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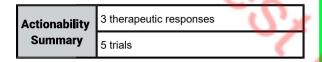
Summit™ Report

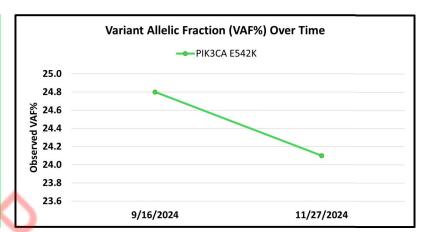
| Patient | Specimen | Physician Information |
|--|--------------------------|-----------------------------------|
| Name:Jane Doe | Type:CSF | Institution:Belay Diagnostics |
| DOB:01/01/1990 Sex Assigned at Birth: Female | Collected:11/27/2025 | Referring Physician:Provider Test |
| MRN:11xx22xx33 | Received:12/02/2024 | |
| Diagnosis:Metastatic Breast Carcinoma | Specimen ID:SumPos-TL-A1 | |

RESULT SUMMARY

POSITIVE







Comments

While no aneuploidy of clinical significance was detected, a high level of chromosomal loss and gain was observed in this sample.

ALTERATION DETAILS

| Clinically Significant Genomic Variants | | | | 70 |
|---|-------|-------------------------|----------------|---|
| Alteration | VAF | Type of Alteration | Classification | Clinical Implications |
| PIK3CA p.E542K c.1624G>A | 24.1% | Substitution - Missense | Tier 1A | Associated with Drug Response; Potentially Relevant Clinical Trials |

Aneuploidy Variants (Chromosome Arm Level Loss or Gain): None

Clinical Implications: Associated with drug response = related to drug sensitivity or resistance as described in Drug Response section of this report; Potentially relevant clinical trials = gene is related to a trial in the Clinical Trials section of this report; Prognostic Implications = related to prognosis as described in Prognostic Implications section of this report; Diagnostic Implications = related to diagnosis as described in Diagnostic Implications

VAF: Variant Allelic Frequency

| Variants of Unknown | Variants of Unknown Significance (Tier 3) | | | | | |
|---------------------|---|-------------|------------|--|--|--|
| chr11q Gain | chr17p Loss | chr21q Loss | chr8p Loss | | | |
| chr12q Loss | chr17q Loss | chr6p Loss | chr8q Gain | | | |
| chr16q Loss | chr19q Gain | chr6q Loss | chr9q Loss | | | |

ACTIONABILITY SUMMARY

| FDA / NCCN Therapies for the Patient's Tumor Type (Tier 1A) | | | | |
|---|-----------|---------|-----------|--|
| Biomarker | Therapies | Setting | Source(s) | |



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| PIK3CA E542K | capivasertib + fulvestrant | Advanced or metastatic; HR+/HER2- | FDA (Approved), NCCN |
|--------------|--|--|----------------------|
| PIK3CA E542K | inavolisib + palbociclib + fulvestrant | Recurrent and advanced/metastatic; HR+/HER2- | FDA (Approved), NCCN |
| PIK3CA E542K | alpelisib + fulvestrant | Subsequent line; HR+/HER2- | FDA (Approved), NCCN |

FDA / NCCN Therapies with Resistance / Decreased Response (Tier 1A): None

Prognostic Implications per NCCN: None

Diagnostic Implications per WHO: None

CLINICAL TRIALS / INVESTIGATIONAL THERAPIES

| PIK3CA E542K | IK3CA E542K | | | | | |
|---|---|---|--|--|--|--|
| Therapy | Clinical Trial | Location/Sponsor | | | | |
| afuresertib + fulvestrant | NCT04851613 (Phase 3) Study Evaluating Efficacy & Safety of Afuresertib Plus Fulvestrant in Patients w/ Locally Advanced or Metastatic HR+/HER2- Breast Cancer | Santa Monica, California Laekna Limited wenwen.he@laekna.com | | | | |
| alpelisib + (aromatase inhibitor or fulvestrant) | NCT04762979 (Phase 2) Alpelisib (BYL719) in Combination With Continued Endocrine Therapy Following Progression on Endocrine Therapy in Hormone Receptor Positive, HER2 Negative, PIK3CA Mutant Metastatic Breast Cancer | Chicago, Illinois Marina N Sharifi msharifi@wisc.edu | | | | |
| gedatolisib + fulvestrant; gedatolisib + palbociclib + fulvestrant | NCT05486143 (Phase 3) Gedatolisib Plus Fulvestrant With or Without Palbociclib vs Standard-of-Care for the Treatment of Patients With Advanced or Metastatic HR+/HER2- Breast Cancer | Winter Haven, Florida Celcuity, Inc. nzack@celcuity.com | | | | |
| gedatolisib + palbociclib + fulvestrant; gedatolisib + fulvestrant | NCT05501886 (Phase 3) Gedatolisib Plus Fulvestrant With or Without Palbociclib vs Standard-of-Care for the Treatment of Patients With Advanced or Metastatic HR+/HER2- Breast Cancer (VIKTORIA-1) | Birmingham, Alabama Celcuity Inc VIKTORIA-1_TRIAL@CELCUITY.COM | | | | |
| inavolisib + fulvestrant | NCT05646862 (Phase 3) A Study Evaluating the Efficacy and Safety of Inavolisib Plus Fulvestrant Compared With Alpelisib Plus Fulvestrant in Participants With HR-Positive, HER2-Negative, PIK3CA Mutated, Locally Advanced or Metastatic Breast Cancer Post CDK4/6i and Endocrine Combination Therapy | York, Pennsylvania Hoffmann-La Roche global-roche-genentech-trials@gene.com | | | | |

TIER 1A THERAPY DETAILS

| PIK3CA E542K | /), | |
|-------------------------|--|---|
| Therapy | Approval / Guideline Summary | Underlying Evidence |
| alpelisib + fulvestrant | FDA approved for HR-positive, HER2-negative, PIK3CA-mutated, advanced or metastatic breast cancer following progression on or after an endocrine-based regimen. NCCN recommended as Category 1/Preferred intervention. | The FDA approval for alpelisib + fulvestrant was supported by data from the randomized, double-blind, placebo-controlled phase-III trial SOLAR-1 (NCT02437318). SOLAR-1 demonstrated that subsequent-line alpelisib + fulvestrant (n = 169), compared with placebo + fulvestrant (n = 172), improved median PFS (HR = 0.65, p = 0.0013; 11.0 mo. vs. 5.7 mo.) in patients with advanced or metastatic, HR-positive, HER2-negative, PIK3CA-mutated breast carcinoma. The additional endpoints were ORR (35.7% vs. 16.2%) and OS (HR = 0.86, n.s.). |



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| capivasertib + fulvestrant | FDA approved for adults with HR-positive, HER2-negative, locally advanced or metastatic breast cancer with one or more PIK3CA/AKT1/PTEN alterations, following progression on at least one endocrine-based regimen in the metastatic setting or recurrence on or within 12 months of completing adjuvant therapy. NCCN recommended as Category 1/Preferred intervention. | The FDA approval for capivasertib + fulvestrant was supported by data from the placebo-controlled, double-blind, randomized, phase-III trial CAPItello-291 (NCT04305496). CAPItello-291 demonstrated that subsequent-line capivasertib + fulvestrant (n = 155), compared with placebo + fulvestrant (n = 134), improved PFS (HR: 0.50, p <0.0001; 7.3 mo. vs. 3.1 mo.) in patients with locally advanced or metastatic HR-positive, HER2-negative breast cancer with with one or more PIK3CA/AKT1/PTEN-alterations. Additional endpoints were ORR (26% vs. 8%) and median DoR (10.2 mo. vs. 8.6 mo.). |
|--|--|---|
| inavolisib + palbociclib + fulvestrant | FDA approved for adults with endocrine-resistant, PIK3CA-mutated, HR-positive, HER2-negative, locally advanced or metastatic breast cancer, following recurrence on or after completing adjuvant endocrine therapy. NCCN recommended as Category 1/Useful in certain circumstances. | The FDA approval for inavolisib + palbociclib + fulvestrant was supported by data from the double blind, randomized, placebo-controlled, phase-III trial INAVO120 (NCT04191499). INAVO120 demonstrated that inavolisib + palbociclib + fulvestrant (n = 161), compared to placebo + palbociclib + fulvestrant (n = 164), improved PFS (15.0 mo. vs. 7.3 mo., HR = 0.43; p < 0.0001) in patients with endocrine-resistant, PIK3CA-mutated, HR-positive, HER2-negative, locally advanced or metastatic breast cancer following recurrence on or after completing adjuvant endocrine therapy. Additional endpoints include an ORR of 58%, median DoR (18.4 mo.) and OS (not mature with 30% deaths in the overall population). |

TEST DETAILS

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| SNVs and Indels (32 genes, 112 amplicons) | | | | | | |
|---|----------------------|--------------------|---|--|--|--|
| AKT1 NM_001014432.1 | ERBB2 NM_004448.2 | GNAS NM_000516.4 | NRAS NM_002524.4 PIK3CA NM_006218.2 PTEN NM_000314.4 RAF1 NM_002880.3 SMAD4 NM_005359.5 TERT NM_198253.2 TP53 NM_000546.5 VHL NM_000551.3 | | | |
| APC NM_000038.5 | ERBB3 NM_001982.3 | H3-3A NM_002107.4 | | | | |
| BRAF NM_004333.4 | ERCC2 NM_000400.3 | HRAS NM_005343.2 | | | | |
| CD79B NM_000626.2 | FBXW7 NM_033632.3 | IDH1 NM_005896.2 | | | | |
| CDH1 NM_004360.3 | FGFR2 NM_000141.4 | IDH2 NM_002168.2 | | | | |
| CDKN2A NM_000077.4 | FGFR3 NM_000142.4 | KRAS NM_004985.3 | | | | |
| CTNNB1 NM_001904.3 | FUS NM_004960.3 | MYD88 NM_002468.4 | | | | |
| EGFR NM_005228.3 | GATA3 NM_001002295.1 | NFE2L2 NM_006164.4 | | | | |

| Aneuploidy (chrom | Aneuploidy (chromosome arm level loss and gain) | | | | | | | | |
|----------------------------------|---|----------------------------------|-------|------------------------------------|--------|------------------|--------------------------------------|--------------------------------------|----------------------------|
| chr1p chr1q chr2p chr2q | chr3p chr3q chr4p chr4q | chr5p chr5q chr6p chr6q | chr8p | chr9p chr9q chr10p chr10q | chr11q | chr14q chr15q | chr16q chr17p chr17q chr18p | chr18q chr19p chr19q chr20p | chr20q chr21q chr22q |

Methods and Limitations

The Summit™ next-generation sequencing (NGS) test investigates tumor DNA (tDNA) extracted from cerebrospinal fluid (CSF) for clinically relevant single/multi nucleotide variants (SNVs, MNVs, indels) and aneuploidy events associated with primary and metastatic central nervous system (CNS) cancers. Methodology involves targeted duplex sequencing of 32 key genes (SNVs, MNVs and Indels) and low pass whole genome sequencing (>0.1x) for the detection of chromosomal arm level loss or gain, aneuploidy (PMID: 37014860). Post target enrichment libraries, generated from 20-40ng of tDNA, are sequenced on the Illumina NovaSeq XPlus, generating 100 bp paired-end sequence reads. The LOD (limit of detection) for SNVs, MNVs and Indels was determined as 0.3% variant allelic fraction (VAF). Variants (mutations and aneuploidy) are called against the human genome build reference hg19 using the Summit™ Genome Analytics (SGA) pipeline (SNVs, MNVs, and Indels - version 1.0.0 and aneuploidy - version 0.6.0), developed at Belay Diagnostics.

Tertiary analysis is performed using the precision oncology workbench (GenomOncology) based on the joint AMP/ASCO/CAP consensus guidelines for interpretation of sequence variants in cancer (PMID: 27993330). Please reach out to contact@belaydiagnostics.com for additional information or queries.

Disclaimers

This test was developed, and its performance characteristics determined by Belay Diagnostics Laboratory (CLIA# 14D2302605), which is certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA) as qualified to perform high complexity testing. This test has not been cleared or approved by the U.S. Food and Drug Administration (FDA). This test may be used for clinical purposes. However, the results of this test do not establish a diagnosis and should not be used alone for diagnosis or patient care decisions or otherwise replace the judgment of a treating physician and must always be interpreted in the context of all relevant clinical and pathological data.



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This test is performed only to evaluate for somatic (i.e., tumor-specific) variants within the genes listed and cannot distinguish between germline and somatic alterations with absolute certainty. This test therefore does not report on incidental findings as defined by the American College for Medical Genetics and Genomics (ACMG) (PMID: 37347242). If a germline variant is suspected, follow-up germline testing using non-neoplastic (normal) tissue should be performed by a laboratory permitted to perform germline genetic testing along with genetic counseling. It is possible for a genomic variant to be present yet go undetected by our assay either due to the heterogeneous nature of the specimen or the limits of detection of our assay. Therefore, to the extent a particular genomic variant is not reported, Belay Diagnostics LLC does not guarantee that the variant does not exist in the specimen provided. Likely benign, and benign variants are not reported. For any reported variant of uncertain significance (VUS), if the classification changes, there is no obligation to send out a new report updating this information.

The information presented in the clinical trials and therapeutic sections of this report is compiled from public sources which are continuously updated. While we strive to ensure this information is accurate and complete, we cannot guarantee the accuracy or completeness of this information. This public sourced information is not ranked in order of potential or predicted efficacy and may not be complete. Specific eligibility criteria should be reviewed as applicable. This information may include associations between a genomic variant (or lack of a variant) and one or more therapeutic agents with potential clinical benefit (or lack of clinical benefit), including agents that are being studied in clinical research. The finding of a genomic variant does not necessarily indicate or demonstrate pharmacologic effectiveness (or lack thereof) of any agent or treatment regimen found in public source information. Similarly, the finding of "no clinically significant variant" does not necessarily indicate or demonstrate lack of pharmacologic effectiveness (or lack of effectiveness) of any agent or treatment regimen found in public source information. Belay Diagnostics expressly disclaims, and makes no representation of or warranty of, the accuracy or completeness with respect to the publicly available information included herein or reviewed or collected during creation of this report.

ACTIONABILITY REFERENCES

FDA: U.S. Food & Drug Administration (fda.gov)

NCCN: National Comprehensive Cancer Network® (NCCN®). Referenced with permission from the NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines®). © National Comprehensive Cancer Network, Inc. 2024. All rights reserved. The NCCN Guidelines® and illustrations herein may not be reproduced in any form for any purpose without the express written permission of the NCCN. To view the most recent and complete version of the guideline, go online to NCCN.org. NCCN makes no warranties of any kind whatsoever regarding their content, use or application and disclaims any responsibility for their application or use in any way.

WHO: World Health Organization Classification of Tumours online (tumourclassification.iarc.who.int)

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