# **TECHNICAL SHEET IDYLLA™ GENEFUSION PANEL**

## Intended Use

The Biocartis Idylla™ GeneFusion Panel is a fully automated in vitro diagnostic test intended for the qualitative detection of specific gene fusions of ALK, ROS1, RET as well as MET exon 14 skipping. The Idylla™ GeneFusion Panel is intended for use with formalin-fixed, paraffinembedded (FFPE) tumor tissue sections from patients with non-small cell lung cancer (NSCLC). The Idylla™ GeneFusion Panel covers the entire process from sample to result, including fully integrated RNA and DNA extraction, reverse transcription of mRNA, real-time PCR amplification and detection, data analysis, and result reporting.

### Indications for Use

The Idylla<sup>™</sup> GeneFusion Panel is for use by healthcare professionals for identifying the panel gene rearrangements for patients with NSCLC, to predict the most appropriate treatment options. The Idylla<sup>™</sup> GeneFusion Panel is not intended to diagnose NSCLC.

# **FEATURES**

Fusion specific detection				
	EML4-ALK	EML4 exon 2; ALK exon 20		
		EML4 exon 6a; ALK exon 20		
		EML4 exon 6b; ALK exon 20		
		EML4 exon 13; ALK exon 20		
		EML4 exon 15; ALK exon 20		
		EML4 exon 17; ALK exon 20		
		EML4 exon 18; ALK exon 20		
		EML4 exon 20; ALK exon 20		
ALK fusions (17)	KIF5B-ALK	KIF5B exon 15; ALK exon 20		
		KIF5B exon 17; ALK exon 20		
		KIF5B exon 24; ALK exon 20		
	HIP1-ALK	HIP1 exon 28; ALK exon 20		
KL TF		HIP1 exon 30; ALK exon 20		
	KLC1-ALK	KLC1 exon 9; ALK exon 20		
	TPR-ALK	TPR exon 15; ALK exon 20		
	TFG-ALK	TFG exon 4; ALK exon 20		
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Fusion specific detection - continued				
		CD74 exon 6; ROS1 exon 32		
	UJ/4-KUSI	CD74 exon 6; ROS1 exon 34		
		SDC4 exon 2; ROS1 exon 32		
	SDC4-ROS1	SDC4 exon 4; ROS1 exon 32		
		SDC4 exon 4; ROS1 exon 34		
		SLC34A2 exon 4; ROS1 exon 32		
ROS1 fusions (13)	SLC34A2-ROS1	SLC34A2 exon 4; ROS1 exon 34		
		SLC34A2 exon 13; ROS1 exon 32		
	EZR-ROS1	EZR exon 10; ROS1 exon 34		
	TPM3-ROS1	TPM3 exon 8; ROS1 exon 35		
		GOPC exon 4; ROS1 exon 36		
	UUFC-KUSI	GOPC exon 8; ROS1 exon 35		
	LRIG3-ROS1	LRIG3 exon 16; ROS1 exon 35		
		KIF5B exon 15; RET exon 11		
RET fusions (7)		KIF5B exon 15; RET exon 12		
	KIESR-DET	KIF5B exon 16; RET exon 12		
		KIF5B exon 22; RET exon 12		
		KIF5B exon 23; RET exon 12		
		KIF5B exon 24; RET exon 11		
	CCDC6-RET	CCDC6 exon 1; RET exon 12		
MET exon 14 skipping	MET exon 14 skipping transcript detection at the exon 13-exon 15 iunction			

# Expression imbalance detection

ALK expression imbalance
ROS1 expression imbalance
RET expression imbalance

Expression imbalance measures the difference between the 3' gene expression level and the 5' gene expression of the kinase gene. Expression imbalance results are only reported in case the specific fusion is not detected. A 'detected' expression imbalance result is indicative for the presence of a fusion and the result should be used in combination with an alternative gene fusion test method (e.g. IHC, FISH or NGS).

Internal GeneFusion controls	
RNA Housekeeping gene 1	ERCC3
RNA Housekeeping gene 2	TMUB2
RNA MET Wild Type	Detection of the Wild Type MET isoform mRNA containing the MET exon 14 sequence
DNA control	KIF11

Minimum specimen requirements			
Sample type	1 x 5 μm FFPE tissue section if tissue area $\ge$ 20 mm <sup>2</sup> 3 x 5 μm FFPE tissue sections if tissue area < 20 mm <sup>2</sup>		
Neoplastic cells	≥ 10%, if less macrodissection is required		
Total turnaround time			
Time	Approx. 180 minutes		
Analytical performance			
Analytical sensitivity	Gene rearrangement ALK ROS1 RET METex14 skipping	LoD in copies / Cartridge 3 000 – 10 000 3 000 5 000 3 000	
Between laboratory reproducibility (648 results at 3 sites with 6 artificial FFPE samples)	99% agreement (645/648)		
Between lot reproducibility (96 results on 3 lots with 4 clinical FFPE samples)	98.9% agreement (95/96)		

### Clinical performance

The clinical performance evaluation compared the Idylla<sup>™</sup> GeneFusion Panel with IHC (VENTANA ALK (D5F3) Assay, Roche Diagnostics GmbH) for ALK. ROS1, RET and METex14 were compared with NGS (Oncomine<sup>™</sup> Focus Assay, Thermo Fisher Scientific). PPA, NPA and OPA for ALK, ROS1 and RET were calculated based on fusion specific results only as well as by combining the fusion specific results with the expression imbalance results.

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### Concordance of the Idylla<sup>™</sup> GeneFusion Panel versus IHC for ALK.

ALK	Fusion Specific Results Only		Including Confirmed	I* Expression Imbalance
	Rate	Agreement (%)	Rate	Agreement (%)
PPA	34/38	89.5%	38/38	100.0%
NPA	118/119	99.2%	118/119	99.2%
OPA	152/157	96.8%	156/157	99.4%

\* Confirmed = samples that are expression Imbalance positive using the Idylla<sup>™</sup> GeneFusion Panel and that were confirmed with the reference method. Expression imbalance results are indicative for the presence of a fusion and should be confirmed with another technology.

### Concordance of the Idylla<sup>™</sup> GeneFusion Panel versus NGS for ROS1.

ROS1	Fusion Specific Results Only		Including Confirmed	a* Expression Imbalance
	Rate	Agreement (%)	Rate	Agreement (%)
PPA	12/15**	80.0%	12/15	80.0%
NPA	187/187	100.0%	187/187	100.0%
ОРА	199/202	98.5%	199/202	98.5%

\* Confirmed = samples that are expression Imbalance positive using the Idylla<sup>™</sup> GeneFusion Panel and that were confirmed with the reference method. Expression imbalance results are indicative for the presence of a fusion and should be confirmed with another technology.

\*\* Of the 3 discordant ROS1 positive samples, the Oncomine™ Focus Assay indicated that there was a low read count for 2 of the 3 samples. All 3 discordant samples tested negative for ROS1 with IHC.

### Concordance of the Idylla<sup>™</sup> GeneFusion Panel versus NGS for RET.

RET	Fusion Specific Results Only		Including Confirmed	* Expression Imbalance
	Rate	Agreement (%)	Rate	Agreement (%)
PPA	13/14	92.9%	14/14	100.0%
NPA	188/188	100.0%	188/188	100.0%
OPA	201/202	99.5%	202/202	100.0%

\* Confirmed = samples that are expression Imbalance positive using the Idylla™ GeneFusion Panel and that were confirmed with the reference method. Expression imbalance results are indicative for the presence of a fusion and should be confirmed with another technology.

### Concordance of the Idylla™ GeneFusion Panel versus NGS for METex14 skipping.

METex14	Rate	Agreement (%)
PPA	48/53*	90.6%
NPA	149/149	100.0%
OPA	197/202	97.5%

\* Of the 5 discordant METex14 positive samples, the Oncomine™ Focus Assay indicated that there was a low read count for 4 of the 5 samples.

LoD: Limit of Detection - PPA: Positive Percent Agreement - NPA: Negative Percent Agreement - OPA: Overall Percent Agreement

# Catalog number Idylla™ GeneFusion Panel A0120/6

The Idylla<sup>™</sup> GeneFusion Panel contains SuperScript<sup>™</sup> III Reverse Transcriptase and is provided subject to a license under patents or patent applications owned by or licensed to Life Technologies Corporation, which license is limited to the human diagnostic field and research field and specifically excludes applications in forensics (including human identity testing). The SuperScript<sup>™</sup> III trademark is owned by Life Technologies Corporation. Patents US 7,700,339, 8,168,383, 8,481,279, 8,486,645, 8,232,060, 8,288,102, 8,377,642, 9,988,688, 9,523,130, 9,096,855, 10,526,661, 9,364,477, 9,539,254, 10,551,383 and pending US applications and all their respective foreign equivalents under license from Cell Signaling Technology, Inc. Idylla<sup>™</sup> Platform and Idylla<sup>™</sup> GeneFusion Panel are available as an IVD in Europe and many other countries outside the US. Please check availability with a local Biocartis representative. © Biocartis NV, February 2023 OUS.FL051.EN.R1.03/2023