AMPINEXT™ NGS Barcode (Index) Set12

Designed to construct multiplex DNA/RNA libraries used for next-generation

The AMPINETY WIGS Bracopte (Index) Sata 12 is designed to construct multiplex DNA/RNA libraries used for next-generation sequencing with the Illumina platform including GAIIx, HiSeq, MiSeq, and NextSeq. This product includes 12 DNA barcodes, each of which contain an index and flow cell binding sequence and can be attached to the sample insert during library preparation. Multiplexing several samples in a single flow cell can be achieved by pooling with the AMPINEXT™ NGS Barcodes. Pooling barcoded samples into a single flow cell significantly reduces hands-on time and provides robust data quality in NGS.

Ordering Information

Order Online »

ENZ-GEN507-0144

144Reactions

Manuals, SDS & CofA

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- Compatible with various library preparation workflows.
- High throughput and cost reduction: Up to 144 multiplexed samples can be processed.
- The indices are designed to be error resistant and allow for proper differentiation between samples.
- Minimizes PCR amplificationbased bias and prevents poor reads from single base errors introduced during PCR.

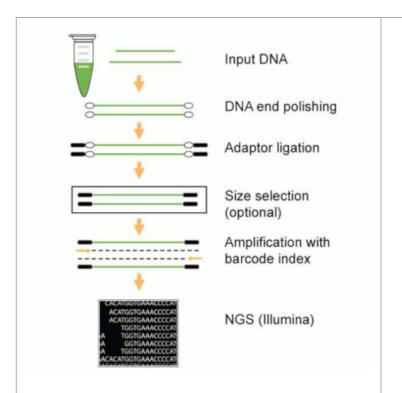


Figure 1. Schematic Procedure for using the AMPINEXT NGS Barcode (Index) Set-12

Handling & Storage

Long Term Storage -20°C

Shipping Blue Ice

Regulatory Status RUO - Research Use Only

Product Details

Application Notes Designed to construct multiplex DNA/RNA libraries used for next-generation sequencing

with the Illumina platform

AMPINEXT™ Barcodes 1-12 (10 µM) **Contents**

AMPINEXT™ Universal Primer (10 µM)

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