



# ACCELERATE YOUR HI-C STUDIES

## with the Dovetail® TopoLink™ v2 Assay

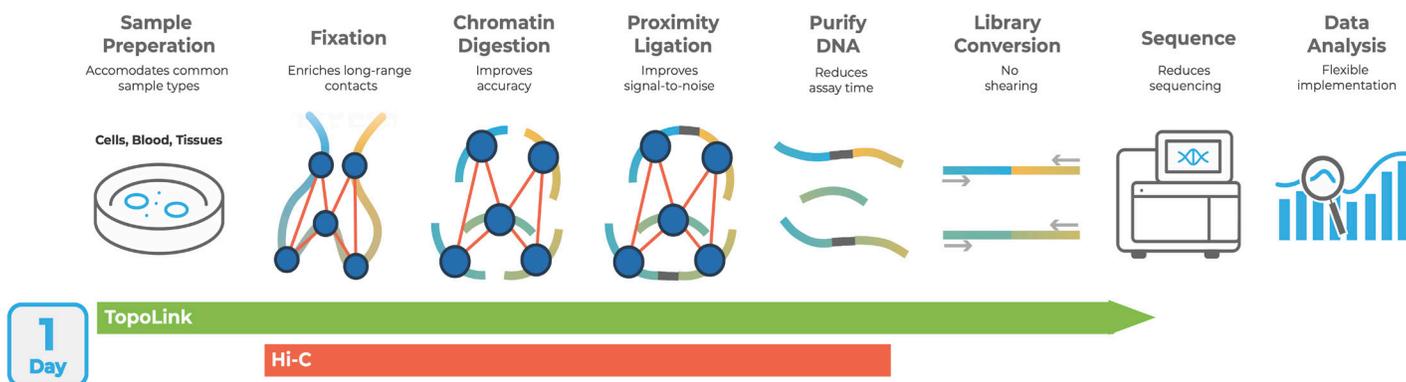
**FASTEST HI-C  
WORKFLOW**

**REDUCED  
SEQUENCING**

**IMPROVED  
ACCURACY**

### ACCELERATE YOUR HI-C STUDIES

Powered by novel LinkPrep™ chemistry, the Dovetail® TopoLink™ Assay addresses one of the most challenging hurdles in Hi-C data generation – assay length and difficulty. The Dovetail® TopoLink™ workflow enables users to go from sample to sequence-ready library in less than a day.

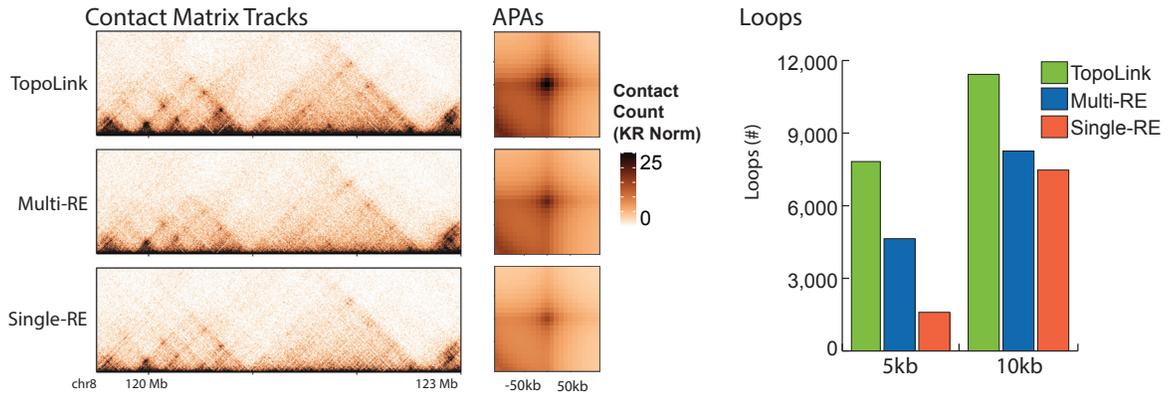


### SEE MORE - SEQUENCE LESS

Detect more TADs and loops compared to traditional Hi-C. Moreover, detected features have higher read support resulting in high-confidence feature calling even at reduced sequencing depths. As a result, fewer libraries and less sequencing are required simplifying experimental design and execution.

	TopoLink	Hi-C
Required Input (cells)	1 Million	2-5 Million
Total Assay Time	1 Day	2-3 Days
Required Proximity Ligation Reactions	1	3 - 4
Required # Libraries	3	6 - 8
Required Read Pairs For Loop Detection	800 Million	1.2 - 1.6 Billion
TADs Called	5,179	2,386
Loops Called	19,253	9,071 - 12,899

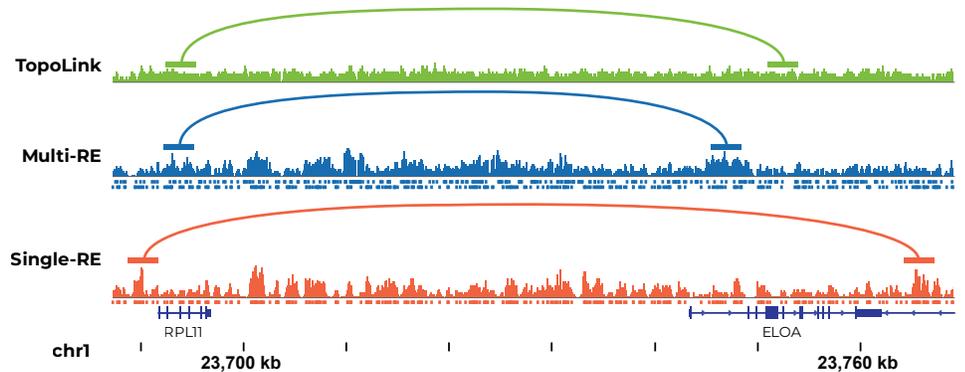
# GENERATE HIGH QUALITY CONTACT MATRICES AND IMPROVE FEATURE DETECTION



## UNBIASED GENOMIC COVERAGE EQUALS IMPROVED ACCURACY

Our uniform sequence coverage produces the most accurate chromatin interaction map possible. Where the non-uniform restriction enzyme site distribution associated with traditional Hi-C methods often skews signals, you can be confident in the position of interactions detected by the Dovetail TopoLink Assay.

### Uneven Sequence Coverage Shifts Loop Position



## ORDER NOW

The Dovetail TopoLink Kit supports all steps from sample preparation through to sequencing library generation – there is no need to purchase a separate library preparation kit.

Part Number	21010
# Reactions	8
# Libraries per sample	3 (24 total library conversions)
Supported Samples	Mammalian Cells, Tissue, Blood

