

fastGEN

A breakthrough system for a one-step NGS library preparation

5NE



Meet the fastGEN technology

Introducing our cutting-edge sequencing solution fastGEN, built on the advanced NGS technology for unparalleled efficiency and user-friendliness.

With an easy-to-use software and diagnostic panels tailored for DNA and RNA analysis, coupled with prompt service and expertise from our application specialists, we are driving a revolution in laboratory diagnostics speed and the availability of personalized medicine.

BioVendor Group developed user-friendly diagnostic NGS kits with excellent analytical parameters and high-speed processing.

Meet our NGS team



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It's faster. Way faster!

Quick & easy detection of mutations in oncomarkers

The fastGEN technology is based on a rapid and easy-handling workflow what significantly lowers the probability of mistakes. The principle of fastGEN is ultra-deep sequencing of short amplicons obtained by a single polymerase chain reaction with specially labeled hybrid primers. FastGEN technology strategically focuses on codons with established treatment options, marking a revolutionary advancement in personalised medicine.

Unique

Simple, ultra-sensitive, specific, and effective technology

Reliable

Significantly lower probability of mistakes, uniform coverage for different amplicons and different samples

Fast

Excellent speed (<30 min hands-on time)

Robust

Robustness achieved by using short amplicons obtained by a single polymerase chain reaction with special tagged hybrid primers

Type of samples suitable for fastGEN technology



DNA isolated from tumor tissue

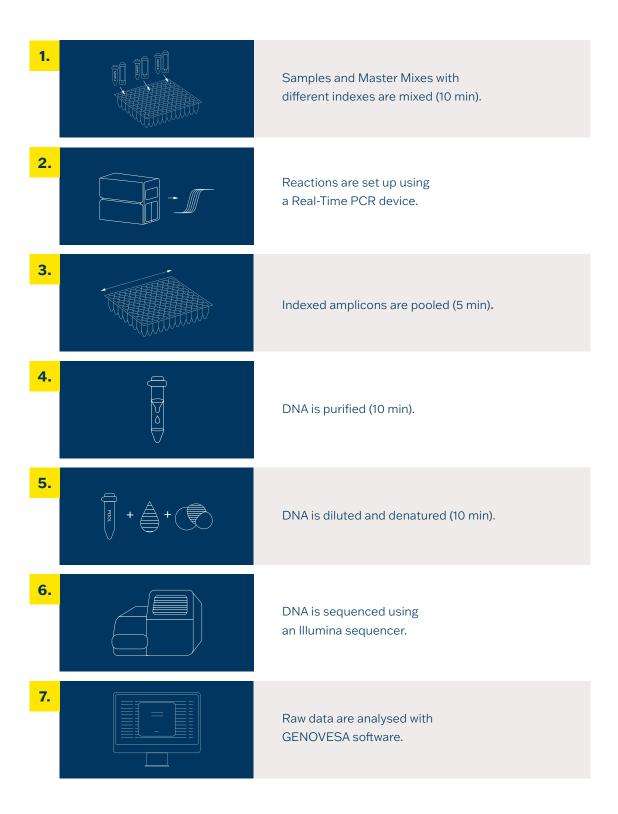


DNA isolated from FFPE tissue samples



Circulating tumor DNA isolated from blood

State-of-the-art prototyping workflow





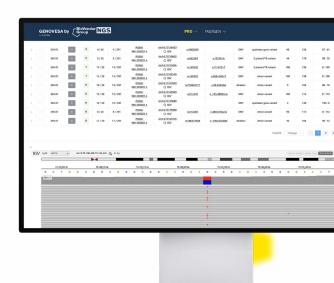
Total time: 4 hours | Hands-on-time: 30 minutes

GENOVESA

Cloud-based database bioinformatic and interpretation system

Key features

- User-friendliness
- Automatized bioinformatic analysis
- Advanced quality control of sequencing data
- Easy variants filtration
- Local clinical variants database
- Visualization of NGS data
- Internal interpretation of variants and sharing between clinics
- Clinical report generation
- Custom baseline creation



From FASTQ to clinical report

GENOVESA enables evaluation of data from small panels, through clinical exome (CES) whole exome data (WES) to whole genome sequencing (WGS) data. It includes a wide range of annotation databases and the possibility of individual customization. Advanced data quality control with the capability of own interpretation of variants and comments. Complete clinical management with integration of International Classification of Diseases (ICD) and HPO terminology.

Sequencing technology

GENOVESA supports analysis of conventional sequencing data (FASTQ, BAM, VCF) regardless of what sequencing technology was used – Illumina, MGI, PacBio, Oxford Nanopore Technologies, Genapsys and many others.

Databases and data sharing

GENOVESA also serves as a database for storing variants, with the possibility of sharing data between individual clinics.

Security

GENOVESA prioritizes data security through HTTPS encryption. We do not rely on third-party providers like Google or Amazon; instead, we host your data in our secure data center located in Prague. We do not collect any sensitive patient data, GENOVESA is fully GDPR compliant.



fastGEN BCR::ABL1 Cancer Kit for hemato-oncology

Molecular monitoring of patients with chronic myeloid leukemia (CML) is a key part of the treatment protocol during the use of tyrosine kinase inhibitors (TKIs). The *BCR::ABL1* fusion gene produces the pathological, constitutively activated tyrosine kinase Bcr-Abl, which is responsible for the uncontrolled proliferation of hematopoietic cells and their reduced response to proapoptotic signals.

The fastGEN technology provides a method of sensitive, specific and time-efficient examination of the mutational status of *BCR::ABL1* using NGS technology.

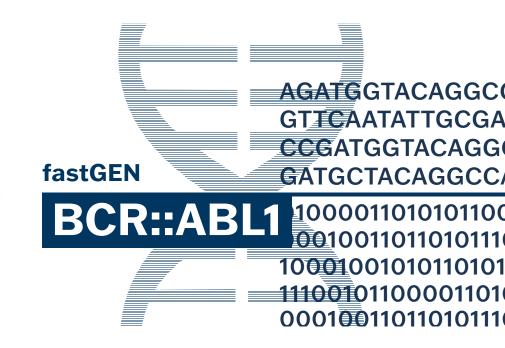


The kit covers codons 237-510

Extraordinary sensitivity

Major, minor and micro break-point

Monitoring the level of minimal residual disease (MRD)



Single technology, multiple opportunities

Kit and data analysis in one package

Kits

Cat. No.	Product	Gene	<u>Detail</u>	<u>Status</u>
RDNGS0001	fastGEN SOLID Cancer Kit	NRAS, KRAS, BRAF	NRAS, KRAS: codons 12, 13, 59, 60, 61, 117 and 146; BRAF: codons 600	CE IVD
RDNGS0002	fastGEN LUNG Cancer Kit	EGFR	exons 18, 19, 20 and 21	CE IVD
RDNGS0003	fastGEN BRAIN Cancer Kit	IDH1, IDH2	IDH1: codon 132 IDH2: codons 140 and 172	CE IVD
RDNGS0007	fastGEN PIK3CA Cancer Kit	PIK3CA	exons 2, 3, 5, 7, 8, 10, 14 and 21	RUO
RDNGS0008	fastGEN TERT Cancer Kit	TERT	promotor and mutations C228T a C250T	RUO
RDNGS0009 RDNGS0009-32	fastGEN TP53 Cancer Kit	TP53	exons 2-11, 2 non-canonical exons (between exon 9 and 10) and adjacent introns (min 6 nt)	RUO
RDNGS0010	fastGEN POLE/ CTNNB1 Cancer Kit	POLE, CTNNB1	POLE: exons 9, 11, 13 and 14 CTNNB1: exons 3, 7 and 8	RUO
RDNGS0011	fastGEN BCR::ABL1 Cancer Kit	ABL1, BCR	BCR::ABL1: minor and major breakpoint	RUO
RDNGS0016	fastGEN H3F3A/IDH1/2 Cancer Kit	H3F3A, IDH1, IDH2	IDH1 - codon 132 IDH2 - codons 140 and 172 H3F3A - codons 28 and 35	RUO
RDNGS0018	fastGEN Intolerance Kit I	ALDOB, AOC1, MCM6	ALDOB1: rs118204429, rs387906225, rs1800546, rs76917243, rs78340951, rs77718928, rs370793608, rs764826805 AOC1: rs10156191, rs1049742, rs2268999, rs1049793, rs2052129 MCM6: rs4988235, rs182549	RUO
RDNGS0019	fastGEN MSI Kit	12 genes	Microsatelite stability	RUO
RDNGS0020-32	fastGEN EGFR/HER2 Cancer Kit	EGFR, HER2	EGFR: exons 18, 19, 20 and 21 HER2: exons 7, 8, 17, 19, 20 and 21	RUO
RDNGS0021	fastGEN MPN Cancer Kit	CALR, JAK2, MPL	CALR: exon 9 JAK2: 12, 13, 14, 16 MPL: 4, 10, 12	RUO
RDNGS1001	fastGEN SOLID II Cancer Kit	NRAS, KRAS, BRAF	NRAS, KRAS: codons 12, 13, 59, 60, 61, 117 and 146; BRAF: codons 600 and 601	RUO



The kits are CE-IVD certified and intended for professional use.

Kit components

The kit contains ready-to-use Master Mixes with 16 sample indexes, read1, read2 seq primers and an index seq primer.









Contacts

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