

COLORECTAL CANCER (CRC)



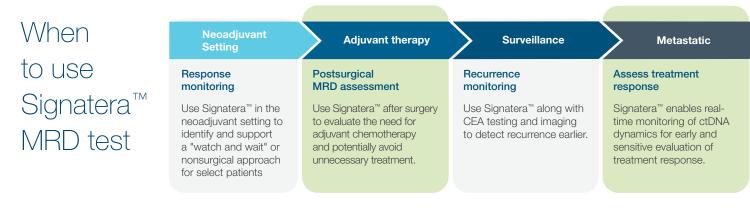
Know Cancer's Next Move

- > Does my patient need adjuvant chemotherapy?
- > Is the treatment working?
- > Is the cancer recurring?

A personalized, tumor-informed approach for molecular residual disease (MRD) detection





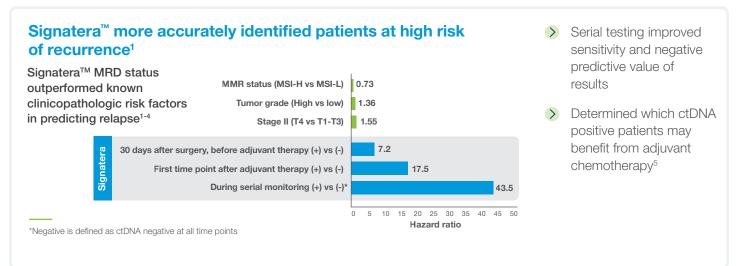


In the neoadjuvant setting

Tailor treatment or surgical strategies based on MRD status using primary tumor biopsy

In the adjuvant setting

Use after surgery to evaluate the benefit of adjuvant chemotherapy



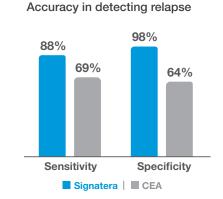
In the surveillance setting

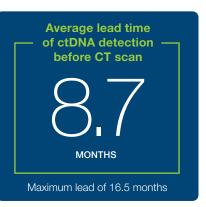
Use along with CEA testing and other surveillance tools to detect recurrence earlier, to inform surgical and / or other early intervention

Signatera[™] was shown to detect relapse more accurately than CEA with clinically meaningful lead times over CT scans¹

- Get clarity when evaluating patients with indeterminate CEA levels or CT scans
- Signatera[™] facilitates shared decision-making and confident treatment planning

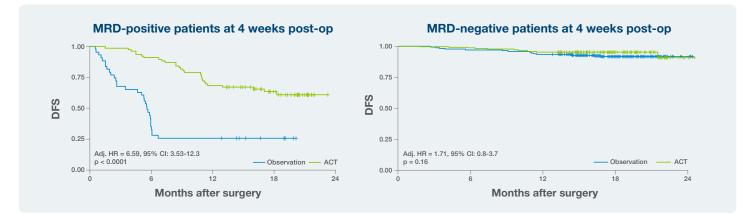
CEA = carcinoembryonic antigen **CT** = computed tomography **ctDNA** = circulating-tumor DNA





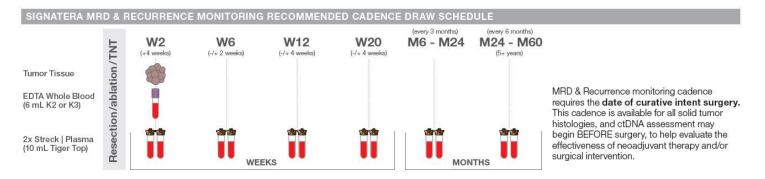
A large-scale prospective, MRD-guided study evaluated the clinical utility of ctDNA analysis in colorectal cancer (CRC)

MRD-positive CRC patients at 4 weeks post-op benefited significantly from chemotherapy while MRD-negative patients at 4 weeks post-op *did not* demonstrate any significant trend in treatment benefit



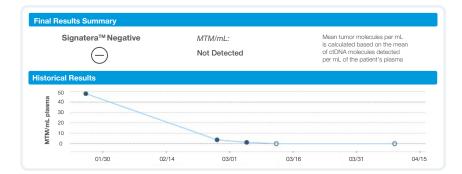
Ordering Signatera[™] for patients with colorectal cancer

Signatera™ is custom-designed for each patient using their own tumor tissue



Track ctDNA dynamics to enable longitudinal monitoring

Signatera[™] reports presence/absence of ctDNA and ctDNA quantity in terms of MTM/mL for longitudinal assessment



 $\textbf{MTM} = mean \ tumor \ molecules$

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Just like no two tumors are alike—Signatera™ is personalized for each patient



Tumor-informed MRD assay for individualized care

• Customized for each patient's unique tumor signature using WES to target the top clonal mutations

High sensitivity and specificity for accurate MRD assessment

- By only tracking tumor-specific variants, sensitivity is optimized with a LOD down to 0.01% VAF⁶
- Filters out germline and CHIP mutations to reduce background noise and to minimize false positives

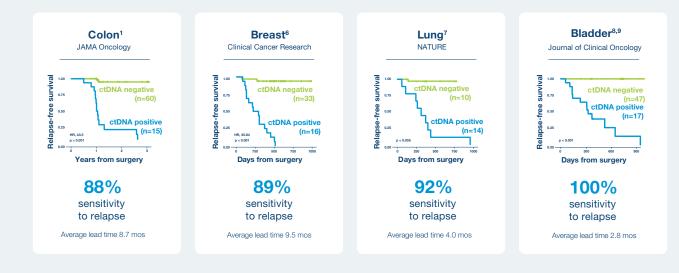


Reliable longitudinal monitoring for confident decision-making

- Tracks ctDNA dynamics by MTM/mL to enable longitudinal monitoring with a simple blood draw
- Follows clonal mutations that should persist as the tumor evolves

LOD = limit of detection; CHIP = clonal hematopoiesis of indeterminate potential; VAF = Varient allele frequency; WES = whole exome sequencing

Signatera[™] is validated across multiple tumor types^{1,6-8}



References

1. Reinert T, Henriksen TV, Christensen E, et al. Analysis of Plasma Cell-Free DNA by Ultradeep Sequencing in Patients With Stages I to III Colorectal Cancer. *JAMA Oncol.* 2019.

 Sinicrope FA, Foster NR, Thibodeau SN, et al. DNA Mismatch Repair Status and Colon Cancer Recurrence and Survival in Clinical Trials of 5-Fluorouracil-Based Adjuvant Therapy. J Natl Cancer Inst. 2011;103(11):863–875.

3. Aoyama, Oba K, Honda M, et al. Impact of postoperative complications on the colorectal cancer survival and recurrence: analyses of pooled individual patients' data from three large phase III randomized trials. *Cancer Med.* 2017;6(7):1573–1580.

4. Yothers G, O'Connell MJ, Lopatin M, et al. Validation of the 12-gene colon cancer recurrence score in NSABP C-07 as a predictor of recurrence in patients with stage II and III colon cancer treated with fluorouracil and leucovorin (FU/LV) and FU/LV plus oxaliplatin. *J Clin Oncol.* 2013;31(36):4512-4519.

5. Kotani D. et al., Molecular residual disease and efficacy of adjuvant chemotherapy in patients with colorectal cancer, *Nature Medicine* v29 Issue 1 Jan 2023

 Coombes RC, Page K, Salari R, et al. Personalized Detection of Circulating Tumor DNA Antedates Breast Cancer Metastatic Recurrence. *Clin Cancer Res.* 2019;25(14):4255-4263.

 Abbosh C, Birkbak NJ, Wilson GA, et al. Phylogenetic ctDNA analysis depicts early-stage lung cancer evolution. *Nature*. 2017;545(7655):446-451.

 Christensen E, Birkenkamp-Demtroder K, Sethi H, et al. Early Detection of Metastatic Relapse and Monitoring of Therapeutic Efficacy by Ultra-Deep Sequencing of Plasma Cell-Free DNA in Patients With Urothelial Bladder Carcinoma. J Clin Oncol. 2019;37(18):1547-1557.
Data on file

Learn more about Signatera[™]: Tel: +1.650.489.9050 | Email: signateraquestions@natera.com | Visit: natera.com/oncology

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