



Innovating Epigenetics Solutions

D-Plex 24 DNBSEQ™ Barcodes

Barcodes modules for D-Plex Small RNA DNBSEQ™ Kit

Cat. No. C05030060 (Set A: [4, 4, 8, 8])

C05030061 (Set B: [24])

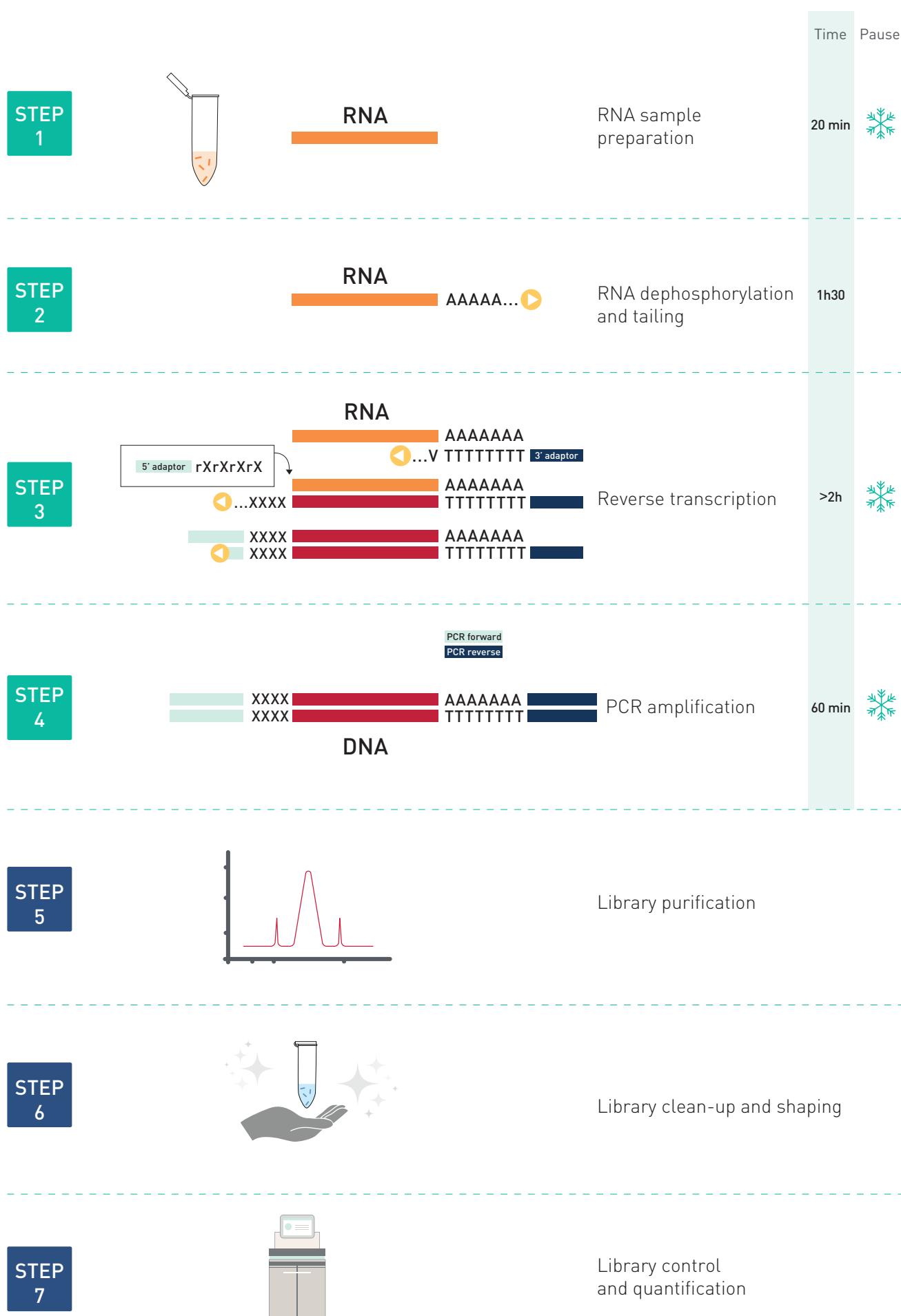
USER GUIDE

v1 12_2020

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Kit Method Overview



Introduction

The Diagenode D-Plex Small RNA DNBSEQ™ Library Preparation Kit is a tool designed for the study of the **small non-coding transcriptome**. The present kit incorporates the unique **D-Plex technology** to generate double-stranded DNA libraries from RNA samples ready to be used for the DNA single-strand circularization step required for **DNBSEQ sequencing** on MGI sequencers.

The D-Plex technology utilizes the innovative **capture and amplification by tailing and switching, a ligation-free method** for library preparation and offers key advantages such as:

- **Ultra-low input capability**
- Ease of use in a **one day, one tube protocol**
- **High library complexity**

The library preparation protocol works on either total intact RNA ($\text{RIN} \geq 8$) extracted and purified from a given sample or a small RNA fraction (<200nt), that might very well represent the circulating content of a **liquid biopsy-type of sample** (blood serum and plasma). The input requirements of the method are flexible and allow the user to perform the method within a wide range of RNA quantities going **from 10 pg** of circulating RNA **up to 100 ng** for total RNA.

The core of the technology relies on **ligation-free reactions** to attach the MGI adaptors to both ends of the library construct. Therefore, the results generated with the D-Plex Small RNA DNBSEQ kit will vastly differ from those produced with a ligase-based approach. For instance, the results generated with a D-Plex kit will encompass a **vast spectrum of small non-coding RNAs** (miRNAs, snoRNAs, snRNAs, piRNAs) whereas a ligase-based approach will enrich the sequencing library in 5'-P – 3'-OH RNAs, mainly mature miRNAs.

Diagenode therefore recommends having a **clear understanding of the scientific question** being asked in a given experiment before proceeding to a small RNA-seq library preparation as the choice of technology will strongly impact the end result.

Kit Materials

Table 1. D-Plex 24 DNBSEQ Barcode sequences – Set A [4, 4, 8, 8]

x-Barcode Set	D-Plex DNBSEQ Barcode #	PCR Reverse Primer sequence	Barcode
4	1	TGTGAGCCAAGGGAGTTGATCGGACCTATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	TAGGCCGAT
	2	TGTGAGCCAAGGGAGTTGGATTCCGTCTTGTCCTAAGACCGCTTGGCCTCCGACTT	GGACGGAATC
	3	TGTGAGCCAAGGGAGTTGCCAGTAAGTTGTCTCCTAAGACCGCTTGGCCTCCGACTT	CTTACTGCCG
	4	TGTGAGCCAAGGGAGTTGTCATTAGGTGGTCTCCTAAGACCGCTTGGCCTCCGACTT	ACCTAATTGA
4	13	TGTGAGCCAAGGGAGTTGCCATTGCCATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	CGGCAATCCG
	14	TGTGAGCCAAGGGAGTTGGAATCCTGATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	ATCAGGATT
	15	TGTGAGCCAAGGGAGTTGCTGGAAATGATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	TCATTCCAGA
	16	TGTGAGCCAAGGGAGTTGATCCAGCATTTGTCTCCTAAGACCGCTTGGCCTCCGACTT	GATGCTGGAT
8	57	TGTGAGCCAAGGGAGTTGCCGTTGAATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	ATTCAACGGA
	58	TGTGAGCCAAGGGAGTTGCCAGTACAGTTGTCTCCTAAGACCGCTTGGCCTCCGACTT	AACTGTACTG
	59	TGTGAGCCAAGGGAGTTGATTGAGGTACTTGTCCTAAGACCGCTTGGCCTCCGACTT	GTACCTCAAT
	60	TGTGAGCCAAGGGAGTTGATTAGAAGTCTTGTCCTAAGACCGCTTGGCCTCCGACTT	GACTTCAAT
	61	TGTGAGCCAAGGGAGTTGCAACGCTTCAATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	TGAAGCGTTG
	62	TGTGAGCCAAGGGAGTTGGGATCGCACGTTGTCTCCTAAGACCGCTTGGCCTCCGACTT	CGTGCATCC
	63	TGTGAGCCAAGGGAGTTGTCCTCCGATTGTCTCCTAAGACCGCTTGGCCTCCGACTT	TCGGAAGGCA
	64	TGTGAGCCAAGGGAGTTGGCGACATCGGTTGTCTCCTAAGACCGCTTGGCCTCCGACTT	CCGATGTCGC
8	65	TGTGAGCCAAGGGAGTTGCATTCTAACGTTGTCTCCTAAGACCGCTTGGCCTCCGACTT	ACTTAGAATG
	66	TGTGAGCCAAGGGAGTTGCCAGGCTTGGATTGTCTCCTAACGACCGCTTGGCCTCCGACTT	TCCAAGCCTG
	67	TGTGAGCCAAGGGAGTTGATCATCGTCTTGTCCTAAGACCGCTTGGCCTCCGACTT	AGACGATGAT
	68	TGTGAGCCAAGGGAGTTGGCTTGAGTTGTCTCCTAACGACCGCTTGGCCTCCGACTT	CTCACAAAGAC
	69	TGTGAGCCAAGGGAGTTGAGTAGGAACGTTGTCTCCTAACGACCGCTTGGCCTCCGACTT	CGTTCCACT
	70	TGTGAGCCAAGGGAGTTGTCACAACCATTGTCTCCTAACGACCGCTTGGCCTCCGACTT	GTGGTTGTGA
	71	TGTGAGCCAAGGGAGTTGCCAGGCCTTGTCTCCTAACGACCGCTTGGCCTCCGACTT	GAAGGCCTGC
	72	TGTGAGCCAAGGGAGTTGGCAAGCTATTGTCTCCTAACGACCGCTTGGCCTCCGACTT	TAGCTGCCA

Table 2. D-Plex 24 DNBSEQ Barcode sequences – Set B [24]

x-Barcode Set	D-Plex DNBSEQ Barcode #	PCR Reverse Primer sequence	Barcode
24	25	TGTGAGCCAAGGAGTTGTTGCCTCTATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	TAGAGGACAA
	26	TGTGAGCCAAGGAGTTGATTGCTAGGTTGCTTCCAAGACCGCTTGGCCTCCGACTT	CCTAGCGAAT
	28	TGTGAGCCAAGGAGTTGACAGCTCAGCTTGCTTCCAAGACCGCTTGGCCTCCGACTT	GCTGAGCTGT
	29	TGTGAGCCAAGGAGTTGATCTAGGTTTGCTTCCAAGACCGCTTGGCCTCCGACTT	AACCTAGATA
	30	TGTGAGCCAAGGAGTTGGAGATGCAATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	TTGCCATCTC
	32	TGTGAGCCAAGGAGTTGCCGATAGCGTTGCTTCCAAGACCGCTTGGCCTCCGACTT	CGCTATCGGC
	33	TGTGAGCCAAGGAGTTGCCATCGTTGCTTCCAAGACCGCTTGGCCTCCGACTT	GCAACGATGG
	34	TGTGAGCCAAGGAGTTGTAACGATTATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	TAATCGTTCA
	35	TGTGAGCCAAGGAGTTGAGAGCGAACATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	GTTCGCTCTA
	36	TGTGAGCCAAGGAGTTGATGTGAGATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	TCTCACACAT
	37	TGTGAGCCAAGGAGTTGATCCTAACAGTTGTCTTCCAAGACCGCTTGGCCTCCGACTT	CTGTTAGGAT
	38	TGTGAGCCAAGGAGTTGCGCGTCTGCGTTGTCTTCCAAGACCGCTTGGCCTCCGACTT	CGCAGACGCG
	39	TGTGAGCCAAGGAGTTGGATGATCCTTTGTCTTCCAAGACCGCTTGGCCTCCGACTT	AAGGATCATC
	49	TGTGAGCCAAGGAGTTGCTGCGTACATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	ATGTACGCAG
	50	TGTGAGCCAAGGAGTTGATCTCATTAATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	TTAATGAGAT
	51	TGTGAGCCAAGGAGTTGAAGTGGCGCATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	TGCGCCACTT
	52	TGTGAGCCAAGGAGTTGGGCCTTAATGTTGCTTCCAAGACCGCTTGGCCTCCGACTT	CATTAAGGCC
	53	TGTGAGCCAAGGAGTTGCTGAGCGGTTGTCTTCCAAGACCGCTTGGCCTCCGACTT	CCGCCTCAGA
	55	TGTGAGCCAAGGAGTTGGATAACCGGCTTGCTTCCAAGACCGCTTGGCCTCCGACTT	GCCGGTTATC
	56	TGTGAGCCAAGGAGTTGCAATATTCTTGTCTTCCAAGACCGCTTGGCCTCCGACTT	GGAATATTGA
	114	TGTGAGCCAAGGAGTTGCTGACTCTGGTTGCTTCCAAGACCGCTTGGCCTCCGACTT	CCAGAGTCAG
	115	TGTGAGCCAAGGAGTTGACTGCCGTGTTGTCTTCCAAGACCGCTTGGCCTCCGACTT	AACAGGCAGT
	116	TGTGAGCCAAGGAGTTGGCATGGAGCTGTCTTCCAAGACCGCTTGGCCTCCGACTT	GCTCCATGAC
	117	TGTGAGCCAAGGAGTTGGATAGACATTGTCTTCCAAGACCGCTTGGCCTCCGACTT	ATGTCTATCC

Table 3. Kit content

Reagent	Quantity	Storage temperature
D-Plex Forward Primer DNBSEQ (FP) (x1)	240 µl	-20°C/4°F
D-Plex Reverse Primer DNBSEQ (RP) (x24)	30 µl each	-20°C/4°F

Multiplexing Advices

The listed PCR Reverse Primers in Tables 1 and 2 are bearing the MGIEasy barcodes that can be used for library multiplexing up to 48. In case of a multiplexing scenario, it is then recommended to follow MGI's library pooling guidelines that are summarized in Table 4.

Table 4. Multiplexing recommendations for the D-Plex Small RNA DNBSEQ kit

Level of multiplexing	Instruction (Mixtures need to be prepared with equal volumes of each barcode to use them in a final volume of 10 µl for the PCR reaction)
1	1. Mix barcoded Reverse Primers of a 4-Barcode Set and give it to the sample. 2. Mix barcoded Reverse Primers of a 8-Barcode Set and give it to the sample.
2	1. Prepare 2 different mixtures with 2 different barcoded Reverse Primers of a 4-Barcode Set and give one mixture to one sample. 2. Prepare 2 different mixtures with 4 different barcoded Reverse Primers of a 8-Barcode Set and give one mixture to one sample.
3	For sample 1 & 2, use the method for 2-sample multiplexing above. For sample 3, use the method for 1-sample multiplexing above. Note that you should use different Barcode Sets for samples 1, 2 and 3.
4	1. Use a different barcoded Reverse Primers of a 4-Barcode Set for each sample. 2. Prepare 4 different mixtures with 2 different barcoded Reverse Primers of a 8-Barcode Set and give one mixture to one sample.
5	For samples 1-4, use the method for 4-sample multiplexing above. For sample 5, use the method for 1-sample multiplexing above. Note that you should use different Barcode Sets for samples 1-4 and 5.
6	For samples 1-4, use the method for 4-sample multiplexing above. For samples 5-6, use the method for 2-sample multiplexing above. Note that you should use different Barcode Sets for sample 1-4 and 5-6.

Level of multiplexing	Instruction (Mixtures need to be prepared with equal volumes of each barcode to use them in a final volume of 10 µl for the PCR reaction)
7	<ol style="list-style-type: none"> 1. For samples 1-4, use the method for 4-sample multiplexing above. (Use 1st Barcode Set). 2. For samples 5-6, use the method for 2-sample multiplexing above. (Use 2nd Barcode Set). 3. For sample 7, use the method for 1-sample multiplexing above. (Use 3rd Barcode set) <p>Note that you should use different Barcode Sets for samples 1-4, 5-6 and 7.</p>
8	Use a different barcoded Reverse Primers of a 8-Barcode Set for each sample.
8n+X (1≤n≤2; X=1-8; total 17-24)	<ol style="list-style-type: none"> 1. For samples 1-8, use the method for 8-sample multiplexing above. 2. For samples 9-8n, use the method for 8-sample multiplexing above. 3. For samples 8n+1 - 8n+X, according to the value of X, use the methods above for X-sample multiplexing accordingly. Remember to use different Barcode Sets. <p>Note that you should use different Barcode sets for steps 1), 2) and 3).</p>
8n+X (3≤n≤5; X=1-8; total 25-48)	<ol style="list-style-type: none"> 1. For samples 1-24 use a different barcoded Reverse Primer of the 24-Barcode Set for each sample. 2. For samples 25-8n, separate the samples into groups of 8, and use the method for 8-sample multiplexing above. Remember to use different Barcode Sets. 3. For samples 8n+1 - 8n+X, according to the value of X, use the methods above for X-sample multiplexing accordingly. Remember to use different Barcode Sets. <p>Note that you should use different Barcode Sets for steps 1), 2) and 3).</p>

Related products

Indexes Module Reference	Reference
D-Plex Small RNA DNBSEQ™ Kit	C05030059
DiaMag 0.2 mL tube magnetic rack	B04000001
DiaMag 1.5 mL tube magnetic rack	B04000003
MicroChIP DiaPure columns	C03040001

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