



## PATIENT INFORMATION

<b>PATIENT:</b> john Doe	<b>DOB:</b> 15 Feb 1966	<b>SEX:</b> M	<b>LAB ID:</b> L123	<b>MRN:</b> M123
<b>COLLECTION DATE</b> 27 Feb 2023	<b>FACILITY NAME</b> Production Test Clinic - DemoData			
<b>RECEIVED DATE</b> 01 Mar 2023	<b>SUBMITTING PHYSICIAN</b> Jane Doe	<b>PHONE</b> ---		
<b>REPORT DATE</b> 06 Mar 2023	<b>TREATING PHYSICIAN/CC</b> ---	<b>PHONE</b> ---		

**CLINICAL HISTORY:** No clinical history provided on the Afirma test requisition form

## RESULTS

**Nodule:** **A** Thyroid, Upper Right, 1.1 cm

## CYTOPATHOLOGY

I Non Diagnostic	II Benign	III Atypia of Undetermined Significance	IV Suspicious for Follicular Neoplasm	V Suspicious for Malignancy	VI Malignant
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**Cytopathology Diagnosis:** Indeterminate - Atypia of Undetermined Significance (AUS - Bethesda Category III)

**Diagnostic Comments:** These features are best classified as atypia of undetermined significance.

**Microscopic Description:** The cytologic and cell block preparations are sparsely cellular and show a few clusters of follicular cells in crowded or microfollicular groups and some colloid.

## AFIRMA GENOMIC SEQUENCING CLASSIFIER

## Ensemble Classifier

Suspicious

## Xpression Atlas

BRAF:p.K601E c.1801A&gt;G

## Other Classifiers

 BRAF p. V600E c. 1799T>A: Negative  
 RET/PTC1, RET/PTC3: Not Detected

 MTC: Negative  
 Parathyroid: Negative

## Clinical Relevance

Potential clinical significance in thyroid cancer

## Risk of Malignancy

~50%<sup>11</sup>

## Associated Neoplasm Type

Follicular Neoplasms (FA, NIFTP, FVPTC, FTC)

FDA Approved Therapy<sup>#</sup>

No alteration-specific therapy currently approved

## TERT PROMOTER REGION

TERT c.-124C>T (C228T): Not Detected  
 TERT c.-146C>T (C250T): Not Detected

## NODULE A RESULTS SUMMARY

The result of this 1.1cm Bethesda III nodule A is Afirma GSC Suspicious and BRAF p.K601E positive which suggests a risk of cancer of ~50%<sup>11</sup>. This genomic alteration is associated with follicular neoplasms (FA, NIFTP, FVPTC, FTC) and a RAS-like profile, which includes rates of lymph node metastases and extrathyroidal extension that are lower than BRAF V600E-like neoplasms, but higher than Non-BRAF-Non-RAS-like neoplasms<sup>9,10</sup>. Clinical correlation and surgical resection should be considered. Consider visiting [clinicaltrials.gov](https://clinicaltrials.gov) to see if there are any available clinical trials relevant to the described molecular variant/fusion discovered on Afirma testing.

## GROSS DESCRIPTION

A: Received 1 vial(s) of sample(s) in CytoLyt, 1 vial(s) of FNAProtect

**CYTOPATHOLOGY REVIEWED AND E-SIGNED ON 01 Mar 2023 06:49 PM BY:**

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**CLINICAL HISTORY:** No clinical history provided on the Afirma test requisition form

TEST PERFORMANCE

Afirma GSC - Ensemble Classifier <sup>1,5</sup>	Cytopathology Diagnosis Indeterminate*
Risk of Malignancy: Afirma GSC Benign	~4%
Risk of Malignancy: Afirma GSC Suspicious	~50%
Sensitivity:	91%
Specificity:	68%
Limit of Detection:	5%

	MTC <sup>3,5</sup>	BRAF V600E <sup>1,2,4,5,11</sup>	RET/PTC <sup>2,5,7,11</sup>	Parathyroid <sup>5,6</sup>	XA Nucleotide Variant Panel**	XA Fusion Panel***	TERT <sup>8</sup>
Sensitivity	>99%			>99%			
Specificity	>99%			>99%			
PPA		>99%			74%	82%	>99%
NPA		>99%	>99%		>99%	>99%	>99%
Confirmation Rate			>99%		>98%	>99%	>99%
Limit of Detection	20%	5%	10%	15%	5%	10%	5%

**References:** 1. Patel KN, et al. *JAMA Surg* 2018. 2. Haugen BR, et al. *Thyroid* 2016. 3. Randolph G, et al. *ATA* 2017. 4. Angell TE, et al. *ATA* 2017. 5. Hao, et al. *Frontiers in Endo* 2019. 6. Sosa JA, et al. *ATA* 2017. 7. Angell, et al. *Frontiers in Endo* 2019. 8. Data on file. 9. TCGA Research Network. *Cell* 2014 10. Yoo, et al. *PLoS Genetics* 2016 11. Goldner, et al. *Thyroid* 2019. 12. Stack, et al. *ATA* 2019. 13. Whitmer D, et al. *Frontiers in Endo* 2022.

\* Indeterminate includes Atypia of Undetermined Significance / Follicular Lesion of Undetermined Significance and (suspicious for) Follicular Neoplasm / Hürthle Cell Neoplasm.  
 † Analytical sensitivity studies demonstrated the test's ability to detect malignant cells in a background of benign cells.  
 ‡ BRAF classifier performance is based on a comparison to a castPCR DNA assay for the BRAF V600E mutation.  
 \*\* Nucleotide variant performance, excluding BRAF V600E, is based on a comparison to a DNA AmpliSeq assay that measures variants using a 5% variant allele frequency threshold.  
 \*\*\* Fusion performance is based on a comparison to an RNA AmpliSeq fusion assay and TaqMan assays.  
 § Confirmation rate is the proportion of positive calls that are confirmed positive by the reference method.  
 ¶ Analytical sensitivity studies demonstrate the test's ability to detect a positive variant in a background of wild type.  
 # FDA approved therapies for thyroid cancer, both specific for genomic alterations and non-specific, may be found at <https://www.cancer.gov/about-cancer/treatment/drugs/thyroid> and <https://www.cancer.gov/about-cancer/treatment/drugs/solid-tumors>. See <https://clinicaltrials.gov> for potentially relevant clinical trials. Afirma XA is not a companion diagnostic and is not conclusive for any therapy.

Associated Neoplasm Type abbreviations - FA, Follicular Adenoma; FTC, Follicular Thyroid Carcinoma; FVPTC, Follicular Variant of Papillary Thyroid Carcinoma; NIFTP, Noninvasive Follicular Thyroid Neoplasm with Papillary-Like Nuclear Features; PTC, Papillary Thyroid Carcinoma.

This NGS assay cannot differentiate somatic and germline variants. Further testing and/or genetic counseling may be warranted depending on the patient's clinical findings, family history and/or variant identified.

Afirma Thyroid FNA Analysis is a diagnostic service provided by Veracyte, Inc. for the assessment of thyroid nodules that includes cytopathology and molecular testing. The Ensemble Classifier, Parathyroid Classifier, MTC Classifier, BRAF V600E Classifier, XA, DNA assay of the TERT promoter region, and their performance characteristics were determined by Veracyte. The Ensemble Classifier measures the expression profile of RNA isolated from the nodule and classifies the sample as benign or suspicious for malignancy. The Parathyroid Classifier determines if the FNA specimen is positive or negative for parathyroid tissue. The Medullary Thyroid Carcinoma (MTC) Classifier determines if the nodule is positive or negative for MTC. The BRAF V600E Classifier measures RNA isolated from the nodule and classifies the sample as positive or negative for the BRAF V600E mutation. The RET/PTC assay sequences the RET and PTC genes to detect RET/PTC1 and RET/PTC3 fusions and reports them as detected or not detected. XA evaluates 593 genes included in Afirma GSC for 905 specific variants and 235 specific fusion pairs. The DNA analysis evaluates the two TERT promoter variants, C228T and C250T.