



the European Union

A precision medicine trial leveraging tissue and bloodbased tumor genomics to optimize treatment in resected stage III and high-risk stage II colon cancer patients: The SAGITTARIUS trial

Background

- Post-surgical circulating tumor DNA (ctDNA) testing can detect molecular residual disease (MRD) [1,2].
- Post-surgical and post-adjuvant chemotherapy (ACT) ctDNA detection is associated with worse recurrence-free survival (RFS) in stage I-III colorectal cancer (CRC) patients. [2-6]
- ctDNA is a prognostic biomarker with predictive potential of ACT efficacy and disease recurrence in CRC. [4-7]
- Prospective studies and trials assess ctDNA for guiding treatment decisions in stage II-III colon cancer patients [8-11]. Ongoing randomized clinical trials are adjusting ACT intensity based on MRD status [12,13]. Unfortunately, intensified ACT is curative in only a proportion of MRD-positive cases [9]. New personalized treatment strategies are needed to increase adjuvant efficacy in molecularly defined colon cancer patient subgroups.

KEY FEATURES

- 1. Diagnosing molecular residual disease after surgery with curative intent
- 2. Establishing the efficacy of a ctDNA-based tailored adjuvant strategy compared to **one-size-fits-all** conventional adjuvant chemotherapy in molecularly defined colon cancer patients
 - Curbing the risk of relapse and improving survival in ctDNA-positive patients

Endpoints

- Primary Secondary
- 2-year RFS in ctDNA-positive patients
- 2-year RFS in ctDNA-negative patients
- 3- and 5-years OS
- Safety and tolerability
- Seroconversion (ctDNA clerarance) rate
- Assesment of FACT-C and EQ-5D-5L
- New biomarkers in ctDNA+ vs ctDNA-



Post-surgery liquid biopsy detects Minimal Residual Disease (MRD) in colon cancer patients, distinguishing two tumors subtypes with different clinical features.

Study Design

SAGITTARIUS is a Phase III randomized clinical trial (RCT) aiming at proving the efficacy of using ctDNA detection to guide the adjuvant clinical management in resected stage III and high-risk stage II colon cancer patients, using a tissue-based personalized treatment approach. Patients are stratified based on post-surgery ctDNA status and molecular biomarkers identified in tumor tissue specimens and then randomized accordingly in two embedded RCTs: Trial-1 for ctDNA-positive patients, and Trial-2 for ctDNA-negative patients.

Reducing toxicity and improving quality of life in ctDNA-negative patients



MSI-H	H/MMRd				
) MSI status MSS/MMRp			MOLECULAR BIOMARKERS	TAILORED	
				TREATMENT	CHEMOTHERAPY
2) POLE status	OLE mut TMB-high	STRATA 1	MSS RAS/RAF mut	CAPOX	A) TEMIRI B) FOLFIRI
3) RAS/RAF status		STRATA 2	MSI-H	nivolumab+ipilimumab	CAPOX
	RAS/RAF mut		MSS POLE mut	nivolumab+ipilimumab	CAPOX
) HER2 status			MSS HER2 ampl	trastuzumab+pertuzumab	CAPOX
	HER2 ampl		MSS Multiple WT	FOLFOX+panitumumab	FOLFIRI
	HER2 wt				

Exploratory

Sample size

ssumptions	 Combined HR=0.63 in overall Trial-1 Expected HR=0.80 in Strata-1 Expected HR=0.50 in Strata-2 80% power; 2-sided alpha=0.05
ample size	 156 events are required > 200 ctDNA-positive patients randomized in Trial-1

Key Eligibility Criteria

- Main Inclusion Histologically confirmed diagnosis of stage III or high-risk stage II colon cancer criteria Availability of primary tumor FFPE block ECOG performance status 0-1 Normal organ functions Main Exclusion History of another neoplastic disease (unless in remission > 5 years) criteria Evidence of metastatic disease
 - Macroscopic or microscopic evidence of residual tumor (R1 or R2 resection)
 - Acute or subacute intestinal occlusion
 - Clinically relevant cardiovascular disease
 - Known history of HIV, Hepatitis B/C, TB

Methods

MRD status is determined using the tumor-informed, personalized ctDNA test (Signatera[™], Natera, Inc.).

Tumor tissue comprehensive genomic profiling (TruSight[™] Oncology Comprehensive EU, Illumina, Inc.) is used to determine molecular biomarkers as microsatellite status (MSI), tumor mutational burden (TMB), gene mutations (mut) and amplification (amp).

Tumor imaging assessment (CT-scan) is performed before randomization and subsequently: every 3 (Trial-1) or 6 (Trial-2) months for 2 years, then every 6 (Trial-1) or 12 (Trial-2) months for 3 years.

Quality of life questionnaires (FACT-C and EQ-5D-5L) and health costs data are longitudinally collected for Quality of Life and Cost Effectiveness analyses.

Biospecimens, including archival tumor tissue, serial blood samples, buccal swabs and stool specimens, are collected for exploratory analyses.

The SAGITTARIUS trial design schema (upper panel), and the molecular biomarker-based stratification factors with hierarchical biomarker priorities consort (lower panel, left) and tailored treatment and chemotherapy associated per each tumor genetic subtype (lowe panel, right).

Study Sites



The SAGITTARIUS clinical networks includes 26 Institutions within 3 different Eurpean Countries (Spain, Italy and Germany)

Pre-existing neuropathy > grade 1

Other Embedded Analysis



References



, Trial SAGITTARIUS

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