UNITED STATES PATENT AND TRADEMARK OFFICE

BEFORE THE PATENT TRIAL AND APPEAL BOARD

INTEGRATED DNA TECHNOLOGIES, INC.,
Petitioner,

v.

TECAN GENOMICS, INC.,
Patent Owner

Case No. IPR2025-00016
U.S. Patent No. 10,876,108

PATENT OWNER'S UPDATED EXHIBIT LIST

Ex.	Short Name	Description
2001	Evans 2009	Mark F. Evans, *The Polymerase Chain Reaction and Pathology Practice*, 15 DIAGN. HISTOPATHOL. 344 (2009)
2002	N/A	*The Cost of Sequencing a Human Genome*, NAT'L INST. OF HEALTH NAT'L HUMAN GENOME RESEARCH INST., https://www.genome.gov/about-genomics/fact-sheets/Sequencing-Human-Genome-cost (last visited July 18, 2025)
2003	Goodwin 2016	Sara Goodwin et al., *Coming of Age: Ten Years of Next-Generation Sequencing Technologies*, 17 NAT. REV. 333 (2016).
2004	Mardis 2017	Elaine R. Mardis, *DNA Sequencing Technologies: 2006-2016*, 12 NAT. PROTOC., 213 (2017).
2005	Fuller 2009	Carl W. Fuller et al., *The Challenges of Sequencing by Synthesis*, 27 NAT. BIOTECHNOL. 1013 (2009).
2006	Marguiles 2005	Marcel Margulies et al., *Genome Sequencing in Open Microfabricated High Density Picoliter Reactors*, 437 NATURE 376 (2005).
2007	Luo 2012	Chengwei Luo et al., *Direct Comparisons of Illumina vs. Roche 454 Sequencing Technologies on the Same Microbial Community DNA Sample*, 7 PLOS ONE 1 (2012).
2008	Hert 2008	Daniel G. Hert et al., *Advantages and Limitations of Next-Generation Sequencing Technologies: A Comparison of Electrophoresis and Non-Electrophoresis Methods*, 29 ELECTROPHORESIS 4618 (2008).
2009	N/A	*Six Years After Acquisition, Roche Quietly Shutters 454*, BioIT World (Oct. 16, 2013), https://www.bio-itworld.com/news/2013/10/16/six-years-after-acquisition-roche-quietly-shutters-454.
2010	Mamanova 2010	Lira Mamanova et al., *Target-Enrichment Strategies for Next-Generation Sequencing*, 7 NAT. METHODS 111 (2010).

Ex.	Short Name	Description
2011	Treangen 2011	Todd J. Treangen et al., *Repetitive DNA and Next-Generation Sequencing: Computational Challenges and Soluitions*, 29 NAT. REV. GENET. 36 (2012).
2012	Brownie 1997	Jannine Brownie et al., *The Elimination of Primer-dimer Accumulation in PCR*, 25 NUCLEIC ACIDS RES. 3235 (1997).
2013	Tewhey 2009	Ryan Tewhey et al., *Microdroplet-based PCR Enrichment for Large-scale Targeted Sequencing*, 27 NAT. BIOTECHNOL. 1025 (2009).
2014	Link 2012	U.S. Patent Application No. 2012/0309002 (published Dec. 6, 2012).
2015	Franck 1998	Sophia M. Franck et al., *Multiplex PCR for Enterotoxigenic, Attaching and Effacing, and Shiga Toxin-Producing Escherichia coli Strains from Calves*, 36 J. CLIN. MICROBIOL. 1795 (1998).
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2017	Fool 2002	University of Hawaii, *The Simple Fool's Guide to PCR* (2002).
2018	Ying 2019	Hua Ying et al., *The Whole-Genome Sequence of the Coral Acropora Millepora*, 1 GENOME BIOL. Genome Biology 1374 (2019).
2019		Office Action Summary, Application No. 13/750,768 (U.S. Patent & Trademark Office July 15, 2015).
2020	Singleton 2011	A. Singleton, *Exome sequencing: a transformative technology*, 10 Lancet Neurol. 942-46 (2011).
2021	Former Bustamante Declaration	Declaration of Carlos D. Bustamente, Ph.D. for Patent Owner's Preliminary Response
2022-2035		*Reserved*

Ex.	Short Name	Description
2036	Sims Deposition 1	Transcript for the June 26, 2025 Deposition of Dr. Peter A. Sims
2037	Sims Deposition 2	Transcript for the June 27, 2025 Deposition of Dr. Peter A. Sims
2038	Bustamante Declaration	Declaration of Carlos D. Bustamente, Ph.D.
2039	Matz Declaration	Declaration of Mikhail V. Matz, Ph.D.
2040	Mullis 1986	K.Mullis et al., *Specific Enzymatic Amplification o fDNA In Vitro: The Polymerase Chain Reaction*, 51 COLDSPRING HARBOR SYMPOSIA QUANTITATIVE BIOL. 1(1986)
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2049	Bower 2010	Neil I. Bower & Ian A. Johnston, *Targeted rapid amplification of cDNA ends (T-RACE)—an improved RACE reaction through degradation of non-target sequences*, 38 NUCLEIC ACIDS RES. 21 (2010)
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2051	Almomani 2011	Rowida Almomani et al., *Experiences with array-based sequence capture; toward clinical applications*, 19 EURO. J. HUMAN GENETICS (2011)
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2053	Quail 2008	Michael A. Quail et al., *A large genome center's improvements to the Illumina sequencing system*, 5 NAT. METHODS 12 (2008)
2054	Kozarewa 2009	Iwanka Kozarewa et al., *Amplification-free Illumina sequencing-library preparation facilitates improved mapping and assembly of (G+C)-biased genomes*, 6 NAT. METHODS 4 (2009)
2055	Syvänen 2001	Ann-Christine Syvänen, *Accessing Genetic Variation: Genotyping Single Nucleotide Polymorphisms*, 2 NAT. REV. (2001)
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2057	Glenn 2011	Travis C. Glenn, *Field guide to next-generation DNA sequencers*, MOLECULAR ECOLOGY RES. (2011)
2058	Zhang 2021	Lu Zhang, *Advances in Metagenomics and Its Application in Environmental Microorganisms*, 12 FRONTIERS IN MICROBIOL. (2021)

Ex.	Short Name	Description
2059	2008 GS-Flx Operating Manual	*Genome Sequencer FLX Operator's Manual*, ROCHE (2008)
2060	Niedringhaus 2011	Thomas P. Niedringhaus, *Landscape of Next-Generation Sequencing Technologies*, 83 ANALYTICAL CHEMISTRY (2011)
2061	Thoma 2011	Generation of Stable mRNA Fragments and Translation of N-Truncated Proteins Induced by Antisense Oligodeoxynucleotides
2062	Ernani 2009	Fred Ernani et al., *Bringing Cost and Process Efficiency to Next Generation Sequencing*, DDW (2009)
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2068	Scaglione 2019	Davide Scaglione et al., *Single primer enrichment technology as a tool for massive genotyping: a benchmark on black poplar and maize*, 124 ANNALS OF BOTANY (2019)
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2070	Knierim 2011	Ellen Knierim, *Systematic comparison of three methods for fragmentation of long-range PCR products for next generation sequencing*, 6 PLoS ONE 11 (2011)
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2072	Whiley 2005	David M. Whiley et al., *Sequence variation in primer targets affects the accuracy of viral quantitative PCR*, 34 J. CLINICAL VIROLOGY (2005)
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2076	JPMorgan Report	Tycho W. Peterson et al., *Next Gen Sequencing Survey*, J.P. MORGAN (2010)
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2079	Additional File 3	Additional File 3 to EX1006
2080	Lewin 1994	Benjamin Lewin, *The eukaryotic gene: conserved exons and unique introns*, in GENES V 677 (1994)
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2082	Nugen Nature	*Accelerating Genomics with Advanced Sample Preparation Solutions*, NATURE RESEARCH CUSTOM MEDIA

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2083	Scolnick 2015	Jonathan A. Scolnick et al., *An Efficient Method for Identifying Gene Fusions by Targeted RNA Sequencing from Fresh Frozen and FFPE Samples*, PLoS One (2015)
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2085	Allegro V2	*Allegro® Targeted Genotyping V2 Product Sheet*, Tecan Genomics, Inc. (2019)
2086	Feng 2016	Weixing Feng et al., *Improving Alignment Accuracy on Homopolymer Regions for Semiconductor-Based Sequencing Technologies*, 17 BMC Genomics 521 (2016)
2087	Nakamura 2011	Kensuke Nakamura et al., *Sequence-Specific Error Profile of Illumina Sequencers*, 39 Nucleic Acids Res. 90 (2011)
2088	Sambrook 2006	Joseph Sambrook et al., *Fragmentation of DNA by Nebulization*, Cold Spring Harb. Protoc. (2006)
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2090	Han 2006	Sangjo Han et al., *AtRTPrimer: Database for Arabidopis Genome-Wide Homogeneous and Specific RT-PCR Primer-Pairs*, 7 BMC Bioinformatics 179 (2006).
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2100	Infinitives	Evin de Leon Sanchez, *Infinitives*, SAN JOSE STATE UNIVERSITY WRITING CENTER, https://www.sjsu.edu/writingcenter/docs/handouts/Infinitives.pdf (last visited July 18, 2025)
2101	Meyer Comparison	Redline document comparing Exhibits 2103 and 2039
2102	Matz Email 1	Email thread between counsel and Petitioner's counsel
2103	Meyer Declaration	Declaration of Eli Meyer, Ph.D.

Ex.	Short Name	Description
2104	Matz Email 2	Email thread between counsel and Petitioner's counsel
2105	Matz Email 3	Email thread between counsel and Petitioner's counsel
2106	Matz Email 4	Email thread between counsel and Petitioner's counsel
2107-2199		*Reserved*
2200		Privileged & Confidential Correspondence (submitted *in camera*)
2201		Privileged & Confidential Correspondence (submitted *in camera*)
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2283		Privileged & Confidential Correspondence (submitted *in camera*)
2284		Privileged & Confidential Correspondence (submitted *in camera*)
2285		Privileged & Confidential Declaration and Timeline Submitted Under Seal for *In Camera* Review

Dated: September 26, 2025

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The undersigned hereby certifies that a copy of the foregoing **PATENT OWNER'S UPDATED EXHIBIT LIST** was served on September 26, 2025 via email to Petitioner by serving the correspondence addresses of record as follows:

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