BECAUSE TIME MATTERS IDYLLA™ PIK3CA-AKT1 MUTATION ASSAY





THINK IDYLLA™ BECAUSE TIME MATTERS

INTRODUCING IDYLLA[™] PIK3CA-AKT1 MUTATION ASSAY

FOR THE MOLECULAR CHARACTERIZATION OF PIK3CA AND AKT1 IN FFPE TUMOR TISSUE SECTIONS



Qualitative detection of **13 PIK3CA** mutations and **1 AKT1** mutation in a single cartridge



Automated result interpretation and test reporting



Less than 2 minutes hands-on time (HOT) Assay turnaround time (TAT) of approx. 150 minutes



Directly from FFPE tumor tissue sections

SPECIMEN REQUIREMENTS

+ 50-600 mm² tissue area for 5 μ m FFPE tissue sections

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- + 25-300 mm² tissue area for 10 μm FFPE tissue sections
- ≥ 20% neoplastic cells

PIK3CA AND AKT1 FACTS

PIK3CA activating mutations have been identified in a broad spectrum of solid tumors¹. In Hormone Receptor (HR)+ breast cancer samples, the prevalence of PIK3CA mutations is 30% to 40%². The Idylla[™] PIK3CA-AKT1 Mutation Assay detects the most prevalent PIK3CA mutations in breast cancer.

| | Distribution of mutations in PIK3CA mutated breast cancer (%) | ldylla™ PIK3CA-AKT1 Mutation Assay |
|--------|---|---|
| H1047R | 35.0 | |
| E545K | 17.5 | ✓ |
| E542K | 10.7 | ✓ |
| N345K | 5.5 | 1 |
| H1047L | 4.0 | ✓ |
| C420R | 1.9 | 1 |
| Q546R | 1.1 | Image: A start of the start of |
| Q546K | < 1.0 | Image: A start of the start of |
| E545A | < 1.0 | ✓ |
| E545G | < 1.0 | 1 |
| E545D | < 1.0 | 1 |
| Q546E | < 1.0 | Image: A start of the start of |
| H1047Y | < 1.0 | 1 |

AKT1-E17K mutations account for 3% of the mutations in HR+ breast cancer samples and are an emerging biomarker in the development of targeted therapies³.



IDYLLA[™] PIK3CA-AKT1 MUTATION ASSAY SHOWS EXCELLENT PERFORMANCE AGAINST NGS

The Idylla[™] PIK3CA-AKT1 Mutation Assay provides a rapid actionable solution which can be seamlessly integrated into virtually any laboratory workflow. The Assay demonstrated 100% concordance with NGS for the presence or absence of PIK3CA and AKT1 mutations in 31 FFPE breast cancer samples with three replicates. The comparison of the results is shown below.

| ldylla™ | Next Generation Sequencing | | |
|---------------------|---|--|--|
| | N345K E542K E545K H1047R H1047L E17K WT TOTAL | | |
| N345K | 15 | | |
| E542K | 12 | | |
| E545K | 18 | | |
| H1047R | 30 | | |
| H1047L | 3 | | |
| E17K | 3 | | |
| WT | 15 | | |
| TOTAL | 96 | | |
| E17K WT TOTAL | 3 15 96 | | |

Positive Percent Agreement (PPA): 100%

Negative Percent Agreement (NPA): 100%

REFERENCES

(1) Ligresti et al. PIK3CA mutations in human solid tumors. Cell Cycle. 2009; 8(9): 1352-1358.

- (2) Martínez-Sáez et al. Frequency and spectrum of PIK3CA somatic mutations in breast cancer. Breast Cancer Res. 2020 May 13;22(1):45.
- (3) Turner et al. Capivasertib in Hormone Receptor-Positive Advanced Breast Cancer. NEJM. 2023; 388:2058-70.

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