

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: 23 Nov 2022 1 of 3

Sample Information

Patient Name: 許怡儂 Gender: Female ID No.: A225623842 History No.: 39928002

Age: 37

Ordering Doctor: DOC2589K 曾令民

Ordering REQ.: 0CCHPPW Signing in Date: 2022/11/23

Path No.: M111-00017 **MP No.:** BR22062

Assay: Oncomine BRCA1/2 Assay

Sample Type: Blood

Date of blood drawing: 2022/11/10

Reporting Doctor: DOC5424G 彭昱璟 (Phone: 8#5424)

Note:

Sample Cancer Type: Breast Cancer

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Relevant Biomarkers

No clinically significant biomarkers found in this sample.

Variant Details

DNA Sequence Variants

Amino Acid Change	Codina						
	County	Locus	Frequency	Transcript	Variant Effect	ClinVar1	Coverage
p.(?)	c26G>A	chr13:32890572	50.06%	NM_000059.3	unknown	Benign	3999
p.(K1132=)	c.3396A>G	chr13:32911888	99.75%	NM_000059.3	synonymous	Benign	3999
p.(L1521=)	c.4563A>G	chr13:32913055	99.92%	NM_000059.3	synonymous	Benign	3994
p.(V2171=)	c.6513G>C	chr13:32915005	99.97%	NM_000059.3	synonymous	Benign	4000
p.(S2414=)	c.7242A>G	chr13:32929232	99.70%	NM_000059.3	synonymous	Benign	3996
p.(V2466A)	c.7397T>C	chr13:32929387	99.82%	NM_000059.3	missense	Benign	3999
p.(S1613G)	c.4837A>G	chr17:41223094	49.30%	NM_007294.4	missense	Benign	3990
	p.(?) p.(K1132=) p.(L1521=) p.(V2171=) p.(S2414=) p.(V2466A)	p.(?) c26G>A p.(K1132=) c.3396A>G p.(L1521=) c.4563A>G p.(V2171=) c.6513G>C p.(S2414=) c.7242A>G p.(V2466A) c.7397T>C	p.(?) c26G>A chr13:32890572 p.(K1132=) c.3396A>G chr13:32911888 p.(L1521=) c.4563A>G chr13:32913055 p.(V2171=) c.6513G>C chr13:32915005 p.(S2414=) c.7242A>G chr13:32929232 p.(V2466A) c.7397T>C chr13:32929387	p.(?) c26G>A chr13:32890572 50.06% p.(K1132=) c.3396A>G chr13:32911888 99.75% p.(L1521=) c.4563A>G chr13:32913055 99.92% p.(V2171=) c.6513G>C chr13:32915005 99.97% p.(S2414=) c.7242A>G chr13:32929232 99.70% p.(V2466A) c.7397T>C chr13:32929387 99.82%	p.(?) c26G>A chr13:32890572 50.06% NM_000059.3 p.(K1132=) c.3396A>G chr13:32911888 99.75% NM_000059.3 p.(L1521=) c.4563A>G chr13:32913055 99.92% NM_000059.3 p.(V2171=) c.6513G>C chr13:32915005 99.97% NM_000059.3 p.(S2414=) c.7242A>G chr13:32929232 99.70% NM_000059.3 p.(V2466A) c.7397T>C chr13:32929387 99.82% NM_000059.3	p.(?) c26G>A chr13:32890572 50.06% NM_000059.3 unknown p.(K1132=) c.3396A>G chr13:32911888 99.75% NM_000059.3 synonymous p.(L1521=) c.4563A>G chr13:32913055 99.92% NM_000059.3 synonymous p.(V2171=) c.6513G>C chr13:32915005 99.97% NM_000059.3 synonymous p.(S2414=) c.7242A>G chr13:32929232 99.70% NM_000059.3 synonymous p.(V2466A) c.7397T>C chr13:32929387 99.82% NM_000059.3 missense	p.(?) c26G>A chr13:32890572 50.06% NM_000059.3 unknown Benign p.(K1132=) c.3396A>G chr13:32911888 99.75% NM_000059.3 synonymous Benign p.(L1521=) c.4563A>G chr13:32913055 99.92% NM_000059.3 synonymous Benign p.(V2171=) c.6513G>C chr13:32915005 99.97% NM_000059.3 synonymous Benign p.(S2414=) c.7242A>G chr13:32929232 99.70% NM_000059.3 synonymous Benign p.(V2466A) c.7397T>C chr13:32929387 99.82% NM_000059.3 missense Benign

¹ Based on Clinvar version 20200329

Variant Details (continued)

DNA Sequence Variants (continued)

Gene	Amino Acid Change	Coding	Locus	Allele Frequency	Transcript	Variant Effect	ClinVar ¹	Coverage
BRCA1	p.(S1436=)	c.4308T>C	chr17:41234470	51.60%	NM_007294.4	synonymous	Benign	4000
BRCA1	p.(K1183R)	c.3548A>G	chr17:41244000	50.08%	NM_007294.4	missense	Benign	3998
BRCA1	p.(E1038G)	c.3113A>G	chr17:41244435	50.64%	NM_007294.4	missense	Benign	3999
BRCA1	p.(P871L)	c.2612C>T	chr17:41244936	52.29%	NM_007294.4	missense	Benign	3995
BRCA1	p.(L771=)	c.2311T>C	chr17:41245237	49.21%	NM_007294.4	synonymous	Benign	3999
BRCA1	p.(S694=)	c.2082C>T	chr17:41245466	49.86%	NM_007294.4	synonymous	Benign	3999

¹ Based on Clinvar version 20200329

Date: 23 Nov 2022

Signatures

Testing Personnel:

Laboratory Supervisor:

Pathologist: