





<b>REPORTING</b>	<b>PHYSICIAN</b>	 <i>Complete Tumor Response Map on page 2</i>
Report Date:	Teh-Ying Chou	
Receipt Date:	Account: Genconn Biotech Co., LTD	
Collection Date:	Address: F15., No 207-5 Sec 3 , Beixin Rd, New Taipei City, Xindian Dist, 23143, Taiwan	
Specimen:	Ph: +886 963 820 633   Fax: N/A	
Status:	Additional Recipient: N/A	

Summary of Detected Somatic Alterations, Immunotherapy Biomarkers & Associated Treatment Options

KEY  Approved in indication  Approved in other indication  Lack of response

Detected Alteration(s) / Biomarker(s)	Associated FDA-approved therapies	Clinical trial availability (see page 3)	% cfDNA or Amplification
TP53 P278L	None	Yes	0.8%
TP53 H179R	None	Yes	0.1%
EGFR Amplification	None	Yes	Low (+)

Variants of Uncertain Clinical Significance  
CDK12 S590A (0.4%)  
The functional consequences and/or clinical significance of alterations are unknown. Relevance of therapies targeting these alterations is uncertain.

Additional Biomarkers

Biomarker	Additional Details
MSI-High	NOT DETECTED

Alterations or biomarkers that were "NOT DETECTED" have been excluded from the summary table above.

We evaluated this sample for 74 genes, including the following guideline-recommended genes for NSCLC

EGFR(T790M and others)

ALK

ROS1

BRAF

MET

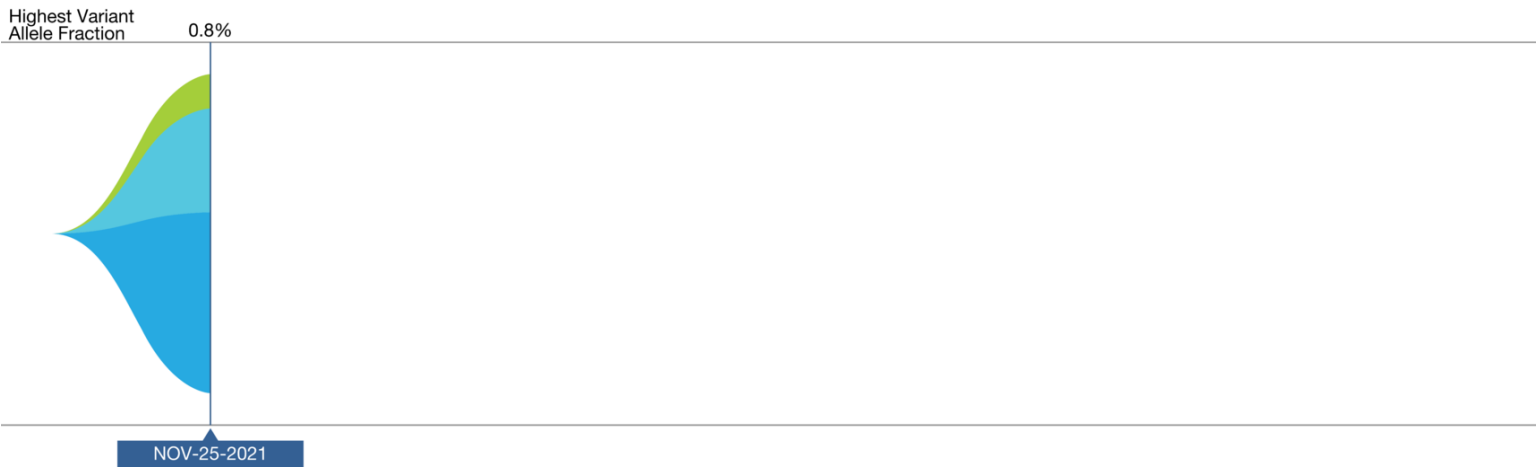
ERBB2(HER2)

RET

NTRK

Guardant360 Tumor Response Map

The Guardant360 Tumor Response Map illustrates the variant allele fraction (% cfDNA) of observed somatic variants at each sample submission. Amplifications are not plotted, and only the first and last five test dates are plotted. Please see the Physician Portal ([portal.guardanthealth.com](https://portal.guardanthealth.com)) for the Tumor Response Map with all test dates.



Detected Alteration(s) / Biomarker(s)	% cfDNA or Amp	
TP53 P278L	0.8%	
CDK12 S590A	0.4%	Variants of Uncertain Clinical Significance §
TP53 H179R	0.1%	
EGFR Amplification Amplifications not graphed above	Low (+) Plasma Copy Number 2.3	

The table above annotates the variant allele fraction (% cfDNA) detected in this sample, listed in descending order.  
§ See definitions section for more detail

Available Clinical Trials (within the same state as the ordering physician)

There may be additional trials not listed here. Visit: [portal.guardanthealth.com](https://portal.guardanthealth.com) or email [clientservices@guardanthealth.com](mailto:clientservices@guardanthealth.com) with A0435653 in the subject line of the email, for additional trials.

Alteration	Trial ID / Contact	Title	Phase	Site(s)
EGFR Amplification	NCT02609776 Use link at the bottom of the page to see if you qualify for an enrolling site (see list). If you still have questions:,JNJ. CT@sylogent.com	Study of Amivantamab, a Human Bispecific EGFR and cMet Antibody, in Participants With Advanced Non-Small Cell Lung Cancer	Phase 1	Kaohsiung, Taiwan Taipei City, Taiwan Taipei, Taiwan Taichung, Taiwan (3)
	NCT04077463 Study Contact,JNJ.CT@sylogent.com,844-434-4210	A Study of Lazertinib as Monotherapy or in Combination With Amivantamab in Participants With Advanced Non-small Cell Lung Cancer	Phase 1	Kaohsiung, Taiwan Taipei City, Taiwan Tainan, Taiwan Taichung, Taiwan
	Visit <a href="https://portal.guardanthealth.com">portal.guardanthealth.com</a> for trials not within the same state as the physician's office			
TP53 P278L	Visit <a href="https://portal.guardanthealth.com">portal.guardanthealth.com</a> for trials not within the same state as the physician's office			
TP53 H179R	Visit <a href="https://portal.guardanthealth.com">portal.guardanthealth.com</a> for trials not within the same state as the physician's office			

More clinical trial options available at [portal.guardanthealth.com](https://portal.guardanthealth.com)

## Definitions

**Variants of Uncertain Clinical Significance:** The functional consequences and/or clinical significance of alterations are unknown. Relevance of therapies targeting these alterations is uncertain.

**Amplification:** Guardant360 detects amplifications in the genes listed in Table 1. Gene amplification results in increased copies of the gene present in the cfDNA. The reported absolute copy number value represents the average copy number for the detected gene that was detected in circulating cfDNA. With the exception of sex-linked genes such as *AR*, 2 copies are expected in the absence of amplification. As the absolute number of copies in circulation is dependent on both tumor fraction and the magnitude of the tumor amplification, amplifications are reported on a semi-quantitative scale.

For *CCNE1*, *EGFR* and *FGFR1*, three levels are reported:

Low (+): Amplification magnitude is below the 50th percentile of amplifications detected by Guardant360.

Medium (++) : Amplification magnitude is between the 50th and 90th percentiles.

High (+++) : Amplification magnitude is above the 90th percentile.

For *BRAF*, *CCND1*, *CCND2*, *CDK4*, *CDK6*, *ERBB2*, *FGFR2*, *KIT*, *KRAS*, *MET*, *PDGFRA*, *RAF1*, *MYC*, *PIK3CA* and *AR*, two levels are reported:

Medium (++) : Amplification magnitude is below the 50th percentile of amplifications detected by Guardant360.

High (+++) : Amplification magnitude is above the 50th percentile.

## Interpretation

Somatic alterations were detected in the circulating cell-free DNA isolated from this patient's blood specimen. These genomic alterations are cancer-associated somatic variants, some of which have been associated with either increased or reduced clinical response to specific treatments. The percentage of altered cell-free DNA circulating (% cfDNA) in blood is related to the unique tumor biology of each patient. Factors that may affect the % cfDNA of detected somatic alterations include tumor growth, turn over, size, heterogeneity, vascularization, disease progression, and treatment.

Amplification was detected in the circulating cell-free DNA isolated from this patient's blood specimen for the annotated gene(s). Unlike tissue-based gene amplification tests (e.g. IHC or FISH), Guardant360 assesses the total representation of a given gene in all circulating cell-free DNA present in the patient's blood sample including material derived from the tumor and healthy tissue alike. As such, the absolute level of amplification present in the blood depends both on the tumor-derived cfDNA content and on the degree of amplification within that fraction and cannot be inferred from bulk cfDNA interrogation. For example, a positive Guardant360 test could represent a small population of cells with extremely high levels of the detected gene amplification. Alternatively, it could represent a large population of cells with low to medium levels of the detected gene amplifications. The exact correlation between amplification detected by Guardant360 compared to IHC or FISH and how each test differentially guides patient management is an area of active investigation.

Method and Limitations

Guardant360 sequences 74 cancer-associated genes to identify somatic alterations. Cell-free DNA (cfDNA) is extracted from plasma, enriched for targeted regions, and sequenced using the Illumina platform and hg19 as the reference genome. All exons are sequenced in some genes; only clinically significant exons are sequenced in other genes. The types of genomic alterations detected by Guardant360 include single nucleotide variants, gene amplifications, fusions, short insertions/deletions (longest detected, 70 base pairs), and splice site disrupting events (see Table 1). Microsatellite Instability (MSI) is assessed for all cancer types by evaluating somatic changes in the length of repetitive sequences on the Guardant360 panel. A "Not Detected" result in samples where the highest % cfDNA is < 0.2% is an inconclusive result because it does not preclude MSI-High status in tissue. MSI status is currently not reported for earlier panel versions. This version of the Guardant360 test is not validated for the detection of other types of genomic alterations, such as complex rearrangements or gene deletions. Certain sample or variant characteristics, such as low cfDNA concentration, may result in reduced analytic sensitivity. Guardant360 cannot discern the source of circulating cfDNA, and for some variants in the range of ~40 to 60% cfDNA, the test cannot easily distinguish germline variants from somatic alterations. Guardant360 is not validated for the detection of germline or de novo variants that are associated with hereditary cancer risk. Tissue genotyping should be considered when plasma genotyping is negative, if clinically appropriate.

Table 1: Genes on the Guardant360 Panel

Guardant360 reports single nucleotide variants, splice site mutations, and insertion and deletion variants (indels) in all clinically relevant exons in 74 genes and reports other variant types in select genes as indicated below.

AKT1	ALK #	APC	AR †	ARAF	ARID1A	ATM	BRAF †	BRCA1
BRCA2	CCND1 †	CCND2 †	CCNE1 †	CDH1	CDK12	CDK4 †	CDK6 †	CDKN2A
CTNNB1	DDR2	EGFR †	ERBB2 †	ESR1	EZH2	FBXW7	FGFR1 †	FGFR2 # †
FGFR3 #	GATA3	GNA11	GNAQ	GNAS	HNF1A	HRAS	IDH1	IDH2
JAK2	JAK3	KIT †	KRAS †	MAP2K1	MAP2K2	MAPK1	MAPK3	MET †
MLH1	MPL	MTOR	MYC †	NF1	NFE2L2	NOTCH1	NPM1	NRAS
NTRK1 #	NTRK3	PDGFRA †	PIK3CA †	PTEN	PTPN11	RAF1 †	RB1	RET #
RHEB	RHOA	RIT1	ROS1 #	SMAD4	SMO	STK11	TERT ‡	TP53
TSC1	VHL							

‡ Guardant360 reports alterations in the promoter region of this gene.

# Guardant360 reports fusion events involving this gene.

† Guardant360 reports amplifications of this gene.

About the Test

The Guardant360 assay was developed and its performance characteristics were determined by Guardant Health, Inc. This test has not been cleared or approved by the U.S. Food and Drug Administration (FDA). The FDA has determined that such clearance or approval is not necessary. This test may be used for clinical purposes and should not be regarded as investigational or for research only. Guardant Health's clinical reference laboratory is certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA) as qualified to perform high complexity clinical laboratory testing. The laboratory report should be interpreted in the context of other clinical information and laboratory, pathology, and imaging studies by a qualified medical professional prior to initiating or changing a patient's treatment plan. The selection of any, all, or none of the drugs associated with potential clinical benefit (or potential lack of clinical benefit) is entirely at the discretion of the treating medical professional. Drug and trial information are based on the diagnosis written on the submitted test request form; this information is not based on any supplemental information provided by the requesting medical professional, including pathology reports or other molecular studies. Some drugs listed in this report may not be approved or cleared by the FDA for the indicated use. Guardant Health makes no endorsement, express or implied, of any product, physician, or procedure contained in this report. This report makes no promises or guarantees that a particular medication will affect (or not affect) the clinical outcome of any patient.

Testing performed at: Guardant Health  
Laboratory Director: Martina Lefterova, MD PhD | CLIA ID: 05D2070300 | CAP #: 8765297 | 505 Penobscot Drive, Redwood City, CA, 94063, USA

### Additional information is available

Any therapeutic annotations are based on publicly available information. This information is described in the "Detailed Therapy Results" and "Relevance of Detected Alterations" sections.

Visit [portal.guardanthealth.com](https://portal.guardanthealth.com) or email [clientservices@guardanthealth.com](mailto:clientservices@guardanthealth.com) with A0435653 in the subject line of the email for:

- Additional clinical trials
- Detailed Therapy Results
- Relevance of Detected Alterations
- References

If you would like to receive this additional information with every Guardant360 report, please call client services at [855.698.8887](tel:855.698.8887) to opt-in.

**Additional information begins on the next page.**

## List of Available Clinical Trials

Alteration	Trial ID / Contact	Title	Phase	Site (number in parenthesis is count of trial sites)
EGFR Amplification	NCT01553942 Lecia V Sequist, MD MPH, lvsquist@partners.org, 617-724-4000	Afatinib With CT and RT for EGFR-Mutant NSCLC	Phase 2	Boston, MA (2)
	NCT02609776 Use link at the bottom of the page to see if you qualify for an enrolling site (see list). If you still have questions:,JNJ. CT@sylogent.com	Study of Amivantamab, a Human Bispecific EGFR and cMet Antibody, in Participants With Advanced Non-Small Cell Lung Cancer	Phase 1	Detroit, MI; Saint Louis, MO; Rochester, MN; Philadelphia, PA; Bethesda, MD; Boston, MA; Chicago, IL; Portland, OR; Tampa, FL; Fairfax, VA; New York, NY (3); CA (5); Canada; Japan (10); China (16); Taiwan (6); Korea, Republic of (8); United Kingdom (3); Italy (3); France (7); Australia (4); Spain (11)
	NCT02795156 Sarah Cannon Development Innovations, LLC, CANN. InnovationsMedical@sarahcannon.com, 844-710-6157	Study Assessing Activity of Molecularly Matched Targeted Therapies in Select Tumor Types Based on Genomic Alterations	Phase 2	Denver, CO; Milwaukee, WI; West Palm Beach, FL; Fort Myers, FL; Saint Petersburg, FL; Nashville, TN; Chattanooga, TN; Kansas City, MO
	NCT02947386 See <a href="https://clinicaltrials.gov/show/NCT02947386">https://clinicaltrials.gov/show/NCT02947386</a>	Nimotuzumab and Nivolumab in Treating Patients With Advanced Non-small Cell Lung Cancer	Phase 1 /Phase 2	Buffalo, NY
	NCT04077463 Study Contact, JNJ.CT@sylogent.com, 844-434-4210	A Study of Lazertinib as Monotherapy or in Combination With Amivantamab in Participants With Advanced Non-small Cell Lung Cancer	Phase 1	Seattle, WA; Detroit, MI; Saint Louis, MO; Philadelphia, PA; Baltimore, MD; Portland, OR; Salt Lake City, UT; Tampa, FL; Fairfax, VA; Boston, MA (3); New York, NY (2); CA (5); Puerto Rico; Japan (7); China (14); Taiwan (4); Korea, Republic of (4); Italy (5); France (7); Germany (11); Spain (8)
	NCT04310007 See <a href="https://clinicaltrials.gov/show/NCT04310007">https://clinicaltrials.gov/show/NCT04310007</a>	Testing the Addition of the Pill Chemotherapy, Cabozantinib, to the Standard Immune Therapy Nivolumab Compared to Standard Chemotherapy for Non-small Cell Lung Cancer	Phase 2	Mount Holly, NJ; Savannah, GA; Phoenix, AZ; Manchester, NH; Worcester, MA; Baltimore, MD; Fargo, ND; Richmond, IN; Bryan, TX; Fort Lauderdale, FL; Fort Smith, AR; Concord, NH; Huntington, WV; Cody, WY; Boston, MA; Birmingham, AL; Greenwood, SC; South Bend, IN; Kingman, AZ; West Haven, CT; Oklahoma City, OK; Glens Falls, NY; Sheridan, WY; Voorhees, NJ; Miami Beach, FL; Albuquerque, NM; Hot Springs, AR; Aventura, FL; Jamestown, ND; Cheyenne, WY; Moorestown, NJ; Bronx, NY (3); Atlanta, GA (2); NV (45); WA (28); WI (34); IA (22); OH (39); ID (13); MI (60); CA (7); MN (38); MO (30); OR (11); IL (49); MT (8); KS (9); AK (9); VA (7); CO (40); KY (10); PA (22); NC (5); LA (6); NE (12)
	NCT04394624 Trial Transparency email recommended (Toll free number for US & Canada), Contact-US@sanofi.com, 800-633-1610 xoption 6	SAR408701 in Combination With Ramucirumab in Pre-treated Patients With Non Squamous Non-small Cell Lung Cancer (NSQ NSCLC)	Phase 2	Detroit, MI; Whittier, CA; Buffalo, NY; Waco, TX; Czechia (3); Korea, Republic of (2); Bulgaria (2); Portugal (2); Spain (5)
	NCT04606381 Study Contact, JNJ.CT@sylogent.com, 844-434-4210	A Study of Amivantamab Subcutaneous (SC) Administration for the Treatment of Advanced Solid Malignancies	Phase 1	West Hollywood, CA; Indianapolis, IN; New York, NY; Portland, OR; Nashville, TN; Canada; Korea, Republic of (2); United Kingdom (2)



## List of Available Clinical Trials

Alteration	Trial ID / Contact	Title	Phase	Site (number in parenthesis is count of trial sites)
	NCT04965090 Helena Yu, MD,yuh@mskcc.org,646-608-3912	A Study of Amivantamab and Lazertinib in People With Non-Small Cell Lung Cancer (NSCLC)	Phase 2	Basking Ridge, NJ; Uniondale, NY; Harrison, NY; Montvale, NJ; Commack, NY; New York, NY; Middletown, NJ
TP53 P278L	NCT04383938 Eyal Attar, MD,info@aprea.com,+1 617 804 6947	Phase 1/2 Study of APR-246 in Combination With Pembrolizumab in Subjects With Solid Tumor Malignancies	Phase 1 /Phase 2	Houston, TX; Saint Louis, MO; Rochester, MN; Phoenix, AZ; Jacksonville, FL; Nashville, TN; Boston, MA (2)
	NCT04555837 Faye M. Johnson,fmjohns@mdanderson.org,713-792-6363	Alisertib and Pembrolizumab for the Treatment of Patients With Rb-deficient Head and Neck Squamous Cell Cancer	Phase 1 /Phase 2	Houston, TX
	NCT04695223 See <a href="https://clinicaltrials.gov/show/NCT04695223">https://clinicaltrials.gov/show/NCT04695223</a>	Arsenic Trioxide for Structural p53 Mutations	Phase 2	China
	NCT04742959 Peng Peng, Ph.D., peng_peng@transtherabio.com,86-25-86901107	Study of TT-00420 Tablet as Monotherapy and Combination Therapy in Patients With Advanced Solid Tumors	Phase 1 /Phase 2	Houston, TX; Duarte, CA; New Brunswick, NJ
	NCT04869475 Min Shi, MD & Ph. D,sm11998@rjh.com.cn,+86-21-64370045	Arsenic Trioxide in Refractory Solid Tumors With Rescuable p53 Mutation	Phase 2	China
TP53 H179R	NCT04383938 Eyal Attar, MD,info@aprea.com,+1 617 804 6947	Phase 1/2 Study of APR-246 in Combination With Pembrolizumab in Subjects With Solid Tumor Malignancies	Phase 1 /Phase 2	Houston, TX; Saint Louis, MO; Rochester, MN; Phoenix, AZ; Jacksonville, FL; Nashville, TN; Boston, MA (2)
	NCT04555837 Faye M. Johnson,fmjohns@mdanderson.org,713-792-6363	Alisertib and Pembrolizumab for the Treatment of Patients With Rb-deficient Head and Neck Squamous Cell Cancer	Phase 1 /Phase 2	Houston, TX
	NCT04695223 See <a href="https://clinicaltrials.gov/show/NCT04695223">https://clinicaltrials.gov/show/NCT04695223</a>	Arsenic Trioxide for Structural p53 Mutations	Phase 2	China
	NCT04742959 Peng Peng, Ph.D., peng_peng@transtherabio.com,86-25-86901107	Study of TT-00420 Tablet as Monotherapy and Combination Therapy in Patients With Advanced Solid Tumors	Phase 1 /Phase 2	Houston, TX; Duarte, CA; New Brunswick, NJ
	NCT04869475 Min Shi, MD & Ph. D,sm11998@rjh.com.cn,+86-21-64370045	Arsenic Trioxide in Refractory Solid Tumors With Rescuable p53 Mutation	Phase 2	China

## Detailed Therapy Results

Alteration	Drug	Trade Name	Target	Current Status
EGFR Amplification	ABBV-221		Anti-Egfr antibody drug conjugate.	Phase 1 (Solid Tumor)
	ABBV-321		Anti-Egfr antibody conjugated to monomethyl auristatin F.	Phase 1 (Solid Tumor) Phase 1 (Glioblastoma, Head and neck squamous cell carcinoma (HNSCC), Brain and Central Nervous System Tumors, Lung squamous cell carcinoma)
	ABT806		Anti-Egfr and EGFRvIII antibody.	Phase 1 (Solid Tumor) Phase 2 (Gastroesophageal junction carcinoma)
	Afatinib	Gilotrif	Irreversible pan-ErbB kinase inhibitor.	Phase 3 (Non-small cell lung carcinoma (NSCLC)) FDA Approved in other indications (Lung squamous cell carcinoma, EGFR-mutant NSCLC)
	Amivantamab	Rybrevant	Bispecific anti-Met/Egfr antibody.	Phase 3 (Non-small cell lung carcinoma (NSCLC)) FDA Approved in other indications (NSCLC with EGFR exon 20 insertion)
	ASP-1929		An antibody-dye conjugate comprised of cetuximab and IRDye700DX acting as photoimmunotherapy.	Phase 3 (Head and neck carcinoma)
	AVID100		Anti-Egfr antibody-drug conjugate with DM1.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 2 (Head and neck squamous cell carcinoma (HNSCC), Breast carcinoma (triple negative))
	BDTX-189		Irreversible Egfr/Her2 inhibitor.	Phase 2 (Solid Tumor)
	BLU-945		Fourth generation Egfr inhibitor targeting T790M and T790M /C797S mutations.	Phase 2 (Non-small cell lung carcinoma (NSCLC))
	CPGJ602		Recombinant human-mouse chimeric anti-Egfr monoclonal antibody.	Phase 2 (Colorectal carcinoma (CRC))
	D2C7-IT		Immunotoxin targeting both wild-type Egfr and Egfr-vIII.	Phase 1 (Glioblastoma)
	Depatuxizumab mafodotin		Anti-Egfr monoclonal antibody drug conjugate.	Phase 1 (Solid Tumor) Phase 3 (Glioblastoma, Glioma)
	DZD9008		Bispecific anti-Egfr/Her2 monoclonal antibody.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 2 (Non-Hodgkin lymphoma (NHL))
	EGFR(V)-EDV-Dox		Doxorubicin-loaded EGFR-targeting nanocells.	Phase 1 (Glioblastoma)
	GC1118		Anti-Egfr monoclonal antibody.	Phase 1 (Solid Tumor) Phase 2 (Gastric carcinoma, Glioblastoma, Gastroesophageal junction carcinoma)
	Hemay022		Egfr tyrosine kinase inhibitor.	Phase 1 (Breast carcinoma (HER2+))
	Lifirafenib		Dual Braf/Egfr inhibitor.	Phase 1 (Non-small cell lung carcinoma (NSCLC)) Phase 1 (Solid Tumor, Brain and Central Nervous System Tumors)
	MM-151		Anti-Egfr monoclonal antibody mixture.	Phase 1 (Non-small cell lung carcinoma (NSCLC)) Phase 1 (Solid Tumor)
	Neratinib	Nerlynx	Egfr/Her2/ErbB4 inhibitor.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) FDA Approved in other indications

## Detailed Therapy Results

Alteration	Drug	Trade Name	Target	Current Status
				(Breast carcinoma (HER2+))
	Nimotuzumab	Theraloc	Anti-Egfr monoclonal antibody.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 3 (Gastric carcinoma, Glioblastoma, Glioma, Pancreatic carcinoma, Head and neck carcinoma, Gastroesophageal junction carcinoma, Esophageal carcinoma, Cervical carcinoma)
	Pirotinib		ErbB family inhibitor.	Phase 1 (Solid Tumor)
	Pozotinib		Egfr/Her2/ErbB4 kinase inhibitor.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 2 (Gastric carcinoma, Head and neck squamous cell carcinoma (HNSCC), Breast carcinoma, Esophageal squamous cell carcinoma, Colorectal carcinoma (CRC))
	Pyrotinib		Egfr/Her2 kinase inhibitor.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 3 (Breast carcinoma)
	Ramucirumab	Cyramza	anti-VEGFR-2 monoclonal antibody.	FDA Approved in this indication (Non-small cell lung carcinoma (NSCLC)) FDA Approved in other indications (Gastric carcinoma, Hepatocellular carcinoma (HCC), Gastroesophageal junction carcinoma, Colorectal carcinoma (CRC))
	SKLB1028		Egfr/Flt3/c-Abl inhibitor.	Phase 2 (Acute myeloid leukemia (AML))
	Sym004		Anti-Egfr antibody mixture.	Phase 2 (Glioma, Head and neck squamous cell carcinoma (HNSCC), Colorectal carcinoma (CRC))
	SYN004		Anti-Egfr monoclonal antibody.	Phase 1 (Solid Tumor) Phase 1 (Lung squamous cell carcinoma)
	Varlitinib		Egfr/Her2 kinase inhibitor.	Phase 2 (Gastric carcinoma, Hepatocellular carcinoma (HCC), Pancreatic carcinoma, Cholangiocarcinoma)
TP53 P278L H179R	Adavosertib		Wee1 tyrosine kinase inhibitor.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 2 (Embryonal tumor with multi-layered rosettes (ETMR), Medulloblastoma, Small cell lung carcinoma (SCLC), Solid Tumor, Primary myelofibrosis (PMF), Ovarian carcinosarcoma, Acute myeloid leukemia (AML), MDS/MPN, unclassifiable, Chronic myelomonocytic leukemia (CMML), Peritoneal papillary serous carcinoma, Myelodysplastic Syndrome (MDS))
	Alisertib		Aurora A small molecule kinase inhibitor.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) Phase 3 (Peripheral T-cell lymphoma (PTCL))
	ALT-801		p53-targeted T-cell receptor-IL2 fusion.	Phase 1 (Solid Tumor) Phase 2 (Melanoma, Urothelial carcinoma, Bladder carcinoma, Urethral melanoma, Urethral carcinoma, Multiple myeloma (MM))
	AMG 900		Aurora A, B, C small molecule kinase inhibitor.	Phase 1 (Solid Tumor) Phase 1 (Acute myeloid leukemia (AML))
	AT9283		Aurora A, B, Jak2, Jak3, Bcr-Abl kinase inhibitor.	Phase 1 (Solid Tumor) Phase 2 (Acute myeloid leukemia (AML), Multiple myeloma (MM), Acute lymphoblastic leukemia (ALL))
	ATO	Trisenox	PML-RARA inhibitor. Inhibits multiple signaling pathways, including the Hedgehog pathway.	Phase 2 (Non-small cell lung carcinoma (NSCLC)) FDA Approved in other indications (Acute myeloid leukemia (AML), Acute promyelocytic leukemia)

Detailed Therapy Results

Alteration	Drug	Trade Name	Target	Current Status
				(APL))
	AZD2811		Nanoparticle formulation of Aurora kinase B inhibitor barasertib (AZD1152).	Phase 1 (Solid Tumor) Phase 2 (Acute myeloid leukemia (AML), Myelodysplastic Syndrome (MDS))
	COTI-2		Reactivates mutant p53.	Phase 1 (Endometrial carcinoma, Head and neck squamous cell carcinoma (HNSCC), Ovarian carcinoma, Cervical carcinoma)
	Debio 0123		Wee1 tyrosine kinase inhibitor.	Phase 1 (Solid Tumor)
	ENMD-2076		Aurora A kinase inhibitor.	Phase 1 (Solid Tumor) Phase 2 (Fibrolamellar hepatocellular carcinoma, Ovarian carcinoma, Breast carcinoma, Fallopian tube adenocarcinoma, Sarcoma)
	Eprenetapopt		Reactivates mutant p53.	Phase 3 (Myelodysplastic Syndrome (MDS))
	Kevetrin		Blocks Mdm2-p53 interaction, restoring transcriptional activity of p53.	Phase 1 (Solid Tumor) Phase 2 (Ovarian carcinoma)
	LY3295668		Aurora kinase A-specific inhibitor.	Phase 1 (Solid Tumor) Phase 2 (Head and neck squamous cell carcinoma (HNSCC), Small cell lung carcinoma (SCLC), Breast carcinoma (triple negative), Breast carcinoma (hormone receptor +, HER2-))
	SGT-53		TP53 gene therapy delivered via transferrin-targeted nanoparticles.	Phase 1 (Solid Tumor) Phase 2 (Glioblastoma, Glioma, Pancreatic carcinoma)
	SNS-314		Aurora kinase A/B inhibitor.	Phase 1 (Solid Tumor)
	TAS-119		Selective Aurora A kinase inhibitor.	Phase 1 (Solid Tumor)
	TT-00420		Aurora kinase A/B inhibitor.	Phase 1 (Solid Tumor) Phase 1 (Breast carcinoma (triple negative))

## Relevance of Detected Alterations

Alteration	Role in Disease	Effect on Drug Sensitivity	Effect on Drug Resistance
TP53 P278L	Loss of tumor suppressor p53, which is encoded by the TP53 gene, is common in aggressive advanced cancers. <sup>(1)</sup> Carriers of a germline mutation in TP53 have Li-Fraumeni Syndrome, an inherited cancer syndrome resulting in multiple tumors in early adulthood, including breast cancer, brain tumors, and leukemias. <sup>(2-4)</sup> Expression of p53 in normal cells is low; however, TP53 alterations, including those that result in loss of p53 tumor suppressor function, may lead to stabilization and increased expression of p53, particularly in the nucleus, and several studies have shown that it may have oncogenic gain-of-function effects. <sup>(5-9)</sup> TP53 alterations are believed to be early events in NSCLC, preceding lymph node metastasis. <sup>(10)</sup> TP53 mutation and expression of p53 have been correlated with the lung squamous cell carcinoma subtype, and p53 expression in lung squamous cell carcinoma has also been associated with disease stage and higher grade tumors. <sup>(11-14)</sup> TP53 mutation has been associated with PD-L1 expression and T-cell infiltration in lung adenocarcinoma samples. <sup>(15-19)</sup>	At present, there are no approved therapies targeting TP53 alterations, despite their high prevalence in cancer. Therapeutic approaches under investigation include gene therapy for TP53 and (dendritic cell-based) TP53 vaccines. <sup>(20-22)</sup> Inhibition of components of the DNA damage checkpoint, including Wee1, has been reported to enhance the activity of DNA-damaging agents in preclinical cancer models with deficiency of p53 function. <sup>(23-25)</sup> Clinical trials of the Wee1 inhibitor adavosertib (MK-1775) are currently underway for patients with solid tumors and hematologic malignancies. Studies have reported Aurora kinase A to be activated in cells harboring TP53 mutation, and Aurora kinase A and B inhibitors have been reported to activate wild-type p53 in cellular assays; thus, tumors retaining a wild-type TP53 allele may benefit from Aurora kinase inhibitors. <sup>(26-31)</sup>	Mutations in TP53 may increase resistance to ionizing radiation therapy. <sup>(32,33)</sup>
TP53 H179R	Loss of tumor suppressor p53, which is encoded by the TP53 gene, is common in aggressive advanced cancers. <sup>(1)</sup> Carriers of a germline mutation in TP53 have Li-Fraumeni Syndrome, an inherited cancer syndrome resulting in multiple tumors in early adulthood, including breast cancer, brain tumors, and leukemias. <sup>(2-4)</sup> Expression of p53 in normal cells is low; however, TP53 alterations, including those that result in loss of p53 tumor suppressor function, may lead to stabilization and increased expression of p53, particularly in the nucleus, and several studies have shown that it may have oncogenic gain-of-function effects. <sup>(5-9)</sup> TP53 alterations are believed to be early events in NSCLC, preceding lymph node metastasis. <sup>(10)</sup> TP53 mutation and expression of p53 have been correlated with the lung squamous cell carcinoma subtype, and p53 expression in lung squamous cell carcinoma has also been associated with disease stage and higher grade	At present, there are no approved therapies targeting TP53 alterations, despite their high prevalence in cancer. Therapeutic approaches under investigation include gene therapy for TP53 and (dendritic cell-based) TP53 vaccines. <sup>(20-22)</sup> Inhibition of components of the DNA damage checkpoint, including Wee1, has been reported to enhance the activity of DNA-damaging agents in preclinical cancer models with deficiency of p53 function. <sup>(23-25)</sup> Clinical trials of the Wee1 inhibitor adavosertib (MK-1775) are currently underway for patients with solid tumors and hematologic malignancies. Studies have reported Aurora kinase A to be activated in cells harboring TP53 mutation, and Aurora kinase A and B inhibitors have been reported to activate wild-type p53 in cellular assays; thus, tumors retaining a wild-type TP53 allele may benefit from Aurora kinase inhibitors. <sup>(26-31)</sup>	Mutations in TP53 may increase resistance to ionizing radiation therapy. <sup>(32,33)</sup>

Relevance of Detected Alterations

Alteration	Role in Disease	Effect on Drug Sensitivity	Effect on Drug Resistance
	tumors. <sup>(11-14)</sup> . TP53 mutation has been associated with PD-L1 expression and T-cell infiltration in lung adenocarcinoma samples. <sup>(15-19)</sup> .		
EGFR Amplification	The presence of an EGFR abnormality (mutation, amplification, or overexpression) can result in an overabundance or overactivity of Egfr protein, which can lead to excessive proliferation. <sup>(34)</sup> . EGFR mutations in NSCLC have been reported to occur more frequently in women, never-smokers, and in patients with adenocarcinoma histology. <sup>(35-44)</sup> .	EGFR amplification or increased copy number may result in elevated Egfr protein expression and thus predict sensitivity to Egfr targeted therapies. The Egfr tyrosine kinase inhibitors (TKIs) erlotinib, afatinib, gefitinib, osimertinib, and dacomitinib have been FDA approved for the treatment of non-small cell lung carcinoma (NSCLC) with specific EGFR mutations; however, only modest clinical benefit for gefitinib or erlotinib has been reported in patients harboring EGFR amplification without concurrent sensitizing mutations. <sup>(45-52)</sup> . Anti-Egfr monoclonal antibodies are also approved in some indications, including cetuximab, which is an approved therapy for head and neck and colorectal cancer, and panitumumab, which is approved in colorectal cancer. <sup>(53-55)</sup> . However, molecular analyses of tumor samples from a Phase 3 study in head and neck squamous cell carcinoma revealed that neither Egfr expression nor EGFR amplification predicted response to cetuximab. <sup>(56,57)</sup> . Studies have reported non-squamous NSCLC patients with metastatic disease and tumors harboring an EGFR exon 19 deletion or L858R mutation to be sensitive to osimertinib, erlotinib, afatinib, gefitinib, dacomitinib, and the combination of erlotinib plus ramucirumab. <sup>(45,58-63)</sup> . Less common activating EGFR mutations have variable sensitivity to EGFR tyrosine kinase inhibitors. <sup>(64)</sup> .	EGFR amplification has been reported as an acquired alteration in 15.5% (16 /103) of EGFR exon19del or L858R-mutant NSCLC patients who were treated with Egfr TKI and in 9.7% (3 /31) of patients treated with Egfr TKI in combination with bevacizumab. <sup>(65)</sup> .



## References

1. Brown C, Lain S, Verma C, Fersht A, Lane D "Awakening guardian angels: drugging the p53 pathway." *Nature reviews. Cancer*(2009): 862-73
2. Malkin D, Li F, Strong L, Fraumeni J, Nelson C, Kim D, Kassel J, Gryka M, Bischoff F, Tainsky M "Germ line p53 mutations in a familial syndrome of breast cancer, sarcomas, and other neoplasms." *Science (New York, N.Y.)*(1990): 1233-8
3. Srivastava S, Zou Z, Pirolo K, Blattner W, Chang E "Germ-line transmission of a mutated p53 gene in a cancer-prone family with Li-Fraumeni syndrome." *Nature*(1991): 747-9
4. Santibáñez-Koref M, Birch J, Hartley A, Jones P, Craft A, Eden T, Crowther D, Kelsey A, Harris M "p53 germline mutations in Li-Fraumeni syndrome." *Lancet (London, England)*(1991): 1490-1
5. Wang Y, Lin R, Tan Y, Chen J, Chen C, Wang Y "Wild-type p53 overexpression and its correlation with MDM2 and p14ARF alterations: an alternative pathway to non-small-cell lung cancer." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2005): 154-64
6. Koga T, Hashimoto S, Sugio K, Yoshino I, Nakagawa K, Yonemitsu Y, Sugimachi K, Sueishi K "Heterogeneous distribution of P53 immunoreactivity in human lung adenocarcinoma correlates with MDM2 protein expression, rather than with P53 gene mutation." *International journal of cancer*(2001): 232-9
7. Kato S, Han S, Liu W, Otsuka K, Shibata H, Kanamaru R, Ishioka C "Understanding the function-structure and function-mutation relationships of p53 tumor suppressor protein by high-resolution missense mutation analysis." *Proceedings of the National Academy of Sciences of the United States of America*(2003): 8424-9
8. Houben R, Hesbacher S, Schmid C, Kauczok C, Flohr U, Haferkamp S, Müller C, Schrama D, Wischhusen J, Becker J "High-level expression of wild-type p53 in melanoma cells is frequently associated with inactivity in p53 reporter gene assays." *PloS one*(2011): e22096
9. Olivier M, Petitjean A, Marcel V, Pétré A, Mounawar M, Plymoth A, de Fromental C, Hainaut P "Recent advances in p53 research: an interdisciplinary perspective." *Cancer gene therapy*(2009): 1-12
10. Chang Y, Wu C, Shih J, Lee Y "Comparison of p53 and epidermal growth factor receptor gene status between primary tumors and lymph node metastases in non-small cell lung cancers." *Annals of surgical oncology*(2011): 543-50
11. Jiang R, Zhang B, Teng X, Hu P, Xu S, Zheng Z, Liu R, Tang T, Ye F "Validating a targeted next-generation sequencing assay and profiling somatic variants in Chinese non-small cell lung cancer patients." *Scientific reports*(2020): 2070
12. Mattioni M, Soddu S, Prodosmo A, Visca P, Conti S, Alessandrini G, Facciolo F, Strigari L "Prognostic role of serum p53 antibodies in lung cancer." *BMC cancer*(2015): 148
13. Bircan A, Bircan S, Kapucuoglu N, Songur N, Ozturk O, Akkaya A "Maspin, VEGF and p53 expression in small biopsies of primary advanced lung cancer and relationship with clinicopathologic parameters." *Pathology oncology research : POR*(2010): 553-61
14. Kim Y, Hammerman P, Kim J, Yoon J, Lee Y, Sun J, Wilkerson M, Pedamallu C, Cibulskis K, Yoo Y, Lawrence M, Stojanov P, Carter S, McKenna A, Stewart C, Sivachenko A, Oh I, Kim H, Choi Y, Kim K, Shim Y, Kim K, Song S, Na K, Choi Y, Hayes D, Kim J, Cho S, Kim Y, Ahn J, Ahn M, Getz G, Meyerson M, Park K "Integrative and comparative genomic analysis of lung squamous cell carcinomas in East Asian patients." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2014): 121-8
15. Dong Z, Zhong W, Zhang X, Su J, Xie Z, Liu S, Tu H, Chen H, Sun Y, Zhou Q, Yang J, Yang X, Lin J, Yan H, Zhai H, Yan L, Liao R, Wu S, Wu Y "Potential Predictive Value of TP53 and KRAS Mutation Status for Response to PD-1 Blockade Immunotherapy in Lung Adenocarcinoma." *Clinical cancer research : an official journal of the American Association for Cancer Research*(2017): 3012-3024
16. Scheel A, Ansén S, Schultheis A, Scheffler M, Fischer R, Michels S, Hellmich M, George J, Zander T, Brockmann M, Stoelben E, Groen H, Timens W, Perner S, von Bergwelt-Baildon M, Büttner R, Wolf J "PD-L1 expression in non-small cell lung cancer: Correlations with genetic alterations." *Oncoimmunology*(2016): e1131379
17. Albitar M, Sudarsanam S, Ma W, Jiang S, Chen W, Funari V, Blocker F, Agersborg S "Correlation of MET gene amplification and TP53 mutation with PD-L1 expression in non-small cell lung cancer." *Oncotarget*(2018): 13682-13693
18. Mansuet-Lupo A, Alifano M, Pécuchet N, Biton J, Becht E, Goc J, Germain C, Ouakrim H, Régnard J, Cremer I, Laurent-Puig P, Dieu-Nosjean M, Blons H, Damotte D "Intratumoral Immune Cell Densities Are Associated with Lung Adenocarcinoma Gene Alterations." *American journal of respiratory and critical care medicine*(2016): 1403-1412
19. Kadara H, Choi M, Zhang J, Parra E, Rodriguez-Canales J, Gaffney S, Zhao Z, Behrens C, Fujimoto J, Chow C, Yoo Y, Kalhor N, Moran C, Rimm D, Swisher S, Gibbons D, Heymach J, Kaftan E, Townsend J, Lynch T, Schlessinger J, Lee J, Lifton R, Wistuba I, Herbst R "Whole-exome sequencing and immune profiling of early-stage lung adenocarcinoma with fully annotated clinical follow-up." *Annals of oncology : official journal of the European Society for Medical Oncology*(2017): 75-82
20. Schuler P, Harasymczuk M, Visus C, Deleo A, Trivedi S, Lei Y, Argiris A, Gooding W, Butterfield L, Whiteside T, Ferris R "Phase I dendritic cell p53 peptide vaccine for head and neck cancer." *Clinical cancer research : an official journal of the American Association for Cancer Research*(2014): 2433-44
21. Vermeij R, Leffers N, van der Burg S, Melief C, Daemen T, Nijman H "Immunological and clinical effects of vaccines targeting p53-overexpressing malignancies." *Journal of biomedicine & biotechnology*(2011): 702146
22. Saito H, Ando S, Morishita N, Lee K, Dator D, Dy D, Shigemura K, Adhim Z, Nibu K, Fujisawa M, Shirakawa T "A combined lymphokine-activated killer (LAK) cell immunotherapy and adenovirus-p53 gene therapy for head and neck squamous cell carcinoma." *Anticancer research*(2014): 3365-70
23. Ma C, Janetka J, Piwnicka-Worms H "Death by releasing the breaks: CHK1 inhibitors as cancer therapeutics." *Trends in molecular medicine*(2011): 88-96
24. Hirai H, Arai T, Okada M, Nishibata T, Kobayashi M, Sakai N, Imagaki K, Ohtani J, Sakai T, Yoshizumi T, Mizuarai S, Iwasawa Y, Kotani H "MK-1775, a small molecule Wee1 inhibitor, enhances anti-tumor efficacy of various DNA-damaging agents, including 5-fluorouracil." *Cancer biology & therapy*(2010): 514-22
25. Bridges K, Hirai H, Buser C, Brooks C, Liu H, Buchholz T, Molkentine J, Mason K, Meyn R "MK-1775, a novel Wee1 kinase inhibitor, radiosensitizes p53-defective human tumor cells." *Clinical cancer research : an official journal of the American Association for Cancer Research*(2011): 5638-48
26. Vilgelm A, Pawlikowski J, Liu Y, Hawkins O, Davis T, Smith J, Weller K, Horton L, McClain C, Ayers G, Turner D, Essaka D, Stewart C, Sosman J, Kelley M, Ecsedy J, Johnston J, Richmond A "Mdm2 and aurora kinase inhibitors synergize to block melanoma growth by driving apoptosis and immune clearance of tumor cells." *Cancer research*(2015): 181-93
27. Li Z, Sun Y, Chen X, Squires J, Nowroozizadeh B, Liang C, Huang J "p53 Mutation Directs AURKA Overexpression via miR-25 and FBXW7 in Prostatic Small Cell Neuroendocrine Carcinoma." *Molecular cancer research : MCR*(2015): 584-91

## References

28. Katayama H, Sen S "Functional significance of Aurora kinase A regulatory interactions with p53-ERα complex in human breast cancer cells." *Hormones & cancer*(2011): 117-24
29. Tentler J, Ionkina A, Tan A, Newton T, Pitts T, Glogowska M, Kabos P, Sartorius C, Sullivan K, Espinosa J, Eckhardt S, Diamond J "p53 Family Members Regulate Phenotypic Response to Aurora Kinase A Inhibition in Triple-Negative Breast Cancer." *Molecular cancer therapeutics*(2015): 1117-29
30. Gully C, Velazquez-Torres G, Shin J, Fuentes-Mattei E, Wang E, Carlock C, Chen J, Rothenberg D, Adams H, Choi H, Guma S, Phan L, Chou P, Su C, Zhang F, Chen J, Yang T, Yeung S, Lee M "Aurora B kinase phosphorylates and instigates degradation of p53." *Proceedings of the National Academy of Sciences of the United States of America*(2012): E1513-22
31. Marxer M, Ma H, Man W, Poon R "p53 deficiency enhances mitotic arrest and slippage induced by pharmacological inhibition of Aurora kinases." *Oncogene*(2014): 3550-60
32. El-Deiry W "The role of p53 in chemosensitivity and radiosensitivity." *Oncogene*(2003): 7486-95
33. Miyasaka A, Oda K, Ikeda Y, Sone K, Fukuda T, Inaba K, Makii C, Enomoto A, Hosoya N, Tanikawa M, Uehara Y, Arimoto T, Kuramoto H, Wada-Hiraike O, Miyagawa K, Yano T, Kawana K, Osuga Y, Fujii T "PI3K/mTOR pathway inhibition overcomes radioresistance via suppression of the HIF1-α/VEGF pathway in endometrial cancer." *Gynecologic oncology*(2015): 174-80
34. Ciardiello F, Tortora G "EGFR antagonists in cancer treatment." *The New England journal of medicine*(2008): 1160-74
35. Li D, Ding L, Ran W, Huang Y, Li G, Wang C, Xiao Y, Wang X, Lin D, Xing X "Status of 10 targeted genes of non-small cell lung cancer in eastern China: A study of 884 patients based on NGS in a single institution." *Thoracic cancer*(2020): 2580-2589
36. Wen L, Wang S, Xu W, Xu X, Li M, Zhang Y, Du X, Liu S "Value of serum tumor markers for predicting EGFR mutations in non-small cell lung cancer patients." *Annals of diagnostic pathology*(2020): 151633
37. Lemine Sow M, El Yacoubi H, Moukafih B, Balde S, Akimana G, Najem S, El Khoyaali S, Abahssain H, Chaibi A, Zeb Khan S, Trapani D, Benzekri A, Ghaouti M, Gamra L, Mestari A, Kettani F, Rahali Y, Mrabti H, Elghissassi I, Errihani H "Frequency and types of EGFR mutations in Moroccan patients with non-small cell lung cancer." *Tumori* (2021): 335-340
38. Si X, Pan R, Ma S, Li L, Liang L, Zhang P, Chu Y, Wang H, Wang M, Zhang X, Zhang L "Genomic characteristics of driver genes in Chinese patients with non-small cell lung cancer." *Thoracic cancer*(2021): 357-363
39. Rizzo S, Petrella F, Buscarino V, De Maria F, Raimondi S, Barberis M, Fumagalli C, Spitaleri G, Rampinelli C, De Marinis F, Spaggiari L, Bellomi M "CT Radiogenomic Characterization of EGFR, K-RAS, and ALK Mutations in Non-Small Cell Lung Cancer." *European radiology*(2016): 32-42
40. Lee S, Bae S, Jung S, Kim C "FDG uptake in non-small cell lung cancer is not an independent predictor of EGFR or KRAS mutation status: a retrospective analysis of 206 patients." *Clinical nuclear medicine*(2015): 950-8
41. Naderi S, Ghorra C, Haddad F, Kourie H, Rassy M, El Karak F, Ghosn M, Abadjian G, Kattan J "EGFR mutation status in Middle Eastern patients with non-squamous non-small cell lung carcinoma: A single institution experience." *Cancer epidemiology*(2015): 1099-102
42. Zhou J, Song X, He H, Zhou Y, Lu X, Ying B "Prevalence and Clinical Profile of EGFR Mutation In Non- Small-Cell Lung Carcinoma Patients in Southwest China." *Asian Pacific journal of cancer prevention : APJCP*(2016): 965-71
43. Lee B, Lee T, Lee S, Choi Y, Han J "Clinicopathologic characteristics of EGFR, KRAS, and ALK alterations in 6,595 lung cancers." *Oncotarget*(2016): 23874-84
44. Zhang Y, Yuan J, Wang K, Fu X, Han X, Threapleton D, Yang Z, Mao C, Tang J "The prevalence of EGFR mutation in patients with non-small cell lung cancer: a systematic review and meta-analysis." *Oncotarget*(2016): 78985-78993
45. Soria J, Ohe Y, Vansteenkiste J, Reungwetwattana T, Chewaskulyong B, Lee K, Dechaphunkul A, Imamura F, Nogami N, Kurata T, Okamoto I, Zhou C, Cho B, Cheng Y, Cho E, Voon P, Planchard D, Su W, Gray J, Lee S, Hodge R, Marotti M, Rukazenzov Y, Ramalingam S "Osimertinib in Untreated EGFR-Mutated Advanced Non-Small-Cell Lung Cancer." *The New England journal of medicine*(2018): 113-125
46. Mok T, Cheng Y, Zhou X, Lee K, Nakagawa K, Niho S, Lee M, Linke R, Rosell R, Corral J, Migliorino M, Pluzanski A, Sbar E, Wang T, White J, Wu Y "Improvement in Overall Survival in a Randomized Study That Compared Dacomitinib With Gefitinib in Patients With Advanced Non-Small-Cell Lung Cancer and EGFR-Activating Mutations." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2018): 2244-2250
47. Cappuzzo F, Finocchiaro G, Grossi F, Bidoli P, Favaretto A, Marchetti A, Valente M, Cseh A, Clementi L, Massey D, Santoro A "Phase II study of afatinib, an irreversible ErbB family blocker, in EGFR FISH-positive non-small-cell lung cancer." *Journal of thoracic oncology : official publication of the International Association for the Study of Lung Cancer*(2015): 665-72
48. Kelly K, Altorki N, Eberhardt W, O'Brien M, Spigel D, Crinò L, Tsai C, Kim J, Cho E, Hoffman P, Orlov S, Serwatowski P, Wang J, Foley M, Horan J, Shepherd F "Adjuvant Erlotinib Versus Placebo in Patients With Stage IB-IIIA Non-Small-Cell Lung Cancer (RADIANT): A Randomized, Double-Blind, Phase III Trial." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2015): 4007-14
49. Ahn M, Park B, Ahn J, Kim S, Kim H, Lee J, Kang J, Cho J, Song H, Park S, Sohn C, Shin S, Choi J, Ki C, Park C, Holmes A, Jänne P, Park K "Are there any ethnic differences in molecular predictors of erlotinib efficacy in advanced non-small cell lung cancer?" *Clinical cancer research : an official journal of the American Association for Cancer Research*(2008): 3860-6
50. Fukuoka M, Wu Y, Thongprasert S, Sunpaweravong P, Leong S, Sriuranpong V, Chao T, Nakagawa K, Chu D, Saijo N, Duffield E, Rukazenzov Y, Speake G, Jiang H, Armour A, To K, Yang J, Mok T "Biomarker analyses and final overall survival results from a phase III, randomized, open-label, first-line study of gefitinib versus carboplatin/paclitaxel in clinically selected patients with advanced non-small-cell lung cancer in Asia (IPASS)." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2011): 2866-74
51. Douillard J, Shepherd F, Hirsh V, Mok T, Socinski M, Gervais R, Liao M, Bischoff H, Reck M, Sellers M, Watkins C, Speake G, Armour A, Kim E "Molecular predictors of outcome with gefitinib and docetaxel in previously treated non-small-cell lung cancer: data from the randomized phase III INTEREST trial." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2010): 744-52
52. Yang J, Sequist L, Geater S, Tsai C, Mok T, Schuler M, Yamamoto N, Yu C, Ou S, Zhou C, Massey D, Zazulina V, Wu Y "Clinical activity of afatinib in patients with advanced non-small-cell lung cancer harbouring uncommon EGFR mutations: a combined post-hoc analysis of LUX-Lung 2, LUX-Lung 3, and LUX-Lung 6." *The Lancet. Oncology*(2015): 830-8
53. Cunningham D, Humblet Y, Siena S, Khayat D, Bleiberg H, Santoro A, Bets D, Mueser M, Harstrick A, Verslype C, Chau I, Van Cutsem E "Cetuximab monotherapy and cetuximab plus irinotecan in irinotecan-refractory metastatic colorectal cancer." *The New England journal of medicine*(2004): 337-45



## References

54. Vermorken J, Mesia R, Rivera F, Remenar E, Kawecki A, Rottey S, Erfan J, Zabolotny D, Kienzer H, Cupissol D, Peyrade F, Benasso M, Vynnychenko I, De Raucourt D, Bokemeyer C, Schueler A, Amellal N, Hitt R "Platinum-based chemotherapy plus cetuximab in head and neck cancer." *The New England journal of medicine*(2008): 1116-27
55. Van Cutsem E, Peeters M, Siena S, Humblet Y, Hendlisz A, Neyns B, Canon J, Van Laethem J, Maurel J, Richardson G, Wolf M, Amado R "Open-label phase III trial of panitumumab plus best supportive care compared with best supportive care alone in patients with chemotherapy-refractory metastatic colorectal cancer." *Journal of clinical oncology : official journal of the American Society of Clinical Oncology*(2007): 1658-64
56. Licitra L, Störkel S, Kerr K, Van Cutsem E, Pirker R, Hirsch F, Vermorken J, von Heydebreck A, Esser R, Celik I, Ciardiello F "Predictive value of epidermal growth factor receptor expression for first-line chemotherapy plus cetuximab in patients with head and neck and colorectal cancer: analysis of data from the EXTREME and CRYSTAL studies." *European journal of cancer (Oxford, England : 1990)*(2013): 1161-8
57. Licitra L, Mesia R, Rivera F, Remenar E, Hitt R, Erfan J, Rottey S, Kawecki A, Zabolotny D, Benasso M, Störkel S, Senger S, Stroh C, Vermorken J "Evaluation of EGFR gene copy number as a predictive biomarker for the efficacy of cetuximab in combination with chemotherapy in the first-line treatment of recurrent and/or metastatic squamous cell carcinoma of the head and neck: EXTREME study." *Annals of oncology : official journal of the European Society for Medical Oncology*(2011): 1078-1087
58. Yang J, Wu Y, Schuler M, Sebastian M, Popat S, Yamamoto N, Zhou C, Hu C, O'Byrne K, Feng J, Lu S, Huang Y, Geater S, Lee K, Tsai C, Gorbunova V, Hirsh V, Bannouna J, Orlov S, Mok T, Boyer M, Su W, Lee K, Kato T, Massey D, Shahidi M, Zazulina V, Sequist L "Afatinib versus cisplatin-based chemotherapy for EGFR mutation-positive lung adenocarcinoma (LUX-Lung 3 and LUX-Lung 6): analysis of overall survival data from two randomised, phase 3 trials." *The Lancet. Oncology* (2015): 141-51
59. Rosell R, Carcereny E, Gervais R, Vergnenegre A, Massuti B, Felip E, Palmero R, Garcia-Gomez R, Pallares C, Sanchez J, Porta R, Cobo M, Garrido P, Longo F, Moran T, Insa A, De Marinis F, Corre R, Bover I, Illiano A, Dansin E, de Castro J, Milella M, Reguart N, Altavilla G, Jimenez U, Provencio M, Moreno M, Terrasa J, Muñoz-Langa J, Valdivia J, Isla D, Domine M, Molinier O, Mazieres J, Baize N, Garcia-Campelo R, Robinet G, Rodriguez-Abreu D, Lopez-Vivanco G, Gebbia V, Ferrera-Delgado L, Bombardieri P, Bernabe R, Bearz A, Artañ A, Cortesi E, Rolfo C, Sanchez-Ronco M, Drozdowskyj A, Queralt C, de Aguirre I, Ramirez J, Sanchez J, Molina M, Taron M, Paz-Ares L "Erlotinib versus standard chemotherapy as first-line treatment for European patients with advanced EGFR mutation-positive non-small-cell lung cancer (EURTAC): a multicentre, open-label, randomised phase 3 trial." *The Lancet. Oncology*(2012): 239-46
60. Wu Y, Cheng Y, Zhou X, Lee K, Nakagawa K, Niho S, Tsuji F, Linke R, Rosell R, Corral J, Migliorino M, Pluzanski A, Sbar E, Wang T, White J, Nadanaciva S, Sandin R, Mok T "Dacomitinib versus gefitinib as first-line treatment for patients with EGFR-mutation-positive non-small-cell lung cancer (ARCHER 1050): a randomised, open-label, phase 3 trial." *The Lancet. Oncology*(2017): 1454-1466
61. Mok T, Wu Y, Thongprasert S, Yang C, Chu D, Saijo N, Sunpaweravong P, Han B, Margono B, Ichinose Y, Nishiwaki Y, Ohe Y, Yang J, Chewaskulyong B, Jiang H, Duffield E, Watkins C, Armour A, Fukuoka M "Gefitinib or carboplatin-paclitaxel in pulmonary adenocarcinoma." *The New England journal of medicine*(2009): 947-57
62. Douillard J, Ostoros G, Cobo M, Ciuleanu T, McCormack R, Webster A, Milenkova T "First-line gefitinib in Caucasian EGFR mutation-positive NSCLC patients: a phase-IV, open-label, single-arm study." *British journal of cancer*(2014): 55-62
63. Nakagawa K, Garon E, Seto T, Nishio M, Ponce Aix S, Paz-Ares L, Chiu C, Park K, Novello S, Nadal E, Imamura F, Yoh K, Shih J, Au K, Moro-Sibilot D, Enatsu S, Zimmermann A, Frimodt-Møller B, Visseren-Grul C, Reck M "Ramucicirumab plus erlotinib in patients with untreated, EGFR-mutated, advanced non-small-cell lung cancer (RELAY): a randomised, double-blind, placebo-controlled, phase 3 trial." *The Lancet. Oncology*(2019): 1655-1669
64. O'Kane G, Bradbury P, Feld R, Leigh N, Liu G, Pisters K, Kamel-Reid S, Tsao M, Shepherd F "Uncommon EGFR mutations in advanced non-small cell lung cancer." *Lung cancer (Amsterdam, Netherlands)*(2017): 137-144
65. Zeng L, Xiao L, Jiang W, Yang H, Hu D, Xia C, Li Y, Zhou C, Xiong Y, Liu L, Liao D, Guan R, Li K, Wang J, Zhang Y, Yang N, Mansfield A "Investigation of efficacy and acquired resistance for EGFR-TKI plus bevacizumab as first-line treatment in patients with EGFR sensitive mutant non-small cell lung cancer in a Real world population." *Lung cancer (Amsterdam, Netherlands)*(2020): 82-88