

HEVA screen



HEVA screen is a kit for the analysis of the ATM, APC, BARD1, BRCA1, BRCA2, BRIP1, CDH1, CHEK2, EPCAM, MLH1, MSH2, MSH6, MUTYH, NBN, PALB2, PMS2, PTEN, RAD50, RAD51C, RAD51D, STK11 and TP53 genes through a molecular protocol based on Next Generation Sequencing (NGS) technologies. The kit is validated for germline analysis (SNPs, indels, CNVs) of DNA extracted from body tissues (blood or others). HEVA screen kit contains all reagents required for the preparation of a specific bidirectional library of amplicons designed for the NGS analysis using Illumina or Ion Torrent sequencers.

KEY FEATURES

- REF: R2010-16 (16 test)
- REF: R2010-48 (48 test)
- Application: germline analysis
- Number of pools: 3
- Panel size: 145 kb
- Input DNA: 20ng/reaction

SAMPLES/RUN*

	Germline
MiSeq Nano Kit v2 (500-cycles)	3
Kit v2 (500-cycles)	40
Kit v3 (600-cycles)	65
Ion 314™ Chip	1
Ion 316™ Chip	4
Ion 318™ Chip/Ion 520™ Chip	7
Ion 530™ Chip	27
Ion PI™ Chip/Ion 540™ Chip	>96

Target genes	Exons	Target genes	Exons
ATM	Full-exons plus 20 bp padding regions	MSH6	Full-exons plus 20 bp padding regions
APC		MUTYH	
BARD1		NBN	
BRCA1		PALB2	
BRCA2		PMS2	
BRIP1		PTEN	
CDH1		RAD50	
CHEK2		RAD51C	
EPCAM		RAD51D	
MLH1		STK11	
MSH2		TP53	

**the maximum number of samples per cartridge/chip estimated assuming an average depth of 300x for germline and 5000x for somatic analysis. The optimal number of samples must be empirically determined on local setups.*

INDEXES/ BARCODES

For Illumina instruments

HR Index set A (6x4) REF: R5001
HR Index set B (6x4) REF: R5002

For Ion Torrent instruments

HR Barcode 1-16 REF: R4001
HR Barcode 17-32 REF: R4002

For ordering info please contact info@4bases.ch

	Kit ID	Cod	Size (n° test)	CLINICAL APPLICATIONS	TARGETS	SAMPLES TYPE
Profiling of HotSpots somatic mutations (SNPs, indels) in cancer tissues	LUNG panel	R1000-16 R1000-48	16 48	NSCLC treatment	EGFR (exons 18, 19, 20, 21) KRAS (exons 2, 3, 4)	Tumor DNA tissue (fresh, frozen, FFPE, FNA, etc.) - Somatic analysis (SNPs, indels)
	COLON panel	R1010-16 R1010-48	16 48	mCRC treatment	KRAS (exons 2, 3, 4) NRAS (exons 2, 3, 4) BRAF (exon 15)	
	BENKit panel	R1020-16 R1020-48	16 48	MultiCancers treatment	KRAS (exons 2, 3, 4) NRAS (exons 2, 3, 4) BRAF (exon 11, 15) EGFR (exons 18, 19, 20, 21) PIK3CA (exons 10, 21)	
	THYRO-ID panel	R1030-16 R1030-48	16 48	Mutations profiling of Papillary Thyroid Carcinoma	KRAS (exons 2, 3, 4) NRAS (exons 2, 3, 4) HRAS (exons 2, 3) BRAF (exon 15) TP53 (exons 4, 5, 6, 7, 8, 9) NOTCH1 (exons 26, 27) PTEN (exons 5, 6, 7, 8) CDKN2A (exons 1, 2) EGFR (exons 18, 19, 20, 21) AKT1 (exon 1) CTNNB1 (exon 1) PIK3CA (exons 10, 21) TSHR (exons 6, 8, 9) hTERT (promoter)	
Full-gene sequencing (all CDS + flanking regions) for germline and/or somatic analysis	BRaCA screen	R2000-16 R2000-48	16 48	Hereditary and Somatic Variants profiling in Breast and Ovary cancer	BRCA1, BRCA2, TP53	Tumor DNA tissue (fresh, frozen, FFPE, FNA, etc.) or other (blood) - Somatic analysis (SNPs, indels) Germline analysis (SNPs, indels, CNVs)
	HECO screen	R2002-16 R2002-48	16 48	Germline Variants profiling in Hereditary nonpolyposis colorectal cancer (HNPCC)	APC, EPCAM, MLH1, MSH2, MSH6, MUTYH, PMS2, STK11	DNA from body tissues (blood or other) - Somatic analysis* (SNPs, indels) Germline analysis (SNPs, indels, CNVs)
	BRaVO screen	R2001-16 R2001-48	16 48	Germline Variants profiling in Hereditary Breast and Ovarian Cancer Syndrome (HBOC)	ATM, BRCA1, BRCA2, CDH1, CHEK2, PALB2, PTEN, TP53	
	HEVA screen	R2010-16 R2010-48	16 48	Hereditary Variants profiling in Breast and Ovary, Lynch Syndrome, and other cancer-related diseases	ATM, APC, BARD1, BRCA1, BRCA2, BRIP1, CDH1, CHEK2, EPCAM, MLH1, MSH2, MSH6, MUTYH, NBN, PALB2, PMS2, PTEN, RAD50, RAD51C, RAD51D, STK11, TP53	
	CFTR screen	R2030-16 R2030-48	16 48	Cystic fibrosis	CFTR	
	NEPHI screen	R2050-16 R2050-48	16 48	Neurofibromatosis (type 1, 2*) Noonan syndrome (type 1, 2)* Legius syndrome* Schwannomatosis*	NF1, SPRED1 NF2*, LZTR1*, SMARCB1*	
	IVF screen	R2040-16 R2040-48	16 48	Hereditary Variants profiling in genetic-related diseases	BDNF, BCHE, ATM, HBB, BLM, ASPA, CHM1, GLA, MEFV, FANCC, G6PC, GALT, GBA, GCDH, GJB2, OTOF, PJKK (DFNB59), HFE2, FPN1, HFE, TFR2, ALDOB, RS1, GALC, GLB1, IDUA, SMPD1, NPC2, NPC1, NBN, FSHR, PAH, PKD2, PKHD1, GAA, M2/ANXA5, ELP1 (IKAP), DHCR7, EPB42, ANK1, HEXA, MPL, MTHFR, F5, F2, ApoE, PAI1, TH, ATP7B, PEX1, AZFa, AZFb, AZFc, MUTYH, BRCA2, BRCA1, APC, SMN2, SMN1, F13A1, CFTR, DMD	
	FUSION screen	R2020-16 R2020-48	16 48	Fusion transcripts and expression imbalances between the 3' and 5' regions of the genes related with Lung cancers	EML4, ALK, ROS1, RET	
Extended screening of cDNA fusion transcripts						