

**LeXonCT Panel v1.0** is a compact panel designed for analysis of solid tumors and liquid biopsy. This panel involves a total number of 69 genes with selected regions covered. With an optimized probe design, this panel offers highly uniform coverage over targeted regions for both FFPE and cell-free DNA.

### Genes with coding regions covered

<i>AKT1</i>	<i>ALK</i>	<b><i>APC</i></b>	<b><i>AR</i></b>	<i>ARAF</i>	<b><i>ATM</i></b>	<b><i>BRAF</i></b>	<b><i>BRCA1</i></b>	<b><i>BRCA2</i></b>	<b><i>CCND1</i></b>	<b><i>CDH1</i></b>	<b><i>CDK12</i></b>
<b><i>CDK4</i></b>	<b><i>CDK6</i></b>	<b><i>CDKN2A</i></b>	<i>CTNNB1</i>	<i>DDR2</i>	<b><i>EGFR</i></b>	<b><i>ERBB2</i></b>	<i>ERCC2</i>	<i>ESR1</i>	<i>EZH2</i>	<b><i>FGFR1</i></b>	<b><i>FGFR2</i></b>
<i>FGFR3</i>	<i>GNAI1</i>	<i>GNAQ</i>	<i>GNAS</i>	<i>HRAS</i>	<i>IDH1</i>	<i>IDH2</i>	<i>JAK2</i>	<i>JAK3</i>	<i>KDM6A</i>	<b><i>KIT</i></b>	<b><i>KRAS</i></b>
<i>MAP2K1</i>	<i>MAP2K2</i>	<b><i>MDM2</i></b>	<b><i>MET</i></b>	<b><i>MLH1</i></b>	<i>MPL</i>	<b><i>MSH2</i></b>	<b><i>MSH6</i></b>	<i>MTOR</i>	<b><i>MYC</i></b>	<b><i>NF1</i></b>	<i>NPM1</i>
<i>NRAS</i>	<i>NTRK1</i>	<i>NTRK2</i>	<i>NTRK3</i>	<b><i>PDGFRA</i></b>	<b><i>PIK3CA</i></b>	<b><i>PMS2</i></b>	<i>PTCH1</i>	<b><i>PTEN</i></b>	<b><i>PTPN11</i></b>	<b><i>RAF1</i></b>	<b><i>RB1</i></b>
<i>RET</i>	<i>ROS1</i>	<i>SMARCB1</i>	<b><i>SMO</i></b>	<b><i>STK11</i></b>	<i>TERT</i>	<b><i>TP53</i></b>	<i>TSC1</i>	<i>TSC2</i>			

Genes with full coding regions shown in bold.

### Genes with selected introns covered

<i>ALK</i>	<i>BRAF</i>	<i>EGFR</i>	<i>FGFR2</i>	<i>FGFR3</i>	<i>MET</i>	<i>NTRK1</i>	<i>NTRK2</i>	<i>PDGFRA</i>	<i>RET</i>	<i>ROS1</i>
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### Microsatellite markers

<i>BAT-25</i>	<i>BAT-26</i>	<i>BAT-40</i>	<i>BAT-RII</i>	<i>NR-21</i>	<i>NR-22</i>	<i>NR-24</i>	<i>NR-27</i>	<i>MONO-27</i>	<i>D2S123</i>	<i>D5S346</i>
<i>D17S261</i>	<i>D17S520</i>	<i>D18S34</i>								

## Performance

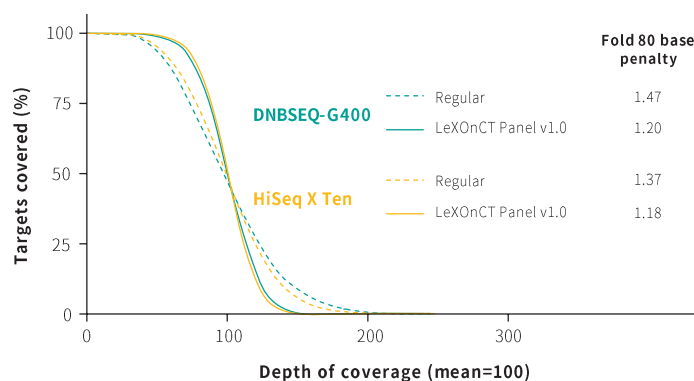
### On-target Rate

**Table 1. On-target rate for gDNA and cfDNA samples**

Platform	gDNA On-target rate (%)	cfDNA On-target rate (%)
HiSeq X Ten, PE150	87.02	87.35
DNBSEQ-G400, PE100	85.47	87.00

The DNA libraries were prepared using LeXPrep library kits with 50 ng of human genomic DNA (Promega, G1512) and 10 ng of plasma cfDNA from a healthy donor, respectively. 1 M read pairs were subsampled for the analysis. On-target rates were calculated as the percentage of mapped read pairs that overlap with probe regions.

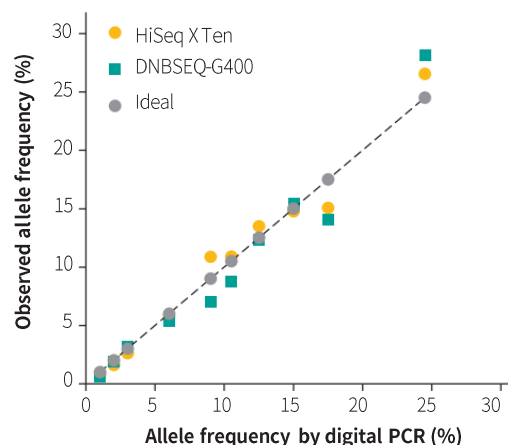
## Coverage Uniformity



**Fig 1. Improved design for cfDNA capture.** The optimized design of the LeXOnCT Panel v1.0 showed improved coverage uniformity for cfDNA libraries on both Illumina® and MGI platforms.

## Variant Analysis

Gene	Variants	Allele frequency by digital PCR	Observed frequency by Vardict	
			HiSeq X Ten	DNBSEQ-G400
NRAS	Q61K	12.5%	13.5%	12.4%
PIK3CA	E545K	9.0%	10.9%	7.0%
PIK3CA	H1047R	17.5%	15.1%	14.0%
EGFR	G719S	24.5%	26.6%	28.2%
EGFR	ΔE746 - A750	2.0%	1.6%	1.9%
EGFR	T790M	1.0%	0.9%	0.6%
EGFR	L858R	3.0%	2.6%	3.2%
BRAF	V600E	10.5%	10.9%	8.8%
KRAS	G13D	15.0%	14.8%	15.4%
KRAS	G12D	6.0%	5.8%	5.4%



**Fig 2. Evaluation of the LeXOnCT Panel v1.0 on the analysis of SNVs and indels.** DNA libraries were prepared from Onco SNV Multiplex 1–25% gDNA (GeneWell, GW-OGTM004) using LeXPrep library kits. The enriched libraries were sequenced either on HiSeq X Ten (PE150) or DNBSEQ-G400 (PE100).

## Ordering Information

Product	Catalog #
LeXOnCT Panel v1.0, 16 rxn	LX01902
LeXOnCT Panel v1.0, 96 rxn	LX01901

For research use only. Not for use in diagnostic procedures.