

Your sequencing data is only as good as
your adapter & index sets



• Is Your Supplier Validating With NGS QC?

Errors in adapter & indexing primer synthesis can
silently compromise your results — wasting valuable time
and budget.

COMMON RISKS

**Without Rigorous QC,
Errors Lead To:**

- ☒ Cross contamination between adapters/index primers
- ☒ Index hopping artefacts
- ☒ Poor functional performance
- ☒ Reduced sequencing data quality

These issues can compromise your
NGS results — and potentially
waste your **time and budget**.
Every batch matters.

LEGISEQ™ GUARANTEE

**Every Batch. Every Time.
NGS QC Verified.**

- ☒ Ultra-low cross-contamination rates
- ☒ High library yield
- ☒ Low dimer percentages
- ☒ High ligation efficiency

Each production batch undergoes
NGS QC validation in addition
to stringent **oligo synthesis QC**
— providing full assurance on
every order.

INTRODUCING

**LegiSeq™ NGS
Ready-to-Use Oligos**
Excellence in Every Read

*Full Length Adapters | Universal Adapters & Index Primers |
Tagmentation compatible Index Primers*

Building the Bases of Discovery

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