

PATIENT

DISEASE
NAME
DATE OF BIRTH
SEX
MEDICAL RECORD #

PHYSICIAN

ORDERING PHYSICIAN
MEDICAL FACILITY
ADDITIONAL RECIPIENT
MEDICAL FACILITY ID
PATHOLOGIST

SPECIMEN

SPECIMEN SITE
SPECIMEN ID
SPECIMEN TYPE
DATE OF COLLECTION
SPECIMEN RECEIVED

NO REPORTABLE ALTERATIONS WITH COMPANION DIAGNOSTIC (CDx) CLAIMS
See professional services section for additional information

OTHER ALTERATIONS & BIOMARKERS IDENTIFIED

Results reported in this section are not prescriptive or conclusive for labeled use of any specific therapeutic product. See professional services section for additional information.

Microsatellite status MS-Stable §
Tumor Mutational Burden 29 Muts/Mb §
ARID1A Y1555fs*8
BRAF G469V
CCND3 amplification §
EP300 EP300(NM_001429) rearrangement exon 12 §
LYN amplification §
MYC amplification §
NTRK1 NTRK1(NM_002529)-BGLAP(NM_199173) fusion (N11; B1*) §
PIK3CB amplification §
RAD21 amplification §
RBM10 P831fs*29
TP53 M237I
VEGFA amplification §

§ Refer to appendix for limitation statements related to detection of any copy number alterations, gene rearrangements, MSI or TMB result in this section.
 Please refer to appendix for Explanation of Clinical Significance Classification and for variants of unknown significance (VUS).

FoundationOne®CDx (FICDx) is a next generation sequencing based in vitro diagnostic device for detection of substitutions, insertion and deletion alterations (indels), and copy number alterations (CNAs) in 324 genes and select gene rearrangements, as well as genomic signatures including microsatellite instability (MSI) and tumor mutational burden (TMB) using DNA isolated from formalin-fixed paraffin embedded (FFPE) tumor tissue specimens. The test is intended as a companion diagnostic to identify patients who may benefit from treatment with the targeted therapies listed in Table 1 in accordance with the approved therapeutic product labeling. Additionally, FICDx is intended to provide tumor mutation profiling to be used by qualified health care professionals in accordance with professional guidelines in oncology for patients with solid malignant neoplasms. The FICDx assay is a single-site assay performed at Foundation Medicine, Inc

TABLE 1: COMPANION DIAGNOSTIC INDICATIONS

INDICATIONS	BIOMARKER	THERAPY
Non-small cell lung cancer (NSCLC)	EGFR exon 19 deletions and EGFR exon 21 L858R alterations	Gilotrif® (Afatinib), Iressa® (Gefitinib), or Tarceva® (Erlotinib)
	EGFR exon 20 T790M alterations	Tagrisso® (Osimertinib)
	ALK rearrangements	Alecensa® (Alectinib), Xalkori® (Crizotinib), or Zykadia® (Ceritinib)
	BRAF V600E	Tafinlar® (Dabrafenib) in combination with Mekinist® (Trametinib)
Melanoma	BRAF V600E	Tafinlar® (Dabrafenib) or Zelboraf® (Vemurafenib)
	BRAF V600E or V600K	Mekinist® (Trametinib) or Cotelliv® (Cobimetinib) in combination with Zelboraf® (Vemurafenib)
Breast cancer	ERBB2 (HER2) amplification	Herceptin® (Trastuzumab), Kadcyla® (Ado-trastuzumab emtansine), or Perjeta® (Pertuzumab)
Colorectal cancer	KRAS wild-type (absence of mutations in codons 12 and 13)	Erbix® (Cetuximab)
	KRAS wild-type (absence of mutations in exons 2, 3, and 4) and NRAS wild type (absence of mutations in exons 2, 3, and 4)	Vectibix® (Panitumumab)
Ovarian cancer	BRCA1/2 alterations	Rubraca® (Rucaparib)

ABOUT THE TEST FoundationOne®CDx is the first FDA-approved broad companion diagnostic for solid tumors.

ABOUT THE TEST FoundationOne®CDx is the first and only FDA-Approved comprehensive companion diagnostic for all solid tumors.

Interpretive content on this page and subsequent pages is provided as a professional service, and is not reviewed or approved by the FDA.

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Biomarker Findings

Tumor Mutational Burden - TMB-High (29 Muts/Mb)
Microsatellite status - MS-Stable

Genomic Findings

For a complete list of the genes assayed, please refer to the Appendix.

BRAF G469V
NTRK1 NTRK1-BGLAP fusion
MYC amplification
PIK3CB amplification
ARID1A Y1555fs*8
CCND3 amplification - equivocal†
EP300 rearrangement exon 12
LYN amplification
RAD21 amplification
RBM10 P831fs*29
TP53 M237I
VEGFA amplification - equivocal†

7 Disease relevant genes with no reportable alterations: **RET, ROS1, ALK, KRAS, ERBB2, MET, EGFR**

† See About the Test in appendix for details.

13 Therapies with Clinical Benefit
0 Therapies with Lack of Response

40 Clinical Trials

BIOMARKER FINDINGS

Tumor Mutational Burden - TMB-High (29 Muts/Mb)

10 Trials see p. 20

Microsatellite status - MS-Stable

GENOMIC FINDINGS

BRAF - G469V

10 Trials see p. 23

NTRK1 - NTRK1-BGLAP fusion

7 Trials see p. 25

MYC - amplification

5 Trials see p. 27

THERAPIES WITH CLINICAL BENEFIT (IN PATIENT'S TUMOR TYPE)	THERAPIES WITH CLINICAL BENEFIT (IN OTHER TUMOR TYPE)
Atezolizumab	Avelumab
Durvalumab	Cemiplimab-rwlc
Nivolumab	
Pembrolizumab	

No therapies or clinical trials. see Biomarker Findings section

THERAPIES WITH CLINICAL BENEFIT (IN PATIENT'S TUMOR TYPE)	THERAPIES WITH CLINICAL BENEFIT (IN OTHER TUMOR TYPE)
Trametinib	Binimetinib
	Cobimetinib
	Regorafenib
	Sorafenib
Crizotinib	none
Larotrectinib	
none	none

The content provided as a professional service by Foundation Medicine, Inc., has not been reviewed or approved by the FDA.

Electronically signed by Jo-Anne Vergilio, M.D. | Jeffrey Ross, M.D., Medical Director | 16 January 2019 | Foundation Medicine, Inc. | 1.888.988.3639

Sample Preparation: 150 Second St., 1st Floor, Cambridge, MA 02141 - CLIA: 22D2027531
Sample Analysis: 150 Second St., 1st Floor, Cambridge, MA 02141 - CLIA: 22D2027531

GRF#

GENOMIC FINDINGS	THERAPIES WITH CLINICAL BENEFIT (IN PATIENT'S TUMOR TYPE)	THERAPIES WITH CLINICAL BENEFIT (IN OTHER TUMOR TYPE)
PIK3CB - amplification	none	none
10 Trials see p. 28		

GENOMIC FINDINGS WITH NO REPORTABLE THERAPEUTIC OR CLINICAL TRIAL OPTIONS

For more information regarding biological and clinical significance, including prognostic, diagnostic, germline, and potential chemosensitivity implications, see the Genomic Findings section.

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CCND3 - amplification - equivocal.....	p. 8	RBM10 - P831fs*29.....	p. 10
EP300 - rearrangement exon 12.....	p. 8	TP53 - M237I.....	p. 10
LYN - amplification.....	p. 9	VEGFA - amplification - equivocal.....	p. 11

NOTE Genomic alterations detected may be associated with activity of certain FDA-approved drugs; however, the agents listed in this report may have varied clinical evidence in the patient's tumor type. Neither the therapeutic agents nor the trials identified are ranked in order of potential or predicted efficacy for this patient, nor are they ranked in order of level of evidence for this patient's tumor type.

SAMPLE