



Prognostic value of post-surgery liquid biopsy cell-free circulating tumor DNA in stage III colon cancer patients - PLCRC-PROVENC3 study

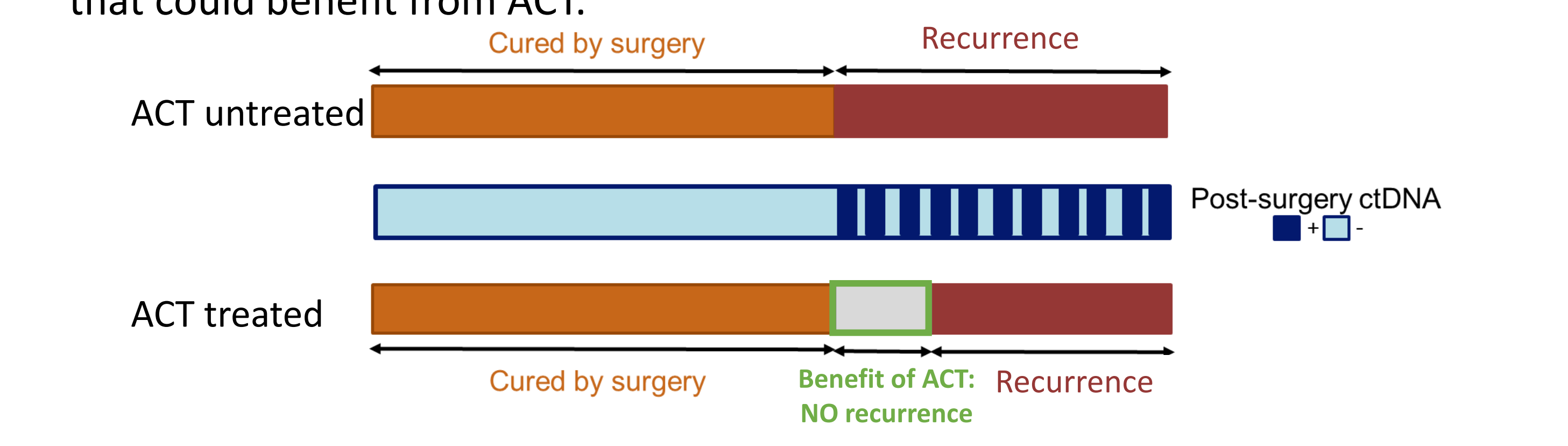
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Background

- Current clinical guidelines in the Netherlands recommend adjuvant chemotherapy treatment (ACT) following resection of the primary tumor for all stage III colon cancer patients.
- Only 15-20% of the patients benefit from ACT: around 55% of stage III colon cancer patients are cured by surgery alone and are being overtreated, and 30% will relapse despite ACT.
- Prognostic biomarkers may improve ACT decisions and reduce futile treatment in this group of patients by identifying the patients at a higher risk of recurrence that could benefit from ACT.



Post-surgery circulating tumor DNA (ctDNA) detection indicates presence of minimal residual disease, and it is a strong prognostic factor in stage II and III colorectal cancer.

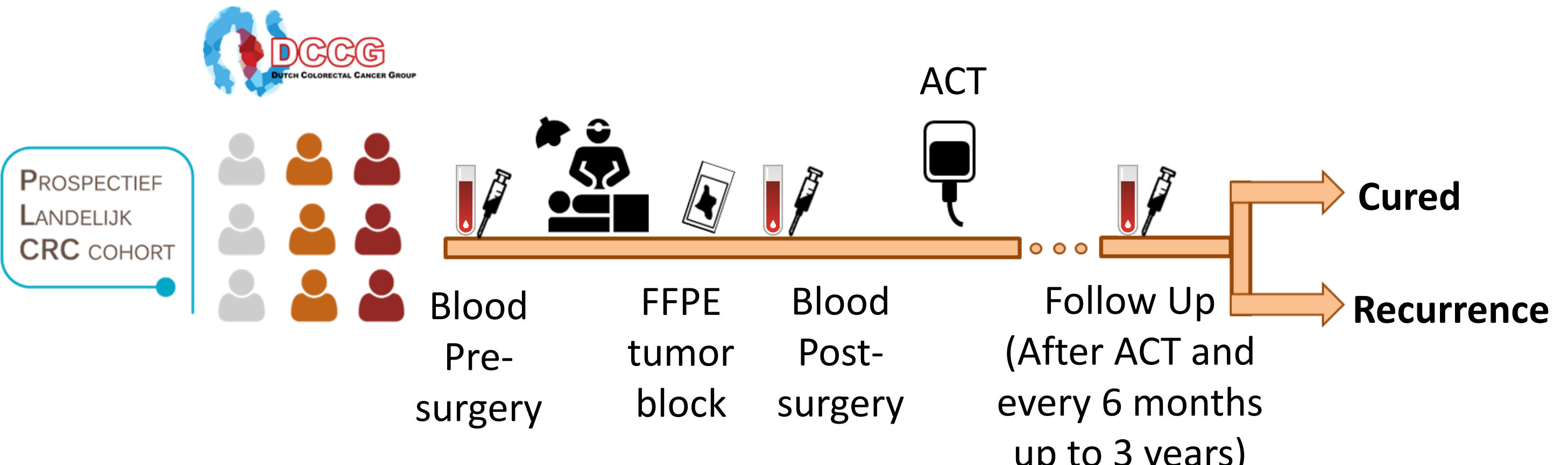
Goal of the study

Investigate the prognostic value of post-surgery ctDNA testing for disease recurrence in ACT treated stage III colon cancer patients

Experimental approach: PROVENC3 study

PROVENC3: (PRO)gnostic Value of Early Notification by Ctdna in Colon Cancer stage 3).

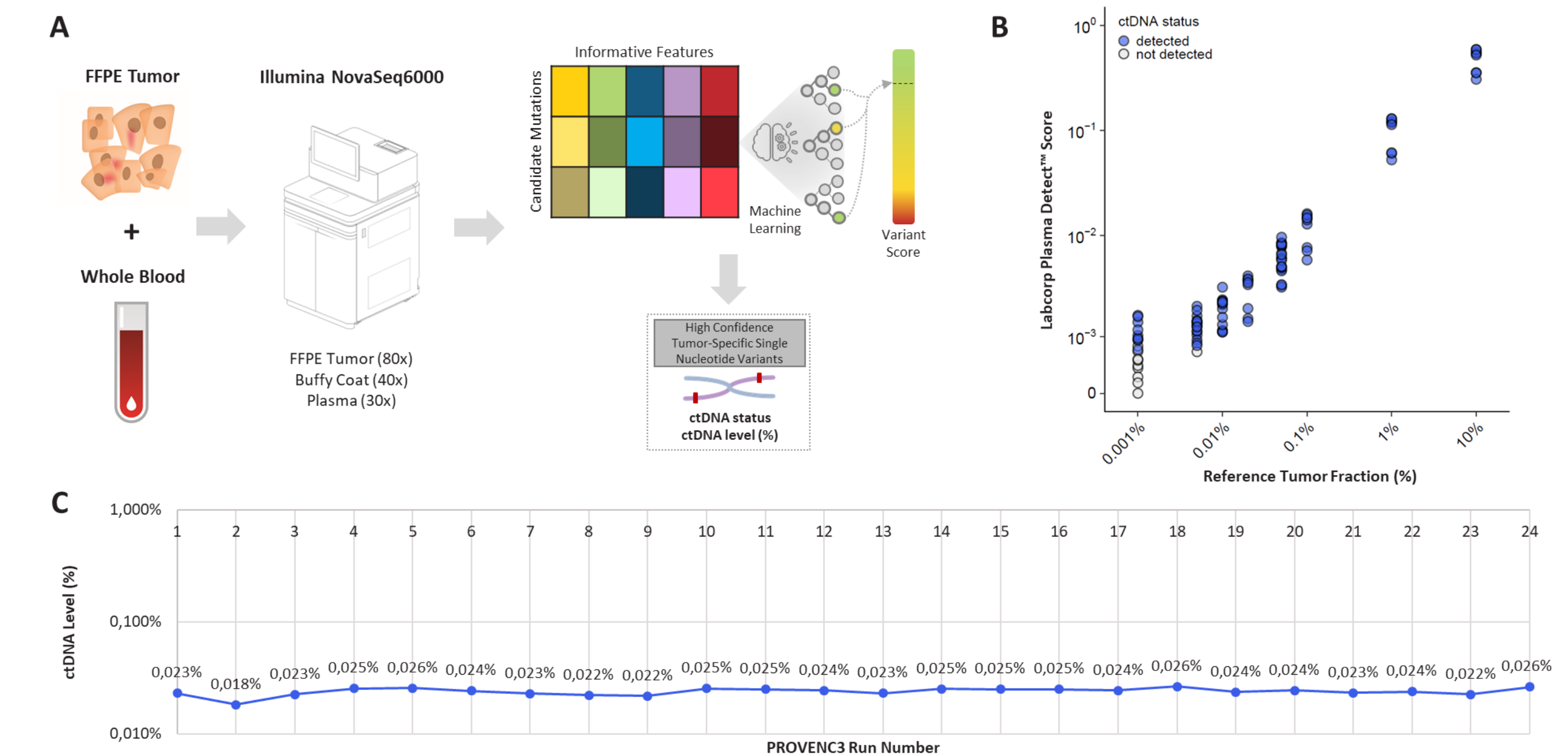
- PROVENC3 is an observational study within the Prospective Dutch Colorectal Cancer Cohort (PLCRC, <https://plcrc.nl/for-international-visitors>).
- 26 Participating hospitals in PROVENC3.
- 236 patients included (Dec2016-Oct2021): Stage III, ACT treated, post-surgery blood available.



- One post-surgery blood sample is collected from day 3-65 after surgery (Median: 10 days, IQR: 14 days), before ACT.
- All biosamples are sent to a central laboratory (The Netherlands Cancer Institute).
- Clinical data is collected in the Netherlands Cancer registry by IKNL.

Experimental approach: ctDNA detection method

Tumor-informed detection of plasma ctDNA through integrated whole genome sequencing (WGS) analyses:



- A) Schematic of the Labcorp Plasma Detect™ assay workflow (adapted from Keefe et al., 2022 Nature Communications and Wood et al., 2018 Science Translational Medicine)
- B) Analytical studies demonstrated a limit of detection (95%) of 0.005% tumor content utilizing contrived reference models derived from commercially available cell lines (including lung cancer, breast cancer, and melanoma), with a specificity of 99.6% (2,015/2,023) observed across 119 noncancerous donor plasma specimens evaluated against 17 reference somatic mutation datasets. The observed tumor fraction was also highly correlated with the reference tumor fraction (Pearson correlation coefficient = 0.96, p<0.001).
- C) Analysis of an external contrived reference control sample demonstrated reproducible results across 24 independent runs evaluated for the PROVENC3 clinical study (CV = 7.2%)

Clinicopathological characteristics and ctDNA MRD results

Results for 114 patients with post-surgery ctDNA analyzed and clinical follow up (FU) information about recurrence available. Median FU 31 months, IQR: 18 months.

	ctDNA positive 22 (18%)	ctDNA negative 92 (82%)	Total 114
Median age (years) (Range)	66 (43-83)	63 (34-79)	63
Gender			
Male	17 (77%)	45 (49%)	62 (54%)
Female	5 (23%)	47 (51%)	52 (46%)
Tumor location			
Right	11 (50%)	34 (37%)	45 (39%)
Left	11 (50%)	58 (63%)	69 (61%)
Differentiation grade			
Well differentiated	0	0	0
Moderately differentiated	17 (77%)	81 (88%)	98 (86%)
Poorly differentiated	3 (14%)	7 (8%)	10 (9%)
Undifferentiated	0	0	0
UNK	2 (9%)	4 (4%)	6 (5%)
T			
T1	0	2 (2%)	2 (2%)
T2	1 (5%)	3 (3%)	4 (4%)
T3	15 (68%)	67 (73%)	82 (72%)
T4	6 (27%)	20 (22%)	26 (23%)
N			
N1	13 (59%)	66 (72%)	79 (69%)
N2	9 (41%)	26 (28%)	35 (31%)
Clinical risk			
Low risk	12 (55%)	56 (61%)	68 (60%)
High risk	10 (45%)	36 (39%)	46 (40%)
MR status			
MSS	21 (95%)	79 (86%)	100 (88%)
MSI	1 (5%)	10 (11%)	11 (10%)
UNK	0	3 (3%)	3 (3%)
Resection			
Radical	20 (91%)	88 (96%)	108 (95%)
Non radical	2 (9%)	1 (1%)	3 (3%)
UNK	0	3 (3%)	3 (3%)
ACT			
3 months CAPOX	15 (68%)	72 (78%)	87 (76%)
6 months capecitabine	2 (9%)	3 (3%)	7 (6%)
6 months CAPOX	3 (14%)	12 (13%)	16 (14%)
3 months capecitabine	2 (9%)	2 (2%)	4 (4%)

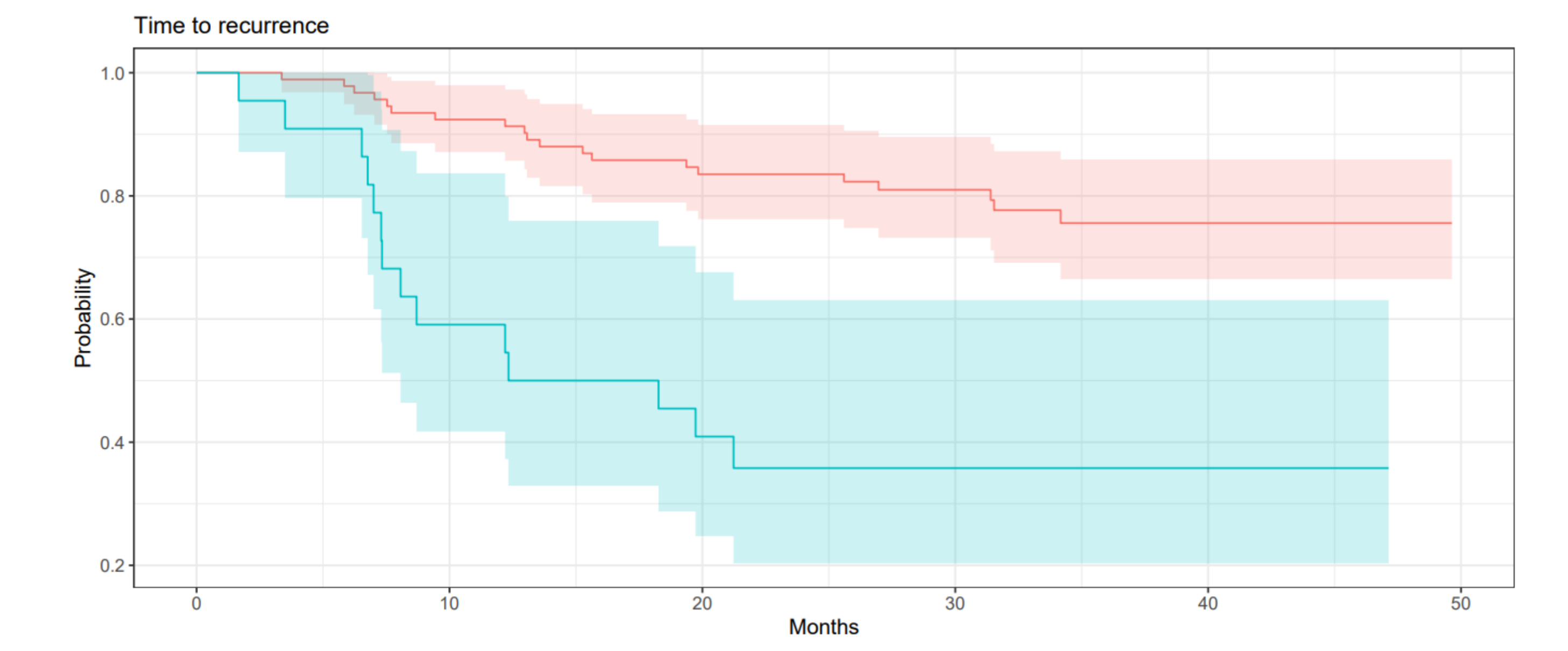
41% of the patients that experienced a recurrence had detectable minimal residual disease (MRD) post-surgery, and 30% of the ctDNA+ patients were likely cured by ACT:

	ctDNA + 22 (18%)	ctDNA - 92 (82%)	Total 114
Recurrence	14 (64%)	20 (22%)	34 (30%)
Non recurrence	8 (36%)	72 (78%)	80 (70%)

	Clinical risk per ctDNA and recurrence status:		Total
	Relapse	No relapse	
ctDNA+ patients	14	8	22
Total	6 (43%)	6 (75%)	12 (55%)
Low risk	8 (57%)	2 (25%)	10 (45%)
High risk			
ctDNA- patients	20	72	92
Total	6 (30%)	50 (70%)	56 (61%)
Low risk	14 (70%)	22 (30%)	36 (39%)
High risk			

Prognostic value of post-surgery ctDNA detection

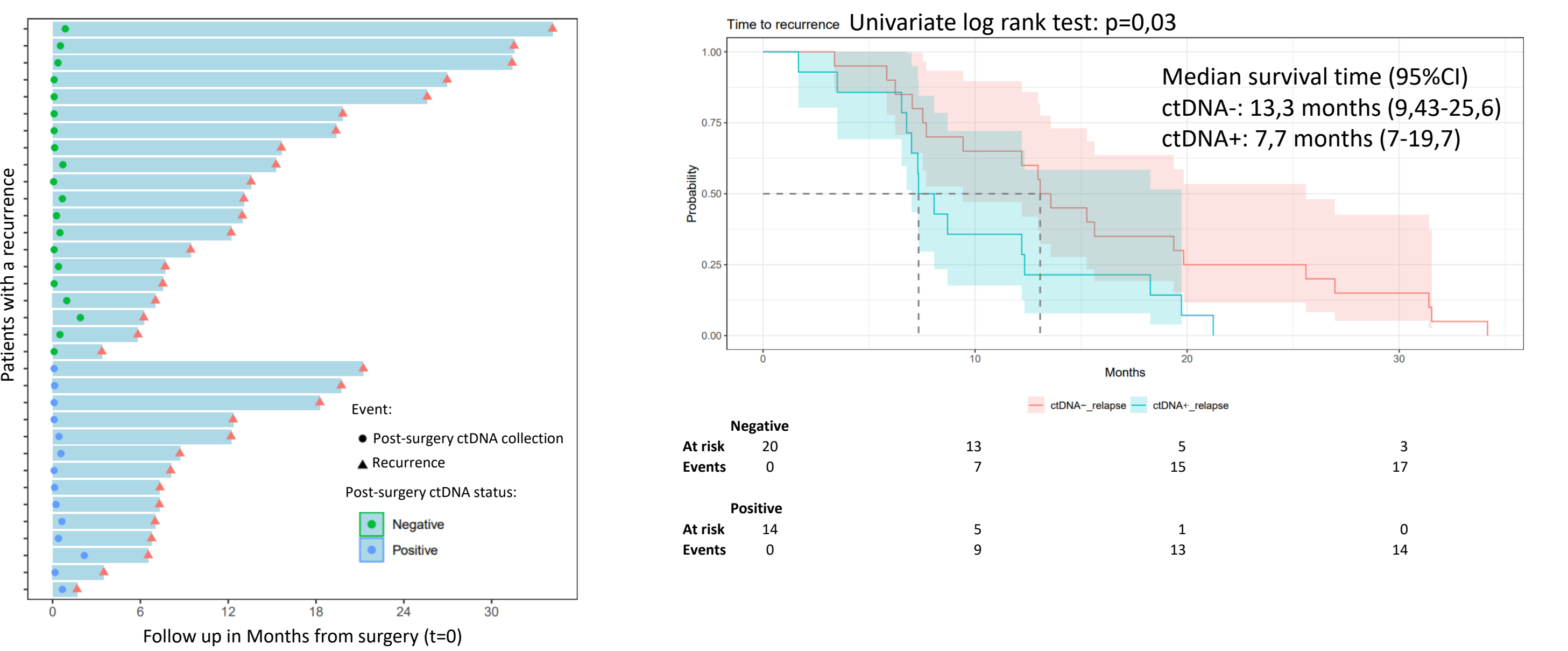
Time to recurrence (TTR) was evaluated for all 114 patients based on post-surgery ctDNA status. Patients with a positive ctDNA status post-surgery are at a higher risk of experiencing a recurrence (HR: 4.52; 95%CI: 2.27-9.01; p<0,001)



	Post-surgery ctDNA status:					
	Negative	Positive				
At risk	92	85	73	55	18	0
Events	0	7	15	17	20	20
At risk	22	13	9	6	2	0
Events	0	9	13	14	14	14

Time to relapse informed by post-surgery ctDNA results

Time to recurrence was evaluated for the 34 patients experiencing a recurrence based on post-surgery ctDNA status. CtDNA positive patients experiencing a recurrence (ctDNA+_relapse) show a shorter time to recurrence than ctDNA negative patients experiencing a recurrence (ctDNA-_relapse).



Conclusions

- Post-surgery ctDNA testing improves the stratification of stage III colon cancer patients for disease recurrence on top of current clinicopathological risk factors
- Approximately one third of the ctDNA-positive patients seem to benefit from ACT.

Next steps

- Complete sample analysis and clinical data collection.
- Based on the results of this study, design of an interventional study towards implementation of ctDNA testing for stage III colon cancer patients.