



Sample Information

Patient Name: 張結坤

Gender: M

ID No.: N101006900

History No.: 42288824

Age: 69

Ordering Doctor: DOC3109L 邱昭華

Ordering REQ.: D4NJ878

Signing in Date: 2020/03/05

Path No.: S109-99207

MP No.: F2003

Assay: Oncomine Focus Assay

Sample Type: FFPE

Block No.: S107-26619A

Percentage of tumor cells: 70%

Note:

Sample Cancer Type: Non-Small Cell Lung Cancer

Table of Contents

	Page
Variant Details	2
Biomarker Descriptions	2
Relevant Therapy Summary	3
Relevant Therapy Details	6

Report Highlights

1 Clinically Significant Biomarkers
 1 Therapies Available
 60 Clinical Trials

Relevant Non-Small Cell Lung Cancer Findings

Gene	Finding	Gene	Finding
ALK	Not detected	MET	Not detected
BRAF	Not detected	NTRK1	Not detected
EGFR	EGFR exon 20 insertion	NTRK3	Not detected
ERBB2	Not detected	RET	Not detected
KRAS	Not detected	ROS1	Not detected

Clinically Significant Biomarkers

Indicated Contraindicated

Genomic Alteration	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
EGFR exon 20 insertion epidermal growth factor receptor Tier: IA Allele Frequency: 33.78%	<div>osimertinib</div> <div>gefitinib²</div>	None	60

Sources included in relevant therapies: FDA1, NCCN, EMA2, ESMO



Prevalent cancer biomarkers without clinical significance based on included data sources

CTNNB1 p.(S37F) c.110C>T

Tier Criteria Met

Genomic Alteration	Tier Classification for Non-Small Cell Lung Cancer
<i>EGFR exon 20 insertion</i> Tier: IA	IA: Biomarker predicts response or resistance to FDA or EMA approved therapies in this cancer type IA: Biomarker is included in NCCN or ESMO guidelines that predict response or resistance to therapies in this cancer type IIC: Biomarker is an inclusion criteria for clinical trials

Reference: Li et al. *Standards and Guidelines for the Interpretation and Reporting of Sequence Variants in Cancer: A Joint Consensus Recommendation of the Association for Molecular Pathology, American Society of Clinical Oncology, and College of American Pathologists.* J Mol Diagn. 2017 Jan;19(1):4-23.

Variant Details

DNA Sequence Variants								
Gene	Amino Acid Change	Coding	Variant ID	Locus	Allele Frequency	Transcript	Variant Effect	Coverage
CTNNB1	p.(S37F)	c.110C>T	COSM5662	chr3:41266113	32.20%	NM_001904.3	missense	2000
EGFR	p.(D770delinsGY)	c.2308_2309insGTT	COSM12427	chr7:55249010	33.78%	NM_005228.4	nonframeshift Insertion	1948
JAK1	p.(=)	c.2199A>G	.	chr1:65310489	80.11%	NM_002227.3	synonymous	1991
ALK	p.(D1529E)	c.4587C>G	.	chr2:29416366	99.95%	NM_004304.4	missense	1998
ALK	p.(I1461V)	c.4381A>G	.	chr2:29416572	99.85%	NM_004304.4	missense	1994
ALK	p.(=)	c.3375C>A	.	chr2:29445458	48.77%	NM_004304.4	synonymous	1997
FGFR3	p.(=)	c.1953G>A	.	chr4:1807894	99.90%	NM_000142.4	synonymous	1994
PDGFRA	p.(=)	c.1701A>G	.	chr4:55141055	99.10%	NM_006206.5	synonymous	1992
FGFR4	p.(P136L)	c.407C>T	.	chr5:176517797	99.40%	NM_213647.2	missense	2000
FGFR4	p.(=)	c.483A>G	.	chr5:176517985	11.61%	NM_213647.2	synonymous	1999
EGFR	p.(G724S)	c.2170G>A	.	chr7:55241722	30.50%	NM_005228.4	missense	2000

Biomarker Descriptions

CTNNB1 (catenin beta 1)

Background: The CTNNB1 gene encodes catenin beta-1 (β -catenin), an integral component of cadherin-based adherens junctions involved in maintaining adhesion and regulating the growth of epithelial cell layers¹. CTNNB1 binds to the APC protein in the cytoplasm and also interacts with TCF and LEF transcription factors in the nucleus to regulate WNT signaling². Steady state levels of CTNNB1 are regulated by ubiquitin-dependent proteolysis^{3,4,5}.



Biomarker Descriptions (continued)

Alterations and prevalence: Recurrent somatic mutations leading to CTNNB1 activation are common in cancer. The most prevalent alterations include missense mutations in exon 3 at codons S33, S37, T41, and S45 that block phosphorylation by GSK- β and inhibit CTNNB1 degradation^{6,7,8,9}. These activating mutations are observed in diverse solid tumors and have a prevalence of 20-30% in hepatocellular carcinoma, 20% of uterine carcinoma, and 15% of adrenocortical carcinoma^{10,11,12,13,14,15,16}.

Potential clinical relevance: Currently, no therapies have been approved for CTNNB1 aberrations. CTNNB1 alterations in EGFR positive lung cancer have been proposed to promote cancer progression and limit the response to EGFR tyrosine kinase inhibitors¹⁷.

EGFR (epidermal growth factor receptor)

Background: The EGFR gene encodes the epidermal growth factor receptor (EGFR) tyrosine kinase, a member of the human epidermal growth factor receptor (HER) family. Along with EGFR/ERBB1/HER1, ERBB2/HER2, ERBB3/HER3, and ERBB4/HER4 make up the HER protein family¹⁸. EGFR ligand induced dimerization results in kinase activation and leads to stimulation of oncogenic signaling pathways including the PI3K/AKT/MTOR and RAS/RAF/MEK/ERK pathways. Activation of these pathways promote cell proliferation, differentiation, and survival^{19,20}.

Alterations and prevalence: Recurrent somatic mutations in the tyrosine kinase domain of EGFR are observed in approximately 10-20% of lung adenocarcinoma and at higher frequencies in never-smoker, female, and in Asian populations with lung cancer^{15,16,21,22}. The most common mutations occur near the ATP-binding pocket of the kinase domain and include short in-frame deletions in exon 19 (EGFR exon 19 deletion) and the L858R amino acid substitution in exon 21²³. These mutations constitutively activate the EGFR kinase resulting in downstream signaling and represent 80% of the EGFR mutations observed in lung cancer. A second group of recurrent activating mutations that are less common include E709K, G719X, S768I, L861Q, and short in-frame insertions in exon 20^{24,25,26,27}. EGFR activating mutations in lung cancer tend to be mutually exclusive to KRAS activating mutations²⁸. Although these variants are common in lung cancer, they are rare in other cancer types. In glioblastoma, recurrent activating EGFR mutations in the extracellular domain include R108K, A289V and G598V^{23,29}. The recurrent focal amplification of the EGFR gene leads to an increase in expression in several cancer types. EGFR is amplified in up to 30% of glioblastoma, 12% of esophageal cancer, 10% of head and neck cancer, 5% of bladder cancer, and 5% of lung squamous cell carcinoma^{15,16,22,29,30}. Deletion of exons 2-7 encoding the extracellular domain of EGFR (EGFRvIII) results in overexpression of a ligand-independent constitutively active protein which is frequently observed in glioblastoma and has been shown to lead to lung cancer development as well as sensitivity to TKIs^{31,32,33}.

Potential clinical relevance: Erlotinib³⁴ (2004), afatinib³⁵ (2013), gefitinib³⁶ (2015), osimertinib³⁷ (2015), and dacomitinib³⁸ (2018) are small molecule TKIs that are FDA approved for non-small cell lung cancer (NSCLC) patients with sensitizing exon 19 deletions and exon 21 L858R mutations. Acquired secondary mutations often confer resistance to first line TKI therapy with the T790M amino acid substitution accounting for 50-60% of cases²³. Osimertinib is also indicated for NSCLC patients harboring EGFR T790M mutations whose disease has progressed on or after treatment with a first line TKI. EGFR targeting antibodies including cetuximab³⁹ (2004), panitumumab⁴⁰ (2006), and necitumumab⁴¹ (2016) are also under investigation in combination with EGFR-targeting TKIs for efficacy against EGFR mutations. The use of cetuximab in combination with afatinib is currently recommended by the NCCN for patients who have progressed after receiving erlotinib, afatinib, dacomitinib, or gefitinib and chemotherapy⁴².

Relevant Therapy Summary

● In this cancer type
 ○ In other cancer type
 ◐ In this cancer type and other cancer types
 ✗ Contraindicated
 ⚠ Both for use and contraindicated
 ✖ No evidence

EGFR exon 20 insertion

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
osimertinib	✗	●	✗	✗	● (II)

* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.



Relevant Therapy Summary (continued)

● In this cancer type ○ In other cancer type ⓘ In this cancer type and other cancer types ⛔ Contraindicated ⚠ Both for use and contraindicated ✕ No evidence

EGFR exon 20 insertion (continued)

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
gefitinib	✕	✕	⛔	✕	● (III)
apatinib + erlotinib, apatinib + gefitinib, apatinib + icotinib hydrochloride	✕	✕	✕	✕	● (IV)
apatinib + gefitinib	✕	✕	✕	✕	● (IV)
gefitinib, radiation therapy	✕	✕	✕	✕	● (IV)
icotinib hydrochloride, radiation therapy	✕	✕	✕	✕	● (IV)
atezolizumab, bevacizumab, chemotherapy	✕	✕	✕	✕	● (III)
bevacizumab (Innovent Biologics), chemotherapy, sintilimab	✕	✕	✕	✕	● (III)
bevacizumab + chemotherapy, bevacizumab (Shanghai Hengrui Pharmaceutical) + chemotherapy	✕	✕	✕	✕	● (III)
chemotherapy, durvalumab	✕	✕	✕	✕	● (III)
chemotherapy, icotinib hydrochloride	✕	✕	✕	✕	● (III)
chemotherapy, nivolumab	✕	✕	✕	✕	● (III)
chemotherapy, toripalimab	✕	✕	✕	✕	● (III)
afatinib, bevacizumab	✕	✕	✕	✕	● (II)
afatinib, chemotherapy, radiation therapy	✕	✕	✕	✕	● (II)
anlotinib hydrochloride + sintilimab	✕	✕	✕	✕	● (II)
apatinib + chemotherapy	✕	✕	✕	✕	● (II)
bevacizumab, osimertinib	✕	✕	✕	✕	● (II)
chemotherapy, ramucirumab	✕	✕	✕	✕	● (II)
chemotherapy, targeted therapy	✕	✕	✕	✕	● (II)
erlotinib	✕	✕	✕	✕	● (II)
erlotinib + chemotherapy	✕	✕	✕	✕	● (II)
erlotinib, radiation therapy	✕	✕	✕	✕	● (II)
gefitinib + chemotherapy	✕	✕	✕	✕	● (II)

* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.



Relevant Therapy Summary (continued)

● In this cancer type ○ In other cancer type ● In this cancer type and other cancer types ⛔ Contraindicated ⚠ Both for use and contraindicated ✕ No evidence

EGFR exon 20 insertion (continued)

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
icotinib hydrochloride	✕	✕	✕	✕	● (II)
ipilimumab, nivolumab	✕	✕	✕	✕	● (II)
KN046	✕	✕	✕	✕	● (II)
poziotinib	✕	✕	✕	✕	● (II)
radiation therapy, tyrosine kinase inhibitors	✕	✕	✕	✕	● (II)
sintilimab	✕	✕	✕	✕	● (II)
sunitinib	✕	✕	✕	✕	● (II)
tarloxotinib	✕	✕	✕	✕	● (II)
afatinib + necitumumab	✕	✕	✕	✕	● (I/II)
bevacizumab + erlotinib + chemotherapy	✕	✕	✕	✕	● (I/II)
DZD-9008	✕	✕	✕	✕	● (I/II)
EMB01	✕	✕	✕	✕	● (I/II)
gefitinib + ningetinib	✕	✕	✕	✕	● (I/II)
icotinib hydrochloride + chemotherapy + radiation therapy	✕	✕	✕	✕	● (I/II)
oleclumab + osimertinib	✕	✕	✕	✕	● (I/II)
TAK788	✕	✕	✕	✕	● (I/II)
APG-1252, osimertinib	✕	✕	✕	✕	● (I)
cetuximab, FATE-NK100	✕	✕	✕	✕	● (I)
durvalumab + oleclumab, oleclumab	✕	✕	✕	✕	● (I)
everolimus + neratinib, neratinib + palbociclib, neratinib + trametinib	✕	✕	✕	✕	● (I)
JNJ-61186372	✕	✕	✕	✕	● (I)
necitumumab, osimertinib	✕	✕	✕	✕	● (I)
osimertinib, osimertinib + radiation therapy	✕	✕	✕	✕	● (I)
pirotinib	✕	✕	✕	✕	● (I)

* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.



Relevant Therapy Summary (continued)

● In this cancer type
 ○ In other cancer type
 ⓘ In this cancer type and other cancer types
 ⛔ Contraindicated
 ⚠ Both for use and contraindicated
 ✕ No evidence

EGFR exon 20 insertion (continued)

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
TP-0903	✕	✕	✕	✕	● (I)
tyrosine kinase inhibitors, tyrosine kinase inhibitors + chemotherapy	✕	✕	✕	✕	● (I)

* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.

Relevant Therapy Details

Current NCCN Information

● In this cancer type
 ○ In other cancer type
 ⓘ In this cancer type and other cancer types
 ⛔ Contraindicated
 🗨 Not recommended
 🛡 Resistance

NCCN information is current as of 2019-08-15. For the most up-to-date information, search www.nccn.org.
 For NCCN International Adaptations & Translations, search www.nccn.org/global/international_adaptations.aspx.

EGFR exon 20 insertion

● osimertinib

Cancer type: Non-Small Cell Lung Cancer

Variant class: EGFR mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- Non-Small Cell Lung Cancer; Brain metastases; Newly diagnosed (Not specified)
- Non-Small Cell Lung Cancer; Leptomeningeal and Spine metastases (Not specified)

Reference: NCCN Guidelines® - NCCN-Central Nervous System Cancers [Version 1.2019]

🗨 pembrolizumab

Cancer type: Non-Small Cell Lung Cancer

Variant class: EGFR mutation

Other criteria: CD274 overexpression

Summary:

NCCN Guidelines® include the following supporting statement(s):

- "A small study suggests that single-agent pembrolizumab is not effective as first-line therapy in patients with metastatic NSCLC and EGFR mutations, even those with PD-L1 levels more than 50%."

Reference: NCCN Guidelines® - NCCN-Non-Small Cell Lung Cancer [Version 7.2019]



EGFR exon 20 insertion (continued)

EGFR tyrosine kinase inhibitor

Cancer type: Non-Small Cell Lung Cancer

Variant class: EGFR exon 20 insertion

Summary:

NCCN Guidelines® include the following supporting statement(s):

- "Patients with EGFR exon 20 insertion mutations are usually resistant to TKIs, although there are rare exceptions."

Reference: NCCN Guidelines® - NCCN-Non-Small Cell Lung Cancer [Version 7.2019]



Current EMA Information

☒ In this cancer type ☐ In other cancer type ☐ In this cancer type and other cancer types ☒ Contraindicated ☒ Not recommended ☒ Resistance

EMA information is current as of 2019-11-25. For the most up-to-date information, search www.ema.europa.eu/ema.

EGFR exon 20 insertion

☒ gefitinib

Cancer type: Non-Small Cell Lung Cancer

Label as of: 2019-05-28

Variant class: EGFR exon 20 insertion

Reference:

https://www.ema.europa.eu/en/documents/product-information/iressa-epar-product-information_en.pdf

Signatures

Testing Personnel:

Laboratory Supervisor:

Pathologist:



References

1. Valenta et al. The many faces and functions of β -catenin. *EMBO J.* 2012 Jun 13;31(12):2714-36. PMID: 22617422
2. Korinek et al. Constitutive transcriptional activation by a beta-catenin-Tcf complex in APC-/- colon carcinoma. *Science.* 1997 Mar 21;275(5307):1784-7. PMID: 9065401
3. Aberle et al. beta-catenin is a target for the ubiquitin-proteasome pathway. *EMBO J.* 1997 Jul 1;16(13):3797-804. PMID: 9233789
4. Winston et al. The SCFbeta-TRCP-ubiquitin ligase complex associates specifically with phosphorylated destruction motifs in IkappaBalpha and beta-catenin and stimulates IkappaBalpha ubiquitination in vitro. *Genes Dev.* 1999 Feb 1;13(3):270-83. PMID: 9990852
5. Kitagawa et al. An F-box protein, FWD1, mediates ubiquitin-dependent proteolysis of beta-catenin. *EMBO J.* 1999 May 4;18(9):2401-10. PMID: 10228155
6. Liu et al. Control of beta-catenin phosphorylation/degradation by a dual-kinase mechanism. *Cell.* 2002 Mar 22;108(6):837-47. PMID: 11955436
7. Miyoshi et al. Activation of the beta-catenin gene in primary hepatocellular carcinomas by somatic alterations involving exon 3. *Cancer Res.* 1998 Jun 15;58(12):2524-7. PMID: 9635572
8. Gao et al. Exon 3 mutations of CTNNB1 drive tumorigenesis: a review. *Oncotarget.* 2018 Jan 12;9(4):5492-5508. PMID: 29435196
9. Morin et al. Activation of beta-catenin-Tcf signaling in colon cancer by mutations in beta-catenin or APC. *Science.* 1997 Mar 21;275(5307):1787-90. PMID: 9065402
10. Schulze et al. Exome sequencing of hepatocellular carcinomas identifies new mutational signatures and potential therapeutic targets. *Nat. Genet.* 2015 May;47(5):505-511. PMID: 25822088
11. Ahn et al. Genomic portrait of resectable hepatocellular carcinomas: implications of RB1 and FGF19 aberrations for patient stratification. *Hepatology.* 2014 Dec;60(6):1972-82. PMID: 24798001
12. Harding et al. Prospective Genotyping of Hepatocellular Carcinoma: Clinical Implications of Next-Generation Sequencing for Matching Patients to Targeted and Immune Therapies. *Clin. Cancer Res.* 2018 Oct 29. PMID: 30373752
13. Cancer et al. Integrated genomic characterization of endometrial carcinoma. *Nature.* 2013 May 2;497(7447):67-73. PMID: 23636398
14. Soumerai et al. Clinical Utility of Prospective Molecular Characterization in Advanced Endometrial Cancer. *Clin. Cancer Res.* 2018 Dec 1;24(23):5939-5947. PMID: 30068706
15. Weinstein et al. The Cancer Genome Atlas Pan-Cancer analysis project. *Nat. Genet.* 2013 Oct;45(10):1113-20. PMID: 24071849
16. Cerami et al. The cBio cancer genomics portal: an open platform for exploring multidimensional cancer genomics data. *Cancer Discov.* 2012 May;2(5):401-4. PMID: 22588877
17. Blakely et al. Evolution and clinical impact of co-occurring genetic alterations in advanced-stage EGFR-mutant lung cancers. *Nat. Genet.* 2017 Dec;49(12):1693-1704. PMID: 29106415
18. King et al. Amplification of a novel v-erbB-related gene in a human mammary carcinoma. *Science.* 1985 Sep 6;229(4717):974-6. PMID: 2992089
19. ErbB Receptors and Cancer. *Methods Mol. Biol.* 2017;1652:3-35. PMID: 28791631
20. Gutierrez et al. HER2: biology, detection, and clinical implications. *Arch. Pathol. Lab. Med.* 2011 Jan;135(1):55-62. PMID: 21204711
21. Pines et al. Oncogenic mutant forms of EGFR: lessons in signal transduction and targets for cancer therapy. *FEBS Lett.* 2010 Jun 18;584(12):2699-706. PMID: 20388509
22. Cancer Genome Atlas Research Network. Comprehensive molecular profiling of lung adenocarcinoma. *Nature.* 2014 Jul 31;511(7511):543-50. doi: 10.1038/nature13385. Epub 2014 Jul 9. PMID: 25079552
23. da et al. EGFR mutations and lung cancer. *Annu Rev Pathol.* 2011;6:49-69. doi: 10.1146/annurev-pathol-011110-130206. PMID: 20887192
24. Arcila et al. EGFR exon 20 insertion mutations in lung adenocarcinomas: prevalence, molecular heterogeneity, and clinicopathologic characteristics. *Mol. Cancer Ther.* 2013 Feb;12(2):220-9. PMID: 23371856



References (continued)

25. Kobayashi et al. EGFR Exon 18 Mutations in Lung Cancer: Molecular Predictors of Augmented Sensitivity to Afatinib or Neratinib as Compared with First- or Third-Generation TKIs. *Clin Cancer Res.* 2015 Dec 1;21(23):5305-13. doi: 10.1158/1078-0432.CCR-15-1046. Epub 2015 Jul 23. PMID: 26206867
26. Yasuda et al. Structural, biochemical, and clinical characterization of epidermal growth factor receptor (EGFR) exon 20 insertion mutations in lung cancer. *Sci Transl Med.* 2013 Dec 18;5(216):216ra177. PMID: 24353160
27. Chiu et al. Epidermal Growth Factor Receptor Tyrosine Kinase Inhibitor Treatment Response in Advanced Lung Adenocarcinomas with G719X/L861Q/S768I Mutations. *J Thorac Oncol.* 2015 May;10(5):793-9. PMID: 25668120
28. Karachaliou et al. KRAS mutations in lung cancer. *Clin Lung Cancer.* 2013 May;14(3):205-14. PMID: 23122493
29. Brennan et al. The somatic genomic landscape of glioblastoma. *Cell.* 2013 Oct 10;155(2):462-77. PMID: 24120142
30. Cancer Genome Atlas Network. Comprehensive genomic characterization of head and neck squamous cell carcinomas. *Nature.* 2015 Jan 29;517(7536):576-82. PMID: 25631445
31. Mitsudomi et al. Epidermal growth factor receptor in relation to tumor development: EGFR gene and cancer. *FEBS J.* 2010 Jan;277(2):301-8. PMID: 19922469
32. Ji et al. Epidermal growth factor receptor variant III mutations in lung tumorigenesis and sensitivity to tyrosine kinase inhibitors. *Proc. Natl. Acad. Sci. U.S.A.* 2006 May 16;103(20):7817-22. PMID: 16672372
33. Gazdar. Activating and resistance mutations of EGFR in non-small-cell lung cancer: role in clinical response to EGFR tyrosine kinase inhibitors. *Oncogene.* 2009 Aug;28 Suppl 1:S24-31. PMID: 19680293
34. https://www.accessdata.fda.gov/drugsatfda_docs/label/2016/021743s025lbl.pdf
35. https://www.accessdata.fda.gov/drugsatfda_docs/label/2019/201292s015lbl.pdf
36. https://www.accessdata.fda.gov/drugsatfda_docs/label/2018/206995s003lbl.pdf
37. https://www.accessdata.fda.gov/drugsatfda_docs/label/2018/208065s011lbl.pdf
38. https://www.accessdata.fda.gov/drugsatfda_docs/label/2018/211288s000lbl.pdf
39. https://www.accessdata.fda.gov/drugsatfda_docs/label/2019/125084s273lbl.pdf
40. https://www.accessdata.fda.gov/drugsatfda_docs/label/2017/125147s207lbl.pdf
41. https://www.accessdata.fda.gov/drugsatfda_docs/label/2015/125547s000lbl.pdf
42. NCCN Guidelines® - NCCN-Non-Small Cell Lung Cancer [Version 7.2019]