



Patient

Name: Patient, Test

Date of Birth: XX/Mon/19XX

Sex: Female

Case Number: TN19-XXXXXX **Diagnosis:** Sarcomatoid carcinoma

Specimen Information

Primary Tumor Site: Anterior wall of bladder

Specimen Site: Bladder, NOS Specimen ID: ABC-1234-XYZ Specimen Collected: XX-Mon-2019 Completion of Testing: XX-Mon-2019

Ordered By

Ordering Physician, MD Cancer Center 123 Main Street Springfield, XY 12345, USA

1 (123) 456-7890

High Impact Results

| BIOMARKER | METHOD | RESULT | THERAPY ASSOCIATION | BIOMARKER LEVEL* |
|-----------|--------------|-------------------|-----------------------|---------------------|
| PD-L1 | 22c3 IHC-CPS | Positive, CPS: 60 | BENEFIT pembrolizumab | Level 1 |

^{*} Biomarker reporting classification: Level 1 - highest level of clinical evidence and/or biomarker association included on the drug label; Level 2 - strong evidence of clinical significance and is endorsed by standard clinical guidelines; Level 3 - potential clinical significance (3A - evidence exists in patient's tumor type, 3B - evidence exists in another tumor type).

Important Note

Please note that PDL1 CPS score is 60 (>=10). CPS is calculated as the number of PD-L1 staining cells (tumor cells, lymphocytes, macrophages) divided by the total viable tumor cells, multiplied by 100. Threshold for positive staining is a CPS >=10. This PD-L1 score is sufficient for use of pembrolizumab in the front-line metastatic setting. Use of pembrolizumab is FDA approved for the treatment of locally advanced or metastatic bladder cancer who are not eligible for cisplatin-containing chemotherapy with a PD-L1 CPS>=10, or in patients who are not eligible for any platinum-containing chemotherapy regardless of PD-L1 status.

Additional Results

| CANCER TYPE RELEVANT BIOMARKERS | | | | | | | |
|---------------------------------|---------|------------------------|--|--|--|--|--|
| Biomarker | Method | Result | | | | | |
| MSI | NGS | Stable | | | | | |
| Mismatch Repair Status | | Proficient | | | | | |
| NTRK1 | RNA-Seq | Fusion Not Detected | | | | | |
| NTRK2 | RNA-Seq | Fusion Not Detected | | | | | |
| NTRK3 | RNA-Seq | Fusion Not Detected | | | | | |
| Tumor Mutational Burd | en | High 19 Mutations/Mb | | | | | |
| ATM | NGS | Mutation Not Detected | | | | | |
| ERBB2 (Her2/Neu) | NGS | Mutation Not Detected | | | | | |
| ERCC2 | NGS | Mutation Not Detected | | | | | |
| FANCC | NGS | Mutation Not Detected | | | | | |
| FGFR1 | RNA-Seq | Fusion Not Detected | | | | | |

| CANCER TYPE RELEVANT BIOMARKERS (cont) | | | | | | |
|--|---------|-----------------------|--|--|--|--|
| Biomarker Metho | | Result | | | | |
| FGFR2 RNA-Se | | Fusion Not Detected | | | | |
| EGER3 | NGS | Mutation Not Detected | | | | |
| rdro | RNA-Seq | Fusion Not Detected | | | | |
| RB1 | NGS | Mutation Not Detected | | | | |
| TSC1 | NGS | Mutation Not Detected | | | | |

| OTHER FINDINGS (see page 2 for additional results) | | | | | | |
|--|--------|------------------------------|--|--|--|--|
| Biomarker | Method | | | | | |
| ARID1A | NGS | Mutated, Pathogenic | | | | |
| | | Exon 3 p.Q479* | | | | |
| FRBB3 | NGS | Mutated, Presumed Pathogenic | | | | |
| LINDUS | CDNI | Exon 3 p.M911 | | | | |

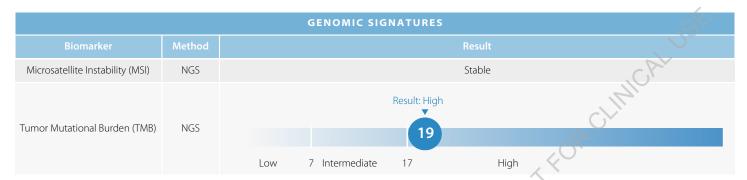
The selection of any, all, or none of the matched therapies resides solely with the discretion of the treating physician. Decisions on patient care and treatment must be based on the independent medical judgment of the treating physician, taking into consideration all available information concerning the patient's condition, the FDA prescribing information for any therapeutic, and in accordance with the applicable standard of care. Whether or not a particular patient will benefit from a selected therapy is based on many factors and can vary significantly. All trademarks and registered trademarks are the property of their respective owners.





Biomarker Results

This summary includes biomarkers most commonly associated with cancer. Complete details of all biomarkers tested can be found in the Appendix.



| | GENES TESTED WITH MUTATIONS/ALTERATIONS | | | | | | | | |
|--------|---|--|--------------------|------|----------------|---------------------|--|--|--|
| Gene | Method | Variant Interpretation | Protein Alteration | Exon | DNA Alteration | Variant Frequency % | | | |
| ARID1A | NGS | Mutated, Pathogenic | p.Q479* | 3 | c.1435C>T | 14 | | | |
| BRCA2 | NGS | Mutated, Variant of Unknown Significance | p.S1667L | 11 | c.5000C>T | 15 | | | |
| ERBB3 | NGS | Mutated, Presumed Pathogenic | p.M91I | 3 | c.273G>A | 38 | | | |
| TP53 | NGS | Mutated, Pathogenic | p.Q331* | 9 | c.991C>T | 38 | | | |

Transcript ID and Variants of Unknown Significance can be found in the Appendix.

Other Findings

| IMMUNOHISTOCHEMISTRY (IHC) | | | | | | | |
|----------------------------|---------------------|--------------|---------------------|--|--|--|--|
| Biomarker | Result | Biomarker | Result | | | | |
| MLH1 | Positive 2+, 90% | PD-L1 (22c3) | Positive, CPS: 60 | | | | |
| MSH2 | Positive 2+, 100% | PMS2 | Positive 1+, 100% | | | | |
| MSH6 | Positive 2+, 95% | | | | | | |

| | GENES TESTED WITHOUT POINT MUTATIONS OR INDELS BY NGS | | | | | | | | | | |
|------|---|-------|-------|--------|-------|-------------------------|-------|--------|--------|------|------|
| ATM | BRAF | BRCA1 | CCND1 | CDKN2A | EGFR | ERBB2 (Her2/ Neu) | ERCC2 | FANCC | FGFR3 | HRAS | IDH1 |
| KIT | KRAS | MET | MTOR | NRAS | NTRK1 | NTRK2 | NTRK3 | PDGFRA | PIK3CA | RB1 | RET |
| TSC1 | | | | | | | | | | | |

Additional results continued on the next page. >

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Clinical Trials Connector™

For a complete list of open, enrolling clinical trials visit MI Portal to access the <u>Clinical Trials Connector</u>. This personalized, real-time web-based service provides additional clinical trial information and enhanced searching capabilities, including, but not limited to:

- · Location: filter by geographic area
- Biomarker(s): identify specific biomarkers associated with open clinical trials to choose from
- Drug(s): search for specific therapies
- Trial Sponsor: locate trials based on the organization supporting the trial(s)

Visit www.CarisMolecularIntelligence.com to view all matched trials. Therapeutic agents listed below may or may not be currently FDA approved for the tumor type tested.

| TARGETED THERAPY CLINICAL TRIALS (211) | | | | | | | | |
|--|-----------|--------|--|--|--|--|--|--|
| Drug Class | Biomarker | Method | Investigational Agent(s) | | | | | |
| Akt inhibitors (4) | ARID1A | NGS | ARQ092, AZD5363 | | | | | |
| Cell cycle inhibitors (3) | TP53 | NGS | LY2606368 | | | | | |
| Immunomodulatory agents (196) | PD-L1 | IHC | MEDI4736, MK-3475, MPDL3280A, MSB0010718C, | | | | | |
| | TMB | NGS | atezolizumab, avelumab, durvalumab, nivolumab, pembrolizumab | | | | | |
| Pan-HER inhibitors (8) | ERBB3 | NGS (S | afatinib, lapatinib, neratinib, pyrotinib | | | | | |

() = represents the total number of clinical trials identified by the Clinical Trials Connector for the provided drug class or table.

All informations and in Please refer to the "Notes of Significance" section that may contain additional information regarding therapy associations.

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