

Product Analysis Certificate

Product: CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro (Packaged)
Cat. #: BCXP10M3VP-V



CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro

Shipment Contents: CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro (Packaged)
Store at -80°C

Description:

Cellecta's CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro is a pooled barcode library that enables the tracking of individual clones derived from a population of cells using either RNA-Seq or PCR/NGS from genomic DNA. The library contains 10 million lentiviral constructs, each expressing a unique barcode sequence. When a cell population is transduced with this barcode library, the barcodes integrate into the genomic DNA of the cells. The result is a founder population where almost every cell has a different DNA-sequenceable barcode that is both integrated into its DNA and expressed on an RNA transcript. Since the barcode is stably integrated, it is passed onto any cell progeny when genomic DNA is replicated. This feature enables identification of all the progeny derived from each individual cell.

When used in conjunction with single-cell RNA-Seq, the expressed barcode may be used to identify expression profiles and activated genes in different cells so that distinct clonal populations of cells from a single progenitor, as well as sub-populations of cells with distinct pathway activations or expression profiles, can be readily identified.

The CloneTracker XP Barcode Library is constructed in a third-generation lentiviral vector that expresses the mVenus fluorescence marker and Puro resistance gene under an EF1a core (EFS) promoter. The specially-designed, optimized barcodes facilitate Next-Gen Sequencing (NGS) data analysis and barcode identification. Using the Illumina NextSeq or HiSeq NGS platform, barcode sequences can be identified either by RNA-Seq or PCR/NGS from genomic DNA and converted to lists with enumerated data.

The titer of packaged libraries provided by Cellecta is functionally determined by transduction of 293T cells at low MOI and FACS of YFP-positive cells.

Biosafety Level: BSL-2
Storage: -80°C
Shelf Life: 1 year from date of receipt
Shipping Conditions: Dry Ice

Product Information (Cellecta Website):

User Manual: <https://www.cellecta.com/resources/product-manuals-and-certificates/>
Vector Info (Sequence, cassette, etc): Please contact Cellecta Technical Support at tech@cellecta.com.

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Contents:

Part #	Description
BCXP10M3VP-V	CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro (packaged) in pScribe5-EFS-mVenus-2A-Puro-bc-3LTR <u>Packaged, >5 × 10⁸ TU:</u> 5.28 × 10⁸ TU , 8.51 × 10 ⁸ TU/ml (620 µl total: 50 µl × 12 vials, 5 µl × 4 vials) Lot# 180815054; Store at -80°C

Library Barcode Sequence Files (sent with PAC):

30nt Barcodes (BC30): Collecta-SEQ-CloneTracker-XP-Barcode-Libraries-100K_x_BC30.xlsx

14nt Barcodes (BC14): Collecta-SEQ-CloneTracker-XP-Barcode-Libraries-100_x_BC14.xlsx

NOTE: The **CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro** uses all 100 of the 14nt barcodes (BC14) in the file above.

Quality Control

Individual Clone Sequencing Data

	Library: CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro
Plasmid Lot #:	180815055
Library Complexity (number of clones):	250 × 10 ⁶
Number of good clones picked:	21
Correct Structure:	>95%
Number of clones with at least one mutation, deletion, or insertion:	0
Mutation / Deletion / Insertion Rate:	0.00%
Estimated % of Inserts without any mutations, deletions, or insertions in <u>barcode</u> portion (Based on Clone QC):	>95%
Estimated % of Inserts without any mutations, deletions, or insertions in <u>barcode</u> portion (Based on NGS QC):	>95%

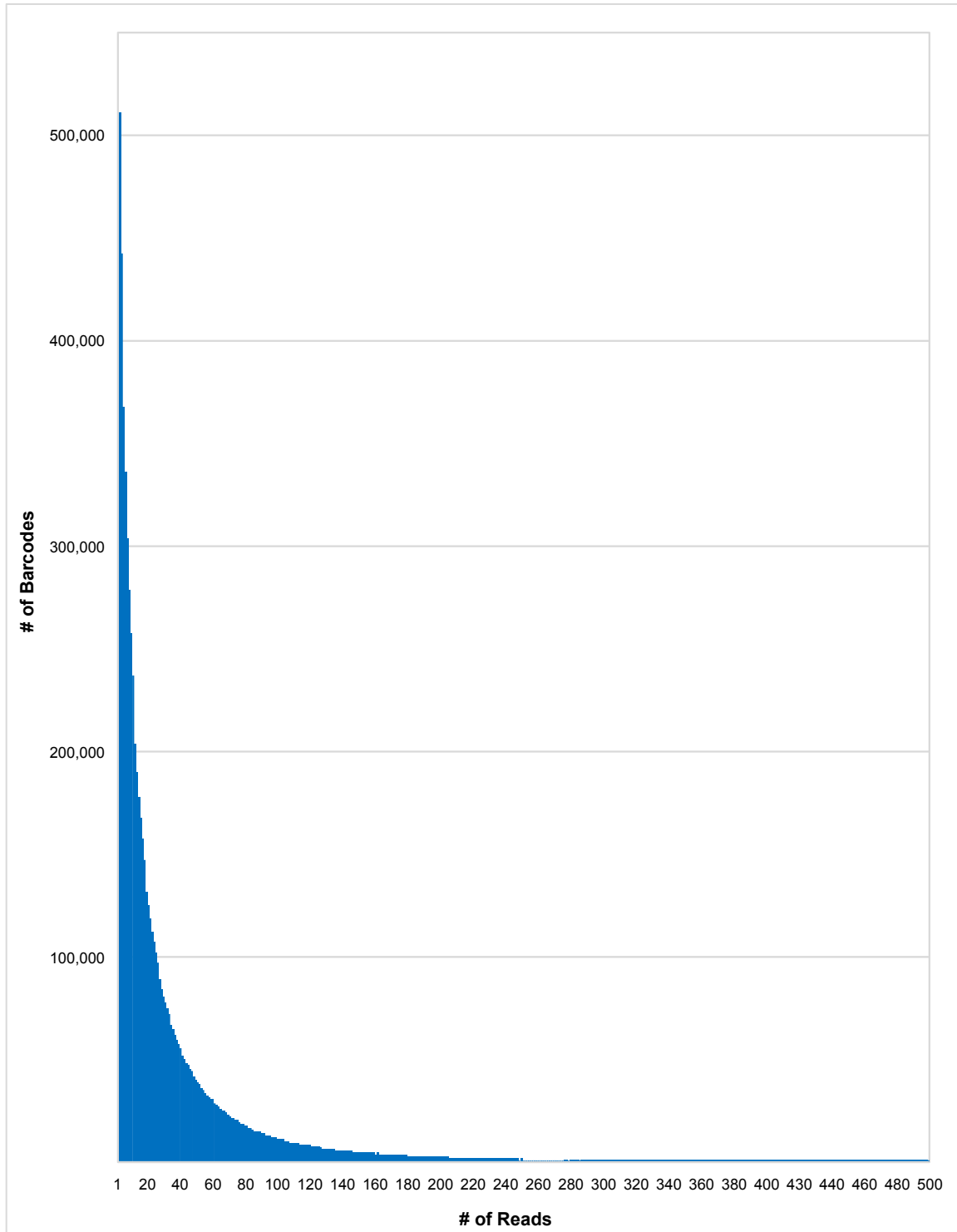
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Next-Gen Sequencing (NGS) Data

Histogram, Barcode Representation (303M total reads)
Number of barcodes (y) with x # of reads



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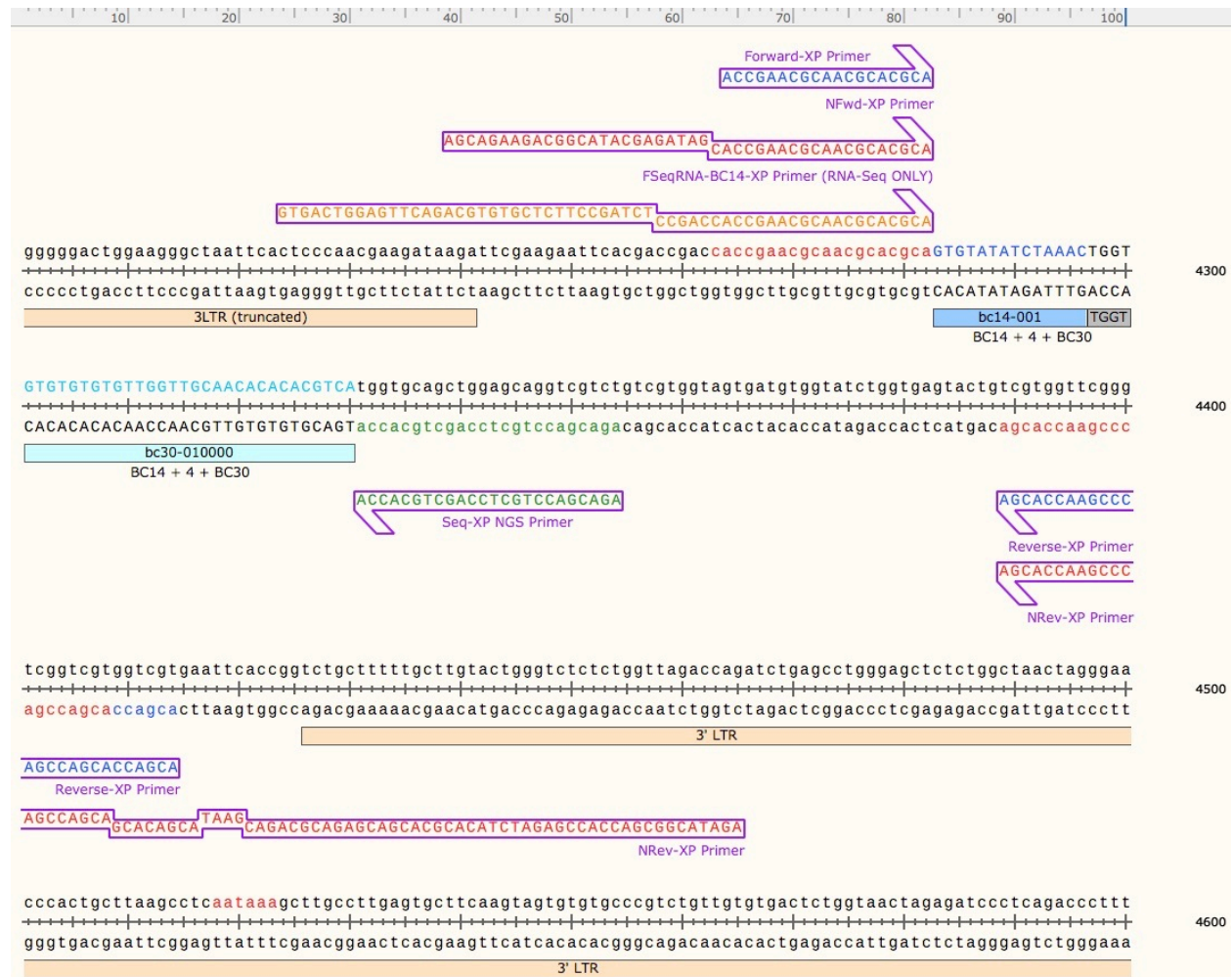
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NGS Barcode Cassette Diagram (in pScribe5 Vector)

Example NGS cassette diagram for the bc14-001_bc30-010000 construct in pScribe5-EFS-Venus-2A-Puro-bc-3LTR, from the CloneTracker XP™ 10M Barcode-3' Library with Venus-Puro. RNA-Seq primer **FSeqRNA-BC14-XP** is also shown and is required in order to read CloneTracker XP barcode sequences on most single-cell Drop-Seq systems.

Expressed barcode structure (sense): BC14-TGGT-BC30-3LTR-Poly(A)signal



Primer*	Sequence	Purpose
Forward-XP Primer	ACCGAACGCAACGCACGCA	1st Round PCR
Reverse-XP Primer	ACGACCAGCACCACCGAACCCGCA	1st Round PCR
P7-NFwd-XP Primer	AGCAGAAGACGGCATAACGAGATAGCACCAGCAACGCACGCA	2nd Round PCR
P5-NRev-XP Primer	AGATACGGCGACCACCGAGATCTACACGACGACGAGACGAGACGAAATACGACACGACGACC GACCCGAACCAGCA	2nd Round PCR
FSeqRNA-BC14-XP Primer	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTCCGACCACCGAACGCAACGCACGCA	scRNA-Seq

* Primers and reagents for multiplexing up to 12 samples also available in Collecta's NGS Prep Kit (See Page 4)

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Diagram of PCR Amplicon

Example Second-Round (Nested) PCR amplicon diagram for a barcode-3' library construct in pScribe5, from the CloneTracker XP™ 10M Barcode-3' Library. The **P7-NFwd-XP Primer** contains the minimal sequence of P7 (5' end, sequence in orange) and **P5-NRev-XP Primer** contains the minimal sequence of P5 (3' end, sequence in orange) required for compatibility with all Illumina flow cells.

- Size of Second-Round (Nested) PCR amplicon: **227 bp**
- Starting library concentration (see NGS Sample Purification step): **10 nM**



NGS Primer*	Sequence	Purpose
Seq-XP NGS Primer	AGACGACCTGCTCCAGCTGCACCA	NGS of barcode (reverse)

* Primers and reagents for multiplexing up to 12 samples also available in Collecta's NGS Prep Kit (See below)

NGS Prep of Screening Samples

Collecta provides the following options for preparing and sequencing your screening samples. For pricing, please inquire.

Cat.#	Description	Quantity
LNGS-300	NGS Prep Kit for Barcode Libraries in pScribe (CloneTracker XP)	Kit for 6-48 samples (48 preps of 50ug each, 12 multiplex)
LNGS-300-SP	Supplementary Primer Set for LNGS-300 (12 Additional Index Primers)	48 Reactions (12 Index Primers x 4 reactions each)
CANA-SQD	NGS of DNA from Genetic Screen	Service (per DNA sample)

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Terms and Conditions

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