

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

AKT1	ALK	APC	AR	ARAF	ATM	BRAF	BRCA1	BRCA2	CCND1	CDH1	CDK12
CDK4	CDK6	CDKN2A	CTNNB1	DDR2	EGFR	ERBB2	ERCC2	ESR1	EZH2	FGFR1	FGFR2
FGFR3	GNA11	GNAQ	GNAS	HRAS	IDH1	IDH2	JAK2	JAK3	KDM6A	KIT	KRAS
MAP2K1	MAP2K2	MDM2	MET	MLH1	MPL	MSH2	MSH6	MTOR	MYC	NF1	NPM1
NRAS	NTRK1	NTRK2	NTRK3	PDGFRA	PIK3CA	PMS2	PTCH1	PTEN	PTPN11	RAF1	RB1
RET	ROS1	SMARCB1	SMO	STK11	TERT	TP53	TSC1	TSC2			

Genes with full coding regions shown in bold.

Genes with selected introns covered

ALK	BRAF	EGFR	FGFR2	FGFR3	MET	NTRK1	NTRK2	PDGFRA	RET	ROS1
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Microsatellite markers

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

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

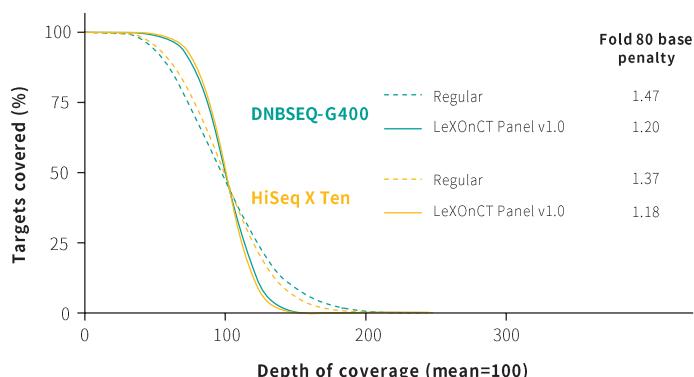


Fig 1. Improved design for cfDNA capture. The optimized design of the LeXOnCT Panel v1.0 showed improved coverage uniformity for cfDNA libraries on both Illumina® and MGI platforms.

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%

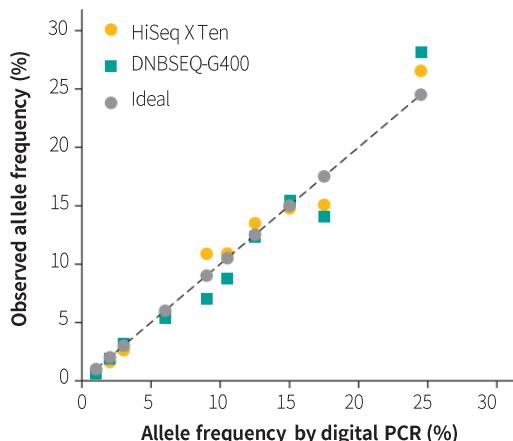


Fig 2. Evaluation of the LeXOnCT Panel v1.0 on the analysis of SNVs and indels. DNA libraries were prepared from Onco SNV Multiplex 1–25% gDNA (GeneWell, GW-OGTM004) using LeXPrep library kits. The enriched libraries were sequenced either on HiSeq X Ten (PE150) or DNBSEQ-G400 (PE100).

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.