



Sample Information

Patient Name: 黃瑤笙**Gender:** Female**ID No.:** Q222294397**History No.:** 41211818**Age:** 41**Ordering Doctor:** DOC5390F 高冠鈞**Ordering REQ.:** 0AZKZPY**Signing in Date:** 2020/12/10**Path No.:** S109-96821**MP No.:** TM20009**Assay:** Oncomine Tumor Mutation Load Assay**Sample Type:** FFPE**Block No.:** S109-38423C**Percentage of tumor cells:** 80%**Note:**

Sample Cancer Type: Colorectal Cancer

Table of Contents	Page
Variants (Exclude variant in Taiwan BioBank with >1% allele frequency)	2
Biomarker Descriptions	3
Relevant Therapy Summary	5
Relevant Therapy Details	6
Clinical Trials Summary	10

Report Highlights

2 Relevant Biomarkers
3 Therapies Available
14 Clinical Trials

Relevant Colorectal Cancer Findings

Gene	Finding
BRAF	Not detected
KRAS	Not detected
NRAS	Not detected
NTRK1	Not detected
NTRK3	Not detected



Relevant Biomarkers

Tier	Genomic Alteration	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
IIC	FBXW7 p.(R479Q) c.1436G>A F-box and WD repeat domain containing 7 Allele Fraction: 0.711	None	None	3
	Tumor Mutational Burden 4.2 Mut/Mb measured	pembrolizumab ¹	ipilimumab + nivolumab nivolumab pembrolizumab	12

Public data sources included in relevant therapies: FDA¹, NCCN, EMA², ESMO

Tier 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.

Although no fusion transcript can be detected, there is high imbalance of the number of 3' reads and 5' reads in the RET gene (3'/5' imbalance value: 25.86). A high 3'/5' imbalance value is suggestive of the presence of gene fusion. The possibility of RET fusion involving partners other than those targeted by the panel cannot be excluded. Further confirmation with other methodologies is suggested.

Variants (Exclude variant in Taiwan BioBank with >1% allele frequency)

DNA Sequence Variants								
Gene	Amino Acid Change	Coding	Variant ID	Locus	Allele Fraction	Transcript	Variant Effect	Coverage
FBXW7	p.(R479Q)	c.1436G>A	COSM22974	chr4:153247366	0.711	NM_033632.3	missense	558
PDE4DIP	p.(E1081K)	c.3241G>A	.	chr1:144882778	0.175	NM_001198834.3	missense	1339
ABL2	p.(G727E)	c.2180G>A	.	chr1:179078177	0.037	NM_005158.4	missense	54
SOX11	p.(=)	c.462G>A	.	chr2:5833315	0.486	NM_003108.3	synonymous	463
LTF	p.(R23dup)	c.68_69insAAG	.	chr3:46501284	1.000	NM_002343.5	nonframeshift Insertion	120
EPHB1	p.(=)	c.1392C>T	.	chr3:134873088	0.275	NM_004441.4	synonymous	2000
PDGFRA	p.(=)	c.1701A>G	.	chr4:55141055	0.998	NM_006206.5	synonymous	1306
GP3M3	p.(?)	c.-3640C>G	.	chr6:32163930	0.737	NM_022107.2	unknown	1392
HSP90AB1	p.(=)	c.504T>C	.	chr6:44217561	0.337	NM_001271970.1	synonymous	2000
PKHD1	p.(Y617H)	c.1849T>C	.	chr6:51918951	0.353	NM_138694.3	missense	1999
DST	p.(=)	c.13692A>G	.	chr6:56362699	0.775	NM_001144769.2	synonymous	227
ADGRB3	p.(=)	c.361C>T	.	chr6:69348928	0.505	NM_001704.2	synonymous	1269
SYNE1	p.(L5015M)	c.15043T>A	.	chr6:152647681	1.000	NM_182961.3	missense	1114
AKAP9	p.(=)	c.10845G>A	.	chr7:91729132	0.486	NM_005751.4	synonymous	621
SAMD9	p.(=)	c.216A>G	.	chr7:92735195	0.489	NM_001193307.1	synonymous	747
UBR5	p.(R2117W)	c.6349C>T	.	chr8:103289360	0.435	NM_015902.5	missense	1848
CSMD3	p.(S1207L)	c.3620C>T	.	chr8:113649141	0.140	NM_198123.1	missense	2000
TAF1L	p.(A38V)	c.113C>T	.	chr9:32635465	0.852	NM_153809.2	missense	799



Variants (Exclude variant in Taiwan BioBank with >1% allele frequency) (continued)

DNA Sequence Variants (continued)

Gene	Amino Acid Change	Coding	Variant ID	Locus	Allele Fraction	Transcript	Variant Effect	Coverage
ABL1	p.(P901A)	c.2701C>G	.	chr9:133760378	0.913	NM_005157.5	missense	1590
BRD3	p.(=)	c.723C>T	.	chr9:136913568	0.067	NM_007371.3	synonymous	463
NUMA1	p.(R1413W)	c.4237C>T	.	chr11:71724312	0.663	NM_006185.3	missense	2000
NUMA1	p.(R723H)	c.2168G>A	.	chr11:71726381	0.665	NM_006185.3	missense	1134
CBL	p.(L857F)	c.2569C>T	.	chr11:119170339	0.300	NM_005188.3	missense	1349
ERBB3	p.(H1165R)	c.3494A>G	.	chr12:56495137	0.908	NM_001982.3	missense	1395
MAP2K1	p.(=)	c.870A>G	.	chr15:66777504	0.505	NM_002755.3	synonymous	1223
TSC2	p.(E1490G)	c.4469A>G	.	chr16:2134692	0.071	NM_000548.4	missense	548
MYH11	p.(N312S)	c.935A>G	.	chr16:15865545	0.064	NM_001040114.1	missense	870
CDH11	p.(=)	c.945G>A	.	chr16:65022114	1.000	NM_001797.3	synonymous	1503
CDH5	p.(R410H)	c.1229G>A	.	chr16:66429973	0.921	NM_001795.4	missense	1236
PER1	p.(=)	c.3024C>T	.	chr17:8046632	0.100	NM_002616.2	synonymous	1874
CDK12	p.(E1345K)	c.4033G>A	.	chr17:37687129	0.268	NM_016507.3	missense	1998
CDK12	p.(R1477T)	c.4430G>C	.	chr17:37687526	0.250	NM_016507.3	missense	1999
TCF3	p.(G431S)	c.1291_1293delGGCinsAGT	.	chr19:1619348	0.943	NM_001136139.3	missense	300
KEAP1	p.(R553W)	c.1657C>T	.	chr19:10599919	0.946	NM_203500.1	missense	56
AMER1	p.(S279G)	c.835A>G	.	chrX:63412332	0.428	NM_152424.3	missense	855
TAF1	p.(V802M)	c.2404G>A	.	chrX:70607228	0.096	NM_004606.4	missense	52
ATRX	p.(D1378E)	c.4134T>G	.	chrX:76912130	0.654	NM_000489.4	missense	786

Biomarker Descriptions

FBXW7 (F-box and WD repeat domain containing 7)

Background: The FBXW7 gene encodes a member of the F-box protein family that functions as the substrate recognition component of the SCF complex, which is responsible for protein ubiquitination and subsequent degradation by the proteasome¹. FBXW7 is a tumor suppressor gene that plays a crucial role in the degradation and turnover of various proto-oncogenes. Aberrations such as mutations or deletions that alter the tumor suppression function can lead to the deregulation of downstream genes, including MYC, MTOR, and NOTCH1, thereby promoting cell proliferation and survival^{1,2,3,4,5,6,7}.

Alterations and prevalence: Mutations in FBXW7 occur at high frequencies in various malignancies, including 40% of uterine carcinoma and 10-15% of stomach, bladder, cervical, and colorectal cancers^{8,9,10,11,12}.

Potential relevance: Currently, no therapies are approved for FBXW7 aberrations. Missense mutations in FBXW7 are associated with poor prognosis and worse overall survival (OS) in comparison to FBXW7 wild-type metastatic colorectal cancer⁸. In a clinical case



Biomarker Descriptions (continued)

report, a patient with FBXW7 R465H-mutated, EGFR/ALK-wildtype lung adenocarcinoma demonstrated tumor shrinkage after treatment with the mTOR inhibitor temsirolimus. In a phase I clinical trial of sirolimus, one hepatocellular fibrolamellar carcinoma patient with the FBXW7 E192A mutation demonstrated stable disease for over 6 months⁷.

Tumor Mutational Burden

Background: Tumor mutational burden (TMB), also known as tumor mutational load (TML), is the count of somatic mutations in the DNA of cancer cells. TMB is determined by next-generation sequencing and is expressed as the number of mutations per megabase (mut/Mb) of DNA coding sequence¹³. Errors in DNA repair, including mutations in the POLE gene and in mismatch repair (MMR) genes, are associated with increased TMB^{14,15,16,17,18}. High TMB is associated with increased neo-antigen burden and has been linked to response to immune checkpoint inhibitors (ICIs) that target the cytotoxic T lymphocyte antigen-4 (CTLA4), programmed death protein 1 (PD1), and programmed death-ligand 1 (PD-L1) inhibitors^{19,20,21,22}.

Alterations and prevalence: In one study of over 100,000 tumor samples, the median TMB value was 3.6 mut/Mb although TMB values vary widely across cancers²³. Certain childhood cancers, leukemia, glioblastoma, and neuroblastoma typically have low mutation burden and median TMB values <1 mut/Mb^{20,23}. In comparison, cancers that experience genotoxic insults including skin cancer and lung cancer have higher median TMB values of approximately 10 mut/Mb^{20,23}. For example, within non-small cell lung cancer (NSCLC), higher TMB was observed in former/current smokers (10.5 mut/Mb) relative to never smokers (0.6 mut/Mb)^{20,23,24}. There is no consensus around the definition of high and low TMB that could be applied universally to all tumor types, instead multiple sources suggest that TMB status is a cancer type specific attribute^{23,25,26}. In NSCLC, several studies have suggested establishing a threshold between low and high TMB of 10 +/- 1 mut/Mb^{27,28,29,30}.

Potential relevance: ICIs stimulate a patient's own T-cells to kill tumors and have exhibited benefits in some patients. The first ICI to be approved by the FDA was ipilimumab (2011), an anti-CTLA4 antibody indicated for the treatment of metastatic melanoma. In 2014, anti-PD-1 antibodies nivolumab (2014) and pembrolizumab (2014) were subsequently approved, for the treatment of metastatic melanoma, and pembrolizumab also for advanced esophageal squamous cell carcinoma. In 2020, the indication for pembrolizumab³¹ was expanded to include TMB-H (>= 10 mut/Mb) solid tumors that have progressed on prior therapy. Indications have been expanded for these ICIs to include several other cancer types including NSCLC, advanced renal cell carcinoma, classical Hodgkin lymphoma, recurrent or metastatic squamous cell carcinoma of the head and neck, urothelial carcinoma, microsatellite instability (MSI)-High or mismatch repair deficient colorectal cancer, and hepatocellular carcinoma. Atezolizumab (2016), avelumab (2017), and durvalumab (2017), that target programmed death-ligand 1 (PD-L1) were subsequently approved by the FDA. However, the predictive biomarkers that underlie the clinical benefits of these approved immunotherapies, including TMB, are under active investigation. Several published studies including the CheckMate 586 and CheckMate 817 clinical trials have concluded that high TMB was associated with improved response to FDA approved checkpoint inhibitors^{28,32,33}. In contrast, several previously promising trials have failed to show an improvement in survival outcomes between high and low TMB including CheckMate 227 (ipilimumab + nivolumab vs. chemotherapy), CheckMate 026 (nivolumab vs. chemotherapy), KEYNOTE 189 (pembrolizumab vs. chemotherapy), KEYNOTE 021 (pembrolizumab vs. pembrolizumab + chemotherapy), and Lung-MAP (nivolumab + ipilimumab vs. nivolumab). In response, suggestions to combine TMB score with PD-L1 expression as a way to increase the predictive power for patient stratification have been reported³⁴. Nivolumab alone or in combination with ipilimumab is recommended for use in NSCLC with evidence of high TMB³⁵. TMB score estimation is affected by the utilized assays, therefore efforts are underway to develop a standardized approach to calculate with the aim to support consistent reporting of TMB values across laboratories^{36,37,38,39}.



Relevant Therapy Summary

☒ In this cancer type
 ☐ In other cancer type
 ☒ In this cancer type and other cancer types
 ☒ No evidence

FBXW7 p.(R479Q) c.1436G>A

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
berzosertib	×	×	×	×	● (II)
palbociclib	×	×	×	×	● (II)
temsirolimus	×	×	×	×	● (II)

Tumor Mutational Burden

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
pembrolizumab	●	○	×	×	● (II)
ipilimumab + nivolumab	×	○	×	×	● (II)
nivolumab	×	○	×	×	● (II)
atezolizumab	×	×	×	×	● (II)
durvalumab, tremelimumab	×	×	×	×	● (II)
ipilimumab, nivolumab	×	×	×	×	● (II)
pembrolizumab, ipilimumab + nivolumab	×	×	×	×	● (II)
entinostat, nivolumab	×	×	×	×	● (I/II)
BAY1905254	×	×	×	×	● (I)
zimberelimab	×	×	×	×	● (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 FDA Information

☒ In this cancer type
 ☐ In other cancer type
 ☒ In this cancer type and other cancer types

FDA information is current as of 2020-10-14. For the most up-to-date information, search www.fda.gov.

Tumor Mutational Burden

● pembrolizumab

Cancer type: Solid Tumor

Label as of: 2020-10-14

Variant class: Tumor Mutational Burden

Indications and usage:

KEYTRUDA® is a programmed death receptor-1 (PD-1)-blocking antibody indicated:

Melanoma

- for the treatment of patients with unresectable or metastatic melanoma.
- for the adjuvant treatment of patients with melanoma with involvement of lymph node(s) following complete resection.

Non-Small Cell Lung Cancer (NSCLC)

- in combination with pemetrexed and platinum chemotherapy, as first-line treatment of patients with metastatic nonsquamous NSCLC, with no EGFR or ALK genomic tumor aberrations.
- in combination with carboplatin and either paclitaxel or paclitaxel protein-bound, as first-line treatment of patients with metastatic squamous NSCLC.
- as a single agent for the first-line treatment of patients with NSCLC expressing PD-L1 [Tumor Proportion Score (TPS) $\geq 1\%$] as determined by an FDA-approved test, with no EGFR or ALK genomic tumor aberrations, and is:
 - stage III where patients are not candidates for surgical resection or definitive chemoradiation, or
 - metastatic.
- as a single agent for the treatment of patients with metastatic NSCLC whose tumors express PD-L1 (TPS $\geq 1\%$) as determined by an FDA-approved test, with disease progression on or after platinum-containing chemotherapy. Patients with EGFR or ALK genomic tumor aberrations should have disease progression on FDA-approved therapy for these aberrations prior to receiving KEYTRUDA®.

Small Cell Lung Cancer (SCLC)

- for the treatment of patients with metastatic SCLC with disease progression on or after platinum-based chemotherapy and at least one other prior line of therapy.¹

Head and Neck Squamous Cell Cancer (HNSCC)

- in combination with platinum and FU for the first-line treatment of patients with metastatic or with unresectable, recurrent HNSCC.
- as a single agent for the first-line treatment of patients with metastatic or with unresectable, recurrent HNSCC whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 1] as determined by an FDA-approved test.
- as a single agent for the treatment of patients with recurrent or metastatic HNSCC with disease progression on or after platinum-containing chemotherapy.

Classical Hodgkin Lymphoma (cHL)

- for the treatment of adult patients with relapsed or refractory cHL.
- for the treatment of pediatric patients with refractory cHL, or cHL that has relapsed after 2 or more lines of therapy.

Primary Mediastinal Large B-Cell Lymphoma (PMBCL)

- for the treatment of adult and pediatric patients with refractory PMBCL, or who have relapsed after 2 or more prior lines of therapy.



Tumor Mutational Burden (continued)

- Limitations of Use: KEYTRUDA® is not recommended for treatment of patients with PMBCL who require urgent cytoreductive therapy.

Urothelial Carcinoma

- for the treatment of patients with locally advanced or metastatic urothelial carcinoma who are not eligible for cisplatin-containing chemotherapy and whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 10] as determined by an FDA-approved test, or in patients who are not eligible for any platinum-containing chemotherapy regardless of PD-L1 status.¹
- for the treatment of patients with locally advanced or metastatic urothelial carcinoma who have disease progression during or following platinum-containing chemotherapy or within 12 months of neoadjuvant or adjuvant treatment with platinum-containing chemotherapy.
- for the treatment of patients with Bacillus Calmette-Guerin (BCG)-unresponsive, high-risk, non-muscle invasive bladder cancer (NMIBC) with carcinoma in situ (CIS) with or without papillary tumors who are ineligible for or have elected not to undergo cystectomy.

Microsatellite Instability-High Cancer or Mismatch Repair Deficient Cancer

- for the treatment of adult and pediatric patients with unresectable or metastatic, microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR)
 - solid tumors that have progressed following prior treatment and who have no satisfactory alternative treatment options,¹ or
 - colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan.¹
- Limitations of Use: The safety and effectiveness of KEYTRUDA® in pediatric patients with MSI-H central nervous system cancers have not been established.

Microsatellite Instability-High or Mismatch Repair Deficient Colorectal Cancer (CRC)

- for the first-line treatment of patients with unresectable or metastatic MSI-H or dMMR colorectal cancer (CRC).

Gastric Cancer

- for the treatment of patients with recurrent locally advanced or metastatic gastric or gastroesophageal junction adenocarcinoma whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 1] as determined by an FDA-approved test, with disease progression on or after 2 or more prior lines of therapy including fluoropyrimidine- and platinum-containing chemotherapy and if appropriate, HER2/neu-targeted therapy.¹

Esophageal Cancer

- for the treatment of patients with recurrent locally advanced or metastatic squamous cell carcinoma of the esophagus whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 10] as determined by an FDA-approved test, with disease progression after one or more prior lines of systemic therapy.

Cervical Cancer

- for the treatment of patients with recurrent or metastatic cervical cancer with disease progression on or after chemotherapy whose tumors express PD-L1 [Combined Positive Score (CPS) ≥ 1] as determined by an FDA-approved test.¹

Hepatocellular Carcinoma (HCC)

- for the treatment of patients with HCC who have been previously treated with sorafenib.¹

Merkel Cell Carcinoma (MCC)

- for the treatment of adult and pediatric patients with recurrent locally advanced or metastatic Merkel cell carcinoma.¹

Renal Cell Carcinoma (RCC)

- in combination with axitinib, for the first-line treatment of patients with advanced RCC.

Endometrial Carcinoma

- in combination with lenvatinib, for the treatment of patients with advanced endometrial carcinoma that is not MSI-H or dMMR, who have disease progression following prior systemic therapy and are not candidates for curative surgery or radiation.¹

Tumor Mutational Burden-High (TMB-H) Cancer



Tumor Mutational Burden (continued)

- for the treatment of adult and pediatric patients with unresectable or metastatic tumor mutational burden-high (TMB-H) [≥ 10 mutations/megabase (mut/Mb)] solid tumors, as determined by an FDA-approved test, that have progressed following prior treatment and who have no satisfactory alternative treatment options.¹
- Limitations of Use: The safety and effectiveness of KEYTRUDA® in pediatric patients with TMB-H central nervous system cancers have not been established.

Cutaneous Squamous Cell Carcinoma (cSCC)

- for the treatment of patients with recurrent or metastatic cutaneous squamous cell carcinoma that is not curable by surgery or radiation.

Adult Indications: Additional Dosing Regimen of 400 mg Every 6 Weeks

- for use at an additional recommended dosage of 400 mg every 6 weeks for all approved adult indications.²

¹This indication is approved under accelerated approval based on tumor response rate and durability of response. Continued approval for this indication may be contingent upon verification and description of clinical benefit in the confirmatory trials.

²This indication is approved under accelerated approval based on pharmacokinetic data, the relationship of exposure to efficacy, and the relationship of exposure to safety. Continued approval for this dosing may be contingent upon verification and description of clinical benefit in the confirmatory trials.

Reference:

https://www.accessdata.fda.gov/drugsatfda_docs/label/2020/125514s085lbl.pdf



Current NCCN Information

- ☒ In this cancer type
 ☐ In other cancer type
 ☒ In this cancer type and other cancer types

NCCN information is current as of 2020-10-01. For the most up-to-date information, search www.nccn.org.
For NCCN International Adaptations & Translations, search www.nccn.org/global/international_adaptations.aspx.

Tumor Mutational Burden

☐ ipilimumab + nivolumab

Cancer type: Non-Small Cell Lung Cancer

Variant class: Tumor Mutational Burden

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- Non-Small Cell Lung Cancer; Emerging biomarker in metastatic disease (Not specified)

Reference: NCCN Guidelines® - NCCN-Non-Small Cell Lung Cancer [Version 8.2020]

☐ nivolumab

Cancer type: Non-Small Cell Lung Cancer

Variant class: Tumor Mutational Burden

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- Non-Small Cell Lung Cancer; Emerging biomarker in metastatic disease (Not specified)

Reference: NCCN Guidelines® - NCCN-Non-Small Cell Lung Cancer [Version 8.2020]

☐ pembrolizumab

Cancer type: Cervical Cancer

Variant class: Tumor Mutational Burden

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- Recurrent or Metastatic Cervical Cancer; Progressed on prior treatment and does not have satisfactory alternative treatment options; Tumor mutational burden-high (TMB-H) [≥ 10 mutations/megabase (mut/Mb)] as determined by an FDA-approved test (Second-line therapy) (Useful in Certain Circumstances)

Reference: NCCN Guidelines® - NCCN-Cervical Cancer [Version 2.2020]



Tumor Mutational Burden (continued)

○ pembrolizumab

Cancer type: Thyroid Gland Anaplastic Carcinoma, **Variant class:** Tumor Mutational Burden
Thyroid Gland Follicular Carcinoma, Thyroid Gland
Hurthle Cell Carcinoma, Thyroid Gland Medullary
Carcinoma, Thyroid Gland Papillary Carcinoma

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- Thyroid Gland Papillary, Follicular, Hurthle Cell Carcinoma; Unresectable locoregional recurrent/persistent disease not amenable to RAI therapy (Not specified)
- Thyroid Gland Papillary, Follicular, Hurthle Cell Carcinoma; CNS or soft tissue or bone metastases not amenable to RAI therapy (Not specified)
- Thyroid Gland Medullary Carcinoma; Locoregional recurrent/persistent disease (Not specified) (Useful in Certain Circumstances)
- Thyroid Gland Medullary Carcinoma; Recurrent or persistent disease; Distant metastases; Asymptomatic, symptomatic or progression of disease (Not specified) (Useful in Certain Circumstances)
- Thyroid Gland Anaplastic Carcinoma; Metastatic (Not specified) (Useful in Certain Circumstances)

Reference: NCCN Guidelines® - NCCN-Thyroid Carcinoma [Version 2.2020]

○ pembrolizumab

Cancer type: Endometrial Cancer **Variant class:** Tumor Mutational Burden

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- Unresectable or Metastatic Endometrial Carcinoma or Uterine Sarcoma; Progressed on prior treatment and does not have satisfactory alternative treatment options; Tumor mutational burden-high (TMB-H) [≥ 10 mutations/megabase (mut/Mb)] as determined by an FDA-approved test (Not specified) (Useful in Certain Circumstances)

Reference: NCCN Guidelines® - NCCN-Uterine Neoplasms [Version 2.2020]

Clinical Trials Summary

FBXW7 p.(R479Q) c.1436G>A

NCT ID	Title	Phase
NCT03718091	A Phase II Study of M6620 (VX-970) in Selected Solid Tumors	II
NCT03297606	Canadian Profiling and Targeted Agent Utilization Trial (CAPTUR): A Phase II Basket Trial	II
NCT03155620	NCI-COG Pediatric MATCH (Molecular Analysis for Therapy Choice) Screening Protocol	II



Clinical Trials Summary (continued)

Tumor Mutational Burden

NCT ID	Title	Phase
NCT03767075	Basket of Baskets: A Modular, Open-label, Phase II, Multicentre Study To Evaluate Targeted Agents in Molecularly Selected Populations With Advanced Solid Tumours	II
NCT02029001	A Two-period, Multicenter, Randomized, Open-label, Phase II Study Evaluating the Clinical Benefit of a Maintenance Treatment Targeting Tumor Molecular Alterations in Patients With Progressive Locally-advanced or Metastatic Solid Tumors MOST: My own specific treatment	II
NCT03297606	Canadian Profiling and Targeted Agent Utilization Trial (CAPTUR): A Phase II Basket Trial	II
NCT03668119	A Randomized, Open-Label, Phase II Study of Nivolumab in Combination With Ipilimumab or Nivolumab Monotherapy in Participants With Advanced or Metastatic Solid Tumors of High Tumor Mutational Burden (TMB-H)	II
No NCT ID	A Multicenter Phase II Study Of Nivolumab Monotherapy In Recurrent And/Or Metastatic Gastrointestinal Cancer Patients With High Tumor Mutation Burden (TMB-H)	II
No NCT ID	TR Accompanied with the Study Titled A Multicenter Phase II Study of Nivolumab Monotherapy in Recurrent and/or Metastatic Gastrointestinal Cancer Patients with High Tumor Mutation Burden (TMB-H).	II
NCT02628067	A Clinical Trial of Pembrolizumab (MK-3475) Evaluating Predictive Biomarkers in Subjects With Advanced Solid Tumors (KEYNOTE 158)	II
NCT02693535	Targeted Agent and Profiling Utilization Registry (TAPUR) Study	II
NCT03838042	INFORM2 Exploratory Multinational Phase I/II Combination Study of Nivolumab and Entinostat in Children and Adolescents with Refractory High-risk Malignancies	I/II
NCT02992964	Pilot Study of Nivolumab in Pediatric Patients With Hypermutant Cancers	I/II
NCT03666273	An Open-label, Phase I, First-in-human, Dose Escalation and Expansion Study to Evaluate the Safety, Tolerability, Maximum Tolerated or Administered Dose, Pharmacokinetics, Pharmacodynamics and Tumor Response Profile of the ILDR2 Function-blocking Antibody BAY1905254 in Patients With Advanced Solid Tumors	I
NCT04087018	A Phase Ib Study to Evaluate the Safety and Clinical Activity of AB122 in Biomarker-Selected Participants With Advanced Solid Tumors	I



Signatures

Testing Personnel:

Laboratory Supervisor:

Pathologist:



References

1. Yeh et al. FBXW7: a critical tumor suppressor of human cancers. *Mol Cancer*. 2018 Aug 7;17(1):115. doi: 10.1186/s12943-018-0857-2. PMID: 30086763
2. Wang et al. Tumor suppressor functions of FBW7 in cancer development and progression. *FEBS Lett*. 2012 May 21;586(10):1409-18. PMID: 22673505
3. Uhlén et al. Proteomics. Tissue-based map of the human proteome. *Science*. 2015 Jan 23;347(6220):1260419. doi: 10.1126/science.1260419. PMID: 25613900
4. Yada et al. Phosphorylation-dependent degradation of c-Myc is mediated by the F-box protein Fbw7. *EMBO J*. 2004 May 19;23(10):2116-25. PMID: 15103331
5. Hori et al. Notch signaling at a glance. *J. Cell. Sci*. 2013 May 15;126(Pt 10):2135-40. PMID: 23729744
6. Aydin et al. FBXW7 mutations in melanoma and a new therapeutic paradigm. *J. Natl. Cancer Inst*. 2014 Jun;106(6):dju107. PMID: 24838835
7. Jardim et al. FBXW7 mutations in patients with advanced cancers: clinical and molecular characteristics and outcomes with mTOR inhibitors. *PLoS ONE*. 2014;9(2):e89388. PMID: 24586741
8. Korphaisarn et al. FBXW7 missense mutation: a novel negative prognostic factor in metastatic colorectal adenocarcinoma. *Oncotarget*. 2017 Jun 13;8(24):39268-39279. PMID: 28424412
9. Cancer Genome Atlas Network. Comprehensive molecular characterization of human colon and rectal cancer. *Nature*. 2012 Jul 18;487(7407):330-7. PMID: 22810696
10. Cancer Genome Atlas Research Network. Comprehensive molecular characterization of urothelial bladder carcinoma. *Nature*. 2014 Mar 20;507(7492):315-22. doi: 10.1038/nature12965. Epub 2014 Jan 29. PMID: 24476821
11. Weinstein et al. The Cancer Genome Atlas Pan-Cancer analysis project. *Nat. Genet*. 2013 Oct;45(10):1113-20. PMID: 24071849
12. 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
13. Yuan et al. Novel technologies and emerging biomarkers for personalized cancer immunotherapy. 4:3. PMID: 26788324
14. Temko et al. Somatic POLE exonuclease domain mutations are early events in sporadic endometrial and colorectal carcinogenesis, determining driver mutational landscape, clonal neoantigen burden and immune response. *J. Pathol*. 2018 Jul;245(3):283-296. PMID: 29604063
15. Mehnert et al. Immune activation and response to pembrolizumab in POLE-mutant endometrial cancer. 126(6):2334-40. PMID: 27159395
16. Bouffet et al. Immune Checkpoint Inhibition for Hypermutant Glioblastoma Multiforme Resulting From Germline Biallelic Mismatch Repair Deficiency. *J. Clin. Oncol*. 2016 Jul 1;34(19):2206-11. PMID: 27001570
17. Humphris et al. Hypermutation In Pancreatic Cancer. *Gastroenterology*. 2017 Jan;152(1):68-74.e2. PMID: 27856273
18. Bonneville et al. Landscape of Microsatellite Instability Across 39 Cancer Types. *JCO Precis Oncol*. 2017;2017. PMID: 29850653
19. Snyder et al. Genetic basis for clinical response to CTLA-4 blockade in melanoma. *N. Engl. J. Med*. 2014 Dec 4;371(23):2189-2199. PMID: 25409260
20. Alexandrov et al. Signatures of mutational processes in human cancer. *Nature*. 2013 Aug 22;500(7463):415-21. PMID: 23945592
21. Rizvi et al. Cancer immunology. Mutational landscape determines sensitivity to PD-1 blockade in non-small cell lung cancer. *Science*. 2015 Apr 3;348(6230):124-8. PMID: 25765070
22. Van et al. Genomic correlates of response to CTLA-4 blockade in metastatic melanoma. *Science*. 2015 Oct 9;350(6257):207-211. PMID: 26359337
23. Chalmers et al. Analysis of 100,000 human cancer genomes reveals the landscape of tumor mutational burden. 9:34. PMID: 28420421
24. Govindan et al. Genomic landscape of non-small cell lung cancer in smokers and never-smokers. *Cell*. 2012 Sep 14;150(6):1121-34. PMID: 22980976
25. Endris et al. Measurement of tumor mutational burden (TMB) in routine molecular diagnostics: in silico and real-life analysis of three larger gene panels. *Int. J. Cancer*. 2019 May 1;144(9):2303-2312. PMID: 30446996



References (continued)

26. Meléndez et al. Methods of measurement for tumor mutational burden in tumor tissue. *Transl Lung Cancer Res.* 2018 Dec;7(6):661-667. PMID: 30505710
27. Hellmann et al. Nivolumab plus Ipilimumab in Lung Cancer with a High Tumor Mutational Burden. *N. Engl. J. Med.* 2018 May 31;378(22):2093-2104. PMID: 29658845
28. Ready et al. First-Line Nivolumab Plus Ipilimumab in Advanced Non-Small-Cell Lung Cancer (CheckMate 568): Outcomes by Programmed Death Ligand 1 and Tumor Mutational Burden as Biomarkers. *J. Clin. Oncol.* 2019 Apr 20;37(12):992-1000. PMID: 30785829
29. Alborelli et al. Tumor mutational burden assessed by targeted NGS predicts clinical benefit from immune checkpoint inhibitors in non-small cell lung cancer. *J. Pathol.* 2020 Jan;250(1):19-29. PMID: 31471895
30. Heeke et al. In-house Implementation of Tumor Mutational Burden Testing to Predict Durable Clinical Benefit in Non-small Cell Lung Cancer and Melanoma Patients. *Cancers (Basel).* 2019 Aug 29;11(9). PMID: 31470674
31. https://www.accessdata.fda.gov/drugsatfda_docs/label/2020/125514s085lbl.pdf
32. F. et al. CheckMate 817: First-Line Nivolumab + Ipilimumab in Patients with ECOG PS 2 and Other Special Populations with Advanced NSCLC. Volume 14, Issue 10, Supplement, Pages S214–S215. Abstract OA04.02
33. Zhu et al. Association Between Tumor Mutation Burden (TMB) and Outcomes of Cancer Patients Treated With PD-1/PD-L1 Inhibitions: A Meta-Analysis. *Front Oncol*, 9:1161, 04 Nov 2019. PMID: 31258479
34. Castellanos et al. Tumor mutational burden (TMB) and PD-L1 expression as predictors of response to immunotherapy (IO) in NSCLC. *Journal of Clinical Oncology* 37, no. 15_suppl (May 20, 2019) 2630-2630. Abstract 2630
35. NCCN Guidelines® - NCCN-Non-Small Cell Lung Cancer [Version 8.2020]
36. <https://www.focr.org/tmb>
37. <http://www.iqnpath.org/category/tmb>
38. Stenzinger et al. Tumor mutational burden standardization initiatives: Recommendations for consistent tumor mutational burden assessment in clinical samples to guide immunotherapy treatment decisions. *Genes Chromosomes Cancer.* 2019 Aug;58(8):578-588. PMID: 30664300
39. Merino et al. Establishing guidelines to harmonize tumor mutational burden (TMB): in silico assessment of variation in TMB quantification across diagnostic platforms: phase I of the Friends of Cancer Research TMB Harmonization Project. *J Immunother Cancer.* 2020 Mar;8(1). PMID: 32217756