

Illumina Adapter Sequences

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TruSight Amplicon Panels

Includes TruSight Myeloid Sequencing Panel and TruSight Tumor 26

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| A701 | ATCACGAC |
| A702 | ACAGTGGT |
| A703 | CAGATCCA |
| A704 | ACAAACGG |
| A705 | ACCCAGCA |
| A706 | AACCCCTC |
| A707 | CCCAACCT |
| A708 | CACCACAC |
| A709 | GAAACCCA |
| A710 | TGTGACCA |
| A711 | AGGGTCAA |
| A712 | AGGAGTGG |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| A501 | TGAACCTT | AAGGTTCA |
| A502 | TGCTAAGT | ACTTAGCA |
| A503 | TGTTCTCT | AGAGAACA |
| A504 | TAAGACAC | GTGTCTTA |
| A505 | CTAATCGA | TCGATTAG |
| A506 | CTAGAACA | TGTTCTAG |
| A507 | TAAGTTCC | GGAACCTA |
| A508 | TAGACCTA | TAGGTCTA |

TruSight Cardio

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| N701 | TAAGGCGA |
| N702 | CGTACTAG |
| N703 | AGGCAGAA |
| N704 | TCCTGAGC |
| N705 | GGACTCCT |
| N706 | TAGGCATG |
| N707 | CTCTCTAC |
| N708 | CAGAGAGG |
| N709 | GCTACGCT |
| N710 | CGAGGCTG |
| N711 | AAGAGGCA |
| N712 | GTAGAGGA |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| A505 | GTAAGGAG | CTCCTTAC |

TruSight One

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| N701 | TAAGGCGA |
| N702 | CGTACTAG |
| N703 | AGGCAGAA |
| N704 | TCCTGAGC |

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| N705 | GGACTCCT |
| N706 | TAGGCATG |
| N707 | CTCTCTAC |
| N708 | CAGAGAGG |
| N709 | GCTACGCT |
| N710 | CGAGGCTG |
| N711 | AAGAGGCA |
| N712 | GTAGAGGA |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| A502 | CTCTCTAT | ATAGAGAG |
| A503 | TATCCTCT | AGAGGATA |
| A504 | AGAGTAGA | TCTACTCT |
| A505 | GTAAGGAG | CTCCTTAC |

TruSight Rapid Capture

Includes TruSight Autism, TruSight Cancer, and TruSight Inherited Disease

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| N701 | TAAGGCGA |
| N702 | CGTACTAG |
| N703 | AGGCAGAA |
| N704 | TCCTGAGC |
| N705 | GGACTCCT |
| N706 | TAGGCATG |
| N707 | CTCTCTAC |

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| N708 | CAGAGAGG |
| N709 | GCTACGCT |
| N710 | CGAGGCTG |
| N711 | AAGAGGCA |
| N712 | GTAGAGGA |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| E502 | CTCTCTAT | ATAGAGAG |
| E505 | GTAAGGAG | CTCCTTAC |
| E506 | ACTGCATA | TATGCAGT |
| E517 | GCGTAAGA | TCTTACGC |

TruSight Tumor 15

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| R701 | ATCACG |
| R702 | CGATGT |
| R703 | TTAGGC |
| R704 | TGACCA |
| R705 | ACAGTG |
| R706 | GCCAAT |
| R707 | CAGATC |
| R708 | ACTTGA |
| R709 | GATCAG |
| R711 | GGCTAC |
| R712 | CTTGTA |

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| R725 | ACTGAT |
| R726 | ATGAGC |
| R727 | ATTCCT |
| R728 | CAAAAG |
| R729 | CAACTA |
| R730 | CACCGG |
| R731 | CACGAT |
| R732 | CACTCA |
| R733 | CAGGCG |
| R734 | CATGGC |
| R735 | CATTTT |
| R736 | CCAACA |
| R749 | GATGCT |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| A501 | TGAACCTT | AAGGTTCA |
| A502 | TGCTAAGT | ACTTAGCA |

Illumina Nextera Library Prep Kits

Includes Nextera DNA, Nextera XT, Nextera Enrichment (**obsolete**), and Nextera Rapid Capture

Nextera Transposase Adapters

(Used for Nextera tagmentation)

Read 1

5' TCGTCGGCAGCGTCAGATGTGTATAAGAGACAG

Read 2

5' GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAG

Nextera Index Kit – PCR Primers

Index 1 Read

5' CAAGCAGAAGACGGCATACGAGAT [i7] GTCTCGTGGGCTCGG

Index 2 Read

5' AATGATACGGCGACCACCGAGATCTACAC [i5] TCGTCGGCAGCGTC

Nextera Index Kit - Index 1 (i7) Adapters

| i7 Bases in Adapter | i7 Index Name | i7 Bases for Sample Sheet |
|---------------------|---------------|---------------------------|
| TCGCCTTA | N701 | TAAGGCGA |
| CTAGTACG | N702 | CGTACTAG |
| TTCTGCCT | N703 | AGGCAGAA |
| GCTCAGGA | N704 | TCCTGAGC |
| AGGAGTCC | N705 | GGACTCCT |
| CATGCCTA | N706 | TAGGCATG |
| GTAGAGAG | N707 | CTCTCTAC |
| CCTCTCTG | N708 | CAGAGAGG |
| AGCGTAGC | N709 | GCTACGCT |
| CAGCCTCG | N710 | CGAGGCTG |
| TGCCTCTT | N711 | AAGAGGCA |
| TCCTCTAC | N712 | GTAGAGGA |

Nextera Index Kit - Index 2 (i5) Adapters

The i5 index names vary for different Nextera products as follows:

- N50x—Nextera DNA
- S50x—Nextera XT
- E50x—Nextera Enrichment and Nextera Rapid Capture

| i5 Bases in Adapter | i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------------|---------------|--|--|
| TAGATCGC | [N/S/E]501 | TAGATCGC | GCGATCTA |
| CTCTCTAT | [N/S/E]502 | CTCTCTAT | ATAGAGAG |
| TATCCTCT | [N/S/E]503 | TATCCTCT | AGAGGATA |
| AGAGTAGA | [N/S/E]504 | AGAGTAGA | TCTACTCT |
| GTAAGGAG | [N/S/E]505 | GTAAGGAG | CTCCTTAC |
| ACTGCATA | [N/S/E]506 | ACTGCATA | TATGCAGT |
| AAGGAGTA | [N/S/E]507 | AAGGAGTA | TACTCCTT |
| CTAAGCCT | [N/S/E]508 | CTAAGCCT | AGGCTTAG |
| GCGTAAGA | [N/S/E]517 | GCGTAAGA | TCTTACGC |

Nextera XT Index Kit v2 - Index 1 (i7) Adapters

| i7 Bases in Adapter | i7 Index Name | i7 Bases for Entry on Sample Sheet |
|---------------------|---------------|------------------------------------|
| TCGCCTTA | N701 | TAAGGCGA |
| CTAGTACG | N702 | CGTACTAG |
| TTCTGCCT | N703 | AGGCAGAA |
| GCTCAGGA | N704 | TCCTGAGC |
| AGGAGTCC | N705 | GGACTCCT |
| CATGCCTA | N706 | TAGGCATG |
| GTAGAGAG | N707 | CTCTCTAC |
| CAGCCTCG | N710 | CGAGGCTG |
| TGCCTCTT | N711 | AAGAGGCA |
| TCCTCTAC | N712 | GTAGAGGA |
| TCATGAGC | N714 | GCTCATGA |

| i7 Bases in Adapter | i7 Index Name | i7 Bases for Entry on Sample Sheet |
|---------------------|---------------|------------------------------------|
| CCTGAGAT | N715 | ATCTCAGG |
| TAGCGAGT | N716 | ACTCGCTA |
| GTAGCTCC | N718 | GGAGCTAC |
| TACTACGC | N719 | GCGTAGTA |
| AGGCTCCG | N720 | CGGAGCCT |
| GCAGCGTA | N721 | TACGCTGC |
| CTGCGCAT | N722 | ATGCGCAG |
| GAGCGCTA | N723 | TAGCGCTC |
| CGCTCAGT | N724 | ACTGAGCG |
| GTCTTAGG | N726 | CCTAAGAC |
| ACTGATCG | N727 | CGATCAGT |
| TAGCTGCA | N728 | TGCAGCTA |
| GACGTCGA | N729 | TCGACGTC |

Nextera XT Index Kit v2 - Index 2 (i5) Adapters

| i5 Bases in Adapter | i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------------|---------------|--|--|
| CTCTCTAT | S502 | CTCTCTAT | ATAGAGAG |
| TATCCTCT | S503 | TATCCTCT | AGAGGATA |
| GTAAGGAG | S505 | GTAAGGAG | CTCCTTAC |
| ACTGCATA | S506 | ACTGCATA | TATGCAGT |
| AAGGAGTA | S507 | AAGGAGTA | TACTCCTT |
| CTAAGCCT | S508 | CTAAGCCT | AGGCTTAG |
| CGTCTAAT | S510 | CGTCTAAT | ATTAGACG |
| TCTCTCCG | S511 | TCTCTCCG | CGGAGAGA |
| TCGACTAG | S513 | TCGACTAG | CTAGTCGA |
| TTCTAGCT | S515 | TTCTAGCT | AGCTAGAA |
| CCTAGAGT | S516 | CCTAGAGT | ACTCTAGG |

| i5 Bases in Adapter | i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------------|---------------|--|--|
| GCGTAAGA | S517 | GCGTAAGA | TCTTACGC |
| CTATTAAG | S518 | CTATTAAG | CTTAATAG |
| AAGGCTAT | S520 | AAGGCTAT | ATAGCCTT |
| GAGCCTTA | S521 | GAGCCTTA | TAAGGCTC |
| TTATGCGA | S522 | TTATGCGA | TCGCATAA |

TruSeq Amplicon Kits

TruSeq Custom Amplicon 1.5, TruSeq Amplicon Cancer Panel, and TruSeq Custom Amplicon Low Input

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| A701 | ATCACGAC |
| A702 | ACAGTGGT |
| A703 | CAGATCCA |
| A704 | ACAAACGG |
| A705 | ACCCAGCA |
| A706 | AACCCCTC |
| A707 | CCCAACCT |
| A708 | CACCACAC |
| A709 | GAAACCCA |
| A710 | TGTGACCA |
| A711 | AGGGTCAA |
| A712 | AGGAGTGG |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| A501 | TGAACCTT | AAGGTTCA |
| A502 | TGCTAAGT | ACTTAGCA |
| A503 | TGTTCTCT | AGAGAACA |
| A504 | TAAGACAC | GTGTCTTA |
| A505 | CTAATCGA | TCGATTAG |
| A506 | CTAGAACA | TGTTCTAG |
| A507 | TAAGTTCC | GGAACCTA |
| A508 | TAGACCTA | TAGGTCTA |

TruSeq HT Kits

Includes TruSeq DNA PCR-Free HT, TruSeq Nano HT, TruSeq Stranded mRNA HT, and TruSeq Total RNA HT

D501–D508 Adapters

AATGATACGGCGACCACCGAGATCTACAC [i5] ACACTCTTTCCCTACACGACGCTCTTCCGATCT

D701–D712 Adapters

GATCGGAAGAGCACACGTCTGAACTCCAGTCAC [i7] ATCTCGTATGCCGTCTTCTGCTTG

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|
| D701 | ATTACTCG |
| D702 | TCCGGAGA |
| D703 | CGCTCATT |
| D704 | GAGATTCC |
| D705 | ATTCAGAA |
| D706 | GAATTCGT |
| D707 | CTGAAGCT |
| D708 | TAATGCGC |
| D709 | CGGCTATG |
| D710 | TCCGCGAA |
| D711 | TCTCGCGC |
| D712 | AGCGATAG |

Index 2 (i5) Adapters

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| D501 | TATAGCCT | AGGCTATA |
| D502 | ATAGAGGC | GCCTCTAT |
| D503 | CCTATCCT | AGGATAGG |

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| D504 | GGCTCTGA | TCAGAGCC |
| D505 | AGGCGAAG | CTTCGCCT |
| D506 | TAATCTTA | TAAGATTA |
| D507 | CAGGACGT | ACGTCCTG |
| D508 | GTACTGAC | GTCAGTAC |

TruSeq LT Kits and TruSeq v1/v2 Kits

Includes TruSeq DNA PCR-Free LT, TruSeq Nano DNA LT, TruSeq DNA v1/v2/LT (**obsolete**), TruSeq RNA v1/v2/LT, TruSeq Stranded mRNA LT, TruSeq Stranded Total RNA LT, TruSeq RNA Access, and TruSeq ChIP

Index sequences are 6 bases as underlined. Enter the underlined 6 bases on the sample sheet.

TruSeq Universal Adapter

5' AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT

TruSeq Index Adapters (Index 1–27)

Index numbers 17, 24, and 26 are reserved.

TruSeq Adapter, Index 1

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACATCACGATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 2

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACCGATGTATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 3

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCATTAGGCATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 4

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCATGACCAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 5

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACACAGTGATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 6

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGCCAATATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 7

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACCAGATCATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 8

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACACTTGAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 9

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGATCAGATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 10

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACTAGCTTATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 11

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGGCTACATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 12

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 13

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACAGTCAACAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 14

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACAGTTCCGTATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 15

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACATGTCAGAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 16

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACCCGTCCCGATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 18

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGTCCGCACATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 19

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGTGAAACGATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 20

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGTGGCCTTATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 21

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGTTCGGAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 22

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACCGTACGTAATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 23

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACGAGTGGATATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 25

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACACTGATATATCTCGTATGCCGTCTTCTGCTTG

TruSeq Adapter, Index 27

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCACATTTCCTTTATCTCGTATGCCGTCTTCTGCTTG

TruSeq Synthetic Long-Read DNA

Double-stranded DNA adapter containing long-range PCR primer binding site, sequencing primer binding site, and end marker sequence.

Long Reads Adapter

5' CCGTTCTTCCCTGCCGAACCCTATCTTCGTCGGCAGCGTCAGATGTGTATAAGAGACAGTACGCTTGCAT

TruSeq Small RNA

RNA 5' Adapter (RA5)

5' GUUCAGAGUUCUACAGUCCGACGAUC

RNA 3' Adapter (RA3)

5' TGGAATTCTCGGGTGCCAAGG

Stop Oligo (STP)

5' GAAUCCACCACGUUCCCGUGG

RNA RT Primer (RTP)

5' GCCTTGGCACCCGAGAATTCCA

RNA PCR Primer (RP1)

5' AATGATACGGCGACCACCGAGATCTACACGTTTCAGAGTTCTACAGTCCGA

RNA PCR Index Primers (RPI1–RPI48)

Index sequence is 6 bases as underlined. Enter the underlined 6 bases on the sample sheet. Index sequences are read in the reverse complement in TruSeq small RNA libraries.

RNA PCR Primer, Index 1 (RPI1)

5' CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 2 (RPI2)

5' CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 3 (RPI3)

5' CAAGCAGAAGACGGCATACGAGATGCCTAAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 4 (RPI4)

5' CAAGCAGAAGACGGCATACGAGATTGGTCAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 5 (RPI5)

5' CAAGCAGAAGACGGCATACGAGATCACTGTGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 6 (RPI6)

5' CAAGCAGAAGACGGCATAACGAGATATTGGCGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 7 (RPI7)

5' CAAGCAGAAGACGGCATAACGAGATGATCTGGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 8 (RPI8)

5' CAAGCAGAAGACGGCATAACGAGATTCAAGTGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 9 (RPI9)

5' CAAGCAGAAGACGGCATAACGAGATCTGATCGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 10 (RPI10)

5' CAAGCAGAAGACGGCATAACGAGATAAGCTAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 11 (RPI11)

5' CAAGCAGAAGACGGCATAACGAGATGTAGCCGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 12 (RPI12)

5' CAAGCAGAAGACGGCATAACGAGATTACAAGGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 13 (RPI13)

5' CAAGCAGAAGACGGCATAACGAGATTTGACTGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 14 (RPI14)

5' CAAGCAGAAGACGGCATAACGAGATGGAACTGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 15 (RPI15)

5' CAAGCAGAAGACGGCATAACGAGATTGACATGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 16 (RPI16)

5' CAAGCAGAAGACGGCATAACGAGATGGACGGTGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 17 (RPI17)

5' CAAGCAGAAGACGGCATAACGAGATCTCTACGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 18 (RPI18)

5' CAAGCAGAAGACGGCATAACGAGATGCGGACGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 19 (RPI19)

5' CAAGCAGAAGACGGCATAACGAGATTTTCACGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 20 (RPI20)

5' CAAGCAGAAGACGGCATAACGAGATGGCCACGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 21 (RPI21)

5' CAAGCAGAAGACGGCATAACGAGATCGAAACGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 22 (RPI22)

5' CAAGCAGAAGACGGCATAACGAGATCGTACGTGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 23 (RPI23)

5' CAAGCAGAAGACGGCATAACGAGATCCACTCGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 24 (RPI24)

5' CAAGCAGAAGACGGCATAACGAGATGCTACCGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 25 (RPI25)

5' CAAGCAGAAGACGGCATAACGAGATATCAGTGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 26 (RPI26)

5' CAAGCAGAAGACGGCATAACGAGATGCTCATGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 27 (RPI27)

5' CAAGCAGAAGACGGCATAACGAGATAGGAATGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 28 (RPI28)

5' CAAGCAGAAGACGGCATAACGAGATCTTTTGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 29 (RPI29)

5' CAAGCAGAAGACGGCATAACGAGATTAGTTGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 30 (RPI30)

5' CAAGCAGAAGACGGCATAACGAGATCCGGTGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 31 (RPI31)

5' CAAGCAGAAGACGGCATAACGAGATATCGTGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 32 (RPI32)

5' CAAGCAGAAGACGGCATAACGAGATTGAGTGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 33 (RPI33)

5' CAAGCAGAAGACGGCATAACGAGATCGCCTGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 34 (RPI34)

5' CAAGCAGAAGACGGCATAACGAGATGCCATGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 35 (RPI35)

5' CAAGCAGAAGACGGCATAACGAGATAAAATGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 36 (RPI36)

5' CAAGCAGAAGACGGCATAACGAGATTGTTGGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 37 (RPI37)

5' CAAGCAGAAGACGGCATAACGAGATATTCCGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 38 (RPI38)

5' CAAGCAGAAGACGGCATAACGAGATAGCTAGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 39 (RPI39)

5' CAAGCAGAAGACGGCATAACGAGATGTATAGGTGACTGGAGTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 40 (RPI40)

5' CAAGCAGAAGACGGCATAACGAGATTCTGAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 41 (RPI41)

5' CAAGCAGAAGACGGCATAACGAGATGTCGTCTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 42 (RPI42)

5' CAAGCAGAAGACGGCATAACGAGATCGATTAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 43 (RPI43)

5' CAAGCAGAAGACGGCATAACGAGATGCTGTAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 44 (RPI44)

5' CAAGCAGAAGACGGCATAACGAGATATTATAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 45 (RPI45)

5' CAAGCAGAAGACGGCATAACGAGATGAATGAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 46 (RPI46)

5' CAAGCAGAAGACGGCATAACGAGATTCGGGAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 47 (RPI47)

5' CAAGCAGAAGACGGCATAACGAGATCTTCGAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

RNA PCR Primer, Index 48 (RPI48)

5' CAAGCAGAAGACGGCATAACGAGATTGCCGAGTGACTGGAGTTCCTTGGCACCCGAGAATTCCA

TruSeq Targeted RNA Expression

Index 1 (i7) Adapters

| i7 Index Name | i7 Bases for Sample Sheet | i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|---------------|---------------------------|
| R701 | ATCACG | R725 | ACTGAT |
| R702 | CGATGT | R726 | ATGAGC |
| R703 | TTAGGC | R727 | ATTCCT |
| R704 | TGACCA | R728 | CAAAAG |
| R705 | ACAGTG | R729 | CAACTA |
| R706 | GCCAAT | R730 | CACCGG |
| R707 | CAGATC | R731 | CACGAT |
| R708 | ACTTGA | R732 | CACTCA |
| R709 | GATCAG | R733 | CAGGCG |

| i7 Index Name | i7 Bases for Sample Sheet | i7 Index Name | i7 Bases for Sample Sheet |
|---------------|---------------------------|---------------|---------------------------|
| R710 | TAGCTT | R734 | CATGGC |
| R711 | GGCTAC | R735 | CATTTT |
| R712 | CTTGTA | R736 | CCAACA |
| R713 | AGTCAA | R737 | CGGAAT |
| R714 | AGTTCC | R738 | CTAGCT |
| R715 | ATGTCA | R739 | CTATAC |
| R716 | CCGTCC | R740 | CTCAGA |
| R717 | GTAGAG | R741 | GACGAC |
| R718 | GTCCGC | R742 | TAATCG |
| R719 | GTGAAA | R743 | TACAGC |
| R720 | GTGGCC | R744 | TATAAT |
| R721 | GTTTCG | R745 | TCATTC |
| R722 | CGTACG | R746 | TCCCGA |
| R723 | GAGTGG | R747 | TCGAAG |
| R724 | GGTAGC | R748 | TCGGCA |

Index 2 (i5) Adapter

| i5 Index Name | i5 Bases for Sample Sheet HiSeq 2000/2500 and MiSeq | i5 Bases for Sample Sheet NextSeq and HiSeq 3000/4000 |
|---------------|--|--|
| A501 | TGAACCTT | AAGGTTCA |
| A502 | TGCTAAGT | ACTTAGCA |
| A503 | TGTTCTCT | AGAGAACA |
| A504 | TAAGACAC | GTGTCTTA |
| A505 | CTAATCGA | TCGATTAG |
| A506 | CTAGAACA | TGTTCTAG |
| A507 | TAAGTTCC | GGAACCTA |
| A508 | TAGACCTA | TAGGTCTA |

Appendix

Process Controls for TruSeq Kits

Included in TruSeq DNA PCR-Free, TruSeq Nano DNA, TruSeq RNA (v1/v2/LT/HT), and TruSeq Exome Kits

CTE2 - 150bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGATCCTACGTTCCAAATGCAGCGAGCTCGTATAACCCTTT
AAGAGTTGCTCTTTTTGTTTGGTAAGTTGCAAATCGAAGTTTTAGATTGAGTTCTACGTCGAGCGGCCGCGAT

CTE2 - 250bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGATCCTTATCTGTCAAACCGCTAATGTCCGTTCTAAGAC
CGTCTGGAGAACACTTGCCCATCAGTGCTTTTGAACCTTTTTTTCACAGGTCCCTTCCGATTACACTGAGAAGCTGA
CCACACCTGCTAGAAGATGGAGGTATGCAGCCCGTTAGTAGGAGTAATACTACCCAGCTTATAACCCCTCAAACGTAG
GGCAGATGGCGGCCGCGAT

CTE2 - 350bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGATCCTAGAGACCATTTCGCGATTCCATGAGACTCCAAGGG
TTCTGCACAACCTTATGCACCTCTATTAGATCATTGTGTTCTACGAAGCCTGGACTGCATTACATATTCACAACCAAC
ATGAGAAGAGCGGAATAGATGGCCGGATGTTTGGTGGCTTTGATATATTGTGAGGAGCATTGCGAACCCCTAGAGCTG
TCCGGTCAAATAACCCCTCACAATAAGTGTAATGTCATGGGATAATCAAAGACTAAGGGAGGGCTTTTATAGAAG
GCGTGAGGTTCATGCTATCCCCCTCTGAAGACGCGGCCGCGAT

CTE2 - 450bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGATCCGTATACGTTTCTAATTTGTAGTTAACGGTTGGATA
CCACTTTGAGGCATGTAATATGGTACTGAGCTTCGGCACAGGGCTCAAATTGCATCATTAAATGTCTCCGATGTGGC
TATATGTCATGGATAAAGGCAGCCCCCTATATCTTTTTTTGTGGCAGCATGGGTCCATCAAAGCAATTATTCAGGGT
CTTAATGACCTCCACAGCTCTAAACGTAATTCATCTGGCTTTGCCTGTACTTACTTCCTCCATGAAAAAAGTGTTG
ATAATGCTCATAATGCTGCCAGCAATTTCTCCTTCTCAAGACTATTCTGGCTTCTGGGTACTTAAAAACAGGG
CTTAGAGTATGGCTGCTGACAAAATTGCACTCTAAACGCTAGCTTAGGTCTTCTGCGGCCGCGAT

CTE2 - 550bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGATCCGTAGCTATCGTTTCGCGAGAAAGTTAGTAGACACA
CAGGACCCAGGCGTGCAAGTCAATTTAGCTGACTACACCGATTCTGGTTAAAGAGCCTATGGCCACCCTTATTTT
AGAGAAAAAAACCACACCTCTAATGTGTTGGGCACTAGAAAAAGCTAACTACCTAGTCCGTTTCTGGACGACTTCA
TTGGGAATAACATACCCCCCACTGTGATTAAGACTGGCACTGTCCTAATGCTTTCTTCAATAGGTTTGGCTCATGTG
TGATTCCCTCTGGCAAACCTTATAGAGGACAAGCAGAATAAACCAATTCAAGGTCGTTGTAGCTGAAGGCCTGGCCTG
CCTGACAGTTAATTATGAGCATGTCTTGCCCTTCATGGTGGATATTCACAGCTGAAAGTGGTATTGGCATTTTTTTTC
TGAGGACACAACGAGGAAATCTGATAAATACGGCCACCTGAAGTCTAGCTCGGAGTTAACAATTTACCACGTTTAGA
GCGGCCGCGAT

CTE2 - 650bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGATCCGCTCGCACTTAGCCTGTAAAGGGGTTTCGCGCTCGT
CTAGTCTGTGCTGTTGCCTGGATAGTAAATTATCATGGTACAAACTTTTAAAGAGCCAGTTAAATGGAGATGGATTTA
AAAAGAGTTATTGTAAAGTCTCCCCAGGTGTGTCATTAAATATCCCAACAGATTGCCCTGGCCTGACCCCTAAATG
CAATTTTGGGATTCCCTTTTAGTTGCTTTTCAATTAATGTACCAGCGCAGTAAAAAAGCACAAAGTATATTGTTTA
TGTAACCTCACTATCTCATTGCACTGGTTACATGGCAGCTTCAGACTGACTAAACTACACTTTTCCCACCATGGTT
CAAAGATCAACAGAACTGGGCCAACAAAAGCAATTTTTTCATGTGGTCTAACTACCAACTTATTATGAGTTAAGTTA
CTTTTAGGTTTAAATCACAGCAGTTTTTCCCTCCACACCTCCCAGAGATACTTTCAGGGTGGCTAAACTTGGCTAA

AGGCTTCCGGACCAACCCCTTGTTCCTTTATGGTGCTTGTGTCCTGACAACCGCGTAAGGCATGGAAATTCAGCTATT
TATCCGATCGTTTATATGGGCGTGC GGCCGCGAT

CTE2 - 750bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGGATCCTTGGACCGTTAATTCATATATCGAAGTAGCAGGTT
GTTGCCCCGCCTGATGTTGCCACTACTTGCTCATGACAGTTTTTTTTAGGCAATGCAAACACTACTATTTGATATTTTTT
TCCAAGTACAGTTGTAGGGTACTCCTTATACTGATTCTTCTGAGCCTGTACGGGGAGCATTAGGTACTGATGTAGTA
GGAGTTGAGCTTCACAAATTCACCAGGTAAGCCCCAAATTTATTTTCTGCTTGGACAGGTCCACCTCACATGGGTCTG
TCTAATATATTAAAAGAGGGATTTTCTTTGCTGTATTGCAGCCCAGTATATCTGTTACTTACAGTAGTAGTCCATTA
TTGCTGGCCTAGGGGCTTTTGTCTCTACACGAACACCACTCTGTAAAATTTGAGGTCGTCCTTAGAGTCAAACCATT
CATGGAGCGCTCTGTGCATCTACCAACTATCGCTAAGCATTCACTTGGTTGGTTTAAAGTGGAGGCAACTCCATTATC
TTCTAGCATACCTTTCCCAGGCTACATGTAGAAAGAGATCTGTTGGGCCCCACTATTTTTTACCCAGGGAAGCCTA
CTTTAGTTATAGCTTGCCAGAGATTTTCTGTGTATGTAGAAGTCATCCACTTTTAAACACCAGGAGGTGGATGTGGG
GCCAGGAAATATGTCAATAACGATACGGGACTTCTAACAGTGACTCGCGGCCGCGAT

CTE2 - 850bp

ATCCTGCAGATGCATCCAGTACTAGTATGGCCCGGGGGATCCTTAAGTCGTGTCCTTCTCCTACGATCTTGTGAACG
ATGGATATTTTTCTTTCTAAACTTTAAACAAACAGTGAGAGATGTTGTTGTGTGTGGAACGACGCTTAGCCTACCGA
GGAAGATCCAGACTACAATAGAATATGTGGCCAAAACCTCTCCGCAACTTCAGCAGCAAAAAGGATATTATTGACATA
ACCTCCTCACAAAAGTACACAAATGGCTAAATAACAGAGCCCCCTCTTTTTACTAGGGAAATGGTGGATGTGGACTT
TAGAATTTAAGATAATAAAGCTCTTGATCCCAATGTTATTTCCATGTGAGGGACATTAAATTGAGTAACCTTTGCCA
CATACCCTCTCCAGAGTCCATTCTCTAAAACCTTGAAGCTCCGCCCCCTTTTTACGCACATTAGGCTTCCAATTACGG
TCAATGGTCTTGAAGATTGGGAGCTTTTGAAGAGTAATAAGAACCATCACAAAAGGAACCCAGAAGCCGGGAGTGT
CTACCAAAAAAATTCAAGGGTTAAAAAAAGTGACATTTTCTCCTGTTTTTTACACATGATTTTGAATGCTGATGGG
TCCACGTCCAGCTCTAAAGGTAGGTTTATGTTTCTCAAAGTTGCTTTCTTGTGAGAATTGAGCCACATCAGGTAGG
TGGGGAAGTAGATCAGTGAGGATGCTTCACATGTGTGGGCACTGGGAACAGAATGCTTCAATAACACGAGCTGACGA
GGGCCCCGCTATGAAAAAAAGATTCTCTGTGCCCCCTGGCGCCTCCGCACTTAAAGAATTGATGACCGTGCGGCCG
GAT

CTE1 - 123bp

GATCCTACGTTCCAAATGCAGCGAGCTCGTATAACCCTTTAAGAGTTGCTCTTTTTGTTTGGTAAGTTGCAAATCGA
AGTTTTAGATTGAGTTCTACGTCGAGCGGCCGCGATATCCTGCAGATGCA

CTE1 - 223bp

GATCCTTATCTGTCAAACCGCTAATGTCCGTTCTAAGACCGTCTGGAGAACACTTGCCCATCAGTGCTTTTGAACC
TTTTTTTACAGGTCCCTTCCGATTACACTGAGAAGCTGACCACACCTGCTAGAAGATGGAGGTATGCAGCCCGTTA
GTAGGAGTAATACTACCCAGCTTATAACCCTCAAACGTAGGGCAGATGGCGGCCGCGATATCCTGCAGATGCA

CTE1 - 323bp

GATCCTAGAGACCATTGCGGATTCCATGAGACTCCAAGGGTTCTGCACAACCTTATGCACCTCTATTAGATCATTGTG
TTCTACGAAGCCTGGACTGCATTACATATTCACAACCAACATGAGAAGAGCGGAATAGATGGCCGGATGTTTGGTGG
CTTTGATATATTGTGAGGAGCATTGCGAACCCTAGAGCTGTCCGGTCAAATAACCCCTCACATAAGTGTAATGTC
ATGGGATAATCAAAGACTAAGGGAGGGCTTTTATAGAAGGCGTGAGGTCATGCTATCCCCCTCTGAAGACGCGGCC
GCGATATCCTGCAGATGCA

CTE1 - 423bp

GATCCGTATACGTTTCTAATTTGTAGTTAACGGTTGGATACCACTTTGAGGCATGTAATATGGTACTGAGCTTCGGC
ACAGGGCTCAAATTGCATCATTAAATGTCTCCGATGTGGCTATATGTCATGGATAAAGGCAGCCCCCTATATCTTTT
TTTGTGGCAGCATGGGTCCATCAAAGCAATTATTCAGGGTCTTAATGACCTCCACAGCTCTAAACGTAATTCATCTG
GCTTTGCCTGTACTTACTTCCCTCCATGAAAAAAGTGTTGATAATGCTCATAATGCTGCCAGCAATTTCCCTCCCTT
CTCAAGACTATTCTGGCTTCCCTGGGTACTTAAAAACAGGGCTTAGAGTATGGCTGCTGACAAAATTGCACTCTAAAC
GCTAGCTTAGGTCTTCTGCGGCCGCGATATCCTGCAGATGCA

CTE1 - 523bp

GATCCGTTAGCTATCGTTCGCGAGAAAGTTAGTAGACACACAGGACCCAGGCGTGCAAGTCAATTTTCAGCTGACTAC
ACCGATTCTGGTTAAAAGAGCCTATGGCCACCCTTATTTTATAGAGAAAAAACCACACCTCTAATGTGTTGGGCACT
AGAAAAAGCTAACTACCTAGTCCGTTTCTGGACGACTTCATTGGGAATAACATACCCCCCACTGTGATTAAGACTGG
CACTGTCTAATGCTTTCTTCAATAGGTTTGGCTCATGTGTGATTCCCTCTGGCAAACCTTATAGAGGACAAGCAGAA
TAAACCAATTCAAGGTCGTTGTAGCTGAAGGCCTGGCCTGCCTGACAGTTAATTATGAGCATGTCTTGCCCTTCATG
GTGGATATTCACAGCTGAAAGTGGTATTGGCATTCTTCTGAGGACACAACGAGGAAATCTGATAAATACGGCCAC
CTGAAGTCTAGCTCGGAGTTACAATTTACCACGTTTAGAGCGGCCGCGATATCCTGCAGATGCA

CTE1 - 623bp

GATCCGCTCGCACTTAGCCTGTAAAGGGGTTTCGCGCTCGTCTAGTCTGTGCTGTTGCCTGGATAGTAAATTATCATG
GTACAAACTTTTTAAGAGCCAGTTAAATGGAGATGGATTTAAAAAGAGTTATTGTAAAGTCTCCCCAGGTGTGTCATT
AAATATCCCAACAGATTGCCCTGGCCTGACCCCTAAATGCAATTTTGGGATTCCCTTTTAGTTGCTTTTCATTA
TGTACCAGCGCAGTAAAAAAGCACAAAGTATATTGTTTATGTAACCTACTATCTCATTTGCACTGGTTACATGGCA
GCTTCAGACTGACTAAACTACACTTTTCCACCATTGGTTCAAAGATCAACAGAACTGGGCCAACAAAAGCAATTTT
TTCATGTGGTCTAACTACCAACTTATTATGAGTTAAGTTACTTTTAGGTTTAAAATCACAGCAGTTTTTCCCTCCAC
ACCTCCCAGAGATACTTTTCAGGGTGGCTAAACTTGGCTAAAGGCTTCCGGACCAACCCTTGTTTTCTTTATGGTGTCTT
GTGTCCTGACAACCGCGTAAGGCATGGAAATTCAGCTATTTATCCGATCGTTTATATGGGCGTGCGGCCGCGATATC
CTGCAGATGCA

CTE1 - 723bp

GATCCTTGGACCGTTAATTCATATATCGAAGTAGCAGGTTGTTGCCCGCCTGATGTTGCCACTACTTGCTCATGAC
AGTTTTTTTAGGCAATGCAAACTACTATTTGATATTTTTTCCAAGTACAGTTGTAGGGTACTCCTTATACTGATTCT
TTCTGAGCCTGTACGGGGAGCATTAGGTACTGATGTAGTAGGAGTTGAGCTTCACAAATTCACCAGGTAAAGCCCAA
TTTATTTTTCTGCTTGGACAGGTCCACCTCACATGGGTCTGTCTAATATATTAAAGAGGGATTTTCTTTGCTGTATT
GCAGCCAGTATATCTGTTACTTACAGTAGTAGTCCATTATTGCTGGCCTAGGGGCTTTTGCTCCTACACGAACACC
ACTCTGTAAAATTTGAGGTGCTCCTTAGAGTCAAACCATTCATGGAGCGCTCTGTGCATCTACCAACTATCGCTAAG
CATTCACTTGGTTGGTTTAAAGTGGAGGCAACTCCATTATCTTCTAGCATACCCTTCCCAGGCTACATGTAGAAAGAG
ATCTGTTGGGCCCCACTATTTTTTACCCAGGGAAGCCTACTTTAGTTATAGCTTGCCAGAGATTTTCTGTGTCATG
TAGAAGTCATCCACTTTTAAACACCAGGAGGTGGATGTGGGGCCAGGAAATATGTCAATAACGATACGGGACTTCTAA
CAGTGACTCGCGGCCGCGATATCCTGCAGATGCA

CTE1 - 823bp

GATCCTTAAGTCGTGTCTCTTCTCCTACGATCTTGTGAACGATGGATATTTTCTTTCTAAACTTTAAACAAACAGTGG
AGAGATGTTGTTGTGTGTGGAACGACGCTTAGCCTACCGAGGAAGATCCAGACTACAATAGAATATGTGGCCAAAAC
TCTCCGCAACTTCAGCAGCAAAAAGGATATTATTGACATAACCTCCTCACAAAAAGTACACAAATGGCTAAATAACA
GAGCCCCCTCTTTTTACTAGGGAAATGGTGGATGTGGACTTTAGAATTTAAGATAATAAAGCTCTTGATCCCAATGTT
ATTTCCATGTGAGGGACATTAAATTGAGTAACCTTTGCCACATACCCTCTCCCAGAGTCCATTCTCTAAAACCTTGAA
GCTCCGCCCCCTTTTTACGCACATTAGGCTTCCAATTACGGTCAATGGTCTTGAAGATTGGGAGCTTTTGAAGAGTAA
TAAGAACCATCACAAAAGGAACCCAGAAGCCGGGAGTGTCTACCAAAAAAATTCAAGGGTTAAAAAAGTGACAT
TTTCTCCTGTTTTTACACATGATTTTGAATGCTGATGGGTCCACGTCCAGCTCTAAAGGTAGGTTTCATGGTTCTCC
AAAGTTGCTTTCTTGTGAGAATTGAGCCACATCAGGTAGGTGGGGAAGTAGATCAGTGAGGATGCTTCACATGTGTG
GGCACTGGGAACAGAATGCTTCAATAACACGAGCTGACGAGGGCCCGCTATGAAAAAAGATTCTCTGTGCCCCCT
GGCGCCTCCGCACTTAAAGAATTGATGACCGTGCGGCCGCGATATCCTGCAGATGCA

CTA - 150bp

GGGGGATCCTACGTTCCAAATGCAGCGAGCTCGTATAACCCTTTAAGAGTTGCTCTTTTTGTTTGGTAAGTTGCAAA
TCGAAGTTTTAGATTGAGTTCTACGTGAGCGGCCGCGATATCCTGCAGATGCATCCAGTACTAGTATGGCCC

CTA - 250bp

GGGGGATCCTTATCTGTCAAAACCGCTAATGTCCGTTCTAAGACCGTCTGGAGAACACTTGCCCATCAGTGCTTTTG
AACCTTTTTTTTACAGGTCCCTTCCGATTACACTGAGAAGCTGACCACACCTGCTAGAAGATGGAGGTATGCAGCCC

GTTAGTAGGAGTAATACTACCCAGCTTATAACCCCTCAAACGTAGGGCAGATGGCGGCCGCGATATCCTGCAGATGCA
TCCAGTACTAGTATGGCCC

CTA - 350bp

GGGGGATCCTAGAGACCATTTCGCGATTCCATGAGACTCCAAGGGTTCTGCACAACCTTATGCACCTCTATTAGATCAT
TGTGTTCTACGAAGCCTGGACTGCATTACATATTCACAACCAACATGAGAAGAGCGGAATAGATGGCCGGATGTTTG
GTGGCTTTGATATATTGTGAGGAGCATTGCGAACCCCTAGAGCTGTCCGGTCAAATAACCCCTCACAATAAGTGTA
TGTCATGGGATAATCAAAAGACTAAGGGAGGGCTTTTATAGAAGGCGTGAGGTCATGCTATCCCCCTCTGAAGACGC
GGCCGCGATATCCTGCAGATGCATCCAGTACTAGTATGGCCC

CTA - 450bp

GGGGGATCCGTATACGTTTCTAATTTGTAGTTAACGGTTGGATACCACTTTGAGGCATGTAATATGGTACTGAGCTT
CGGCACAGGGCTCAAATTGCATCATTAAATGTCTCCGATGTGGCTATATGTCATGGATAAAGGCAGCCCCCTATATC
TTTTTTTGTGGCAGCATGGGTCCATCAAAGCAATTATTCAGGGTCTTAATGACCTCCACAGCTCTAAACGTAATTCA
TCTGGCTTTGCCTGTACTTACTTCCCTCCATGAAAAAAGTGTTGATAATGCTCATAATGCTGCCAGCAATTTCCCTC
CCTTCTCAAGACTATTCTGGCTTCCCTGGGTACTTAAAAACAGGGCTTAGAGTATGGCTGCTGACAAAATTGCACTCT
AAACGCTAGCTTAGGTCTTCTGCGGCCGCGATATCCTGCAGATGCATCCAGTACTAGTATGGCCC

CTA - 550bp

GGGGGATCCGTTAGCTATCGTTTCGCGAGAAAGTTAGTAGACACACAGGACCCAGGCGTGCAAGTCAATTTAGCTGA
CTACACCGATTCTGGTTAAAAGAGCCTATGGCCACCCTTATTTTAGAGAAAAAAACCACACCTCTAATGTGTTGGG
CACTAGAAAAAGCTAACTACCTAGTCCGTTTCTGGACGACTTCATTGGGAATAACATACCCCCCACTGTGATTAAGA
CTGGCACTGTCTAATGCTTTTCTTCAATAGGTTTGGCTCATGTGTGATTCCCTCTGGCAAACCTTATAGAGGACAAGC
AGAATAAACCAATTCAAGGTCGTTGTAGCTGAAGGCCTGGCCTGCCTGACAGTTAATTATGAGCATGTCTTGCCCTT
CATGGTGGATATTCACAGCTGAAAGTGGTATTGGCATTTTTTTCTGAGGACACAACGAGGAAATCTGATAAATACGG
CCACCTGAAGTCTAGCTCGGAGTTACAATTTACCACGTTTAGAGCGGCCGCGATATCCTGCAGATGCATCCAGTAC
TAGTATGGCCC

CTA - 650bp

GGGGGATCCGCTCGCACTTAGCCTGTAAAGGGGTTTCGCGCTCGTCTAGTCTGTGCTGTTGCCTGGATAGTAAATTAT
CATGGTACAACTTTTAAAGAGCCAGTTAAATGGAGATGGATTTAAAAGAGTTATTGTAAAGTCTCCCCAGGTGTGT
CATTAAATATCCCAACAGATTGCCCTGGCCTGACCCCTAAATGCAATTTTGGGATTCCCTTTTAGTTGCTTTTCATT
AAAATGTACCAGCGCAGTAAAAAAGCACAAAGTATATTGTTTATGTAACCTACTATCTCATTGCACTGGTTACAT
GGCAGCTTCAGACTGACTAAAACCTACTTTTCCCACCATGGTTCAAAGATCAACAGAACTGGGCCAACAAAAGCAA
TTTTTTTCATGTGGTCTAACTACCAACTTATTATGAGTTAAGTTACTTTTAGGTTTAAATCACAGCAGTTTTTCCCT
CCACACCTCCCAGAGATACTTTTCAGGGTGGCTAAACTTGGCTAAAGGCTTCCGGACCAACCCTTGTTTCTTTATGGT
GCTTGTGTCCTGACAACCGCGTAAGGCATGGAATTCAGCTATTTATCCGATCGTTTATATGGGCGTGCGGCCGCGA
TATCCTGCAGATGCATCCAGTACTAGTATGGCCC

CTA - 750bp

GGGGGATCCTTGACCGTTAATTCATATATCGAAGTAGCAGGTTGTTGCCCGCCTGATGTTGCCACTACTTGCTCA
TGACAGTTTTTTTAGGCAATGCAAACTACTATTTGATATTTTTTTTCCAAGTACAGTTGTAGGGTACTCCTTATACTG
ATTCTTCTGAGCCTGTACGGGGAGCATTAGGTACTGATGTAGTAGGAGTTGAGCTTCACAAATTCACCAGGTAAGCC
CAAATTTATTTTCTGCTTGGACAGGTCCACCTCACATGGGTCTGTCTAATATATTAAAAGAGGGATTTTCTTTGCTG
TATTGCAGCCCAGTATATCTGTTACTTACAGTAGTAGTCCATTATTGCTGGCCTAGGGGCTTTTGCTCCTACACGAA
CACCCTCTGTAAAATTTGAGGTGCTCCTTAGAGTCAAACCATTATGAGAGCGCTCTGTGCATCTACCAACTATCGC
TAAGCATTCACTTGGTTGGTTTAAAGTGGAGGCAACTCCATTATCTTCTAGCATACCCTTCCCAGGCTACATGTAGAA
AGAGATCTGTTGGGCCCCACTATTTTTTTCACCCAGGGAAGCCTACTTTAGTTATAGCTTGCCAGAGATTTTCTGTGT
CATGTAGAAGTCATCCACTTTTAAACACCAGGAGGTGGATGTGGGGCCAGGAAATATGTCAATAACGATACGGGACTT
CTAACAGTGACTCGCGGCCGCGATATCCTGCAGATGCATCCAGTACTAGTATGGCCC

CTA - 850bp

GGGGGATCCTTAAGTCGTGTCTTCTCTCTACGATCTTGTGAACGATGGATATTTTCTTTCTAAACTTTAAACAAACA
GTGGAGAGATGTTGTTGTGTGTGGAACGACGCTTAGCCTACCGAGGAAGATCCAGACTACAATAGAATATGTGGCCA
AAACTCTCCGCAACTTCAGCAGCAAAAAGGATATTATTGACATAACCTCCTCACAAAAAGTACACAAATGGCTAAAT
AACAGAGCCCCTCTTTTTACTAGGGAAATGGTGGATGTGGACTTTAGAATTTAAGATAATAAAGCTCTTGATCCCAA
TGTTATTTCCATGTGAGGGACATTAAATTGAGTAACCTTTGCCACATACCCTCTCCCAGAGTCCATTCTCTAAAACCT
TGAAGCTCCGCCCCCTTTTTACGCACATTAGGCTTTCCAATTACGGTCAATGGTCTTGAAGATTGGGAGCTTTTGAAGA
GTAATAAGAACCATCACAAAAAGGAACCCAGAAGCCGGGAGTGTCTACCAAAAAAATTCAAGGGTTAAAAAAAAGTG
ACATTTTCTCCTGTTTTTTACACATGATTTTGAATGCTGATGGGTCCACGTCCAGCTCTAAAGGTAGGTTTCATGGTT
CTCCAAAGTTGCTTTTCTTGTGAGAATTGAGCCACATCAGGTAGGTGGGGAAGTAGATCAGTGAGGATGCTTCACATG
TGTGGGCACTGGGAACAGAATGCTTCAATAACACGAGCTGACGAGGGCCCGCTATGAAAAAAAAGATTCTCTGTGCC
CCCTGGCGCCTCCGCACTTAAAGAATTGATGACCGTGCGGCCGCGATATCCTGCAGATGCATCCAGTACTAGTATGG
CCC

CTL - 150bp

AGTATGGCCCCGGGGGATCCTACGTTCCAAATGCAGCGAGCTCGTATAACCCTTTAAGAGTTGCTCTTTTTGTTTGGT
AAGTTGCAATCGAAGTTTTAGATTGAGTTCTACGTCGAGCGGCCGCGATATCCTGCAGATGCATCCAGTACA

CTL - 250bp

AGTATGGCCCCGGGGGATCCTTATCTGTCAAAACCGCTAATGTCCGTTCTAAGACCGTCTGGAGAACACTTGCCCATC
AGTGCTTTTGAACCTTTTTTTTACAGGTCCCTTCCGATTACACTGAGAAGCTGACCACACCTGCTAGAAGATGGAGG
TATGCAGCCCGTTAGTAGGAGTAATACTACCCAGCTTATAACCCTCAAACGTAGGGCAGATGGCGGCCGCGATATCC
TGCAGATGCATCCAGTACA

CTL - 350bp

AGTATGGCCCCGGGGGATCCTAGAGACCATTTCGCGATTCCATGAGACTCCAAGGGTTCTGCACAACCTTATGCACCTCT
ATTAGATCATTGTGTTCTACGAAGCCTGGACTGCATTACATATTACAAACCAACATGAGAAGAGCGGAATAGATGGC
CGGATGTTTGGTGGCTTTGATATATTGTGAGGAGCATTGCGAACCCTAGAGCTGTCCGGTCAAATAACCCCTCACA
ATAAGTGTAATGTCATGGGATAATCAAAAGACTAAGGGAGGGCTTTTATAGAAGGCGTGAGGTCATGCTATCCCCCT
CTGAAGACGCGGCCGCGATATCCTGCAGATGCATCCAGTACA

CTL - 450bp

AGTATGGCCCCGGGGGATCCGTATACGTTTCTAATTTGTAGTTAACGGTTGGATACCACTTTGAGGCATGTAATATGG
TACTGAGCTTCGGCACAGGGCTCAAATTGCATCATTAAATGTCTCCGATGTGGCTATATGTCATGGATAAAGGCAGC
CCCCATATCTTTTTTTTGTGGCAGCATGGGTCCATCAAAGCAATTATTAGGGTCTTAATGACCTCCACAGCTCTAA
ACGTAATTCATCTGGCTTTGCCTGTACTTACTTCCCTCCATGAAAAAAGTGTTGATAATGCTCATAATGCTGCCCAG
CAATTTCTCCCTTCTCAAGACTATTCTGGCTTCCCTGGGTACTTAAAAACAGGGCTTAGAGTATGGCTGCTGACAAA
ATTGCACTCTAAACGCTAGCTTAGGTCTTCTGCGGCCGCGATATCCTGCAGATGCATCCAGTACA

CTL - 550bp

AGTATGGCCCCGGGGGATCCGTTAGCTATCGTTCGCGAGAAAGTTAGTAGACACACAGGACCCAGGCGTGCAAGTCAA
TTTCAGCTGACTACACCGATTCTGGTTAAAAGAGCCTATGGCCACCCTTATTTTAGAGAAAAAAACCACACCTCTA
ATGTGTTGGGCACTAGAAAAAGCTAACTACCTAGTCCGTTTCTGGACGACTTCATTGGGAATAACATACCCCCCACT
GTGATTAAGACTGGCACTGTCCTAATGCTTTCTTCAATAGGTTTGGCTCATGTGTGATTCCCTCTGGCAAACCTTATA
GAGGACAAGCAGAATAAACCAATTCAAGGTCGTTGTAGCTGAAGGCCTGGCCTGCCTGACAGTTAATTATGAGCATG
TCTTGCCCTTCATGGTGGATATTCACAGCTGAAAGTGGTATTGGCATTTTTTTCTGAGGACACAACGAGGAAATCTG
ATAAATACGGCCACCTGAAGTCTAGCTCGGAGTTAACAATTTACCACGTTTAGAGCGGCCGCGATATCCTGCAGATG
CATCCAGTACA

CTL - 650bp

AGTATGGCCCCGGGGGATCCGCTCGCACTTAGCCTGTTAAGGGGTTTCGCGCTCGTCTAGTCTGTGCTGTTGCCTGGAT
AGTAAATTATCATGGTACAACTTTTAAAGAGCCAGTTAAATGGAGATGGATTTAAAAAGAGTTATTGTAAAGTCTCC
CCAGGTGTGTCATTAAATATCCCAACAGATTGCCCTGGCCTGACCCCTAAATGCAATTTTGGGATTCCCTTTTAGT

TGCTTTTCATTAAATGTACCAGCGCAGTAAAAAAGCACAAAGTATATTGTTTATGTAACCTACTATCTCATTGCA
CTGGTTACATGGCAGCTTCAGACTGACTAAACTACACTTTTCCCACCATGGTTCAAAGATCAACAGAAGTGGGCCA
ACAAAAGCAATTTTTTTCATGTGGTCTAACTACCAACTTATTATGAGTTAAGTTACTTTTAGGTTTAAAATCACAGCA
GTTTTTCCCTCCACACCTCCAGAGATACTTTAGGGTGGCTAAACTTGGCTAAAGGCTTCCGGACCAACCTTGTT
TCTTTATGGTGCTTGTGTCCTGACAACCGCGTAAGGCATGGAAATTCAGCTATTTATCCGATCGTTTATATGGGCGT
GCGGCCGCGATATCCTGCAGATGCATCCAGTACA

CTL - 750bp

AGTATGGCCCCGGGGATCCTTGGACCGTTAATTCATATATCGAAGTAGCAGGTGTTGCCCCGCCTGATGTTGCCAC
TACTTGCTCATGACAGTTTTTTTTAGGCAATGCAAACCTACTATTTGATATTTTTTCCAAGTACAGTTGTAGGGTACT
CCTTATACTGATTCTTCTGAGCCTGTACGGGGAGCATTAGGTACTGATGTAGTAGGAGTTGAGCTTCACAAATTCAC
CAGGTAAGCCCAATTTATTTTCTGCTTGGACAGGTCCACCTCACATGGGTCTGTCTAATATATTAAAAGAGGGATT
TTCTTTGCTGTATTGCAGCCCAGTATATCTGTTACTTACAGTAGTAGTCCATTATTGCTGGCCTAGGGGCTTTTGCT
CCTACACGAACACCACCTCTGTAAATTTGAGGTCGTCCTTAGAGTCAAACCATTATGAGAGCGCTCTGTGCATCTAC
CAACTATCGCTAAGCATTCACTTGGTTGGTTTAAAGTGGAGGCAACTCCATTATCTTCTAGCATACCCTTCCCAGGCT
ACATGTAGAAAGAGATCTGTTGGGCCCCACTATTTTTTACCCAGGGAAGCCTACTTTAGTTATAGCTTGCCAGAGA
TTTTCTGTGTCATGTAGAAGTCATCCACTTTTAAACCAGGAGGTGGATGTGGGGCCAGGAAATATGTCAATAACGA
TACGGGACTTCTAACAGTGACTCGCGCCGCGATATCCTGCAGATGCATCCAGTACA

CTL - 850bp

AGTATGGCCCCGGGGATCCTTAAAGTCGTGTCCTTCTCCTACGATCTTGTGAACGATGGATATTTTCTTTTC
TAAACTTTAAACAAACAGTGGAGAGATGTTGTTGTGTGTGGAACGACGCTTAGCCTACCGAGGAAGATCC
AGACTACAATAGAATATGTGGCCAAAACCTCTCCGCAACTTCAGCAGCAAAAAGGATATTATTGACATAAC
CTCCTCACAAAAAGTACACAAATGGCTAAATAACAGAGCCCCTCTTTTTACTAGGGGAAATGGTGGATGTG
GACTTTAGAATTTAAGATAATAAAGCTCTTGATCCCAATGTTATTTCCATGTGAGGGACATTAAATTGAG
TAACCTTTGCCACATACCCTCTCCCAGAGTCCATTCTCTAAAACCTGAAGCTCCGCCCTTTTTTACGCAC
ATTAGGCTTCCAATTACGGTCAATGGTCTTGAAGATTGGGAGCTTTTGAAGAGTAATAAGAACCATCACA
AAAAGGAACCCAGAAGCCGGGAGTGTCTACCAAAAAAATTCAAGGGTTAAAAAAAAGTGACATTTTCTCC
TGTTTTTTACACATGATTTTGAATGCTGATGGGTCCACGTCCAGCTCTAAAGGTAGGTTTCATGGTTCTCC
AAAGTTGCTTTCTTGTGAGAATTGAGCCACATCAGGTAGGTGGGGAAGTAGATCAGTGAGGATGCTTCAC
ATGTGTGGGCACTGGGAACAGAATGCTTCAATAACACGAGCTGACGAGGGCCCGCTATGAAAAAAAAGAT
TCTCTGTGCCCCCTGGCGCCTCCGCACTTAAAGAATTGATGACCGTGCGGCCGCGATATCCTGCAGATGC
ATCCAGTACA

Legacy Kits

The kits listed in this section are no longer sold.

Nextera DNA Sample Prep Kit (Epicentre Biotechnologies)

(Obsolete)

As a replacement, use catalog # FC-121-1031.

Transposon Sequences

5' -GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAG

5' -GCCTTGCCAGCCCGCTCAGAGATGTGTATAAGAGACAG

Adapters (showing optional bar code)

5' -AATGATACGGCGACCACCGAGATCTACACGCTCCCTCGCGCCATCAG

5' -CAAGCAGAAGACGGCATACGAGAT [**barcode**] CGGTCTGCCTTGCCAGCCCGCTCAG-3'

PCR Primers

5' -AATGATACGGCGACCACCGA

5' -CAAGCAGAAGACGGCATACGA

Oligonucleotide Sequences for Genomic DNA

(Obsolete)

Adapters

5' P-GATCGGAAGAGCTCGTATGCCGTCTTCTGCTTG

5' ACACTCTTTCCCTACACGACGCTCTTCCGATCT

PCR Primers

5' AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT

5' CAAGCAGAAGACGGCATACGAGCTCTTCCGATCT

Genomic DNA Sequencing Primer

5' ACACTCTTTCCCTACACGACGCTCTTCCGATCT

Oligonucleotide Sequences for Paired End DNA

(Obsolete)

PE Adapters

5' P-GATCGGAAGAGCGGTTCAGCAGGAATGCCGAG

5' ACACTCTTTCCCTACACGACGCTCTTCCGATCT

PE PCR Primer 1.0

5' AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT

PE PCR Primer 2.0

5' CAAGCAGAAGACGGCATACGAGATCGGTCTCGGCATTCCTGCTGAACCGCTCTTCCGATCT

PE Read 1 Sequencing Primer

5' ACACTCTTTCCCTACACGACGCTCTTCCGATCT

PE Read 2 Sequencing Primer

5' CGGTCTCGGCATTCCTGCTGAACCGCTCTTCCGATCT

Oligonucleotide Sequences for the Multiplexing Sample Prep Oligo Only Kit

(Obsolete)

Multiplexing Adapters

5' P-GATCGGAAGAGCACACGTCT

5' ACACTCTTTCCCTACACGACGCTCTTCCGATCT

Multiplexing PCR Primer 1.0

5' AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT

Multiplexing PCR Primer 2.0

5' GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT

Multiplexing Read 1 Sequencing Primer

5' ACACTCTTTCCCTACACGACGCTCTTCCGATCT

Multiplexing Index Read Sequencing Primer

5' GATCGGAAGAGCACACGTCTGAACTCCAGTCAC

Multiplexing Read 2 Sequencing Primer

5' GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT

PCR Primer Index Sequences 1–12

PCR Primer, Index 1

5' CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTC

PCR Primer, Index 2

5' CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTC

PCR Primer, Index 3

5' CAAGCAGAAGACGGCATACGAGATGCCTAAGTGACTGGAGTTC

PCR Primer, Index 4

5' CAAGCAGAAGACGGCATACGAGATTGGTCAGTGACTGGAGTTC

PCR Primer, Index 5

5' CAAGCAGAAGACGGCATACGAGATCACTGTGTGACTGGAGTTC

PCR Primer, Index 6

5' CAAGCAGAAGACGGCATACGAGATATTGGCGTGACTGGAGTTC

PCR Primer, Index 7

5' CAAGCAGAAGACGGCATACGAGATGATCTGGTGACTGGAGTTC

PCR Primer, Index 8

5' CAAGCAGAAGACGGCATACGAGATTCAAGTGTGACTGGAGTTC

PCR Primer, Index 9

5' CAAGCAGAAGACGGCATACGAGATCTGATCGTGACTGGAGTTC

PCR Primer, Index 10

5' CAAGCAGAAGACGGCATACGAGATAAGCTAGTGACTGGAGTTC

PCR Primer, Index 11

5' CAAGCAGAAGACGGCATACGAGATGTAGCCGTGACTGGAGTTC

PCR Primer, Index 12

5' CAAGCAGAAGACGGCATACGAGATTACAAGGTGACTGGAGTTC

Oligonucleotide Sequences for the v1 and v1.5 Small RNA Kits

(Obsolete)

RT Primer

5' CAAGCAGAAGACGGCATACGA

5' RNA Adapter

5' GUUCAGAGUUCUACAGUCCGACGAUC

3' RNA Adapter

5' P-UCGUAUGCCGUCUUCUGCUUGUdT

v1.5 Small RNA 3' Adapter

5' /5rApp/ATCTCGTATGCCGTCTTCTGCTTG/3ddC/

Small RNA PCR Primer 1

5' CAAGCAGAAGACGGCATACGA

Small RNA PCR Primer 2

5' AATGATACGGCGACCACCGACAGGTCAGAGTTCTACAGTCCGA

Small RNA Sequencing Primer

5' CGACAGGTTCTACAGTCCGACGATC

Revision History

| Document | Date | Description of Change |
|------------------------------|--------------|--|
| Document # 1000000002694 v00 | October 2015 | <p>Added information for the following TruSight kits: TruSight Cardio, TruSight Myeloid Sequencing Panel, TruSight One, TruSight Rapid Capture, TruSight Tumor 15, and TruSight Tumor 26.</p> <p>Grouped TruSeq Amplicon Kits section for TruSeq Custom Amplicon 1.5, TruSeq Amplicon Cancer Panel, and TruSeq Custom Amplicon Low Input.</p> <p>Marked obsolete kits as obsolete.</p> <p>Grouped legacy kit information in new section titled Legacy Kits.</p> <p>Reformatted and reorganized the contents, and assigned document # 1000000002694.</p> |

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