Enabling personalized cancer care

oncobit

OncobitTM PM for personalized monitoring of melanoma

Imaging data is often difficult to interpret and may not accurately reflect disease progression. Clinical studies have shown that circulating tumor DNA can improve the understanding of patient prognosis and is an appropriate biomarker for monitoring disease recurrence, progression and response to therapy.*

OncobitTM PM is a digital PCR platform technology consisting of optimized PCR reagents and a proprietary analysis software. This complete solution enables the sensitive detection of specific cancer markers in circulating tumor DNA, as well as an unbiased and automatic report generation, providing a quantitative measure of circulating tumor DNA including the mutant allele frequency.

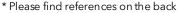
- Fully validated with clinical samples according to European guidelines
- Unbiased and automatic data analysis and reporting
- High accuracy demonstrated with > 1000 patient samples
- Precise quantification, e.g. standard error of 0.23% at 1.5% MAF** for BRAFV600E***
- High sensitivity, detecting as low as 2 copies per 10,000 wild-type copies (equivalent to 0.02% MAF**)

Target gene	Single-plex assays for	Limit of Blank (molecules)	Typical Indication
BRAF	V600E/K/R/E2/D	2.8/2.8/1.4/2.7/2.5	Cutaneous melanoma
NRAS	Q61K/L/R	1.8/1.8/1.8	Cutaneous melanoma
GNAQ	Q209P/L	1.8/1.8	Uveal melanoma
GNA11	Q209L	1.8	Uveal melanoma
SF3B1	R625H/C	5.7/7.5	Uveal melanoma

For more information, please contact

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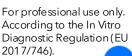




** Mutant allele frequency

*** Input material of 18ng cfDNA







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