xGen™ UDI-UMI ADAPTERS

Overview

xGen[™] UDI-UMI Adapters are full-length adapters for ligating unique dual index (UDI) and unique molecular identifiers (UMI) to library molecules for Illumina® sequencing. A UDI is an 8-nucleotide sequence that is specific to a given sample library. UDI-UMI Adapters include the molecular index/barcode which is a 9-nucleotide sequence that uniquely tags each molecule within a sample library. These sample indexes are used for demultiplexing during data analysis when matching individual sequence reads to the correct sample. Before using UDI-UMI Adapters, refer to the appropriate library prep kit protocol for further information.

Product details

The xGen UDI-UMI Adapters are available at 15 μM concentration, in a single-use (96-well) plate with a pierceable seal. Each well contains one specific index pair for indexing one sample.

- xGen UDI-UMI Adapters, 16 rxn
- xGen UDI-UMI Adapters, 96 rxn

Low level multiplexing

It is not recommended to mix indexes between products unless you first verify color-balancing of the barcodes, and sufficient edit distance. For specific questions about multiplexing, reach out to our Scientific Applications Support team at applicationsupport@idtdna.com.

Handling and storage

- Store the xGen UDI-UMI Adapters at -20°C
- Thaw adapters on ice before use and keep on ice during use
- Do not heat adapters above room temperature (15–25°C)
- If any material remains unused, carefully re-seal the plate with a new adhesive seal to prevent cross-contamination
- Important: Never attempt to heat seal the plate again.

Directions for use

- 1. Before starting, thaw the xGen UDI-UMI Adapters on ice.
 - Important: Keep the xGen UDI-UMI Adapters on ice during use.
- 2. After thawing, briefly vortex the plate to mix, then centrifuge to collect the liquid in the bottom of the well before breaking the seal.
- 3. Before plating the adapters into the ligation reaction, prepierce the seal of the plate with a pipette tip, then directly pipette the required volume of primers.
 - Important: Always use a separate pipette tip for each well to avoid cross-contamination of indexes.
- 4. Prepare the Ligation Master Mix as instructed in the library prep protocol and adding the exact quantity of adapted specified in the protocol.
 - Note: The optimal amount of adapter is dependent on the protocol and input DNA quantity going into library prep.
 - Tip: If you need to dilute the adapter, always use the NGS Adapter Buffer (10 mM Tris, 0.1 mM EDTA and 100 mM NaCl, pH 8.0).
- 5. Return any unused portion of the plate to storage at -20°C.

Sequencing and analysis

- To fill out the index sequences in the sample sheet, utilize the IDT Master Index List.
- The UMI (Figure 1) is in-line with the i7 index. To sequence the UMI, increase the number of cycles by 9 for the i7 read.



Figure 1. xGen UDI-UMI Adapter.



Note: See the Analysis Guidelines for details on use of the UMI for error correction.



Tip: Strand-specific libraries constructed using the xGen UDI-UMI Adapters have the insert strandedness flipped (i.e., to the opposite strand) when compared to standard TruSeq[™] adapters (Illumina). This strand flip must be accounted for in data analysis for strandspecific applications (e.g., stranded RNA-seq, bisulfite sequencing).

Plate layouts

10006914-xGen[™] UDI-UMI Adapters, 16 rxn

	1	2	3	4	5	6	7	8	9	10	11	12
А	1	9	empty									
В	2	10	empty									
С	3	11	empty									
D	4	12	empty									
E	5	13	empty									
F	6	14	empty									
G	7	15	empty									
Н	8	16	empty									

10005903-xGen™ UDI-UMI Adapters, 96 rxn

	1	2	3	4	5	6	7	8	9	10	11	12
А	1	9	17	25	33	41	49	57	65	73	81	89
В	2	10	18	26	34	42	50	58	66	74	82	90
С	3	11	19	27	35	43	51	59	67	75	83	91
D	4	12	20	28	36	44	52	60	68	76	84	92
Е	5	13	21	29	37	45	53	61	69	77	85	93
F	6	14	22	30	38	46	54	62	70	78	86	94
G	7	15	23	31	39	47	55	63	71	79	87	95
Н	8	16	24	32	40	48	56	64	72	80	88	96

Technical support: applicationsupport@idtdna.com

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