Patient

Name: Terri Brennan Date of Birth: 2006-02-24

Sex: Male

Case Number: 3587

Diagnosis: Adenocarcinoma

Specimen Information

Ordered By

Primary Tumor Site: Pancreatic

Specimen Site: Duodenum (for periampullary tumors)

Specimen ID: 8379

Specimen Collected: 2024-02-18

Test Initiated: 2024-02-23

| Biomarker | Method | Analyte | Result | | Biomarker level | |
|-----------|--------|-----------|---------------------|-------------------|---|---------|
| ER | IHC | protein | Positive 3+, 100% | BENEFIT | abemaciclib, palbociclib, ribociclib, endocrine, therapy, everolimus | level 2 |
| PR | IHC | protein | Positive 2+, 95% | BENEFIT | abemaciclib, palbociclib, ribociclib, endocrine therapy | level 2 |
| ТМВ | seq | DNA tumor | 11 m/Mb Low | BENEFIT | pembrolizumab | level 2 |
| ERBB2 | IHC | Protien | Negative 0 | LACK OF BENEFT | trastuzumab, ado-trastuzumab emtansine, pertuzumab, fam-trastuzumab deruxtecan-nxki, lapatinib, neratinib, tucatinib | level 1 |

Cancer-Type Relevant Biomarkers

| BioMarker | Method | Analyte | Result |
|------------------------|--------|-----------|---------------------|
| AR | IHC | Protien | Positive 3+, 43% |
| MUTYH | Seq | RNA-Tumor | Fusion not detected |
| AR | IHC | Protien | Negative 3+, 97% |
| Mismatch repair status | IHC | Protien | Negative 1+, 80% |
| MYO1G | Seq | DNA-Tumor | Stable |

Genomic Signatures

| BioMarker | Method | Analyte | Result |
|--------------------------------------|--------|-----------|--|
| Microsatellite instability | Seq | DNA tumor | Low |
| Tumor mutational burden | Seq | DNA tumor | 11 mutations/Mb Low |
| Genomic loss of heterozygosity (LOH) | Seq | DNA tumor | Low - 10% of tested genmoic segments exhibit LOH |

Genes Tested with Pathogenic Alterations or likely Pathogenic Alterations

| Gene | Method | Analyte | Variant Interpretation | Protien Alteration | Exon | DNA Alteration | Allele Frequency % |
|-------|--------|-----------|------------------------|--------------------|------|----------------|--------------------|
| ERBB2 | Seq | DNA tumor | Benign | p.R80X | 10 | c.322G>C | 24.7 |
| IDH2 | Seq | DNA tumor | Benign | p.A774X | 11 | 1 | 6.74 |
| KIT | Seq | DNA tumor | Benign | p.D463H | 9 | c.1387G>C | 28.75 |
| RB1 | Seq | DNA tumor | Benign | p.I543V | 17 | c.1679T>G | 6.67 |
| TEK | Seq | DNA tumor | Likely Pathogenic | p.M1R | 9 | c.2T>G | 7.8 |
| MAPK1 | Seq | DNA tumor | Benign | p.Q125R | 14 | c.602G>A | 2.46 |

Gene Variants of Unknown Significance

| Gene | Method | Analyte | Variant Interpretation | Protien Alteration | Exon | DNA Alteration | Allele Frequency % |
|-------|--------|-----------|-----------------------------------|--------------------|------|----------------|--------------------|
| FLT3 | Seq | DNA tumor | Variant of uncertain significance | p.G1386D | 10 | 1 | 5.19 |
| CCND3 | Seq | DNA tumor | Variant of uncertain significance | p.M1R | 18 | c.2T>G | 3.27 |
| FOXL2 | Seq | DNA tumor | Variant of uncertain significance | p.K659E | 20 | c.1647T>A | 17.8 |

Immunohistochemistry Results

| Biomarker | Result |
|-----------|-------------------|
| PMS2 | Positive 3+, 98% |
| AR | Negative 3+, 94% |
| MSH2 | Positive 2+, 60% |
| ERBB2 | Positive 1+, 9% |

Genes Tested with Indeterminate Results by Tumor DNA Sequencing

PAX5 SF3B1 FBXW7 NTRK1

Specimen Information

Specimen ID: 8379 Specimen Collected: 2024-02-18 Specimen Recieved: 2024-02-23 Testing Initiated: 2024-02-23

Gross Description: 8379

Pathological Diagnosis:

Left breast, central, 12:00, suspicious mass, 12-gauge core needle biopsy: Infiltrating moderately-differentiated mammary carcinoma, grade 2, Nottingham score 6 (architectural grade 3, nuclear grade 2, mitotic figures 1).

Dissection Information:

Molecular testing of this specimen was performed after harvesting of targeted tissues with an approved manual microdissection technique. Candidate slides were examined under a microscope and areas containing tumor cells (and separately normal cells, when necessary for testing) were circled. A laboratory technician harvested targeted tissues for extraction from the marked areas using a dissection microscope.

Clinical Trials Connector

| CHEMOTHERAPY CLINICAL TRIALS | | | | | | |
|--|--------|-----|------------------------|---|--|--|
| Drug class Biomarker Method Analyte Investigational ag | | | Investigational agents | | | |
| Anti hormonal therapy | ER | IHC | protein | anastrazole, letrozole, exemestane, fulvestrant, tamoxifen, goserelin, leuprolide | | |
| Anti hormonal therapy | PR | IHC | protein | anastrazole, letrozole, exemestane, fulvestrant, tamoxifen, goserelin, leuprolide | | |
| Anti inflammatory agents | PIK3CA | NGS | DNA tumor | aspirin | | |

| TARGETED THERAPY CLINICAL TRIALS | | | | | | |
|----------------------------------|--------|---------|------------------------|--|--|--|
| Drug class Biomarker Method | | Analyte | Investigational agents | | | |
| Akt inhibitors | ARID1A | NGS | DNA tumor | AZD5363, MK-2206, ipataserib | | |
| immunomodulatory agents | TMB | NGS | DNA tumor | avelumab, atezolizumab, durvalumab, ipilimumab, nivolumab, pembrolizumab | | |
| PARP inhibitors | NBN | NGS | DNA tumor | BGB-290, BMN-673, olaparib, rucaparib, talazoparib | | |
| Akt/mTor inhibitors | PIK3CA | NGS | DNA tumor | AZD5363, BYL719, MK-2206, ipataserib, everolimus, temsirolimus | | |