

Circulating tumor DNA to detect minimal residual disease, response to adjuvant therapy and identify patients at high risk of recurrence in stage I-III CRC.

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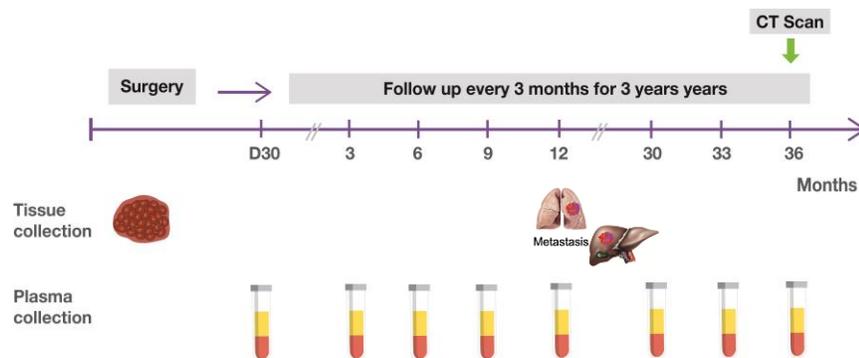
Aims and Study Design

1. To detect MRD and stratify patients based on risk of relapse using Signatera™ ctDNA (bespoke, mPCR NGS) assay.
2. To assess post-therapy relapse risk in ctDNA-positive patients.
3. To determine molecular lead time of recurrence between ctDNA and CT imaging.

Patient Characteristics

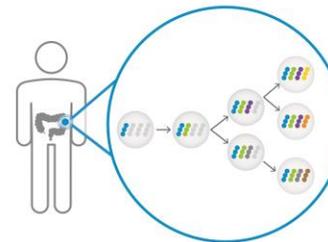
- Patients enrolled = 198
- Patients analyzed = 193
- Median follow-up: 21.9 (1.4-41.9) months
- Stage I: (n=5), Stage II: (n=85), Stage III: (n=103)
- Adjuvant treated: (n=105)

INCLIVA and Aarhus University - Clinical Protocol

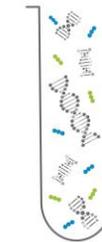


Signatera's Molecular Protocol

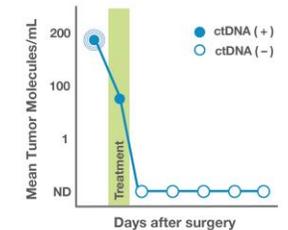
Sequence tumor tissue to identify unique signature of tumor mutations



Custom design and manufacture personalized mPCR assay for each patient, targeting the top 16 clonal mutations found in tumor

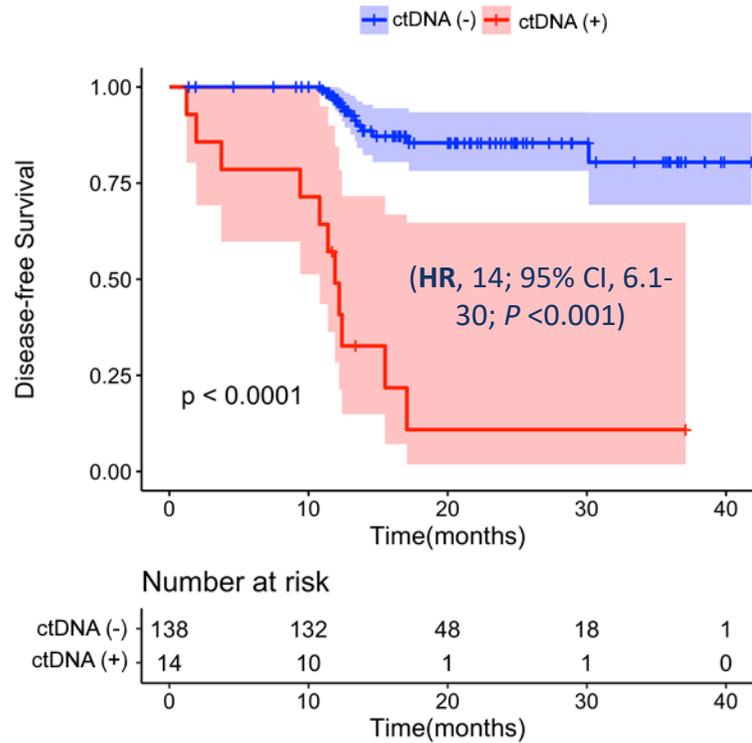


Use personalized assay to test patient's blood for presence of circulating tumor DNA (ctDNA)



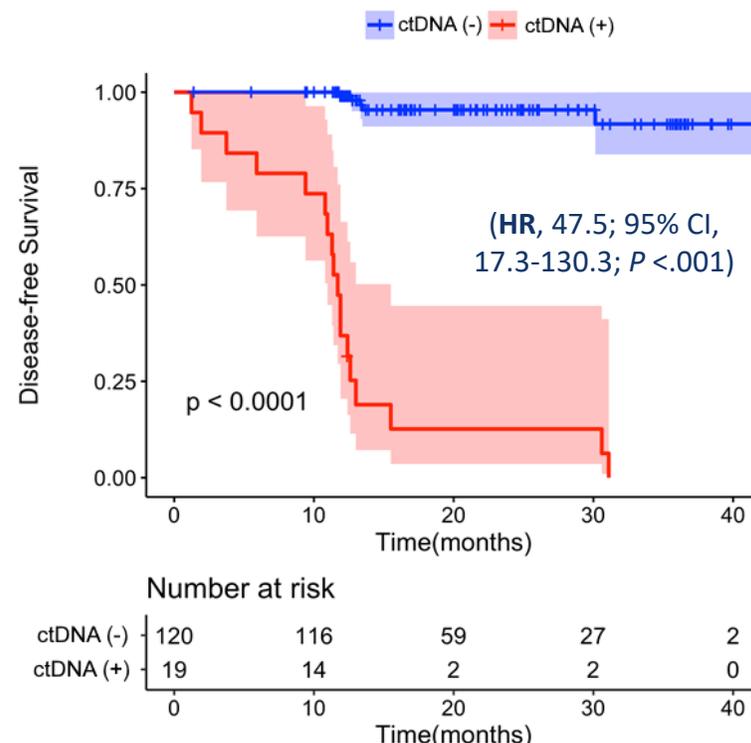
Results | Relapse-Risk Stratification by ctDNA Status

Post-operative setting (single timepoint)



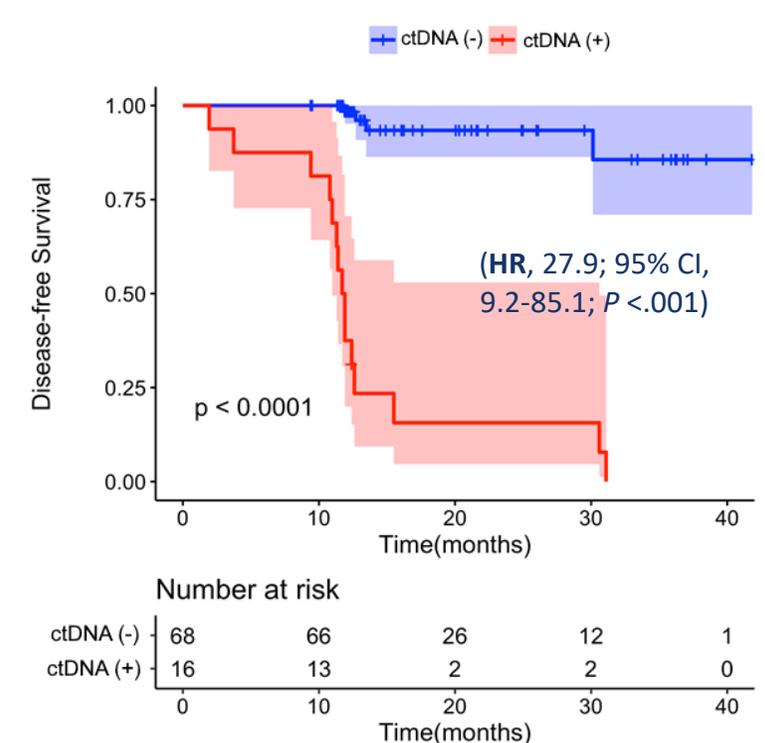
MRD-Positive: 9.2% (14/152)
Patients relapsed: 78.5% (11/14)

Post-definitive treatment setting



ctDNA positivity in longitudinal follow-up was associated with significantly worse DFS

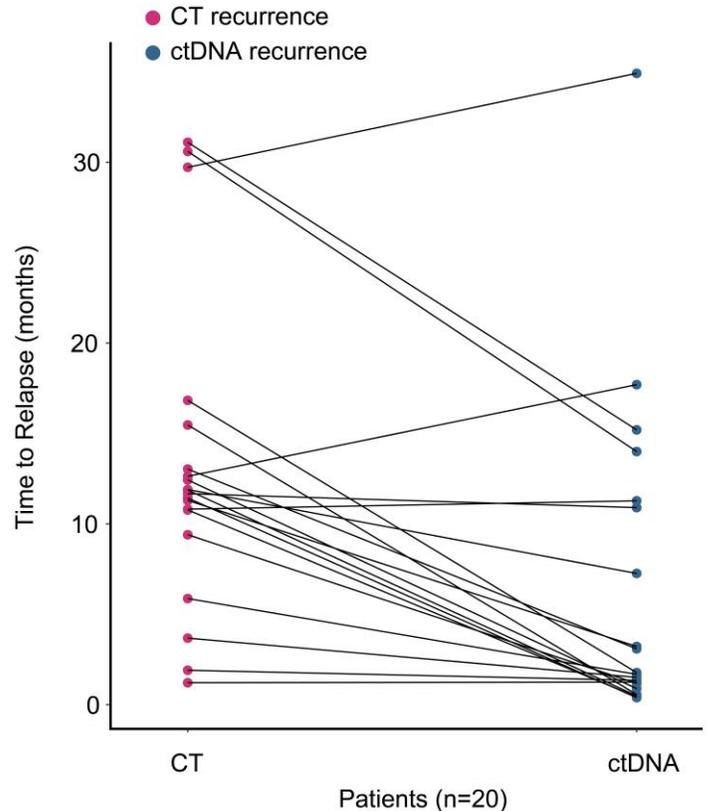
Post-ACT setting



Post-ACT longitudinal ctDNA positivity was strongly associated with significantly worse DFS

Results | Molecular Lead Time and Multivariate Analysis

Molecular Lead Time



Median lead time over radiological recurrence was 8.15 (IQR 0.56-16.6) months, $P < 0.001$.

Multivariable Cox Proportional-Hazards Model

Multivariable analysis included: ctDNA status, age, sex, stage, ACT, MSI, perineural invasion and number of resected lymph nodes

Akaike Information Criterion statistics suggests that the best model includes ctDNA status and age.

N = 137 Events = 23	HR	CI	Pr(> z)
Longitudinal ctDNA status (ref. negative)	53.19	(18.87-149.90)	<0.001
Age (ref. <70 years)	1.99	(0.81-4.86)	0.13

ctDNA was the only significant risk factor

Conclusions

1. Signatera ctDNA assay showed a sample level specificity of over 99.8% in a multi-institutional study.
2. Post-operative ctDNA status was the only significant predictor of DFS than all other risk factors combined.
3. ctDNA-positive status at the end of adjuvant treatment may potentially indicate resistance to standard therapy.
4. Longitudinal ctDNA tracking enables identification of patients at higher risk of relapse with a median lead time of 8.15 months over radiologic identification.
5. ctDNA analysis can help guide treatment decisions both in the adjuvant and post-adjuvant setting.

Disclosures

Prof. Andrés Cervantes

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Shruti Sharma, Svetlana Shchegrova, Derrick Renner, Himanshu Sethi, Bernhard Zimmermann, Alexey Aleshin are employees at Natera and own stock/options to own stock in the company .

All other authors declare no conflicts of interest.