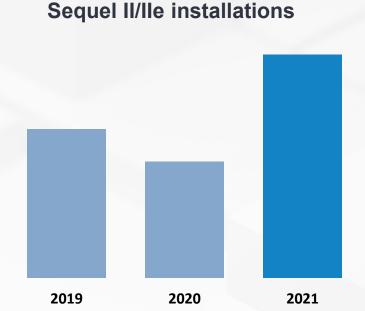
>40
Countries

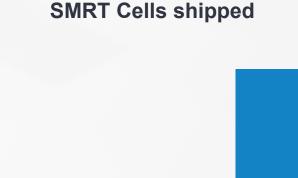
Commercial focus and new products drove breakout performance in 2021



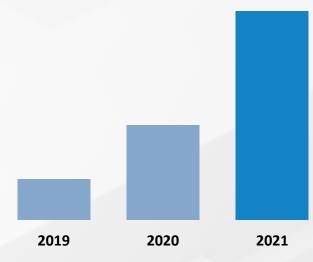


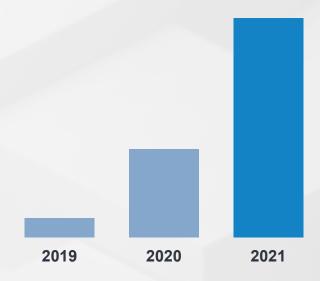
Record performance for Sequel II/IIe platform in 2021











84%

Growth in installed base 220%

Growth in Sequel II/e SMRT Cells 250%

Growth in data generated from Sequel II/IIe

Most productive year in PacBio history



Collaboration to accelerate long-read WGS into routine clinical care

PacBio HUMAN GENOMICS

~25% more HiFi data 3× lower DNA input



Nanobind technology extends our capabilities in extraction + sample prep

OMNIOME

Sequencing by binding adds differentiated short read technology

PacBio MICROBIAL GENOMICS

Launched HiFiViral as PacBio's first kitted solution



Q2

Q3

Q3

Q4

~50 Employees dedicated to research + technology development

~280

Employees dedicated to product development

~80

Employees dedicated to operations/manufacturing

HiFiViral delivering complete SARS-CoV-2 genomes for all circulating variants



> 16K Samples shipped

20 Global customers













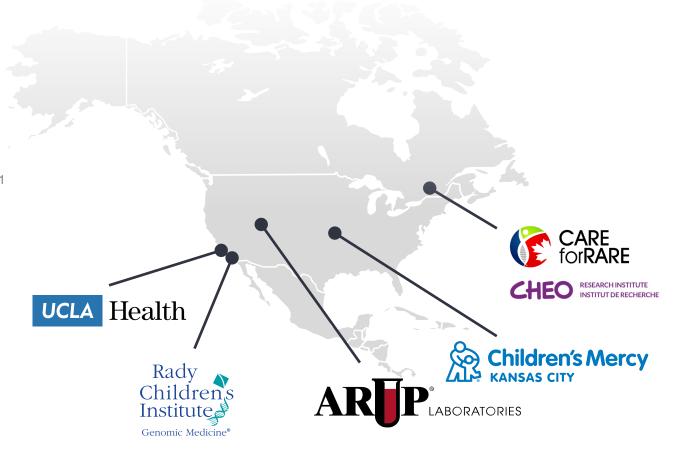
Collaborations demonstrating increased potential of HiFi in clinical research

"Incorporating structural variants (SVs) from genome sequencing (GS) added up to 13% of new diagnoses in previously unsolved cases. HiFi-GS yielded increased discovery rate with >4-fold more rare coding SVs than short-read GS."

- Genomic Answers for Kids/ Children's Mercy Kansas City¹

"Long-read sequencing can identify numerous variants, both small and structural that are not readily detectable by short reads. The number of small coding variants in disease genes missed by short reads is of particular concern and may impact the overall diagnostic rate for critically ill children."

- Matthew Bainbridge, Rady Children's Hospital²



^{1.} doi:10.1101/2021.10.07.21264628

^{2.} https://eventpilotadmin.com/web/page.php?page=IntHtml&project=ASHG21&id=1989

2022 strategic priorities build on our 2021 successes



Execution matters

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



Progress product pipeline

Demonstrate SBB as differentiated short-read offering + drive future HiFi platform development



Delight our customers

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

Significant improvements in efficiency + usability to Sequel IIe in 1H 2022





SMRTbell prep kit 3.0

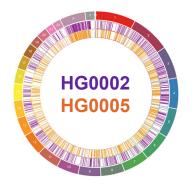
Template binding kits 3.1/3.2

40% less DNA required

>60% reduction in hands-on time

>30% fewer tubes required





SMRT Link v11.0

5-methylcytosine detection

Methylation calling on instrument

Gene editing QC workflow

High-throughput sample setup

Expanding ecosystem enables customers to achieve more with Sequel II/IIe

More samples

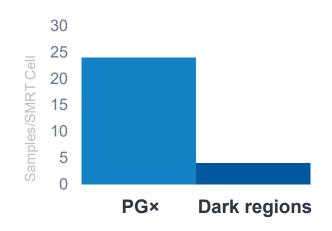
Community panels

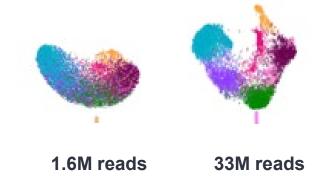
More reads

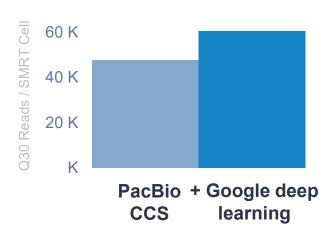
Single-cell Iso-Seq

More data

Deep learning technologies













Partnership to help the 75%¹ of families with unexplained rare disease cases after short-read sequencing



"This study represents our continued commitment to the 100,000 Genomes Project participants, and also to our quest to seek out the benefits of new disruptive technologies."

Parker Moss

Chief Ecosystem & Partnership Officer, Genomics England



Berry Genomics partnership to purchase 50+ desktop systems at launch



Announcing development of first desktop HiFi-based sequencing platform



Makes HiFi sequencing more accessible to new labs for targeted assays and panels



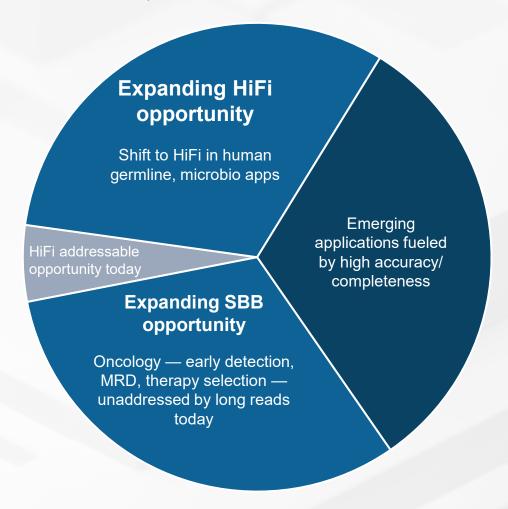
Initial focus on **Chinese distributed market** through NMPA but will be **commercialized globally**





Short-read sequencing is a multi-billion-dollar revenue opportunity for PacBio

SEQUENCING OPPORTUNITY





A more accurate short-read technology gives customers **flexibility**



A more diversified product portfolio drives PacBio scale



Ultimately allows PacBio to participate in highest growth short-read **clinical** applications like oncology

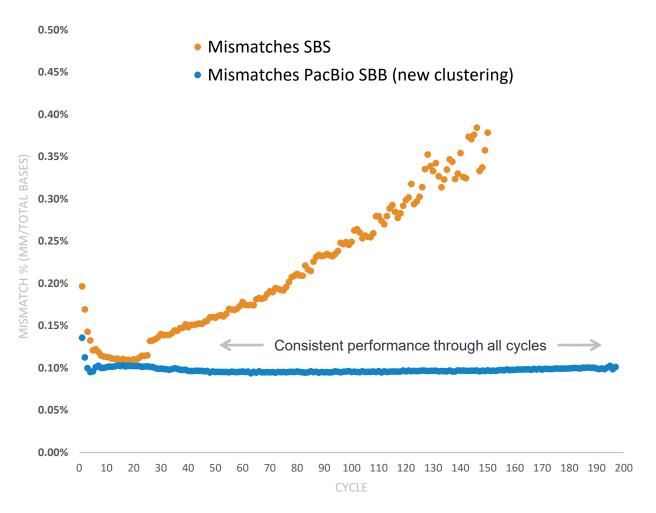
Implemented novel clustering method to vastly improve SBB performance





Even lower error rate than original emulsion PCR method

SBB demonstrates greater accuracy + consistency vs current technologies





Even lower error rate than original emulsion PCR method

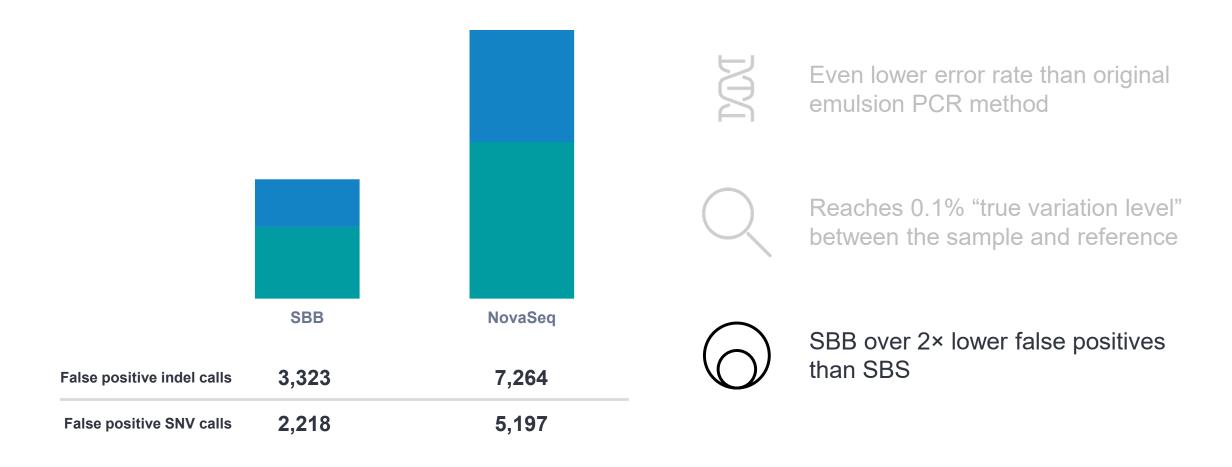


Reaches 0.1% "true variation level" between the sample and reference

Mismatch rate by cycle

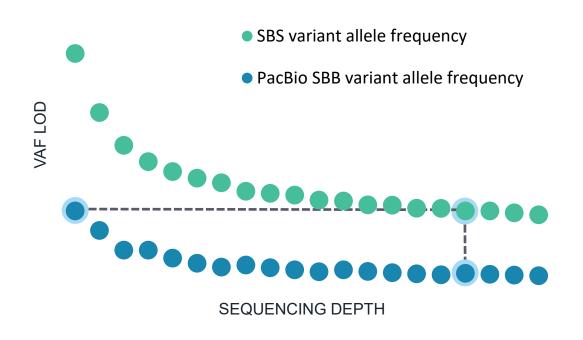


Higher accuracy translates to lower false positive base calls





SBB accuracy will help realize the full potential of liquid biopsy



Higher sensitivity



Detect even lower VAF



Maximize use cases (MRD, etc.)

Lower depth



Lower compute + storage needs



Reduce complexity (UMIs, etc.)



Customer collaborations accelerate SBB entry into liquid biopsy applications







Key takeaways



Transformation in 2021 set the foundation for growth



Only company with both long- and short-read sequencing technologies¹



Our sequencing tech continues to lead the industry in both accuracy + completeness



Multiple platform development programs underway, which are expected to drive growth

PacBio

1. Based on company estimates

PacBio

We create some of the world's most advanced sequencing technologies.

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