

Department of Pathology and Laboratory Medicine No.201, Sec. 2, Shipai Rd., Beitou District, Taipei City, Taiwan 11217, R.O.C.

Tel: 02-2875-7449

Date: 08 Oct 2020 1 of 4

Sample Information

Patient Name: 王文賢 Gender: Male ID No.: F120553642 History No.: 39779550

Age: 53

Ordering Doctor: DOC1373L 陳三奇

Ordering REQ.: 0AWYGSE Signing in Date: 2020/10/07

Path No.: \$109-89712 **MP No.:** F20084

Assay: Oncomine Focus Assay

Sample Type: FFPE Block No.: S109-28920A Percentage of tumor cells: 90%

Note:

Sample Cancer Type: Liver Cancer

Table of Contents	Page
Variants (Exclude variant in Taiwan BioBank with >1% allele frequency)	1
Biomarker Descriptions	2

Report Highlights

0 Relevant Biomarkers0 Therapies Available0 Clinical Trials

Relevant Biomarkers

No clinically significant biomarkers found in this sample.

Prevalent cancer biomarkers without relevant evidence based on included data sources CTNNB1 p.(D32G) c.95A>G

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

DNA Sequence Variants Allele Amino Acid Change Variant ID Variant Effect Coverage Gene Coding Locus Frequency Transcript CTNNB1 p.(D32G) c.95A>G COSM5681 chr3:41266098 17.76% NM_001904.3 missense 1999

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Date: 08 Oct 2020 2 of 4

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}.

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 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.

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Pathologist:

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Signatures		
Testing Personnel:		
Laboratory Supervisor:		

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Date: 08 Oct 2020 4 of 4

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