# Monitor your patients with gynecologic cancer using tumor-informed ctDNA testing to inform management decisions

Current tools have limited sensitivity and specificity; Signatera<sup>™</sup> can help monitor your patients across the treatment continuum.



## Signatera<sup>™</sup> identifies patients at high risk of recurrence and detects recurrence earlier<sup>1</sup>



 $\triangleright$ 

After completion of adjuvant/definitive therapy, ctDNA was detectable in 23% of patients, all of whom experienced disease progression.<sup>1</sup>

Longitudinally, ctDNA was detected with 100% sensitivity and specificity<sup>1</sup>, while CA-125 had much lower sensitivity (43%) and specificity (78%).<sup>2</sup>

# Signatera<sup>™</sup> quantifies ctDNA over time to help answer critical questions during immunotherapy treatment

- > Is the treatment working?
- Should treatment be changed or escalated?
- Is the cancer actually progressing?
- Is the patient an exceptional responder?



Signatera<sup>™</sup> ctDNA dynamics help inform early if single agent IO treatment is working or if a change in therapy should be considered.



Early ctDNA dynamics during immunotherapy have been demonstrated to be a strong predictor of treatment benefit across solid tumors and in gynecologic cancers specifically.<sup>3-5</sup>

### Ordering Signatera<sup>™</sup> for your gynecologic cancer patients:

- Can be run at any time point from diagnosis through recurrence
- Medicare coverage for patients with stage II-IV ovarian cancer to inform treatment decisions in the adjuvant setting and for early
  detection of recurrence in the surveillance/maintenance settings, as well as monitoring response to immunotherapy in patients
  with any solid tumor
- One blood draw, one tumor sample, three results: easily add on hereditary cancer and/or tumor genomic profiling to your Signatera<sup>™</sup> order

#### Natera's Oncology Portfolio:

Empower<sup>™</sup> Hereditary cancer test Empower<sup>™</sup> for hereditary cancer testing provides clear answers you can act on by analyzing up to 81 genes across 12+ common hereditary cancers. Empower includes more commonly known genes associated with gynecologic cancers (including *BRCA1, BRCA2, BRIP1, RAD51C, RAD51D*, and mismatch repair (MMR) genes for Lynch Syndrome), to help guide therapeutic decisions.

Altera<sup>™</sup> Tumor genomic prof Add on Altera<sup>™</sup> Genomic Profiling test which utilizes whole-exome and whole-transcriptome sequencing to identify clinically relevant biomarkers that may help guide treatment selection (including MSI, *BRCA1/2*, HR genes, MMR genes, TMB, *BRAF, RET*, and *NTRK*), with no additional tumor sample needed.

#### References:

- 1. Hou et al. Gynecol Oncol. 2022; 167:334-341.
- 2. Chapman et al. Poster presented at 2021 AACR Annual Meeting.
- 3. Bratman et al. Nature Cancer. 2020; 1(9):873-881.
- 4. Recio et al. Poster presented at 2023 SGO Annual Meeting.
- 5. Backes et al. Poster presented at 2024 SGO Annual Meeting

Learn more at: natera.com/oncology



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Signatera<sup>™</sup> has been developed and its performance characteristics determined by the CLIA-certified laboratory performing the test. The test has not been cleared or approved by the US Food and Drug Administration (FDA). CAP accredited, ISO 13485 certified, and CLIA certified. © 2024 Natera, Inc. All Rights Reserved. SGN\_MD\_OS\_Gyn\_20240312\_NAT-9300001