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**Molecular characterization of human glioblastoma:
longitudinal follow-up of plasmatic microRNAs levels by
droplet digital PCR**

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ABSTRACT

In uno studio longitudinale prospettico caso-controllo sono stati analizzati, mediante droplet digital PCR, microRNA plasmatici, in particolare miR 21-5p, miR 23b-3p e miR 34a-5p, in qualità di biomarcatori periferici di astrocitomi IDH-mutati e glioblastomi IDH-wildtype grado 4 (secondo la quinta edizione della classificazione dei tumori del sistema nervoso centrale dell'Organizzazione Mondiale della Sanità).

L'espressione plasmatica di miR 21-5p, miR 23b-3p e miR 34a-5p è stata documentata sia nei volontari sani sia nei pazienti al momento della diagnosi. Sono stati registrati livelli circolanti di miR 34a-5p significativamente più elevati nei pazienti rispetto ai volontari sani, identificando un cut-off diagnostico assoluto di 1.25 copie/ μ L. Inoltre, è stato possibile riconoscere due gruppi di pazienti, rispettivamente con alte e basse concentrazioni plasmatiche assolute di miRNA al momento della diagnosi.

miR 21-5p, miR 23b-3p e miR 34a-5p hanno mostrato lo stesso andamento longitudinale nel tempo con un progressivo aumento a 1 e 3 mesi dopo l'intervento chirurgico, momento in cui sono state registrate le concentrazioni massime, seguito da una graduale diminuzione a 6 e 12 mesi dall'intervento.

Correlazioni significative dei livelli plasmatici di ciascun miRNA sono state identificate tra il momento della diagnosi e 1 mese dopo l'intervento chirurgico e, più interessante, è stata riscontrata una correlazione reciproca tra le concentrazioni di miR 21-5p e miR 23b-3p a ciascun *timing point* e tra miR 21-5p e miR 34a-5p al momento della diagnosi. Una *network analysis* ha evidenziato alcune possibili vie molecolari di interazione tra questi miRNA nel contesto patologico dei gliomi, come i meccanismi di controllo del ciclo (E2F1, E2F3, RB1, CDK6) e della crescita cellulare (PTEN, PIK3R1, IGF1R). Infine, livelli elevati di miRNA circolanti sono risultati predittori affidabili di sopravvivenza globale e sopravvivenza libera da malattia. Nonostante la novità dei risultati descritti nel nostro lavoro, si rendono necessari ulteriori futuri studi che prendano in analisi più ampie coorti di pazienti per supportare i nostri dati.

1. INTRODUCTION

1.1 Liquid biopsy and glioblastoma

The identification of circulating biomarkers for 'liquid biopsy' is gradually surging in popularity in the oncology research fields due to its potential huge clinical applications for population screening, tumor monitoring and delivery of targeted therapies deriving from tumor molecular profiling. Liquid biopsies are "non-invasive" procedures based on sampling and analysing circulating tumour cells, tumour DNA, cell-free RNA, extracellular vesicles, tumour-educated platelets, proteins and metabolites derived from various biological fluids^{1,2}. MicroRNAs (miRNAs), a family of small non-coding RNAs, are deregulated in patients affected by cancer. Detection of tumour-specific peripheral biomarkers may be also useful in managing patients affected by primary brain tumours³⁻⁷.

Glioblastoma (GBM) is the most frequent primary malignant brain tumour in adults, accounting for approximately 15% of all intracranial neoplasms and 45-50% of all primary malignant brain tumours. The annual incidence of GBM is between 3.19 and 3.55 cases per 100000 people *per year* in Europe and USA. GBM preferentially affects older adults, with a peak of incidence between 55 and 85 years^{8,9}. These malignant brain tumours progress rapidly and clinical presentation largely depends on their location. They may manifest with focal neurological deficits, behavioural alterations, neurocognitive impairment, seizures and signs or symptoms of increased intracranial pressure, such as headaches, nausea and vomiting. The gold standard of treatment for GBM is surgical resection followed by concurrent conformal radiotherapy (RT) and chemotherapy¹⁰⁻¹³. Nevertheless, GBM has a dismal prognosis, with the highest mortality rate within 15-18 months after diagnosis, and a 5-year overall survival (OS) lower than 5%^{8,9}.

Despite growing efforts, no peripheral biomarker has been validated for GBM. Different circulating factors have been explored in the context of GBM, but variable and sometimes conflicting results have been reported^{5,6,14-16}. miRNA levels in serum and plasma are more stable, reproducible and consistent among individuals in comparison with other circulating nucleic acids^{17,18}. Furthermore, there is a lack of studies that take into consideration the major revisions of the fifth edition of the World Health Organization (WHO) Classification of Tumours of the Central Nervous System (CNS)^{13,19}. The revised classification of adult-type, diffuse astrocytic gliomas distinguishes **astrocytoma, IDH-mutant WHO grade 4** from **GBM, IDH-wildtype WHO grade 4**. Molecular parameters have been added as biomarkers of grading in addition to histological features, such as microvascular proliferation and/or necrosis. Astrocytomas IDH-mutant WHO grade 4 are diagnosed in case of *CDKN2A/B* homozygous deletion, and GBM IDH-wildtype WHO grade 4 in case of *TERT* promoter mutation, *EGFR* amplification or +7/-10 signature (combined gain of chromosome 7 and loss of chromosome 10)^{13,19,20}.

1.2 microRNAs and their role in cancer

In the context of circulating biomarkers, our interest focused on miRNAs, small non-coding RNAs, typically 20-23 nucleotides in length²¹⁻²⁵. They are synthesized by RNA pol II as *primary miRNA* (*pri-miRNAs*), then processed into *precursor miRNAs* (*pre-miRNAs*) and *miRNA duplex* by consecutive ribonuclease cleavages – Drosha-Pasha complex, Dicer in the cell nucleus and cytoplasm respectively (**Figure 1**).

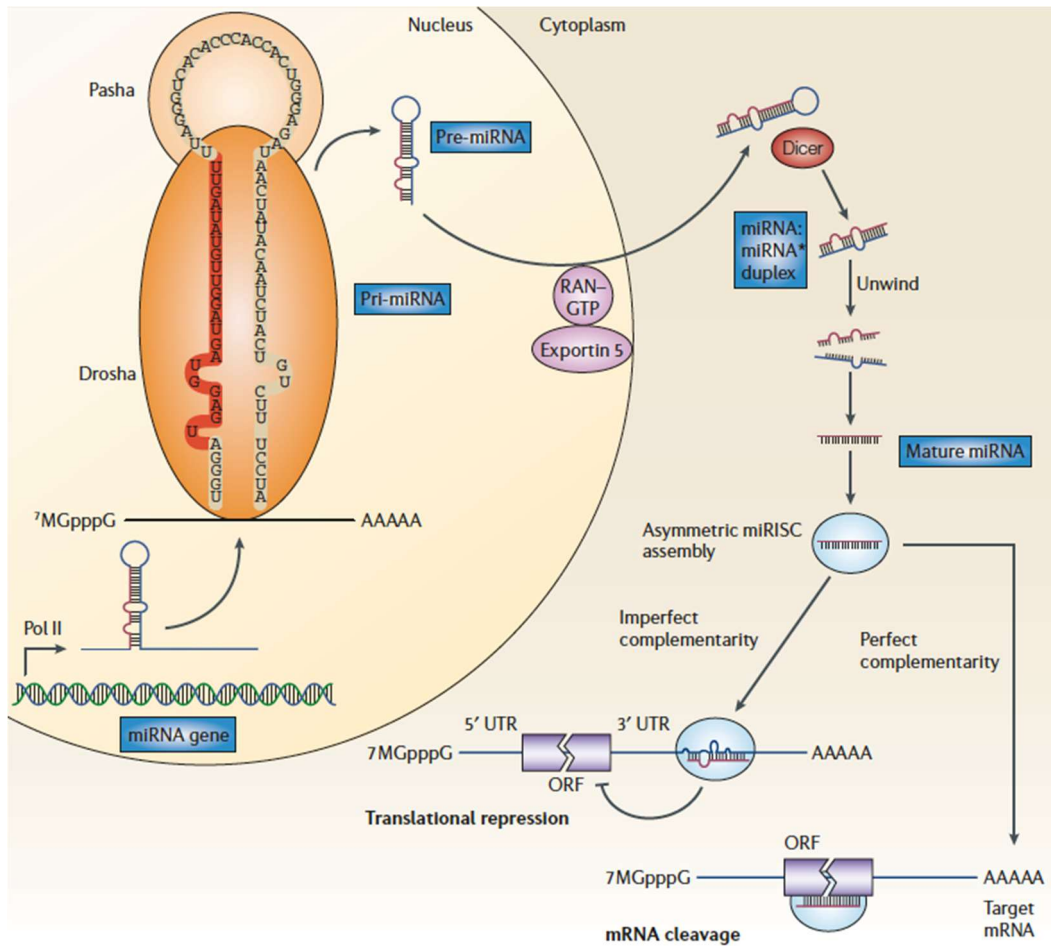


Figure 1 – The biogenesis of miRNAs; from Esquela-Kersher 2006²¹.

miRNAs act mainly as post-transcriptional regulators of gene expression and particularly as ‘silencers’ of messenger RNAs (mRNA)²¹⁻²⁸. The base pairing of miRNAs to target mRNAs depends primarily on 7-8 nucleotides at their 5’ end, known as ‘seed region’. Consequently, each miRNA may target many mRNAs and each mRNA can be targeted by multiple miRNAs altering the cellular phenotype^{25,26,29}. Depending on the degree of complementarity of their pairing, miRNAs exert their regulatory effect in two ways: i) mRNA degradation, and ii) translation inhibition. miRNAs that bind with perfect or nearly perfect complementarity to mRNA sequences induce the cleavage and degradation of mRNA transcripts by ribonucleases in the miRNA-associated induced-silencing complex (miRISC). miRNAs that pair with imperfect complementary sites within the 3’ untranslated

regions of their mRNA targets repress the target-gene expression post-transcriptionally, by blocking translation initiation and/or elongation^{21,29} (**Figure 1**).

miRNAs are expressed both in physiological and pathological conditions. They exhibit a causal role in tumorigenesis of a wide variety of human cancers, acting either as *onco-miRNAs* or *tumour-suppressor miRNAs*. miRNAs are detectable in tumour tissues and biological fluids, such as peripheral blood, because released by tumour and/or microenvironment cells^{17,21,23-25,27,28,30}. Three major potential pathways have been hypothesized for miRNAs to enter the circulation: i) energy-free passive release from damaged cells consequent to apoptosis or necrosis into circulation; ii) active and selective secretion mediated by the RNA-binding protein dependent pathway; iii) active and selective secretion facilitated by extra-cellular vesicles^{18,31} (**Figure 2**).

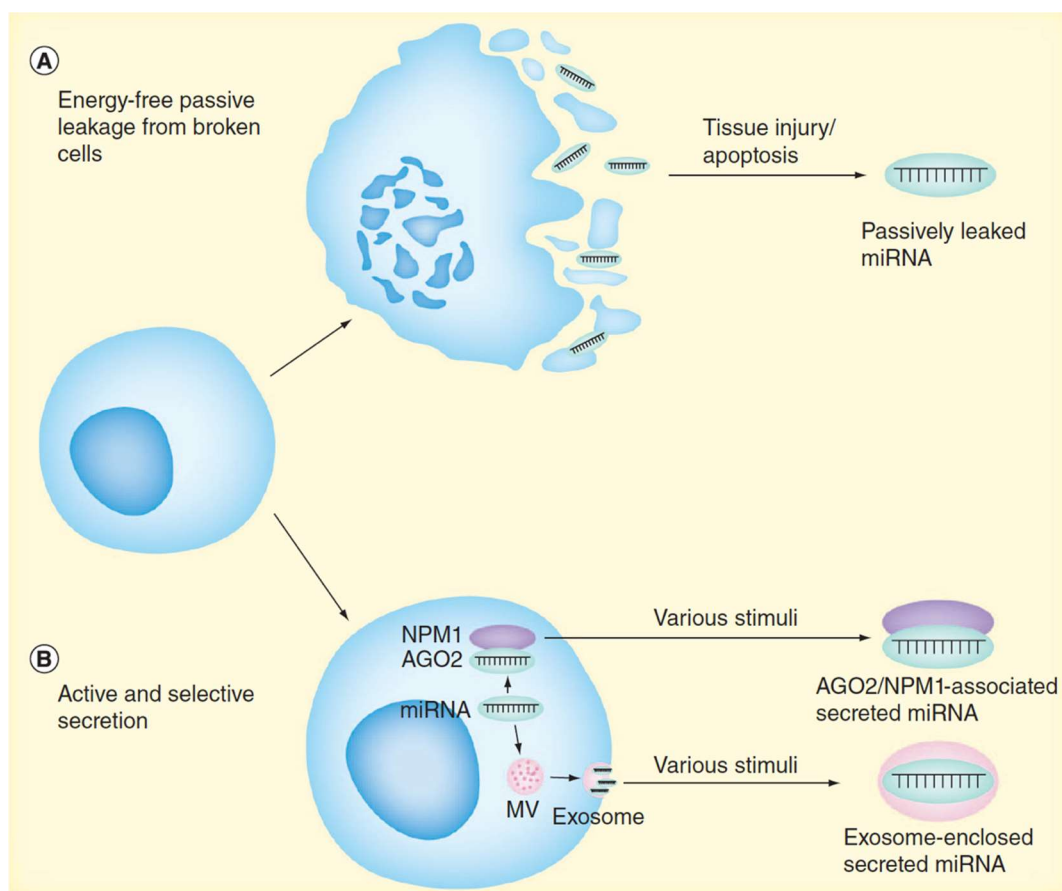


Figure 2 – Sources of circulating miRNAs; from Redova 2013¹⁸.

1.3 Detection of circulating miRNAs by droplet digital polymerase chain reaction (ddPCR)

Measurements of circulating transcripts' expression levels, especially miRNAs, may be challenging because of their low concentrations.

miRNAs can be detected and quantified by reverse transcriptase polymerase chain reaction (PCR) achieved by real-time quantitative PCR (qPCR) or digital PCR.

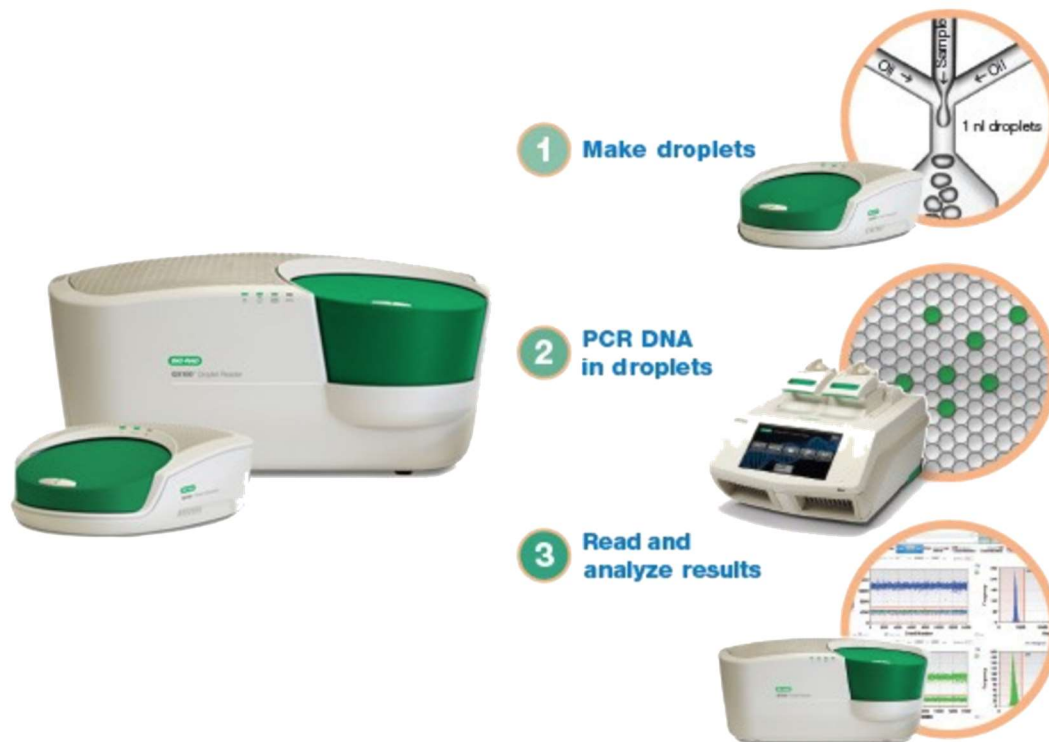


Figure 3 – ddPCR workflow (Bio-Rad Laboratories, Inc.; Hercules, CA, USA).

Digital PCR builds on traditional PCR amplification steps (denaturation, annealing and extension cycles) and intercalating dye and fluorescent probe-based detection methods to provide a highly sensitive absolute quantification of nucleic acids without the need for endogenous controls and standard curves. Particularly, the droplet digital PCR (ddPCR) is a method (**Figure 3**) based on water-oil emulsion droplet technology³². The sample is partitioned into 20,000 nanoliter-sized droplets. Template molecules are distributed randomly among droplets, such that some droplets have no template molecules and others have one or more (at most few). Each droplet undergoes PCR amplification and fluorescence analysis separately. The droplets are then individually counted and scored as *positive* or *negative* by passing in a single stream through a fluorescence detector. Due to the random nature of the partitioning, the fluorescence data after amplification – the ratio between the number of positive events (positive droplets) and the total number of independent events (the total number of droplets) – are well fitted by a Poisson distribution – a function that relates the probability of a given number of events occurring independently in a given sample when the average rate of occurrence is known – to determine the absolute copy number of the target DNA molecule (copies/ μ L) in the input reaction mixture.

2. AIMS OF THE STUDY

This study is aimed to develop a new line of research in neuro-oncology focusing on circulating biomarkers in GBM.

The discovery of miRNAs, both cellular and extracellular miRNAs has revealed their great potential as novel and robust diagnostic and prognostic cancer biomarkers. miRNAs can effectively control tumorigenesis by affecting different signal transduction pathways^{15,33}. Data obtained from tissue biopsies (mainly by RNA-sequencing technology) have been catalogued in the OncomiR Cancer database (OMCD) www.oncomir.umn.edu/omcd, where only 5 cases of GBM have been reported³⁴. Liquid biopsies are not included in the OMCD and their results in patients affected by GBM are extremely variable, in part due to challenges of carrying out accurate measurements of transcripts' expression levels^{5,6,15,16}.

The primary objective of our study was to correlate the circulating levels of non-coding RNAs in patients affected by GBM with clinical, neuroradiological and pathological tumour features in order to identify the possible role of peripheral GBM biomarkers as early and reliable predictors of tumour response to treatments (surgery, radiotherapy and chemotherapy), tumour recurrence and patients' survival. We focused on circulating miRNAs^{4,17,18,30} selected according to literature review, analysing their plasmatic levels both in patients affected by GBM and healthy volunteers:

- **miR 21-5p**, an *onco-miRNA* highly expressed in cell cultures, tumour tissue and serum or plasma derived from patients affected by GBM^{14,15,35-44};
- **miR 23b-3p**, overexpressed in GBM tissue and cell cultures⁴⁵⁻⁴⁸;
- **miR 34a-5p**, a *tumour suppressor miRNA* investigated in cell cultures, tumour tissue and serum or plasma of patients with GBM^{15,37,39,44,49-53}.

3. MATERIALS AND METHODS

3.1 Study population

Twenty-three patients presenting with diffuse high-grade astrocytic gliomas (**Figure 4**) operated at the Neurosurgical Department of the ASST Cremona between April and December 2021 were prospectively included in the study.

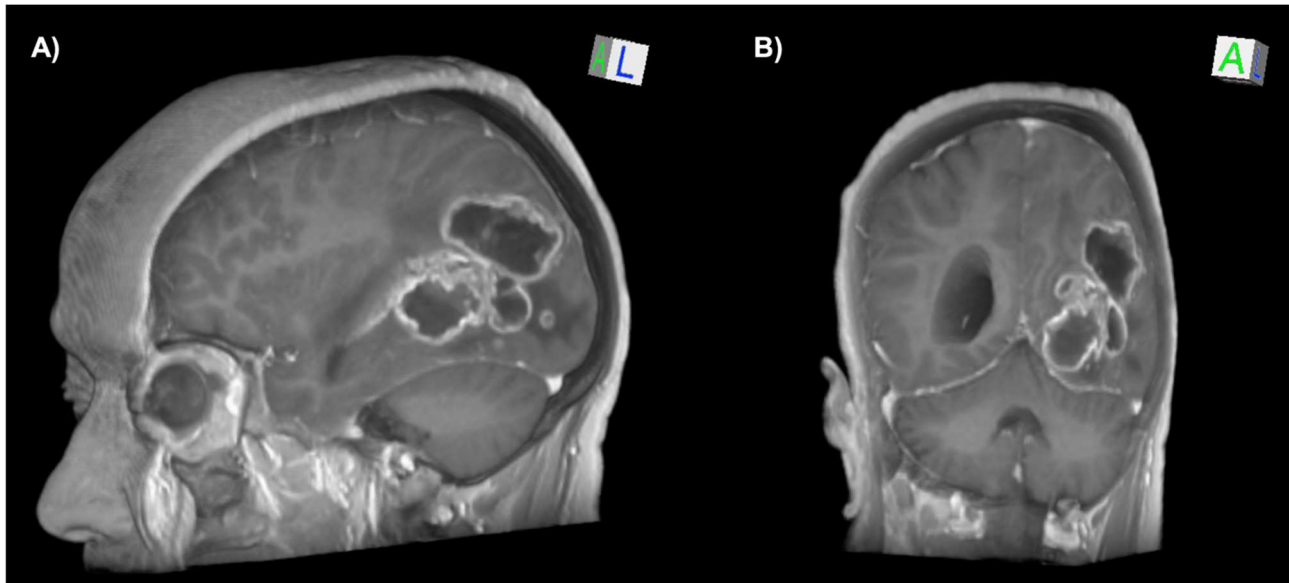


Figure 4 – Diffuse high-grade astrocytic glioma appearance on T1-weighted post-contrast sequences: **A)** 3D sagittal reconstruction; **B)** 3D coronal reconstruction.

The patients' inclusion criteria were: age older than 18 years, prognosis greater than six months, Karnofsky performance status (KPS) ≥ 70 and pathological report consistent with **GBM IDH-wildtype WHO grade 4** or **astrocytoma IDH-mutant WHO grade 4**^{13,19}. Furthermore, 32 healthy volunteers were enrolled in the study.

All the procedures were conducted in accordance with the ethical standards of the institutional research committee and with the 1964 Helsinki declaration and its later amendments. Informed consent was obtained from all patients and healthy volunteers. The local ethics committee approved the study (protocol number 32219, dated 2 October 2019).

3.2 Preoperative clinical, neuroradiological and surgical features

Clinical, neuroradiological and surgical characteristics were collected and recorded in an anonymized database.

The clinical features included: age at diagnosis, gender, KPS, modified Rankin scale (mRS) and medical therapies. According to KPS, used to quantify general well-being and independence in daily life activities, patients were classified as “independent” (KPS \geq 90) or “dependent” (KPS \leq 80). Similarly, they were categorized as “presenting” (mRS > 2) or “non-presenting” (mRS \leq 2) significant disabilities in keeping with mRS, a specific score for neurological pathological conditions⁵⁴. Therapies with steroids (dexamethasone) and anti-epileptic drugs (i.e. levetiracetam, lacosamide) at the time of diagnosis were registered.

Tumour location, volume, contrast enhancement and necrosis were assessed by two independent investigators on preoperative magnetic resonance imaging (MRI) images.

3.3 Treatments

3.3.1 Surgery

All patients underwent microsurgical fluorescence-guided tumour resection by craniotomy or frameless neuronavigated robotic-assisted biopsy, in keeping with the patient’s and tumour’s characteristics. They were administered an oral solution of 5-aminolevulinic acid (5-ALA, 20 mg/Kg of body weight) between 2 and 4 hours prior to surgery. Tumour fluorescence intensity was assessed under microscopic blue light (wavelength: 375-440 nm) and defined as “absent”, “faint” or “strong”⁵⁵ (**Figure 5**).

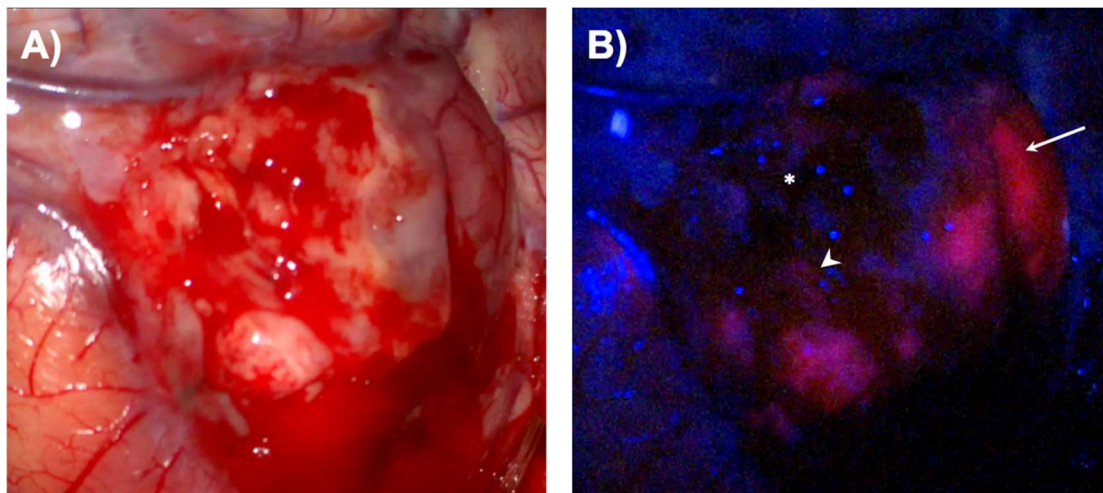


Figure 5 – Fluorescence-guided (5-ALA) microscopic tumour resection (Zeiss; Oberkochen, Germany): **A**) intraoperative microscopic tumour appearance under white light (10x); **B**) intraoperative microscopic tumour appearance under blue light (wavelength: 375-440 nm): “absent” (white asterisk), “faint” (white arrowhead) and “strong” (white arrow) tumour fluorescence intensity (10x).

3.3.2 Adjuvant therapies: radiotherapy and chemotherapy

One month after surgery, patients with newly diagnosed GBM IDH-wildtype WHO grade 4 or astrocytoma IDH-mutant WHO grade 4 underwent conformal RT and concomitant chemotherapy with temozolomide, an alkylating agent that adds methyl groups to the purine and pyrimidine groups particularly inducing the formation of O⁶-methylguanine (O⁶MeG), which leads to DNA double-strand breaks, chromosomal aberrations and ultimately cell death⁵⁶⁻⁵⁸.

RT was delivered following a standard or hypo-fractionated protocol. The standard RT protocol was based on 2 Gy per daily fraction, 5 days per week, over 6 weeks, for a total dose of 60 Gy. The hypo-fractionated RT protocol was reserved to patients older than 70 or with postoperative KPS lower than 70 and was planned to reach a total dose of 40 Gy in 15 fractions over 3 weeks¹². RT was delivered to the gross tumor volume with a 2-to-3-cm margin for the clinical target volume. Concomitant temozolomide was administered at a dose of 75 mg per square meter of body-surface area per day, 7 days per week from the first until the last day of RT, however for no longer than 49 days^{10,11}. After a 4-week break, patients clinically and neurologically fit for chemotherapy were administered up to 12 cycles of adjuvant chemotherapy with temozolomide following a 5-day schedule every 28 days at a dose of 150 mg per square meter of body-surface area for the first cycle, increased to 200 mg per square meter of body-surface area per day, until unacceptable toxicity, tumour progression, recurrence or death^{10,11}.

Patients experiencing tumour recurrence during adjuvant temozolomide and those with recurrent GBM were operated and then started on regorafenib, an oral inhibitor of several kinases involved in tumour angiogenesis (VEGFR1-3, TIE2), oncogenesis (KIT, RET, RAF1, BRAF), interactions between tumour and microenvironment (PDGFR, FGFR), and tumour immunity (CSF1R). Regorafenib was administered at a dose of 160 mg per day, 7 days per week for the first 3 weeks every 4-week cycle, 4 weeks after surgery in case of operation. Dose reduction, to 120 mg or 80 mg per day, or treatment discontinuation were adopted in case of toxicity⁵⁹.

3.4 Pathology

The histological diagnosis followed the fifth edition of the WHO Classification of Tumours of the CNS¹⁹. Tumour tissue was fixed in buffered 10% formalin for 24-48 hours, dehydrated in graded alcohols and paraffin embedded following the standard protocols. Serial 3- μ m thick sections were obtained from paraffin tumour blocks and mounted on glass slides, then underwent deparaffinization (by rinsing into a bath of xylene twice for 5 minutes each) and hydration (by rinsing into ethanol baths at reducing concentrations 100%, 90% and 70% and distilled water three times for 5 minutes each) before staining.

3.4.1 Histology

Serial sections were stained with hematoxylin and eosin (H&E). GBM typically appear as highly cellular tumours, usually composed of poorly differentiated, sometimes pleomorphic, tumour cells with nuclear atypia and brisk mitotic activity. Prominent microvascular proliferation and/or necrosis are essential diagnostic features (**Figure 6**).

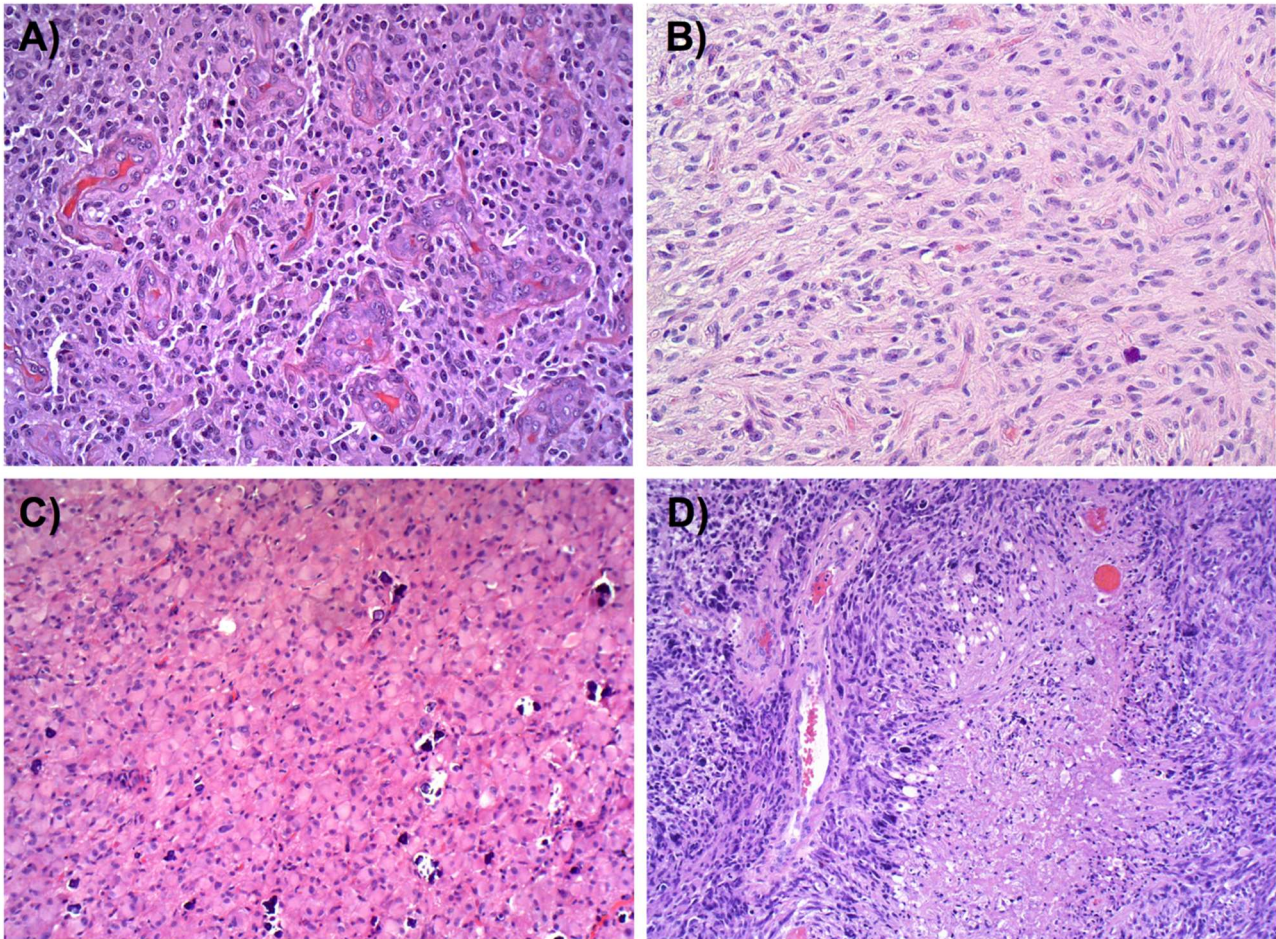


Figure 6 – GBM's morphological features: **A**) microvascular proliferation (white arrows; H&E – 20x); **B**) tumour cells with spindle features (H&E – 20x); **C**) tumour cells with epithelioid features and microcalcifications (H&E – 10x); **D**) necrosis (H&E – 10x).

3.4.2 Immunohistochemistry

Immunohistochemical (IHC) analyses were performed using the BOND Polymer Refine Detection system on the LEICA BOND III staining platform (Leica Biosystem; Wetzlar, Germany) for primary antibodies directed against the following antigens: synaptophysin (clone 27G12; 1:200), neurofilament (clone 2F11; 1:500), NeuN (A60; 1:500), GFAP (clone GA.5; 1:400), SOX10 (clone BC34; 1:200), EMA (clone E29; 1:500), CD34 (clone QBEnd/10; 1:1000), ATRX (polyclonal; 1:400), IDH1 R132H (clone H09; 1:50) and p53 (clone DO-7; 1:100). Otherwise, Roche's antibodies ready

to use were detected on the platform BenchMark ULTRA using the UltraView Universal DAB Detection Kit (Roche; Basel, Switzerland) for: Ki-67 (clone 30-9; prediluted) with the Ultraview Amplification Kit; BRAF (clone VE1; prediluted) and OLIG 2 (clone EP112; prediluted) with the Optiview Amplification Kit.

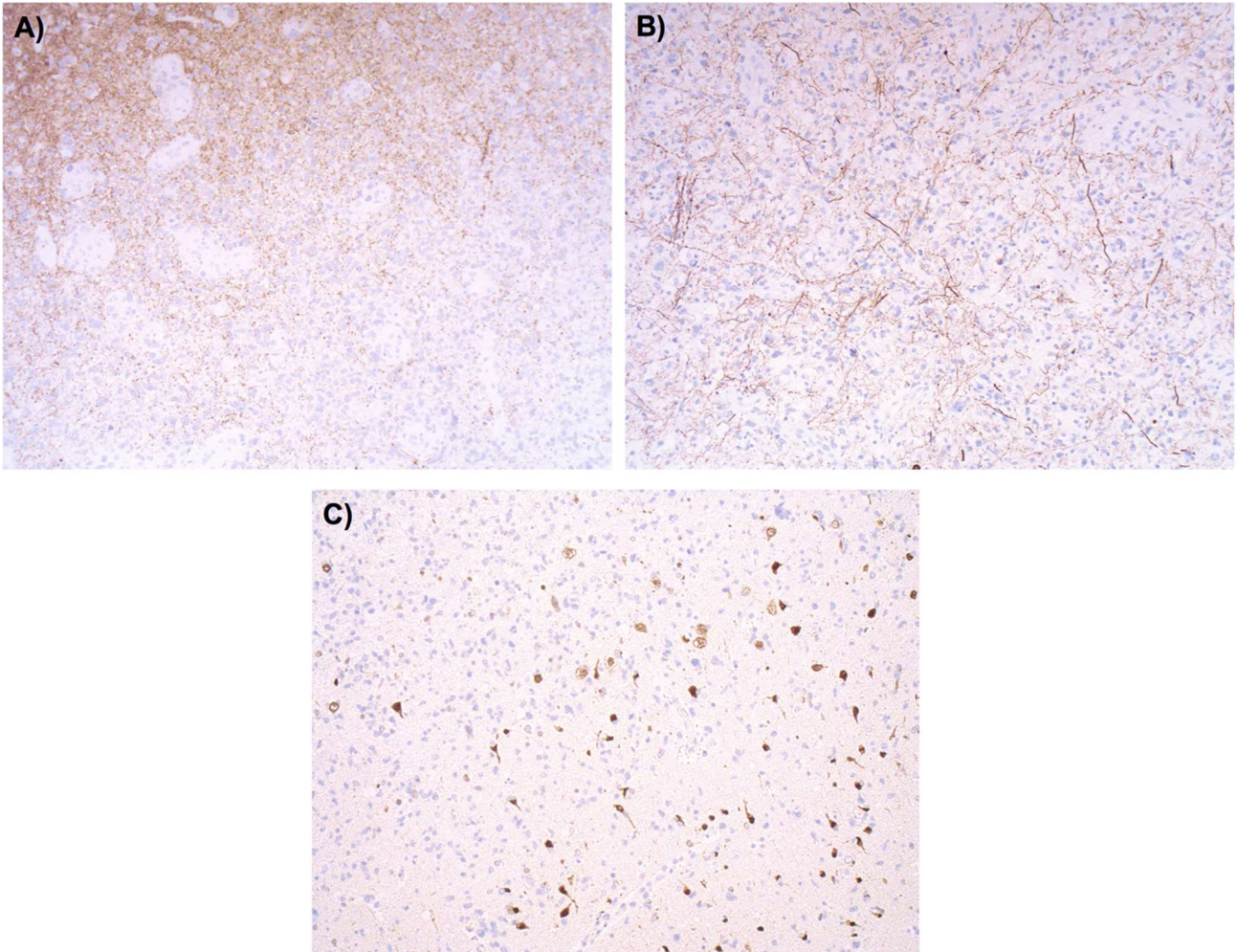


Figure 7 –GBM infiltration (synaptophysin, neurofilament, NeuN negative) of brain tissue (synaptophysin, neurofilament, NeuN positive): **A)** IHC for synaptophysin (10x); **B)** IHC for neurofilament (10x); **C)** IHC for NeuN (10x).

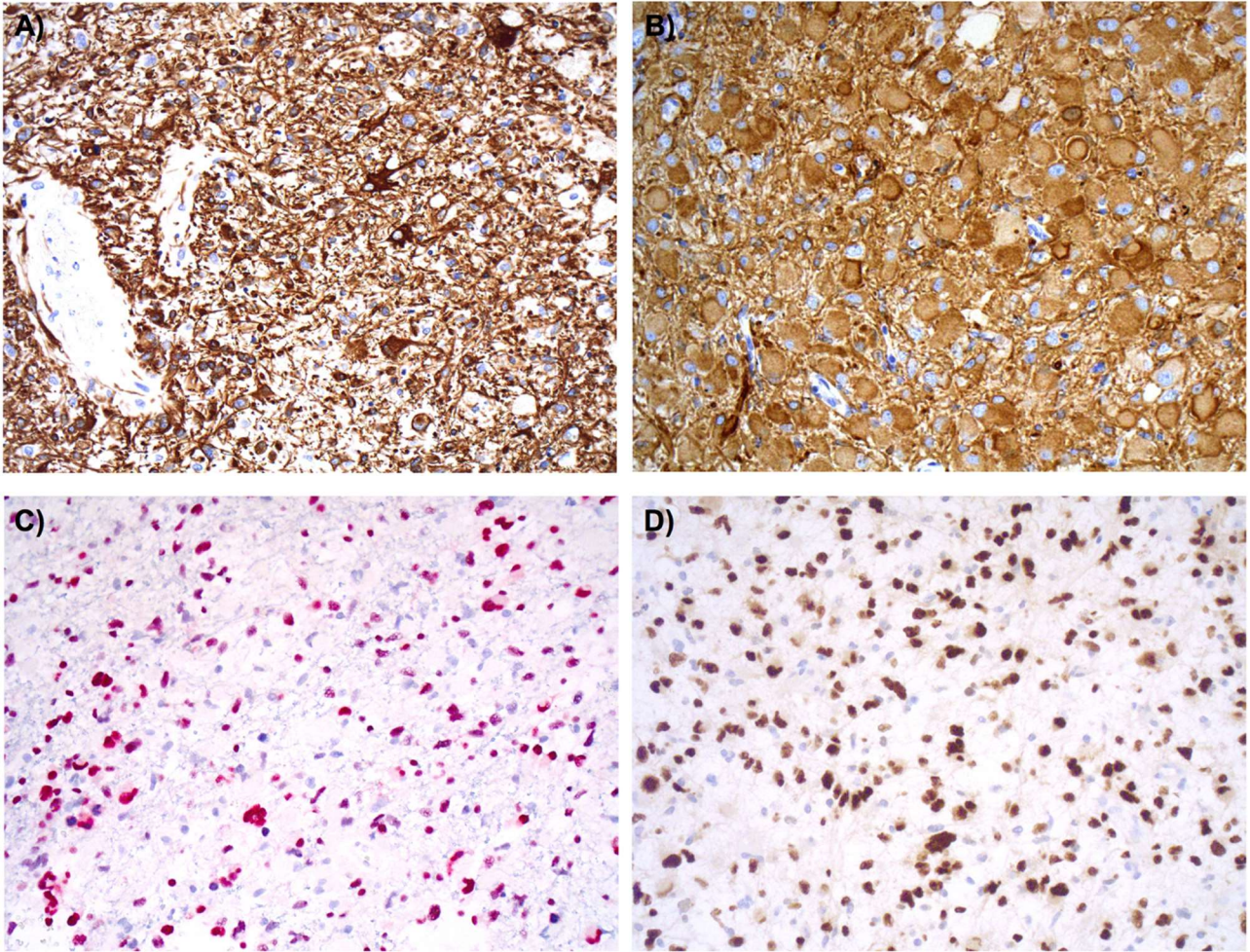


Figure 8 – A, B GFAP expression in GBM: (IHC – 20x); **C** SOX10 positivity in GBM (IHC – 20x); **D** Olig2 positivity in GBM (IHC – 20x).

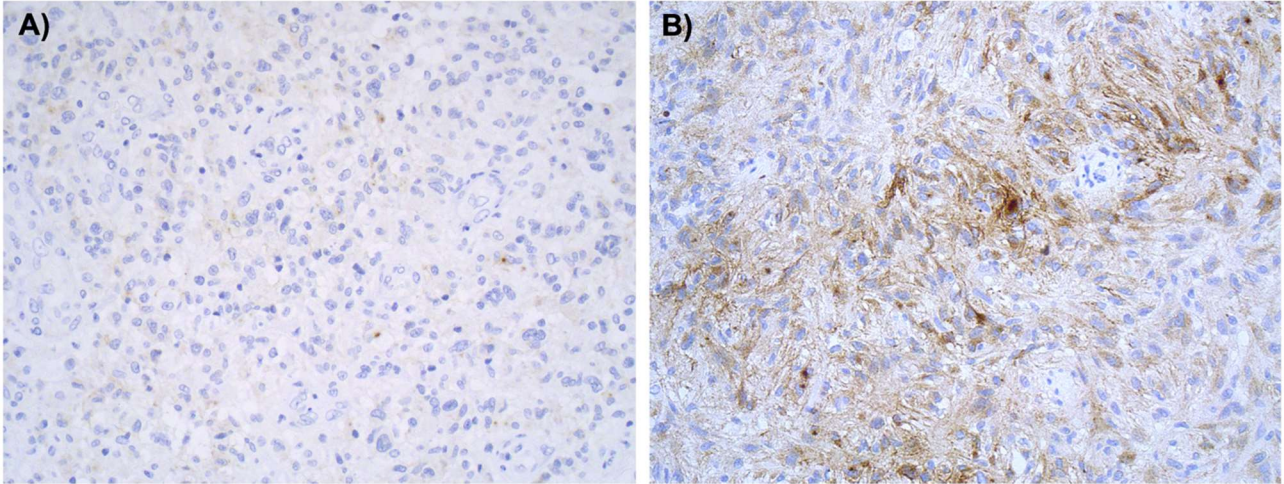


Figure 9 – EMA expression in GBM: **A**) negative (IHC – 20x); **B**) positive (IHC – 20x).

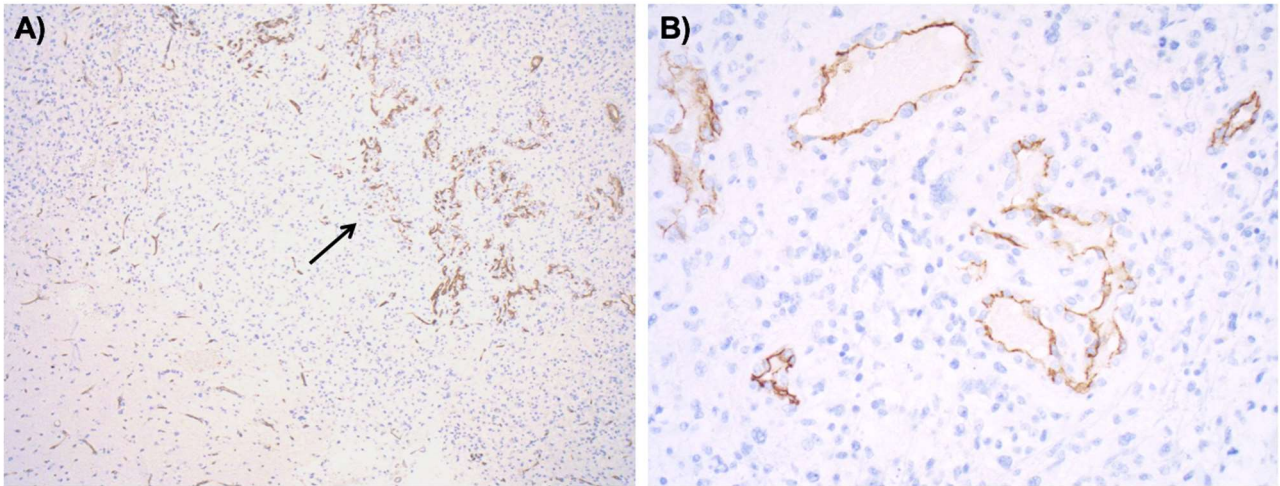


Figure 10 – GBM microvascular proliferation: **A)** vascular patterns in normal brain tissue and tumour (black arrow) (CD34; IHC – 4x); **B)** microvascular tumour proliferation (CD34; IHC – 20x).

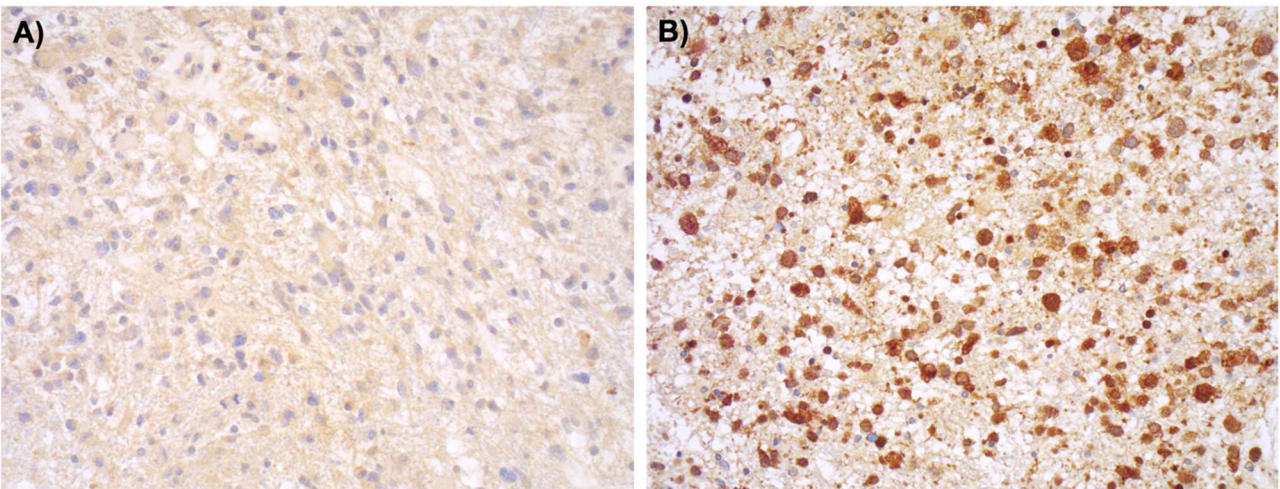


Figure 11 – IDH1 R132H status in GBM: **A)** IDH1 wild-type (negative; IHC – 20x); **B)** IDH1 mutant (positive; IHC – 20x).

The stainings for ATRX and p53 underwent semi-quantification by two expert neuropathologists. ATRX expression was classified as “lost” or “retained” (**Figure 12**), while p53 as “absent”, “intermediate”, or “strong” (**Figure 13**). ATRX is a chromatin remodelling protein and its loss of function causes microsatellite instability and telomere maintenance impairment but also increased tumour sensitivity to alkylating agents and RT⁶⁰.

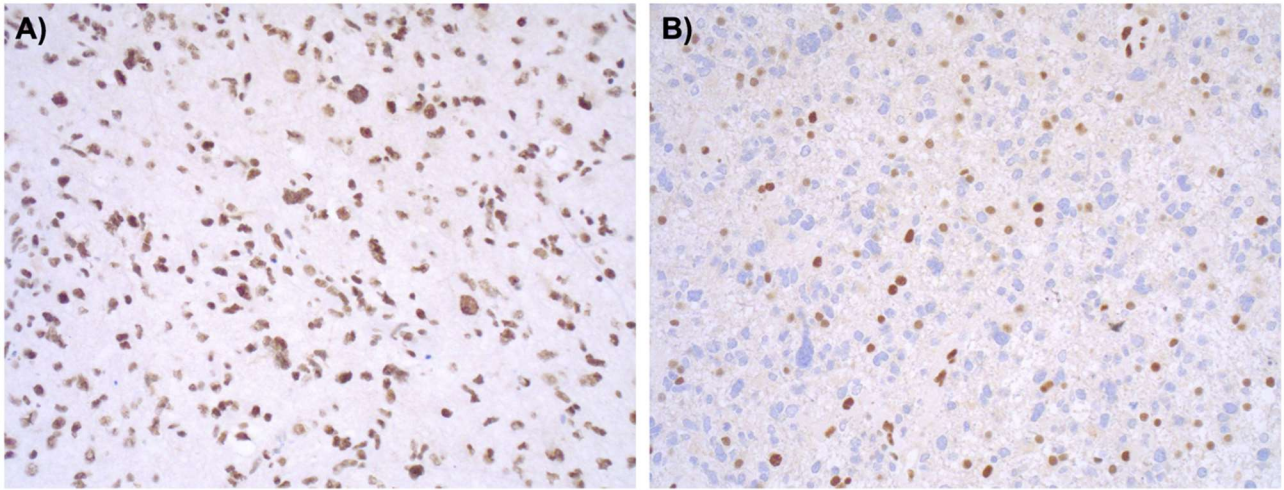


Figure 12 – ATRX expression in GBM: **A)** retained nuclear expression (IHC – 20x); **B)** lost nuclear expression (IHC – 20x).

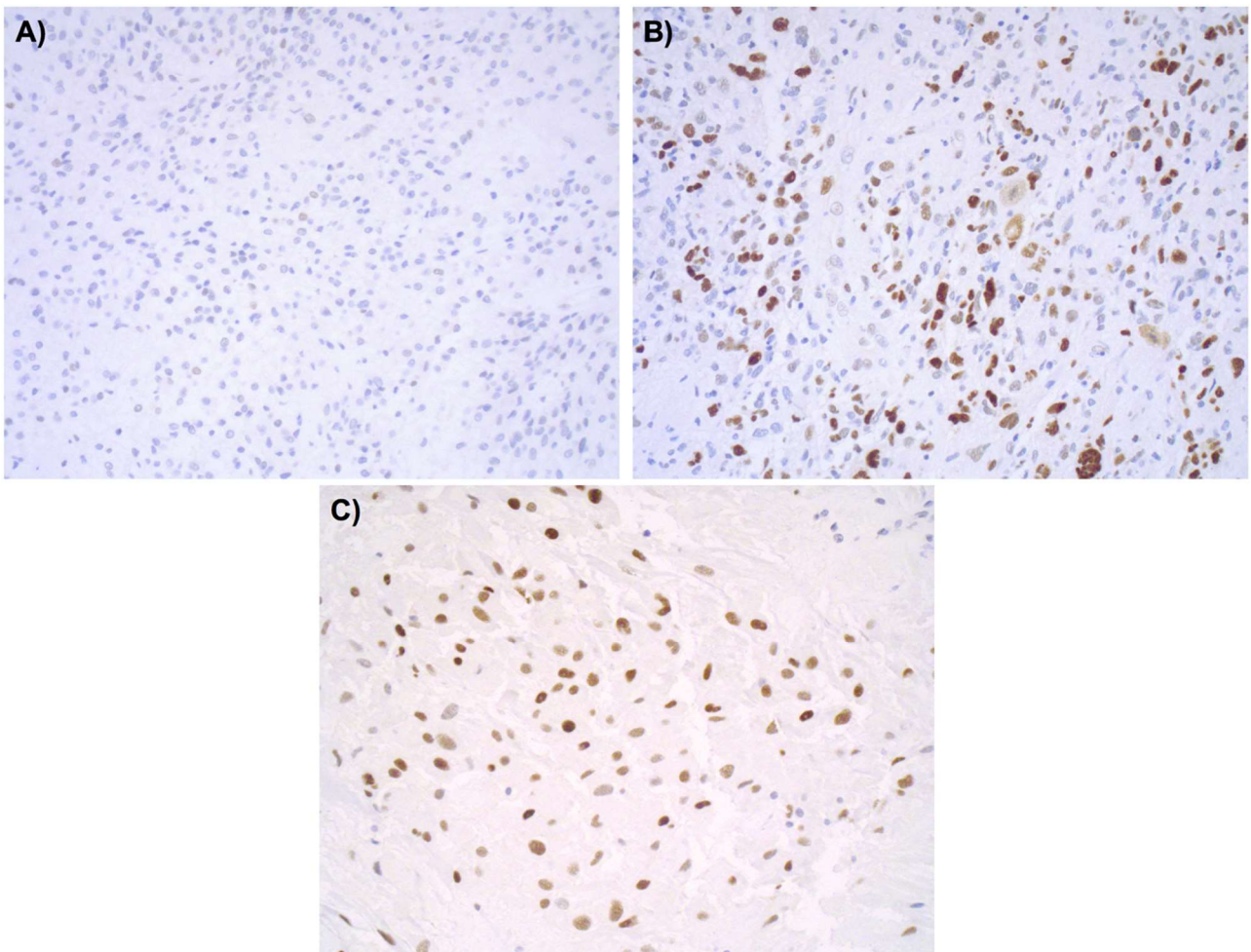


Figure 13 – p53 status in GBM: **A)** negativity (IHC – 10x); **B)** “intermediate” positivity (IHC – 20x); **C)** “strong” positivity (IHC – 20x).

The Ki-67 proliferative index was evaluated by counting the number of positive cells in at least 1000 tumour cells focusing on hot-spot tumour areas (**Figure 14**).

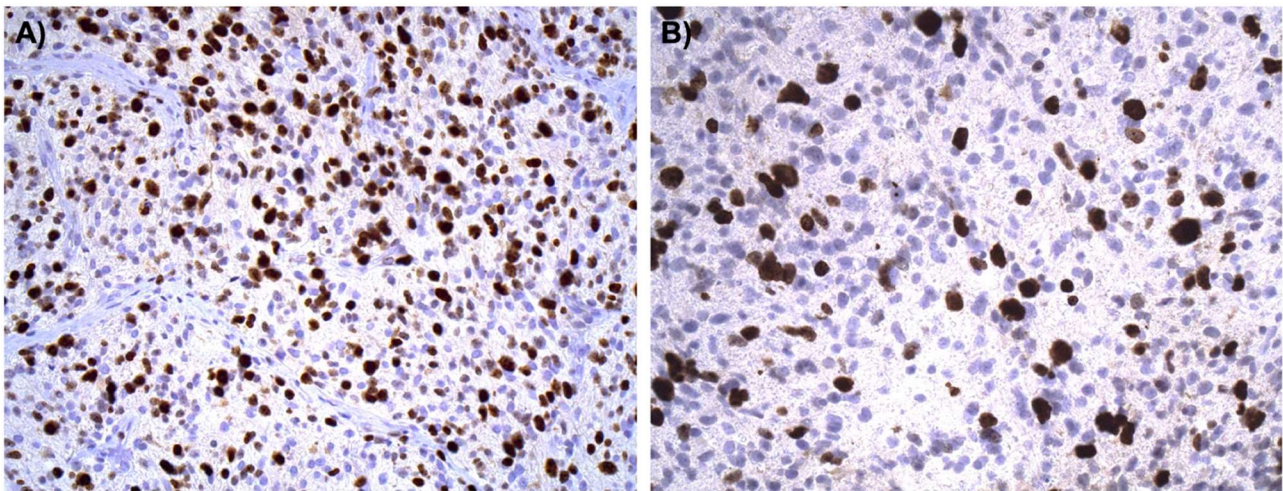


Figure 14 – Ki67 labelling: **A)** IHC – 10x; **B)** IHC – 20x.

3.4.3 Determination of IDH mutations and MGMT promoter methylation status

The most representative tumour area was circled on a reference H&E-stained section. Five serial 5- μ m thick sections were cut from formalin-fixed and paraffin-embedded tissue for DNA extraction. Genomic DNA was isolated from the selected tumour area using the Promega Maxwell DNA FFPE Preparation Kit (Promega; Madison, WI, USA).

A Real Time PCR was performed on the platform RT800-96 (EasyPGX® qPCR instrument 96, Diatech Pharmacogenetics SRL, Jesi, Italy) to detect mutations on *IDH1* (codon 105: G105G; codon 132: R132H, R132C, and R132S/G/L/I/V) and *IDH2* (codon 140: R140G/W/Q/L; codon 172: R172K and R172G/W/T/M/S). *Isocitrate dehydrogenase 1 (IDH1)* on chromosome 2q33 and *IDH2* on chromosome 15q26.1 encode for enzymes that function as homodimers and catalyse a reversible oxidative decarboxylation reaction of isocitrate to alpha-ketoglutarate to generate NADPH from NADP⁺, providing defence from oxidative damage and supporting cell fitness. Glioma-associated *IDH* mutations lead to a gain of function: synthesis of (D)-2-hydroxyglutarate causes epigenetic shift, DNA repair deficiencies, RNA demethylases, histone modification, and metabolism changes that all favour the gliomagenesis process^{61,62}.

The *O*⁶-methylguanine methyltransferase (*MGMT*) promoter methylation status was assessed by pyrosequencing (Pyromark Q961.D, kit MGMT plus, Diatech Pharmacogenetics SRL; Jesi, Italy). According to the percentage of MGMT promoter methylation status, tumours were grouped in unmethylated (< 9%), and methylated (methylated 9-29%; hyper-methylated > 29%)⁶³. *MGMT* sited on chromosome 10q26.3 encodes for an enzyme that transfers methyl groups at the *O*⁶ site of guanine (*O*⁶MeG) to a cysteine residue in its catalytic pocket, thus finally avoiding cell death promoted by alkylating agents, such as temozolomide. The *MGMT* expression is mainly regulated by epigenetic modifications, especially the methylation of CpG island in the promoter region, that reduces the transcription of *MGMT* increasing tumour chemosensitivity^{56,58,64}.

3.5 Extent of tumour resection analysis

Early postoperative MRI was routinely performed within 48 hours of surgery to assess the extent of tumour resection – near total (NTR) or subtotal (STR) if the tumour removal was greater or lesser than 95%, respectively⁶⁵ – and postoperative contrast enhancement of surgical cavity classified as “absent”, “linear” or “nodular” (Figures 15-17).

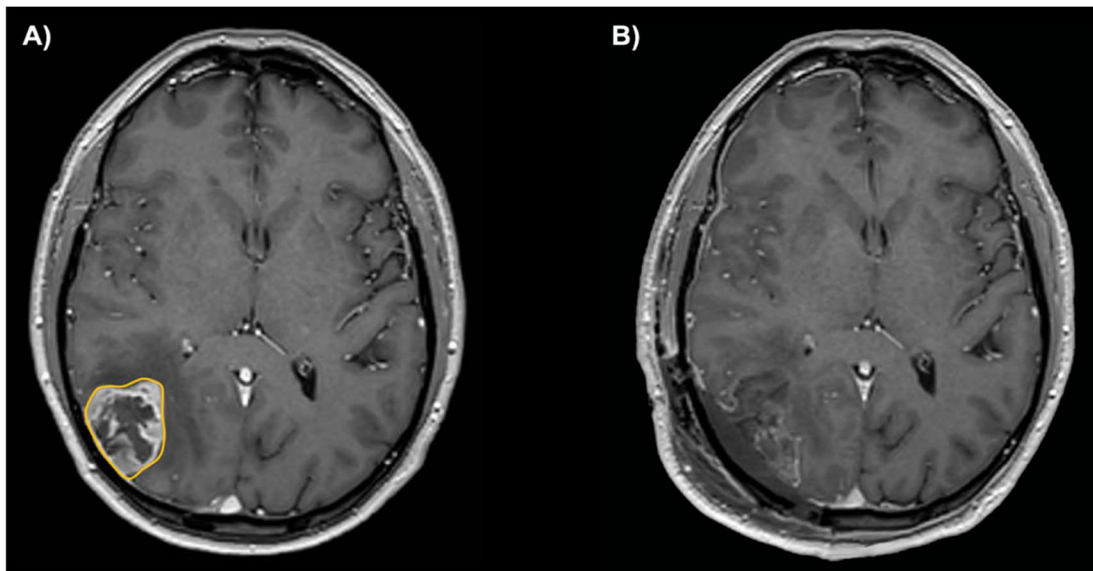


Figure 15 – Preoperative (A) and postoperative (B) T1-weighted post-contrast axial images showing NTR with *absence of pathological enhancement* of surgical cavity.

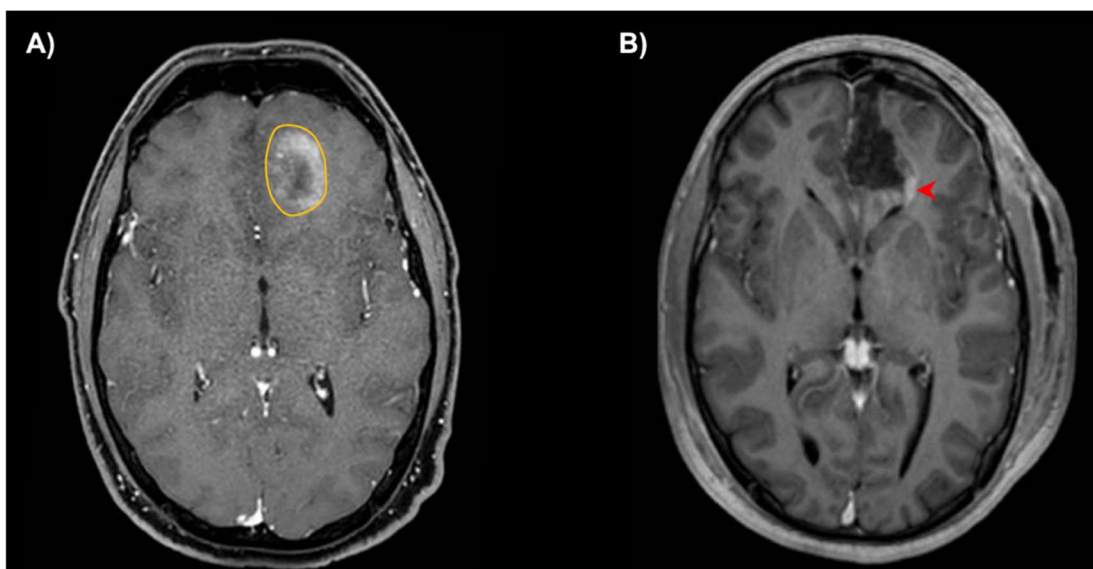


Figure 16 – Preoperative (A) and postoperative (B) T1-weighted post-contrast axial images showing NTR with *linear enhancement* (red arrowhead) of surgical cavity.

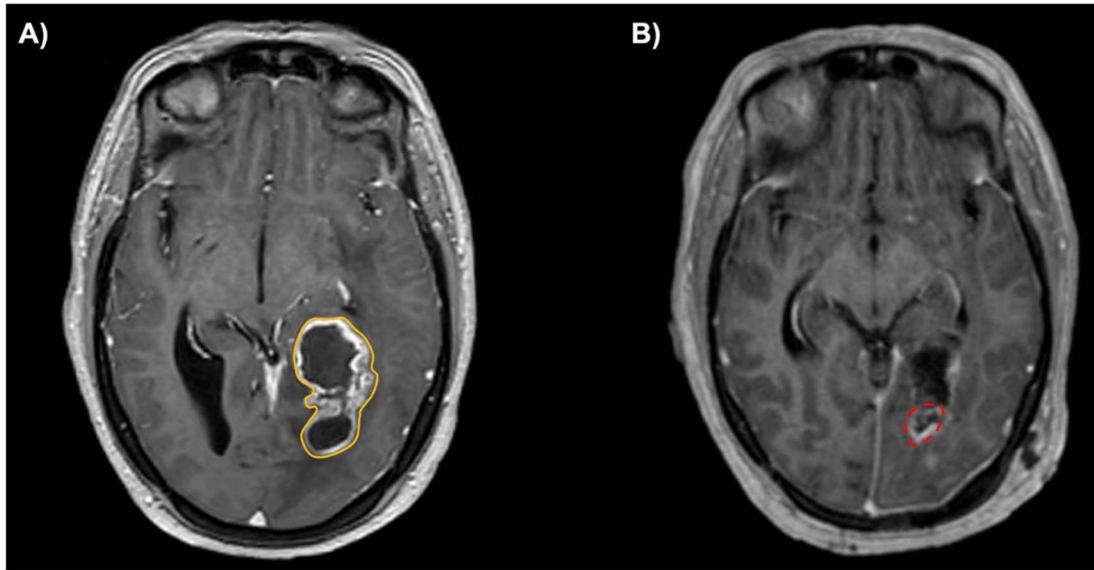


Figure 17 – Preoperative (A) and postoperative (B) T1-weighted post-contrast axial images showing NTR with nodular enhancement (dotted red area) of surgical cavity.

Tumour volumes were assessed on T1-weighted contrast enhanced volumetric images, retrieved from the institution’s digital archive system and stored as DICOM files. The imaging analysis was carried out using OsiriX MD (Osirix MD 9.0®, Pixmeo, Swiss). Tumour volume, expressed in cubic centimetres (cc), was computed with the Region of Interest (ROI) methodology. Tumour and postoperative pathologic enhancement of surgical cavity were segmented on axial images, then multiplied by the slice thickness through the software^{66,67} (Figures 18-19).



Figure 18 – Preoperative tumour volume computation by Osirix (Osirix MD 9.0®, Pixmeo, Swiss).

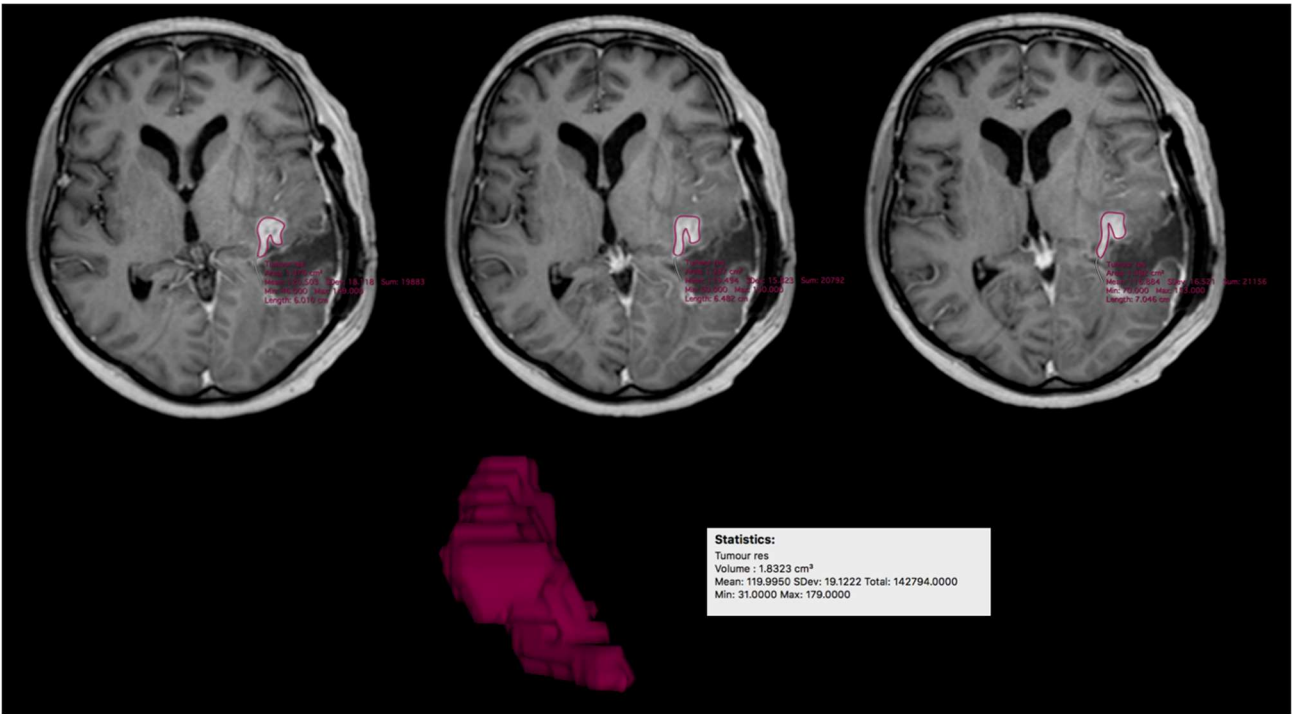


Figure 19 – Postoperative residual tumour volume computation by Osirix (Osirix MD 9.0[®], Pixmeo, Swiss).

The percentage of tumour resection was computed according to the formula:

$$\frac{(\text{preoperative tumour volume} - \text{postoperative tumour remnant volume})}{\text{preoperative tumour volume}} \times 100$$

3.6 Peripheral blood and tumour samples collection

Peripheral blood samples were collected at diagnosis (before surgery) and regularly over the follow-up (1, 3, 6 and 12 months after surgery), specifically in 23 patients at diagnosis, 16 patients 1 month after surgery, 8 patients 3 months after surgery, 9 patients 6 months after surgery and 5 patients 12 months after surgery (**Figure 20**).

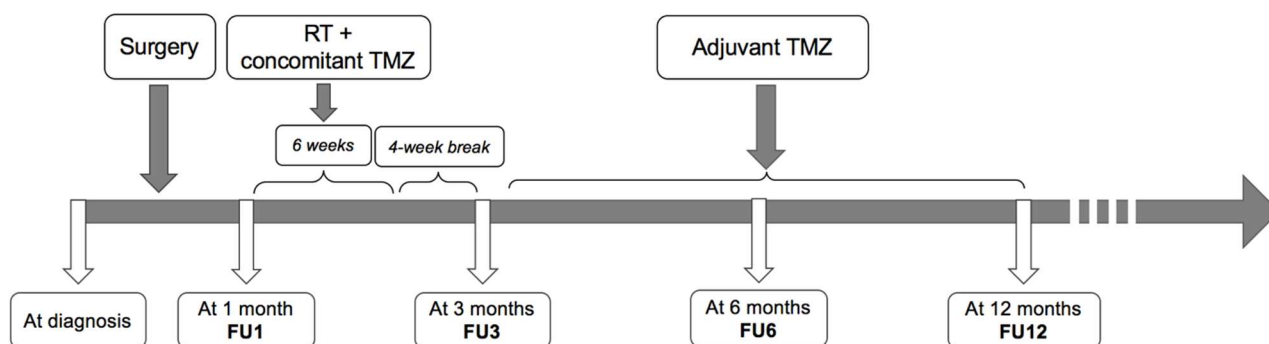


Figure 20 – Timing line of blood samples' collection reporting the standard adjuvant therapy scheme (see paragraph: “*Adjuvant therapies: radiotherapy and chemotherapy*”).

Plasma (5), buffy coat (1) and whole blood (4) aliquots were obtained by processing the peripheral blood samples collected in EDTA-coated collection tubes (BD Vacutainer® K2E 10.8mg), while serum aliquots (4) by processing the peripheral blood samples gathered in serum-specific collection tubes containing an inert gel (BD Vacutainer® SST™ II Advance 8.5 mL). The aliquots were then frozen at -80°C .

Plasma aliquots were obtained within 30 minutes from the withdrawal by: 1) spinning the peripheral blood at 3000 rpm for 10 minutes at 4°C ; 2) collecting and further centrifuging the supernatant at 4000 rpm for 20 minutes at 4°C . Buffy coat was achieved from the white interphase obtained after the first centrifugation of peripheral blood (1) in the plasma processing.

The serum-specific collection tubes were kept at room temperature for 60 minutes after the withdrawal to allow the coagulation process. Serum aliquots were obtained by spinning the peripheral blood at 3000 rpm for 15 minutes at 20°C and collecting the upper phase.

Furthermore, representative tumour tissue samples were taken during surgery and fresh frozen at -80°C .

3.7 Quantification of circulating miRNA levels in plasma

3.7.1 Total RNA isolation from plasma

Total RNA was isolated from 200 μL of plasma by means of the miRNeasy Mini Kit (Qiagen), which combines phenol/guanidine-based lysis of samples and silica membrane-based purification of total RNA, including small RNAs. According to the manufacturer's instructions, plasma samples were lysed in 1 mL of QIAzol Lysis Reagent (Qiagen) and 2.5 μL of 5 nM synthetic cel-miR-39-3p (Integrated DNA Technologies) from *C. elegans* were added to each sample as spike-in. After the addition of 200 μL of chloroform and centrifugation at 12000 g for 15 minutes at 4°C, the lysate was separated into the upper aqueous phase containing RNA, the white interphase containing DNA, and the lower organic phase. The upper aqueous phase was transferred into a new tube and the miRNeasy kit protocol was followed by using the miRNeasy MinElute spin columns, which allow total RNA to bind to the membrane and remove phenol or other contaminants. RNA was eluted from the spin columns in 35 μL of nuclease-free water. Total RNA concentration was measured using a NanoDrop spectrophotometer (Thermo Fisher Scientific, Inc.; Waltham, MA, USA) and RNA quality was assessed using the 260/280 ratio.

3.7.2 Droplet digital PCR (ddPCR) workflow

Circulating levels of miR-21-5p, miR 23b-3p and miR 34a-5p were quantified using the specific TaqMan microRNA assays (Thermo Fisher Scientific, Inc.; Waltham, MA, USA): miR 21-5p assay (ID 000397); miR 23b-3p assay (ID 000400); miR 34a-5p assay (ID 000426). Each assay employs the target-specific stem-loop primer during the reverse transcription to produce a template, which is then quantified in ddPCR using the specific TaqMan probe.

For each miRNA, 2.5 μL of purified total RNA were reversed transcribed using the TaqMan microRNA Reverse Transcription kit (Thermo Fisher Scientific, Inc.; Waltham, MA, USA) in a 7.5 μL reaction containing 0.075 μL of dNTPs, 0.500 μL of 50 U/ μL MultiScribe™ Reverse Transcriptase, 0.750 μL of 10X Reverse Transcription Buffer, 0.095 μL of RNase Inhibitor, 2.08 μL of nuclease-free water and 1.5 μL of specific stem-loop primers. The reverse transcription reaction was performed at 16°C for 30 minutes, followed by incubation at 42°C for 30 minutes and 85°C for 5 minutes.

The ddPCR experiments were performed according to the QX200 TaqMan ddPCR protocol. For each miRNA, 1.33 μL of reverse transcription product, 11 μL of 2X ddPCR Supermix for probes (Bio-Rad Laboratories, Inc.; Hercules, CA, USA) and 1.1 μL of specific 20X TaqMan PCR probe assay (Thermo Fisher Scientific, Inc.; Waltham, MA, USA) were mixed to obtain a final volume of 22 μL . 20 μL of the ddPCR assay mixture and 70 μL of droplet generation oil for probes (Bio-Rad Laboratories, Inc.; Hercules, CA, USA) were loaded into a disposable droplet generator cartridge (Bio-Rad Laboratories, Inc.; Hercules, CA, USA). Once the droplet generation process has been completed by the QX200 droplet generator (Bio-Rad Laboratories, Inc.; Hercules, CA, USA), 45 μL of the mix containing the droplets were transferred to a 96-well PCR plate (Bio-Rad Laboratories,

Inc.; Hercules, CA, USA) using a multichannel pipette. The plate was heat-sealed with foil and placed in a conventional thermal cycler set as follows: 95°C for 10 minutes, 40 cycles of 94°C for 30 seconds followed by 58°C for 1 minute, 98°C for 10 minutes, 4°C for 40 minutes and a 4°C indefinite hold. A negative control, containing all components of the reaction except for the template, was included in each ddPCR experiment for every miRNA assay. Following the PCR amplification step, the plate containing the droplets was loaded into the QX200 Droplet Reader (Bio-Rad Laboratories, Inc.; Hercules, CA, USA) and the results were analysed using QuantaSoft Software, version 1.7 (Bio-Rad Laboratories, Inc.; Hercules, CA, USA), with Absolute Quantification (ABS) set as the type of experiment. For each plate, the total number of droplets (positive and negative) was verified; one-dimension plot (1-D plot) was used to manually set the threshold and distinguishing the positive droplets, which contained at least one copy of the target miRNA and exhibited high fluorescence, from the negative droplets with very low fluorescence (**Figure 21**). The fraction of positive droplets was fitted by the software to a Poisson distribution to determine the concentration of each target miRNA in copies/ μL .

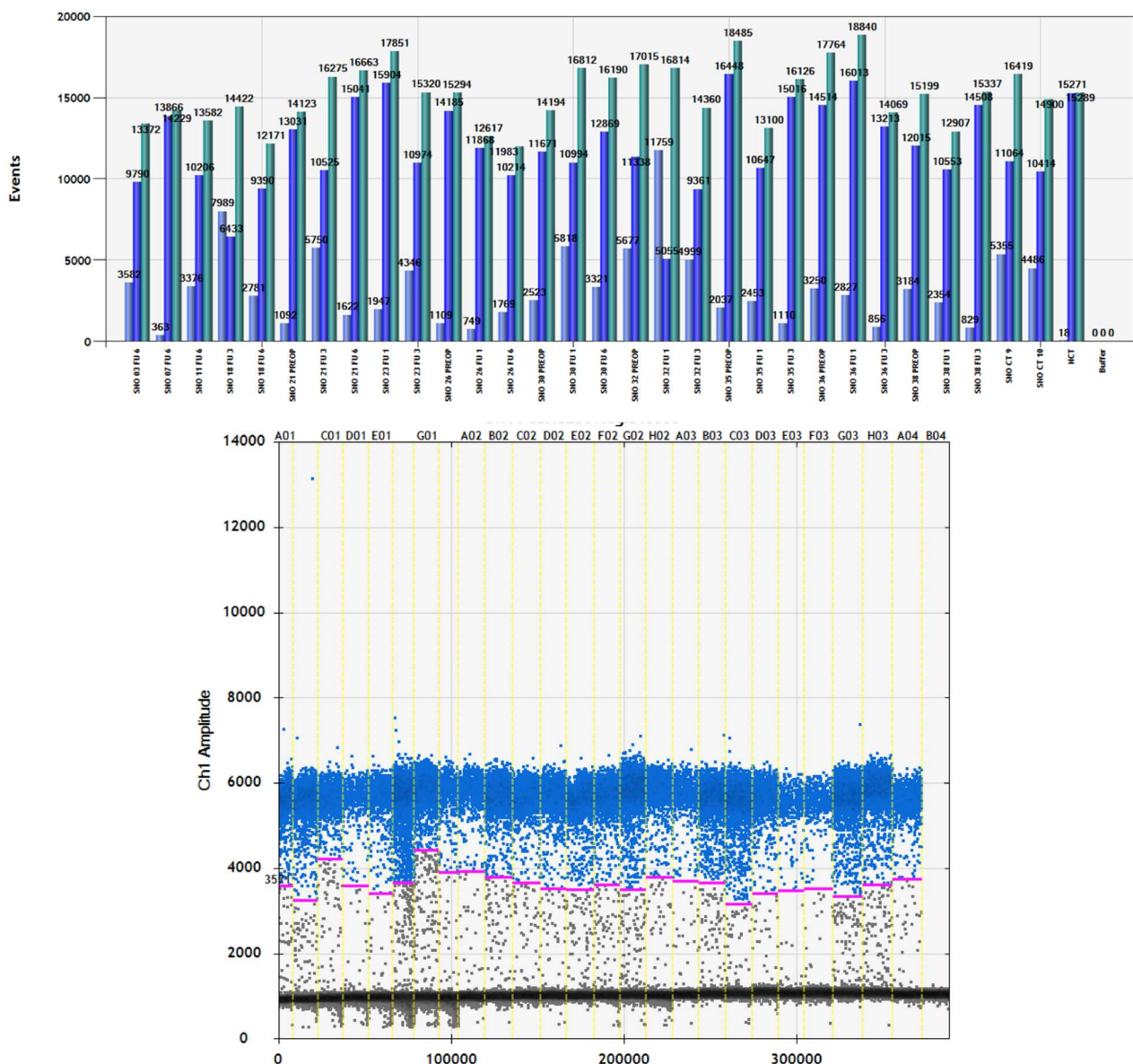


Figure 21 – Illustrative outputs obtained with QuantaSoft Software (version 1.7; Bio-Rad Laboratories, Inc.; Hercules, CA, USA): A) histograms depicting the number of droplets (positive, negative, total); B) 1-D plot.

3.8 Computation of miRNAs' plasmatic concentrations changes

The percentage of change of miRNAs' plasmatic concentrations between different timing points was calculated according to the following formula:

$$\Delta [miR_{T_2} - miR_{T_1}] = \frac{[miR_{T_2}] - [miR_{T_1}]}{[miR_{T_1}]} \times 100$$

The concentrations were defined as *increased* or *stable/decreased* if their variation was greater or lower than 10%.

3.9 miRNA network creation

The miRNet Version 2.0 (<https://www.mirnet.ca/miRNet/home.xhtml>) allowed the creation of miRNA-centric networks for systems-level interpretation of miRNA functions and gene regulations⁶⁸.

The process started selecting “miRNAs” as module of input. Thereafter in the *Upload section* “*H. sapiens*” was selected as *Organism*, “*miRBase ID*” as *ID type* and “*Genes (miRTarBase v8.0)*” as *Targets*; “*hsa-miR-21-5p*; *hsa-miR-23b-3p*; *hsa-miR-34a-5p*” were entered in the *miRNA list*. In the Network Builder we set the *Degree cutoff* at *1.0* and applied it to all the network nodes.

3.10 Statistical analysis

Categorical variables were reported as numbers (percentages) and compared with either the Pearson's chi-square test with Yates' correction or Fisher's exact test as appropriate. Continuous variables were expressed as means (\pm standard deviation, SD) or medians (interquartile range, IQR) and compared with the unpaired Student's *t* test or Mann-Whitney U test as appropriate, after confirmation of normality of distribution by histograms' visual inspection and the Shapiro-Wilk test. The correlations of plasmatic miRNA concentrations with patients' and tumours' variables were reported either as Pearson's or rho Spearman's correlation coefficients (*r*) in case of linear and non-linear associations, respectively. The results of linear regression were reported as adjusted R² estimates on included figures.

Receiver operating characteristics (ROC) curves were generated to assess sensitivity, specificity and predictive power of each miRNA for GBM. The area under curve (AUC) values ranged from 0

to 1: values around 0.5 indicated an equal distribution among healthy volunteers and GBM patients; values between 0.5 and 1 suggested that miRNAs' levels were higher in GBM patients; values between 0 and 0.5 indicated that miRNA were higher in healthy volunteers^{3,35}. The cut-off values were identified according to the Youden index (YI)^{69,70}.

Hierarchical cluster analysis with Ward's methods and K-means cluster analysis were performed to classify patients affected by GBM according to their miRNAs' plasmatic levels.

Multivariate linear regression analysis was used to determine which variables independently predicted plasmatic miRNA levels at different timing points. Results of multivariate modelling were expressed as odds ratios (OR), 95% confidence interval (CI) after adjustment for covariates and *p* for significance.

The overall survival (OS) and recurrence free survival (RFS) analyses were performed according to the Kaplan-Meier method from the date of surgery to the date of death, tumour recurrence/progression or end of observation. Censoring was applied to the last follow-up date for patients who remained alive. The log-rank test was used to compare survivals⁷⁰.

The Cox regression analysis was adopted to determine which variables independently predicted OS and RFS. Results of multivariate modelling were expressed as hazard ratios (HR), 95% confidence interval (CI) after adjustment for covariates and *p* for significance.

A two-tailed probability (*p*) value ≤ 0.05 was considered statistically significant.

Statistical analysis was performed using the package SPSS for Windows, Version 23.0 (SPSS Inc.; Chicago, IL, USA).

4. RESULTS

4.1 Patients' and tumour's features

The mean age at diagnosis was 57.9 years (± 9.6) with a male prevalence (60.9%). Eighteen patients (78.3%) were "independent" in daily life activities (KPS ≥ 90) and 21 (91.3%) did not present significant neurological disabilities (mRS ≤ 2). Steroids and anti-epileptic drugs were preoperatively administered in 21 (91.3%) and 10 cases (43.5%), respectively (see **Appendix – Table 1**).

Lesions were left-sided in 11 cases (47.8%), right-sided in 9 cases (39.1%) and sited along the midline in 3 cases (13.1%). The preoperative brain MRI showed tumour contrast enhancement in all cases and necrosis in 87.0% of them. The mean tumour volume was 33.8 cc (± 19.1); 10 patients (43.5%) presented a tumour volume smaller than 30.0 cc.

Nineteen (82.6%) were newly diagnosed tumours. Intraoperative "strong" tumour fluorescence was recorded in all but one (95.7%) surgeries. The pathological report was consistent with GBM IDH-wildtype WHO grade 4 in 22 cases (95.7%) and astrocytoma IDH-mutant WHO grade 4 in the remaining one (4.3%). Twelve cases (52.2%) showed un-methylated MGMT, 4 (17.4%) methylated MGMT and the remaining 7 (30.4%) hyper-methylated MGMT, with no statistical difference in their distribution ($p=0.119$). ATRX expression was "retained" (in the cell nucleus) in 20 cases (87.0%) and "lost" in 3 cases (13.0%, $p<0.001$). p53 tumour expression was classified as "absent" in 2 cases (8.7%), "intermediate" in 13 cases (56.5%) and "strong" in 8 cases (34.8%, $p=0.028$). The mean Ki67% was 41.3 (± 19.8).

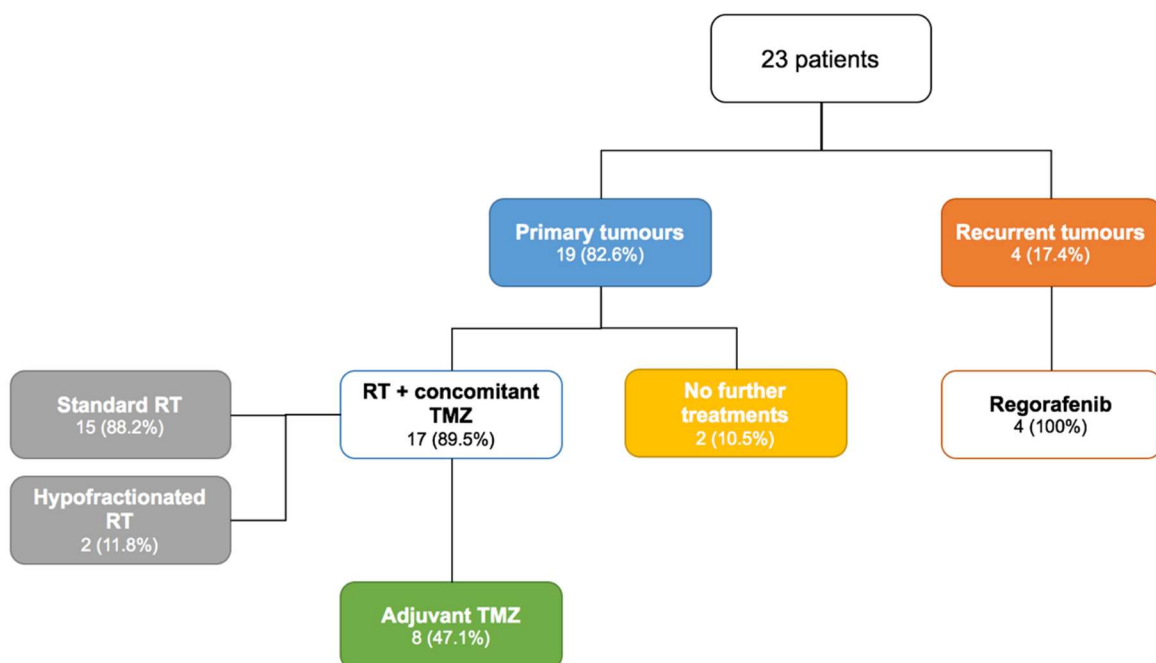


Figure 22 – Flow chart showing patients’ distribution according to adjuvant therapies (RT: radiotherapy; TMZ: temozolomide).

Seventeen out of 19 patients (89.5%) operated for newly diagnosed tumours were treated with RT and concomitant temozolomide. The RT protocol was standard in 15 (88.2%) and hypo-fractionated in the remaining 2 (11.8%). Overall, 12 patients (52.2%) were suitable for adjuvant chemotherapy. Eight of them were started on temozolomide, 3 of which were treated with regorafenib at tumour recurrence, and 4 presenting with recurrent GBM were treated with regorafenib (**Figure 22**).

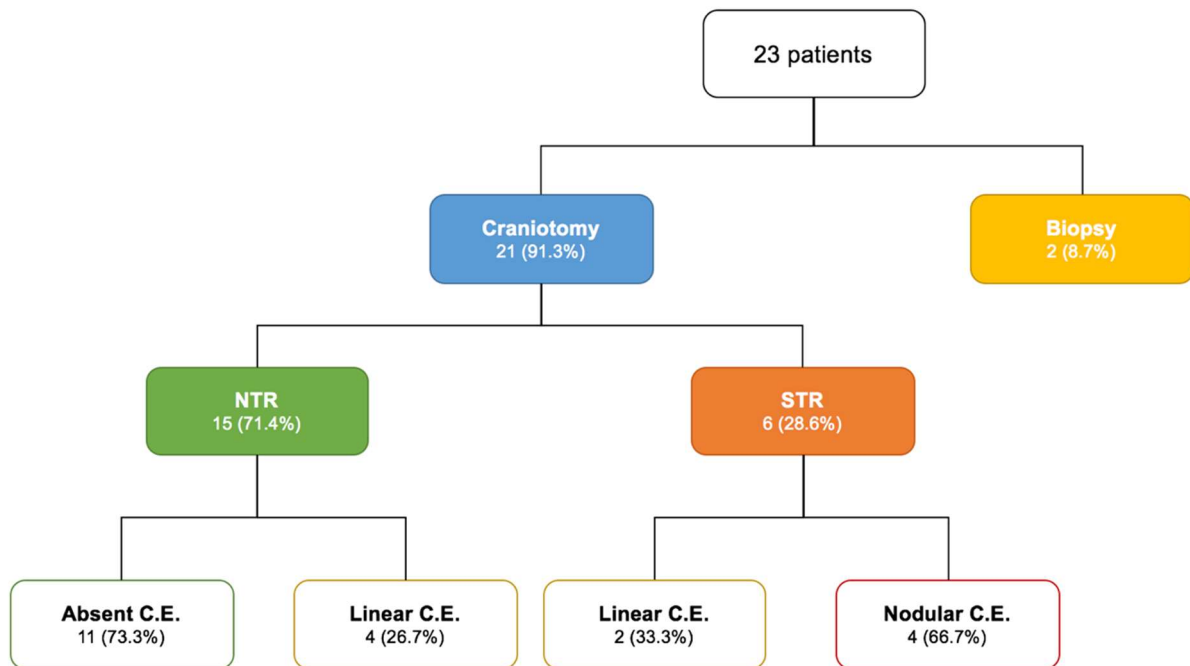


Figure 23 – Flow chart showing patients’ distribution according to the extent of tumour resection (C.E.: contrast enhancement of surgical cavity; NTR: near-total resection; STR: subtotal resection).

Craniotomy was performed in 21 patients (91.3%) and NTR was achieved in 15 of them (71.4%). In case of STR (6, 28.6%), the pathologic contrast enhancement of surgical cavity was “linear” in 2 (33.3%) and “nodular” in 4 (66.7%) cases (**Figure 23**) with a mean tumour remnant volume of 5.88 cc (± 3.74). The mean percentage of tumour resection was 99.3% (± 1.0) and 80.7% (± 7.9) in case of NTR and STR, respectively (see **Appendix – Table 2**).

4.2 Results of ddPCR assays

Overall, the ddPCR assays provided a mean of 16265.48 total droplets (± 2045.53 ; median: 16342.00, IQR 15147.00-17716.00) showing similar results for each miRNA (miR 21-5p: 15643.15 \pm 1808.62; miR 23b-3p: 16728.15 \pm 2134.99; miR 34a-5p: 16429.80 \pm 2041.81; **Figure 24**).

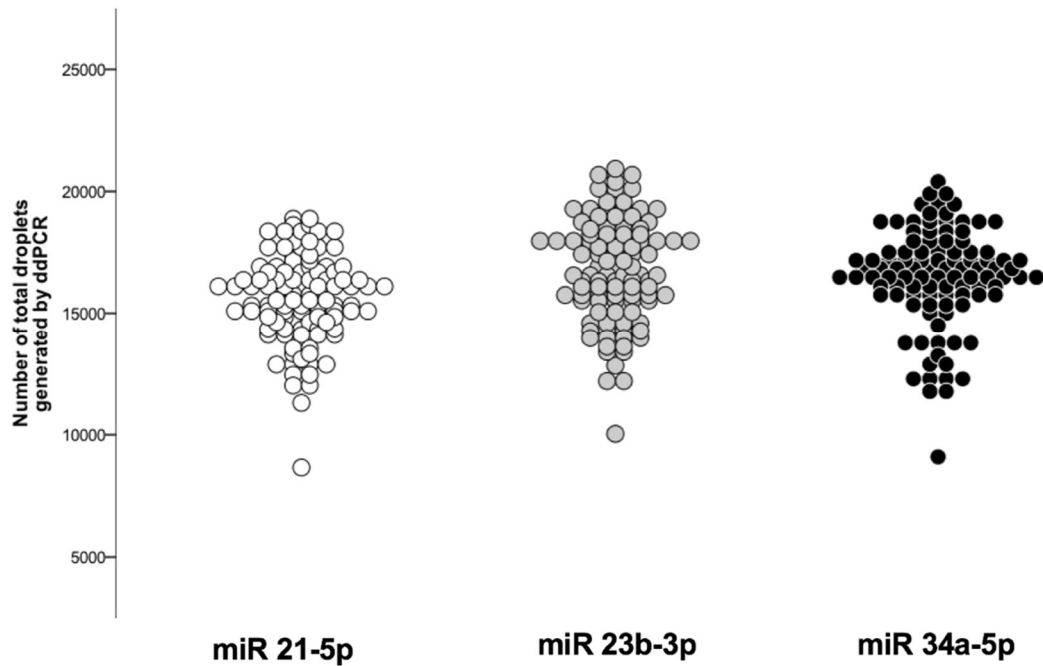


Figure 24 – Dot plots showing the total number of droplets generated by ddPCR for miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C).

The mean number of positive droplets detected by ddPCR was 650.80 (± 1443.36 ; median: 56.00, IQR 13.00-429.00). The mean number of positive droplets was greater for miR 21-5p (1783.18 ± 2054.95) followed by miR 23b-3p (129.29 ± 222.63) and miR 34a-5p (22.14 ± 29.95) (**Figure 25**).

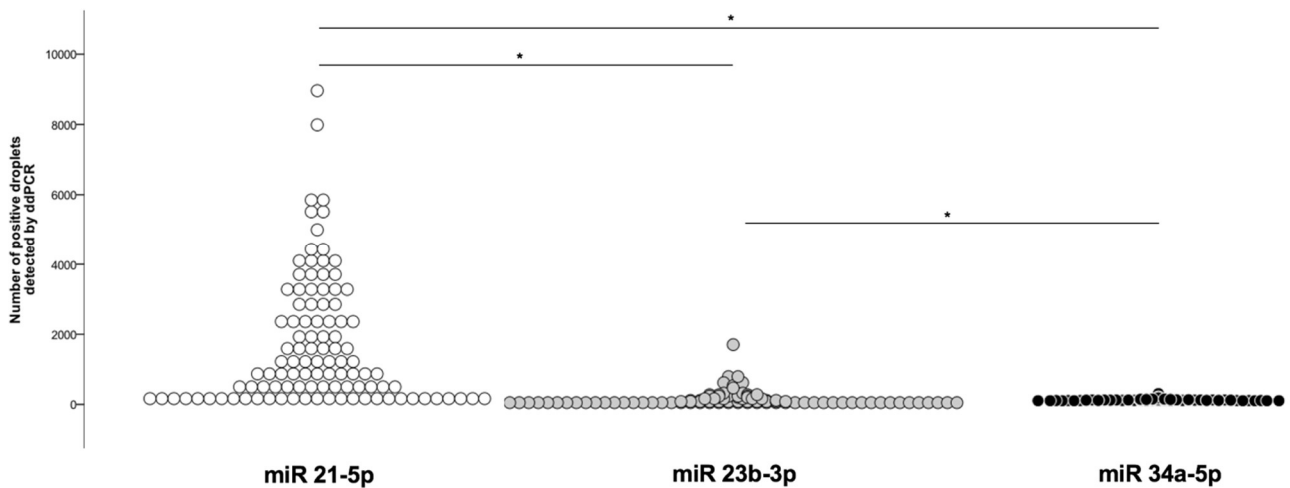


Figure 25 – Dot plots showing the number of positive droplets detected by ddPCR for miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C) (*: $p < 0.001$).

4.3 miRNAs' plasmatic concentrations in healthy volunteers and patients

miR 21-5p, miR 23b-3p and miR 34a-5p were all detectable in healthy volunteers with a mean plasmatic concentration of 114.28 copies/ μL (± 125.41), 4.43 copies/ μL (± 5.40) and 0.68 copies/ μL

(± 0.62), respectively. Patients presented on average 115.49 copies/ μL (± 132.57) of miR 21-5p, 7.18 copies/ μL (± 10.15) of miR 23b-3p and 1.28 copies/ μL (± 1.10) of miR 34a-5p at diagnosis. Healthy volunteers showed significantly lower mean circulating levels of miR 34a-5p than patients at diagnosis (0.68 vs. 1.28 copies/ μL ; $p=0.012$) (**Figure 26**).

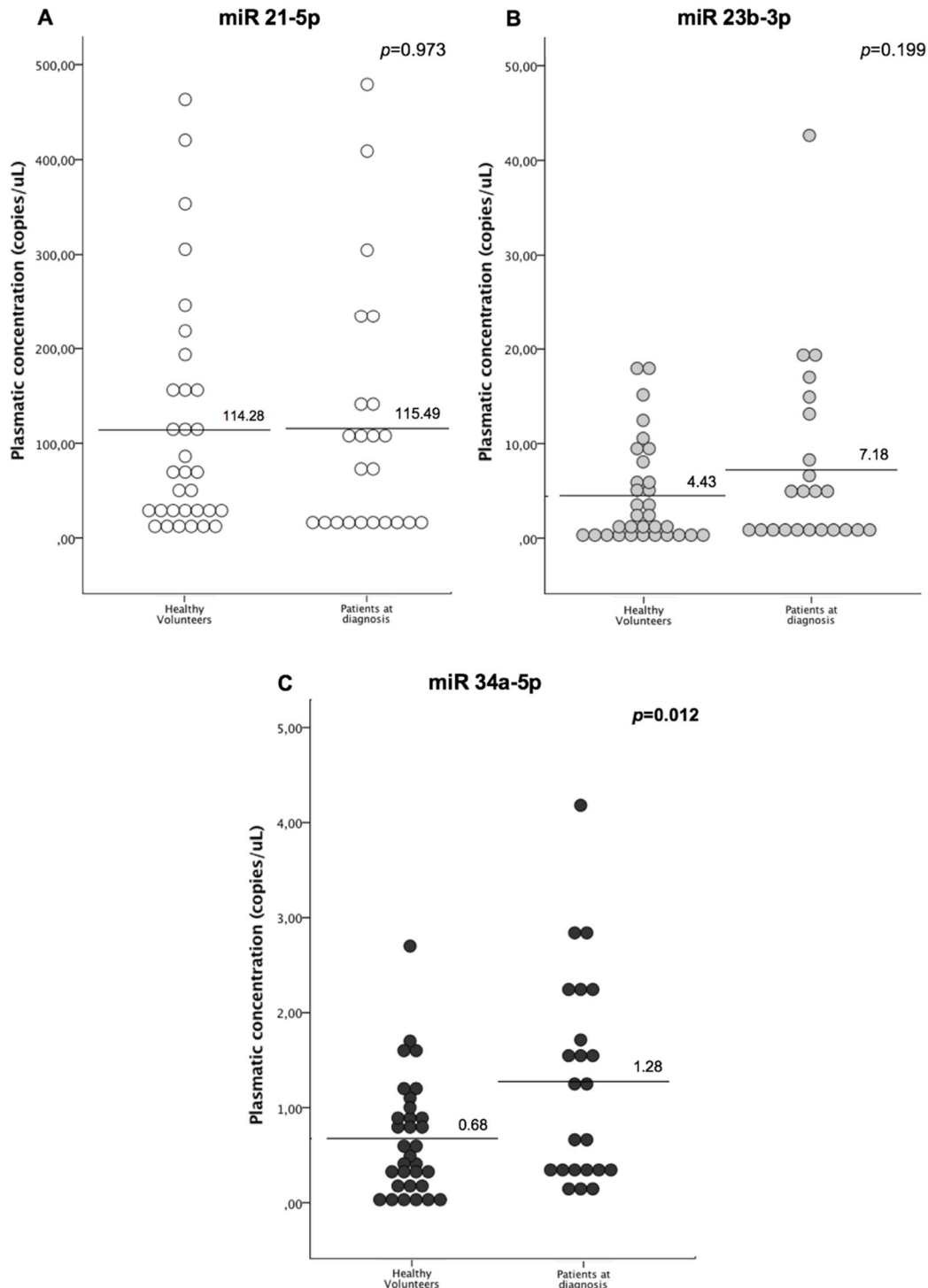


Figure 26 – Dot plots showing plasmatic concentrations of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) in healthy volunteers and patients at diagnosis (mean values represented by lines).

The diagnostic efficacy of the tested miRNAs in distinguishing patients affected by GBM from healthy individuals was evaluated using the ROC curve analysis (**Figure 27**). The AUC was 0.484 for miR 21-5p (95% C.I. 0.326 – 0.643, $p=0.844$), 0.537 for miR 23b-3p (95% C.I. 0.375-0.700, $p=0.639$) and 0.664 for miR 34a-5p (95% C.I. 0.515-0.814, $p=0.039$). The cut-off value for miR 34a-5p was 1.25 copies/ μ L (sensitivity: 47.8%; specificity: 87.5%; YI: 0.353).

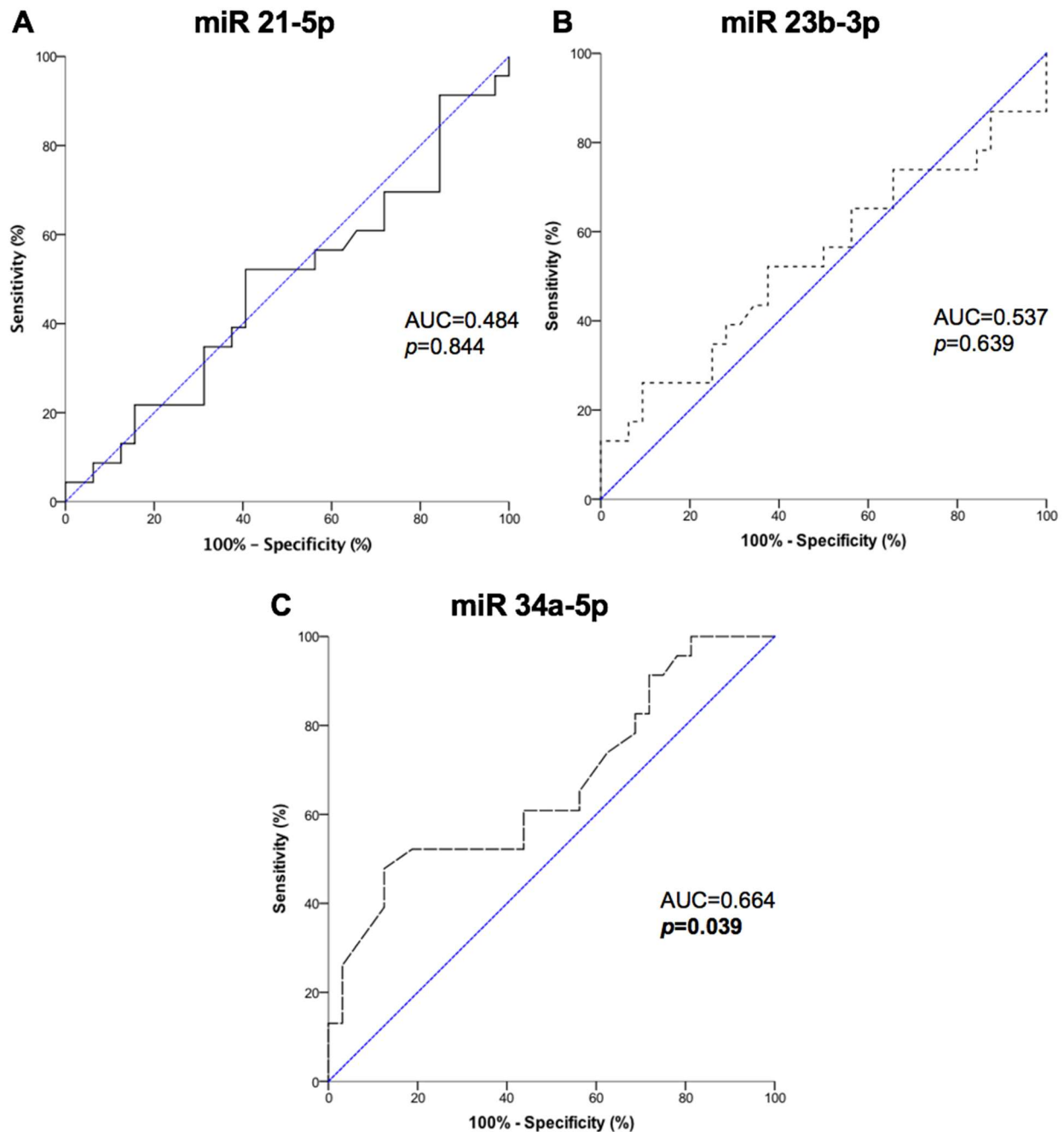


Figure 27 – ROC curves of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between healthy volunteers and patients at diagnosis.

Mean plasmatic concentrations of miR 21-5p (356.16 ± 303.89 vs. 114.28 ± 125.41 copies/ μ L; $p=0.001$) and miR 23b-3p (22.26 ± 19.04 vs. 4.43 ± 5.40 copies/ μ L; $p<0.001$) in patients 3 months

after surgery reached significantly higher levels compared to healthy volunteers. However, mean circulating levels of miR 34a-5p remained significantly higher in patients at each timing point compared to healthy volunteers (**Figure 28**).

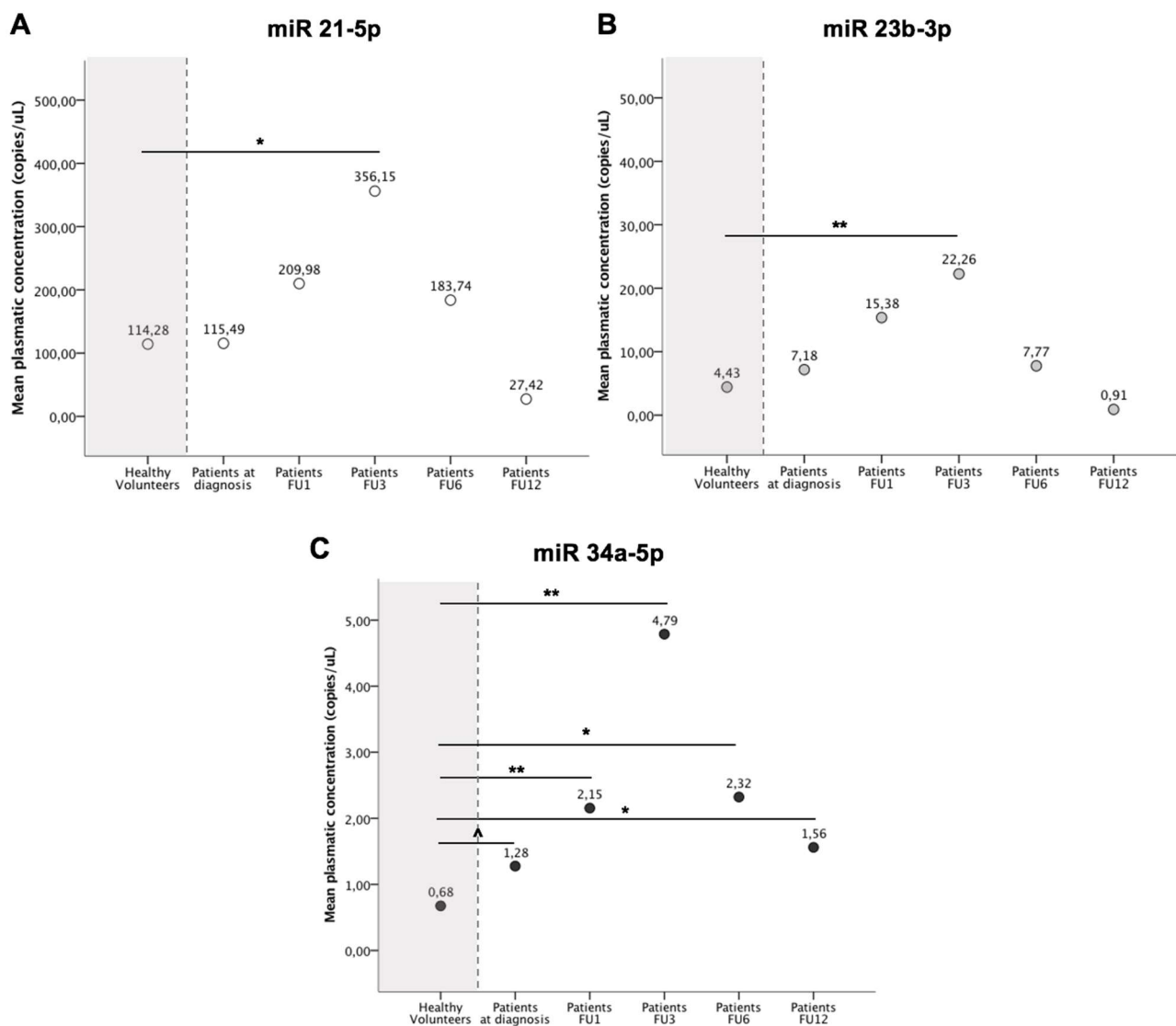


Figure 28 – Comparison of mean circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between healthy volunteers and patients at different timing points (**: $p < 0.001$; *: $p = 0.001$; ^: $p < 0.05$).

4.4 Patients' clustering according to miRNAs' plasmatic concentrations at diagnosis

Patients affected by GBM were clustered according to the similarity of their miRNAs' absolute circulating concentrations at the time of diagnosis (**Figure 29**). Specifically, two main groups of patients were identified: group 1 presented low absolute levels of all miRNAs; group 2 showed high levels of miRNAs. miR 21-5p (mean: 331.60 vs. 55.46 copies/ μ L, $p < 0.001$) and miR 23b-3p (22.66 vs. 2.88 copies/ μ L, $p < 0.001$), but not miR 34a-5p (1.90 vs. 1.10 copies/ μ L, $p = 0.155$), were significantly higher in group 2.

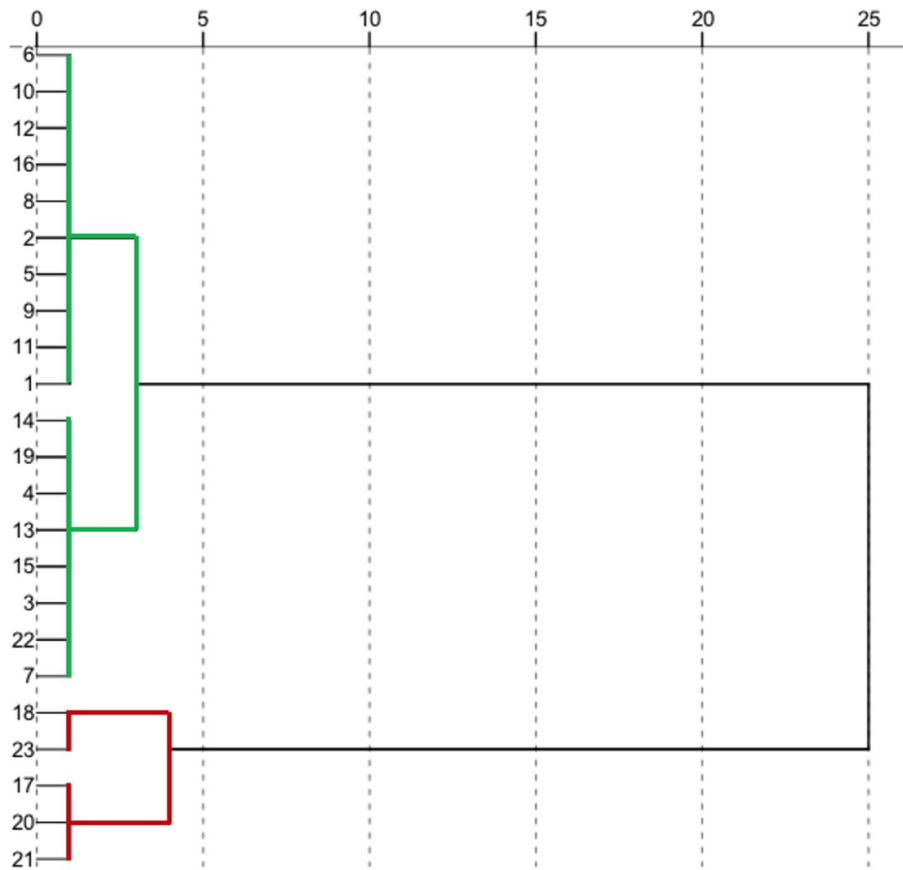


Figure 29 – Dendrogram showing the hierarchical clustering of GBM patients according to their absolute circulating miRNAs' concentrations (miR 21-5p, miR 23b-3p and miR 34a-5p) at the time of diagnosis. Group 1 (green line); group 2 (maroon line).

Furthermore, we performed a hierarchical clustering to group GBM patients according to the similarity of their miRNAs' circulating profile at the time of diagnosis relative to the mean plasmatic levels in healthy volunteers (**Figure 30**). Patients were clustered into four distinct groups: group 1 presented high expression levels for all miRNAs; group 2 showed low levels of miR 21-5p and high levels of miR 23b-3p; group 3 presented low circulating levels of all miRNAs; group 4 mainly low levels of miR 21-5p and miR 23b-3p and high levels of miR 34a-5p.

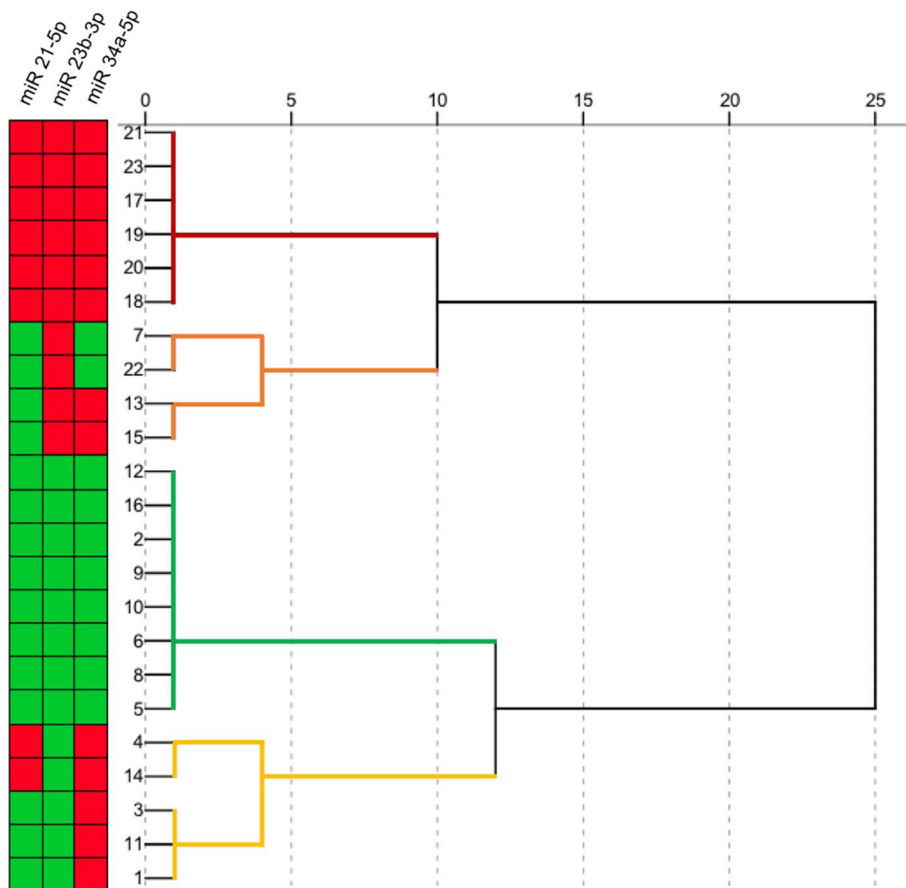


Figure 30 – Hierarchical clustering classifies GBM patients according to their circulating miRNAs’ concentrations at the time of diagnosis (miR 21-5p, miR 23b-3p and miR 34a-5p) relative to the plasmatic levels in healthy volunteers. Group 1 (maroon line); group 2 (orange line); group 3 (green line); group 4 (yellow line). Matrix coding: a) red: above the mean miRNAs’ circulating levels in healthy volunteers; b) green: below the mean miRNAs’ circulating levels in healthy volunteers.

4.5 Longitudinal course of miRNAs’ plasmatic concentrations

4.5.1 Variations of miRNAs’ plasmatic concentrations

Patients experienced increase, decrease or stability of miRNAs plasmatic concentrations over time. Specifically, miR 21-5p: decreased and increased in 8 cases (50%), respectively, between the diagnosis and 1 month after surgery; decreased in 4 cases (66.7%) and increased in 2 cases (33.3%) between 1 and 3 months after surgery; decreased in 4 cases (80%) and increased in 1 case (20%) between 3 and 6 months after surgery; eventually, decreased in 4 cases (100%) between 6 and 12 months after surgery (**Figure 31**).

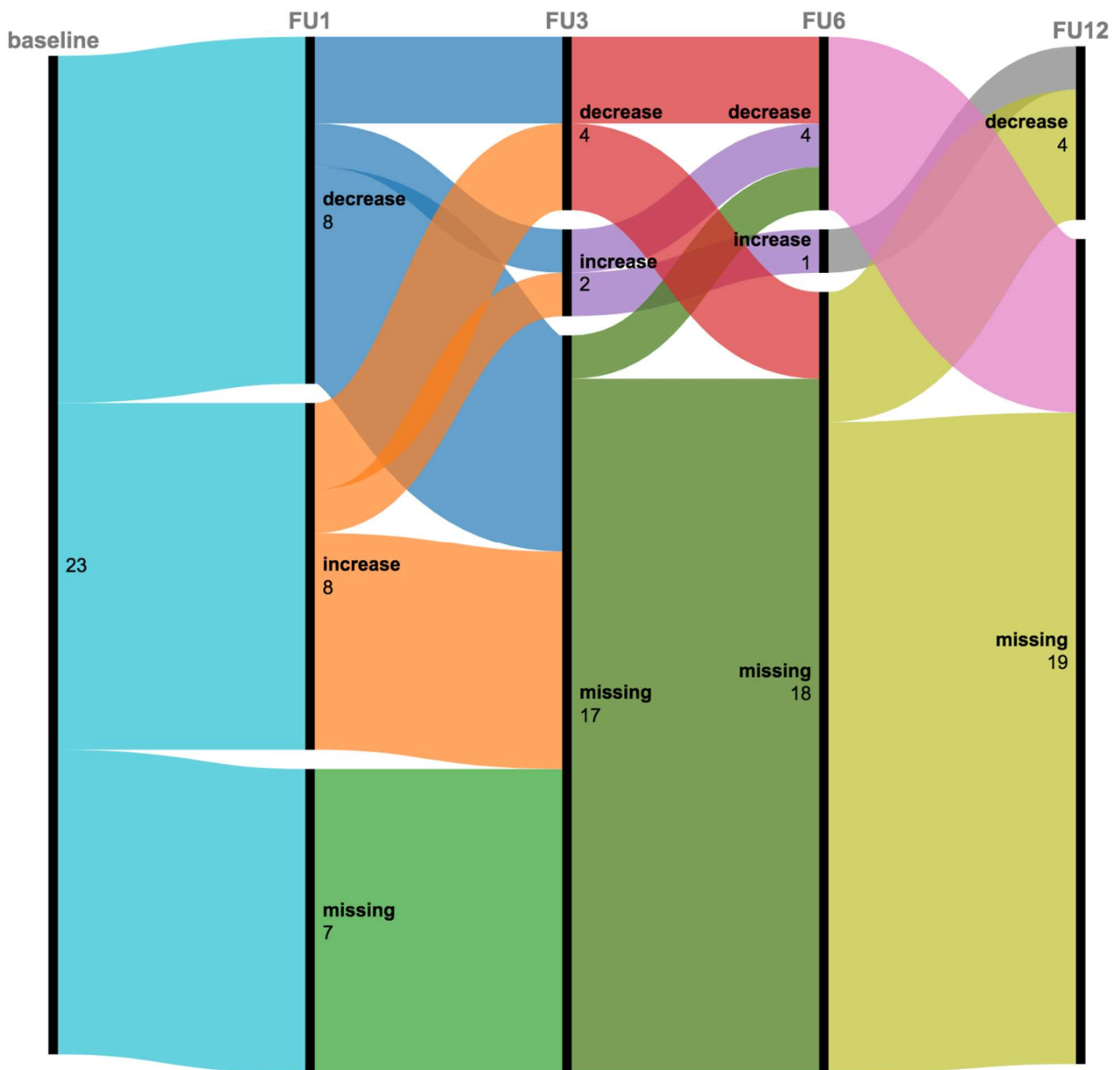


Figure 31 – Alluvial diagram for miR 21-5p plasmatic concentrations stratified by variations (increase, decrease and stability) over time (1, 3, 6 and 12 months after surgery).

miR 23b-3p plasmatic concentrations showed: stability in 2 cases (12.5%), increase in 8 cases (50%) and decrease in 6 cases (37.5%) between the diagnosis and 1 month after surgery; decrease in 4 cases (66.7%) and increase in 2 cases (33.7%) between 1 and 3 months after surgery; decrease in 5 cases (100%) between 3 and 6 months after surgery; decrease in 4 cases (100%) between 6 and 12 months after surgery (**Figure 32**).

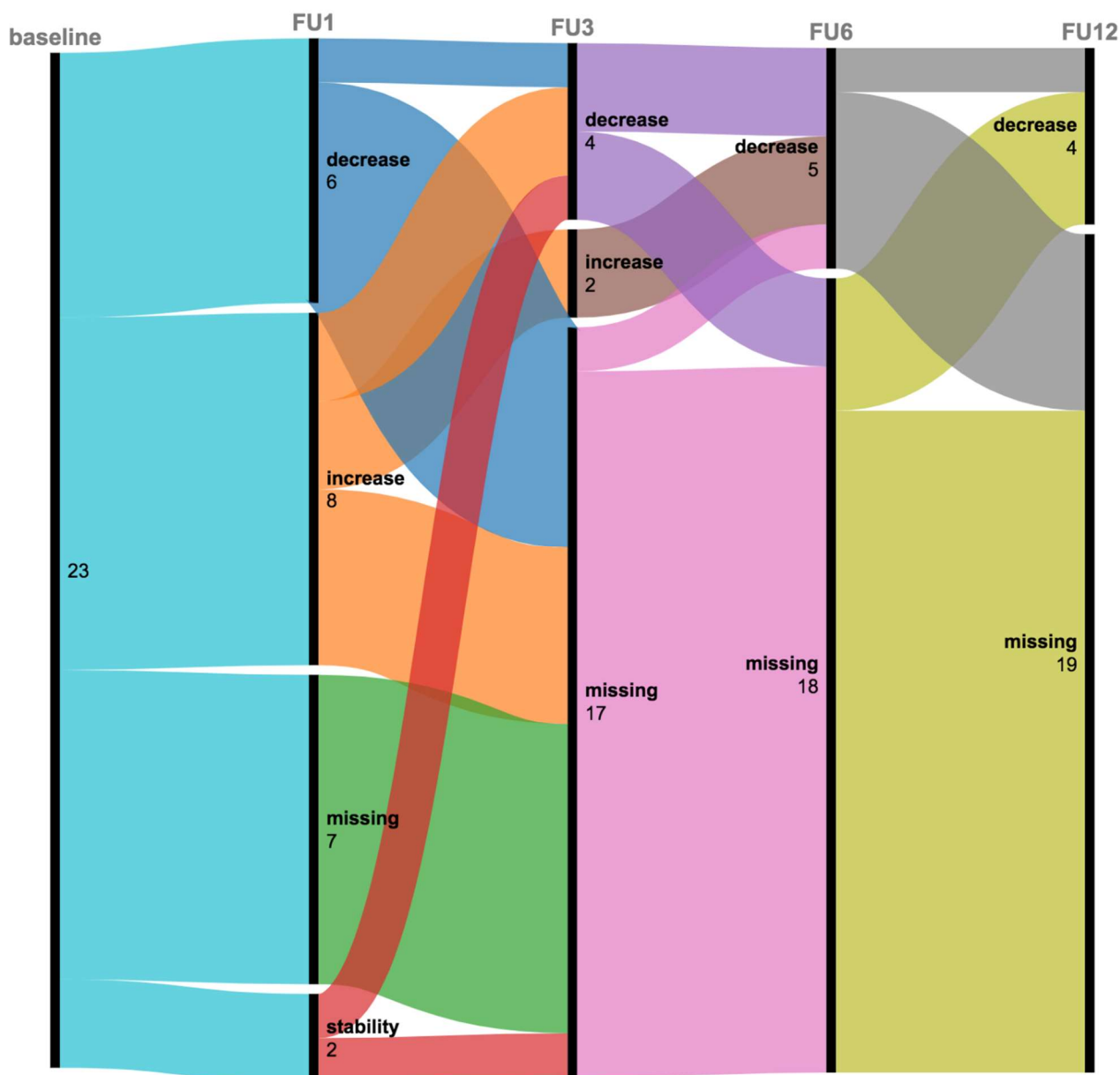


Figure 32 – Alluvial diagram for miR 23b-3p plasmatic concentrations stratified by variations (increase, decrease and stability) over time (1, 3, 6 and 12 months after surgery).

The longitudinal course of miR 34a-5p circulating levels registered: increase in 12 cases (75%) and decrease in 4 cases (25%) between the diagnosis and 1 month after surgery; decrease in 3 cases (50%), increase in 2 cases (33.7%) and stability in 1 case (16.7%) between 1 and 3 months after surgery; decrease in 3 cases (60%) and stability in 2 cases (40%) between 3 and 6 months after surgery; decrease and stability in 2 cases (50%), respectively, between 6 and 12 months after surgery (**Figure 33**).

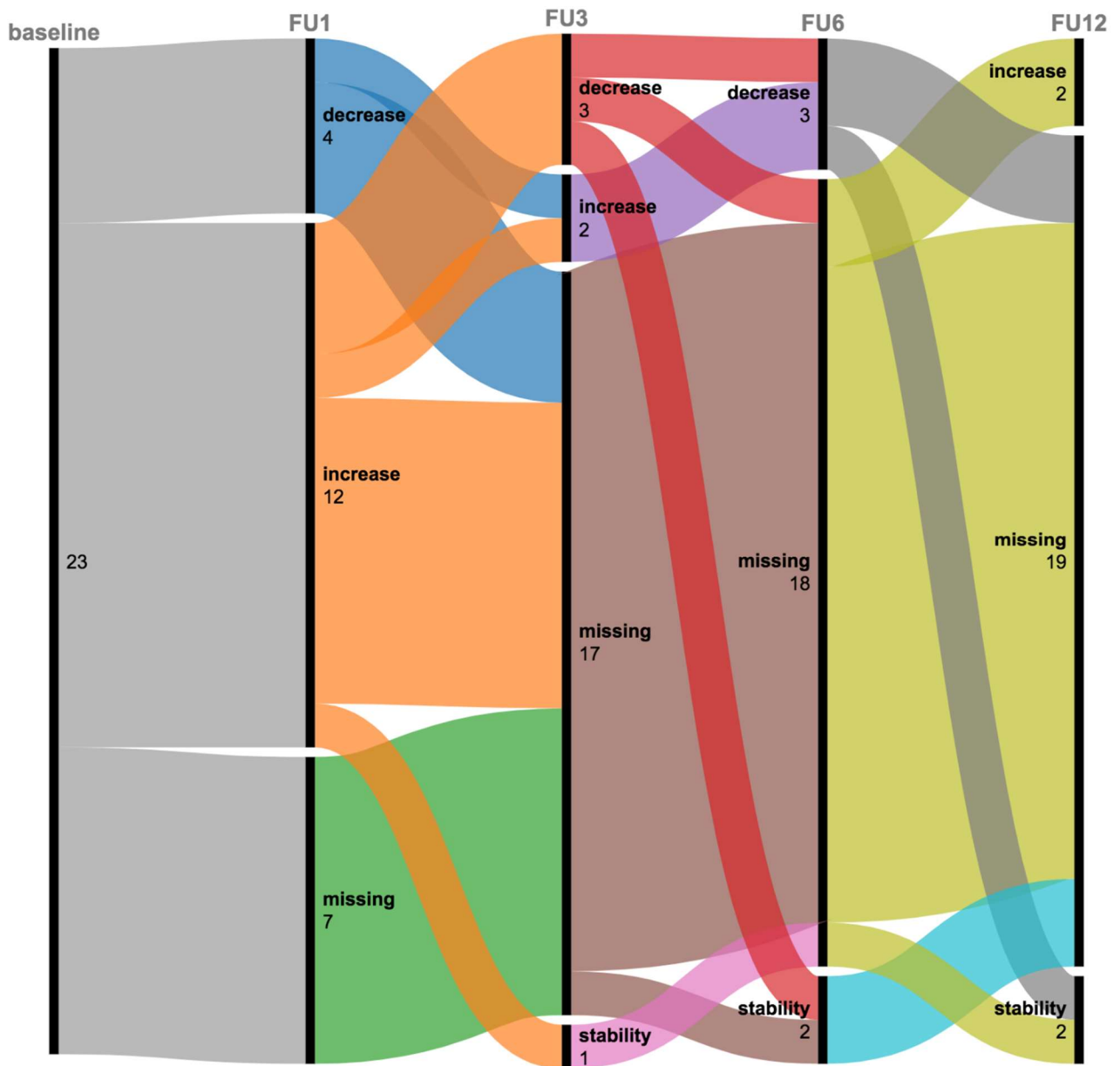


Figure 33 – Alluvial diagram for miR 34a-5p plasmatic concentrations stratified by variations (increase, decrease and stability) over time (1, 3, 6 and 12 months after surgery).

Between the diagnosis and 1 month after surgery the mean rates of increase were 55.5%, 74.7%, 53.7% and the mean rates of decrease were 42.2%, 58.4%, 55.7% for miR 21-5p, miR 23b-3p and miR 34a-5p, respectively (**Figure 34**).

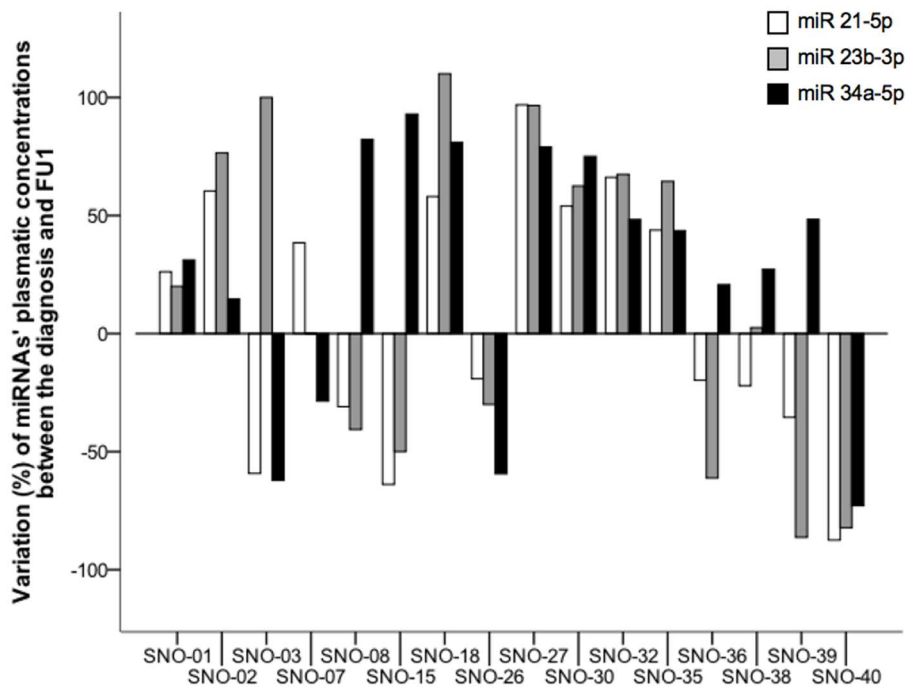


Figure 34 – Variations (%) of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations between the diagnosis and 1 month after surgery.

Between 1 and 3 months after surgery the mean rates of increase were 1621.9%, 15755.68%, 793.3% and the mean rates of decrease were 65.9%, 67.5%, 51.8% for miR 21-5p, miR 23b-3p and miR 34a-5p, respectively (**Figure 35**).

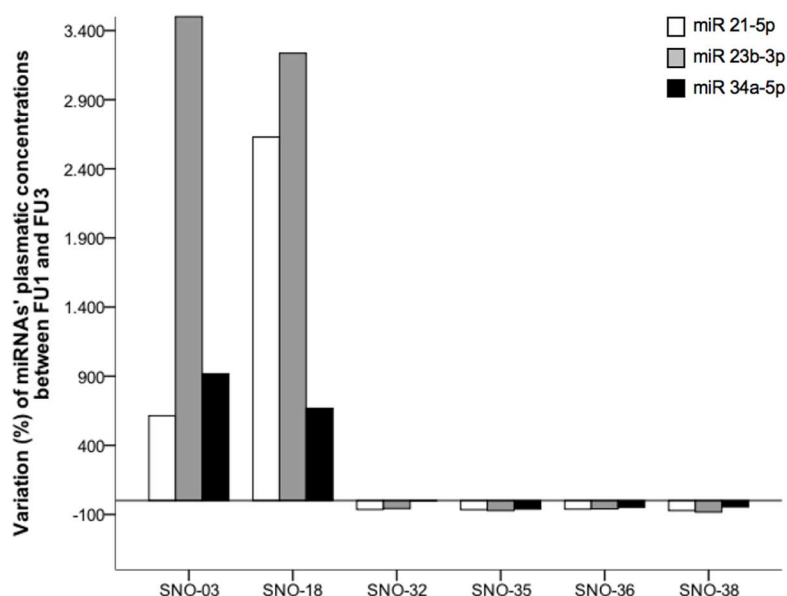


Figure 35 – Variations (%) of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations between 1 and 3 months after surgery.

Between 3 and 6 months after surgery the mean rates of increase were 37.5%, 0%, 0% and the mean rates of decrease were 74.2%, 76.7%, 78.5% for miR 21-5p, miR 23b-3p and miR 34a-5p, respectively (Figure 36).

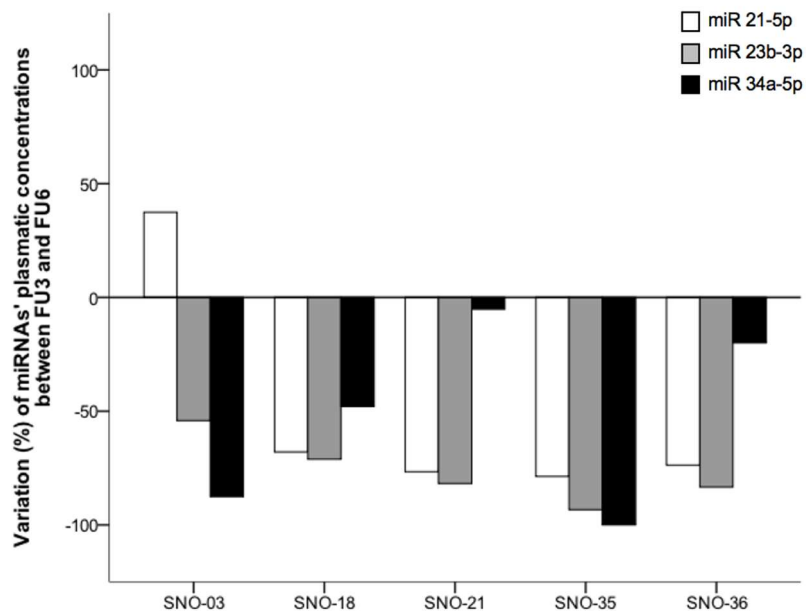


Figure 36 – Variations (%) of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations between 3 and 6 months after surgery.

Eventually, between 6 and 12 months after surgery miR 21-5p, miR 23b-3p and miR 34a-5p increased in mean of 0%, 0%, 125.0% and decreased in mean of 85.9%, 85.8%, 66.7%, respectively (Figure 37).

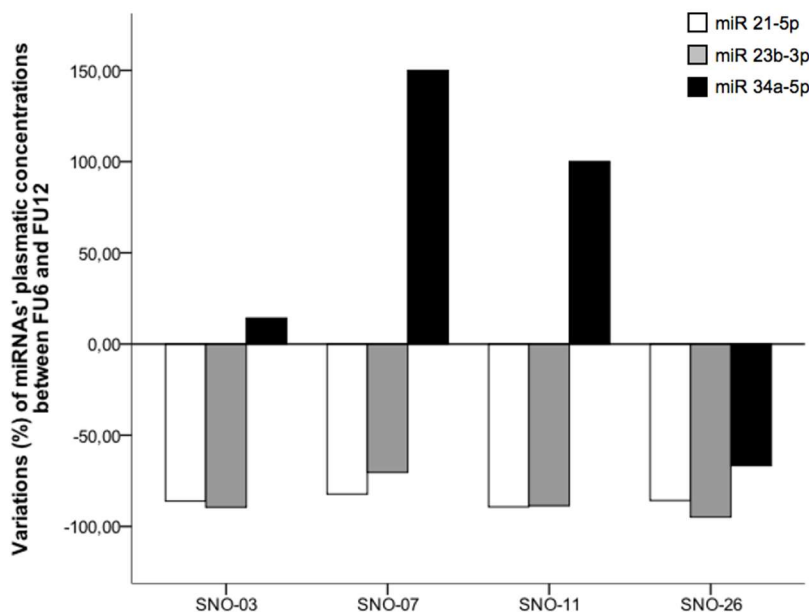
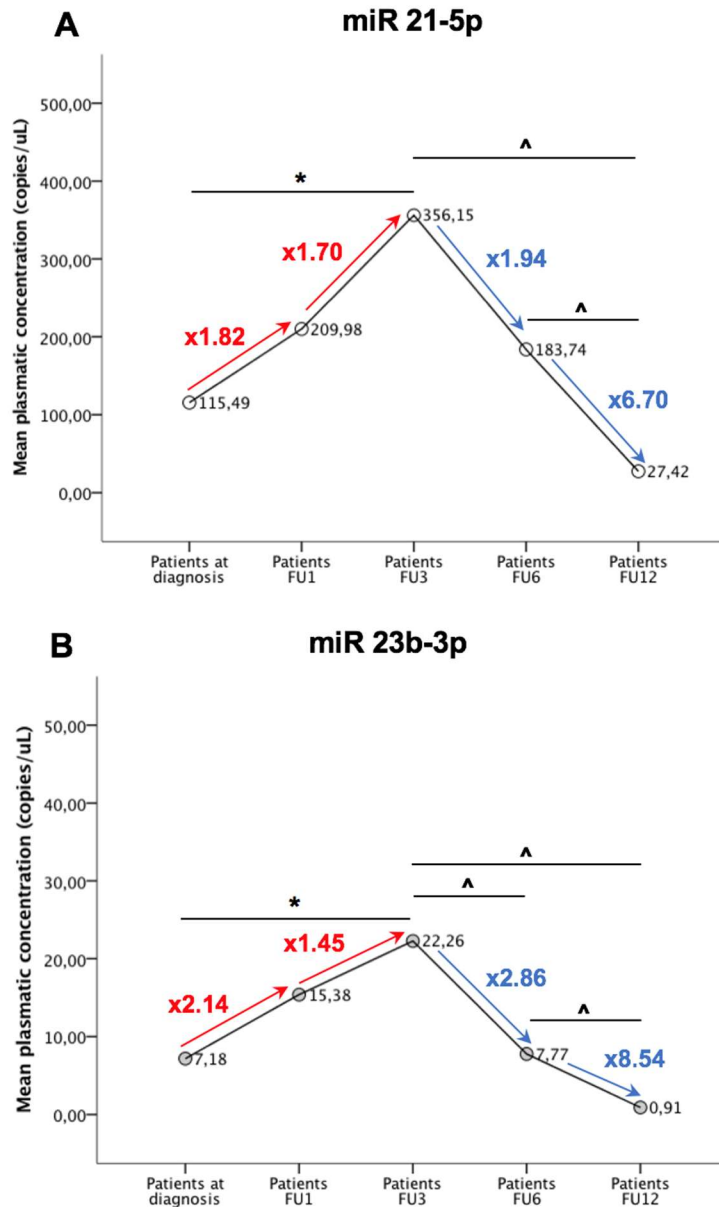


Figure 37 – Variations (%) of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations between 6 and 12 months after surgery.

4.5.2 Mean miRNAs' plasmatic concentrations course

The mean plasmatic concentrations of each miRNA showed the same longitudinal course in patients affected by GBM. The circulating levels progressively increased 1 and 3 months after surgery, then decreased 6 and 12 months after surgery (**Figure 38**).



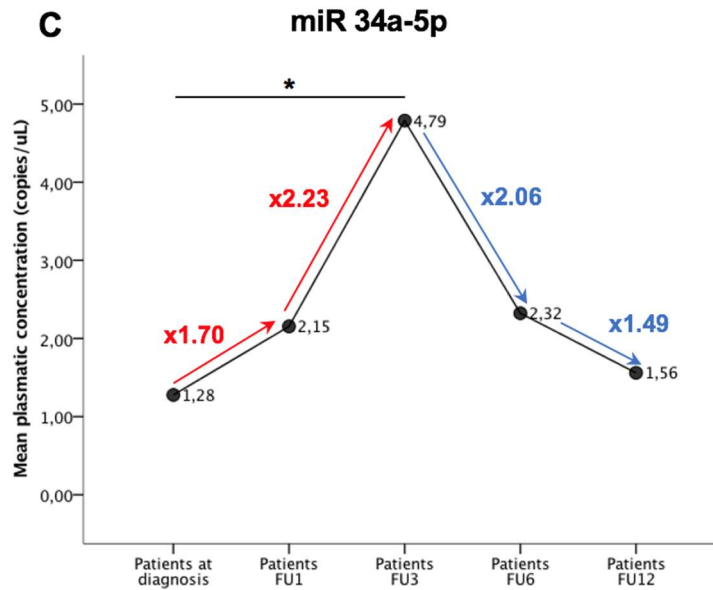


Figure 38 – Longitudinal course of mean plasmatic concentrations of miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C) (*: $p < 0.01$; ^: $p < 0.05$).

In particular, the mean plasmatic concentration of miR 21-5p increased by 1.82 folds between the diagnosis and 1 month after surgery (115.49 ± 132.57 vs. 209.98 ± 356.28 copies/ μ L; $p=0.251$) and by 1.70 folds between the first and the third month after surgery (209.98 ± 356.28 vs. 356.15 ± 303.89 copies/ μ L; $p=0.310$). Then it decreased by 1.94 folds between 3 and 6 months (356.15 ± 303.89 vs. 183.74 ± 141.87 copies/ μ L; $p=0.174$) and by 6.70 folds between 6 and 12 months after surgery (183.74 ± 141.87 vs. 27.42 ± 17.27 copies/ μ L; $p=0.033$).

Similarly, the mean miR 23b-3p levels showed a 2.14 fold increase between the diagnosis and 1 month after surgery (7.18 ± 10.15 vs. 15.38 ± 32.73 copies/ μ L; $p=0.345$) and a 1.45 fold increase between 1 and 3 months after surgery (15.38 ± 32.73 vs. 22.26 ± 19.04 copies/ μ L; $p=0.523$), followed by a 2.86 fold decrease between 3 and 6 months (22.26 ± 19.04 vs. 7.77 ± 6.16 copies/ μ L; $p=0.047$) and a 8.54 fold decrease between 6 and 12 months after surgery (7.77 ± 6.16 vs. 0.91 ± 0.42 copies/ μ L; $p=0.031$).

Eventually, the mean miR 34a-5p plasmatic concentrations were 1.70 folds higher 1 month after surgery than at the time of diagnosis (2.15 ± 1.71 vs. 1.28 ± 1.10 copies/ μ L; $p=0.083$), and 2.23 folds higher 3 months than 1 month after surgery (4.79 ± 5.17 vs. 2.15 ± 1.71 copies/ μ L; $p=0.074$), then 2.06 and 1.49 folds lower 6 months than 3 months after surgery (2.32 ± 2.43 vs. 4.79 ± 5.17 copies/ μ L; $p=0.246$) and 12 months than 6 months after surgery (1.56 ± 0.34 vs. 2.32 ± 2.43 copies/ μ L; $p=0.505$), respectively.

All the investigated miRNAs displayed significantly higher plasmatic concentrations 3 months after surgery compared to the time of diagnosis (miR 21-5p: 356.15 ± 303.89 vs. 115.49 ± 132.57 copies/ μ L, $p=0.004$; miR 23b-3p: 22.26 ± 19.04 vs. 7.18 ± 10.15 copies/ μ L, $p=0.008$; miR 34a-5p:

4.79 ± 5.17 vs. 1.28 ± 1.10 copies/μL, $p=0.004$). Furthermore, miR 21-5p and miR 23b-3p showed significantly lower circulating levels 12 months after surgery (miR 21-5p: 27.42 ± 17.27 copies/μL; miR 23b-3p: 0.91 ± 0.42 copies/μL) compared to 3 months (miR 21-5p: 356.15 ± 303.89 copies/μL, $p=0.037$; miR 23b-3p: 22.26 ± 19.04 copies/μL, $p=0.031$) and 6 months after surgery (miR 21-5p: 183.74 ± 141.87 copies/μL, $p=0.033$; miR 23b-3p: 7.77 ± 6.16 copies/μL, $p=0.031$).

4.6 Correlation analyses between miRNAs' plasmatic concentrations at diagnosis and clinical, neuroradiological and pathological features

There were no significant differences in plasmatic concentration of miRNAs and clinical, neuroradiological and histological features apart from preoperative corticosteroid therapy and ATRX status.

The mean circulating levels of miR 23b-3p (6.14 ± 10.01 vs. 18.10 ± 1.56 copies/μL, $p<0.001$) and miR 34a-5p (1.16 ± 1.07 vs. 2.56 ± 0.35 copies/μL, $p=0.021$) at diagnosis resulted significantly higher in patients who were not administered preoperative corticosteroids (**Table 1**).

| | miR 21-5p at diagnosis (mean, SD) | <i>p</i> value | miR 23b-3p at diagnosis (mean, SD) | <i>p</i> value | miR 34a-5p at diagnosis (mean, SD) | <i>p</i> value |
|------------------------|---|----------------|--|-------------------|--|----------------|
| Gender (n, %) | | | | | | |
| Males (14, 60.9) | 86.62 (92.39) | 0.200 | 4.38 (6.22) | 0.170 | 1.36 (1.25) | 0.638 |
| Females (9, 39.1) | 160.39 (175.30) | | 11.53 (13.62) | | 1.15 (0.86) | |
| mRS (n, %) | | | | | | |
| ≤ 2 (21, 91.3) | 109.67 (122.45) | 0.668 | 7.16 (10.40) | 0.682 | 1.29 (1.08) | 0.737 |
| > 2 (2, 8.7) | 218.45 (268.06) | | 10.40 (12.45) | | 1.75 (1.48) | |
| KPS (n, %) | | | | | | |
| 70-80 (5, 21.7) | 156.04 (163.53) | 0.576 | 8.58 (8.10) | 0.754 | 2.12 (1.39) | 0.182 |
| 90-100 (18, 78.3) | 108.82 (128.13) | | 7.12 (11.06) | | 1.10 (0.91) | |
| Steroids (n, %) | | | | | | |
| Yes (21, 91.3) | 95.72 (117.87) | 0.204 | 6.14 (10.01) | < 0.001 | 1.16 (1.07) | 0.021 |
| No (2, 8.7) | 323.00 (120.21) | | 18.10 (1.56) | | 2.56 (0.35) | |
| AEDs (n, %) | | | | | | |
| Yes (10, 43.5) | 70.18 (69.93) | 0.155 | 3.56 (4.56) | 0.137 | 1.29 (1.33) | 0.969 |
| No (13, 56.5) | 150.34 (159.76) | | 9.96 (12.41) | | 1.27 (0.94) | |

Table 1 – Comparison of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis with patients' clinical features (23 patients).

AEDs: anti-epileptic drugs; KPS: Karnofsky Performance Status; mRS: modified Ranking Score.

Furthermore, the plasmatic concentrations of all miRNAs were significantly higher at the time of diagnosis in tumours showing retained ATRX expression on IHC (miR 21-5p: 136.05 ± 137.44 vs.

8.15 ± 5.87 copies/μL, $p=0.001$; miR 23b-3p: 8.51 ± 10.73 vs. 0.36 ± 0.40 copies/μL, $p=0.004$; miR 34a-5p: 1.47 ± 1.11 vs. 0.37 ± 0.03 copies/μL, $p<0.001$) (**Table 2**).

| | miR 21-5p at diagnosis (mean, SD) | <i>p</i> value | miR 23b-3p at diagnosis (mean, SD) | <i>p</i> value | miR 34a-5p at diagnosis (mean, SD) | <i>p</i> value |
|---------------------------------------|--|----------------|---|----------------|---|-------------------|
| Preoperative MRI | | | | | | |
| Location (n, %) | | | | | | |
| Right (9, 39.1) | 105.18 (121.57) | 0.602 | 5.53 (6.65) | 0.443 | 1.19 (1.04) | 0.807 |
| Left (11, 47.8) | 140.62 (154.59) | | 9.87 (13.12) | | 1.43 (1.22) | |
| Midline (3, 13.1) | 54.27 (71.65) | | 2.24 (2.73) | | 1.00 (1.04) | |
| Necrosis (n, %) | | | | | | |
| Yes (20, 86.7) | 122.78 (143.06) | 0.271 | 8.06 (10.97) | 0.075 | 1.14 (0.92) | 0.486 |
| No (3, 13.3) | 62.43 (64.58) | | 2.57 (2.45) | | 2.16 (2.07) | |
| Tumour volume | | | | | | |
| ≤ 30 cc (10, 43.5) | 105.85 (128.68) | 0.765 | 5.22 (7.27) | 0.403 | 1.71 (1.41) | 0.096 |
| > 30 cc (13, 56.5) | 122.90 (140.22) | | 8.68 (11.99) | | 0.94 (0.65) | |
| Primary vs recurrent GBM | | | | | | |
| Primary (19, 82.6) | 119.38 (140.51) | 0.718 | 7.25 (10.81) | 0.932 | 1.29 (1.14) | 0.863 |
| Recurrent (4, 17.4) | 96.98 (99.27) | | 6.85 (7.40) | | 1.19 (0.99) | |
| MGMT methylation status (n, %) | | | | | | |
| Un-methylated (12, 52.2) | 134.44 (141.68) | 0.484 | 9.83 (12.23) | 0.190 | 1.14 (0.86) | 0.565 |
| Methylated (11, 47.8) | 94.81 (125.22) | | 4.28 (6.67) | | 1.42 (1.39) | |
| ATRX expression (n, %) | | | | | | |
| Lost (3, 13.0) | 8.15 (5.87) | 0.001 | 0.36 (0.40) | 0.004 | 0.37 (0.03) | < 0.001 |
| Retained (20, 87.0) | 136.05 (137.44) | | 8.51 (10.73) | | 1.47 (1.11) | |
| p53 expression (n, %) | | | | | | |
| Absent (2, 8.7) | 165.00 (196.58) | 0.574 | 10.30 (13.01) | 0.782 | 0.84 (1.08) | 0.817 |
| Intermediate (13, 56.5) | 82.20 (80.05) | | 5.55 (6.35) | | 1.35 (1.26) | |
| Diffuse (8, 34.8) | 128.99 (165.81) | | 8.16 (15.39) | | 1.15 (0.86) | |

Table 2 – Comparison of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis with histological and radiological tumour' features (23 patients).

ATRX: alpha thalassemia/mental retardation syndrome X-linked; MGMT: O-6-methylguanine-DNA-methyltransferase.

However, the correlation and multivariate analyses showed no significant results (see **Appendix – Tables 3-6**).

4.7 Correlation analyses between miRNAs' plasmatic concentrations over the follow up, extent of tumour resection and adjuvant therapies

Analysing the miRNAs' levels over the follow up, the only significant difference was found in miR 34a-5p plasmatic concentration 1 month after surgery that was significantly higher in patients who underwent near-total compared to subtotal resection (2.70 ± 1.93 vs. 1.13 ± 0.61 copies/ μ L, $p = 0.040$). The correlation and multivariate analyses did not show any significant results (see **Appendix – Tables 7-17**).

4.8 Mutual correlation analyses between miRNAs' plasmatic concentrations at different timing points

Correlation analyses of miRNA plasmatic concentrations at different timing points were detailed in **Tables 3-7**.

| | miR 21-5p at diagnosis | | | miR 23b-3p at diagnosis | | | miR 34a-5p at diagnosis | | |
|---------------------|---------------------------|----------------|------------------|----------------------------|----------------|------------------|----------------------------|----------------|--------------|
| | R | R ² | <i>p</i> | R | R ² | <i>p</i> | R | R ² | <i>p</i> |
| At diagnosis | | | | | | | | | |
| miR 21-5p | | | | 0.931 | 0.867 | <0.001 | 0.420 | 0.176 | 0.046 |
| miR 23b-3p | 0.931 | 0.867 | <0.001 | | | | 0.253 | 0.064 | 0.244 |
| miR 34a-5p | 0.420 | 0.176 | 0.046 | 0.253 | 0.064 | 0.244 | | | |
| FU1 | | | | | | | | | |
| miR 21-5p | 0.651 | 0.424 | 0.006 | 0.796 | 0.634 | <0.001 | 0.013 | 0.000 | 0.961 |
| miR 23b-3p | 0.672 | 0.452 | 0.004 | 0.819 | 0.671 | <0.001 | 0.018 | 0.000 | 0.947 |
| miR 34a-5p | 0.160 | 0.026 | 0.554 | 0.182 | 0.033 | 0.501 | 0.522 | 0.272 | 0.038 |
| FU3 | | | | | | | | | |
| miR 21-5p | -0.325 | 0.106 | 0.432 | -0.141 | 0.020 | 0.738 | -0.750 | 0.563 | 0.032 |
| miR 23b-3p | 0.227 | 0.052 | 0.589 | 0.386 | 0.149 | 0.345 | -0.424 | 0.180 | 0.295 |
| miR 34a-5p | -0.572 | 0.327 | 0.139 | -0.480 | 0.230 | 0.228 | -0.242 | 0.059 | 0.564 |
| FU6 | | | | | | | | | |
| miR 21-5p | -0.226 | 0.051 | 0.559 | -0.289 | 0.084 | 0.451 | -0.130 | 0.017 | 0.738 |
| miR 23b-3p | 0.029 | 0.001 | 0.940 | 0.041 | 0.002 | 0.917 | -0.282 | 0.080 | 0.462 |
| miR 34a-5p | -0.227 | 0.052 | 0.556 | -0.145 | 0.021 | 0.710 | -0.242 | 0.059 | 0.530 |
| FU12 | | | | | | | | | |
| miR 21-5p | 0.839 | 0.704 | 0.075 | 0.137 | 0.019 | 0.826 | 0.220 | 0.048 | 0.722 |
| miR 23b-3p | -0.007 | 0.000 | 0.991 | -0.001 | 0.000 | 0.999 | -0.354 | 0.125 | 0.559 |
| miR 34a-5p | -0.479 | 0.229 | 0.415 | -0.607 | 0.368 | 0.277 | 0.757 | 0.573 | 0.138 |

Table 3 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/ μ L) at diagnosis and miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/ μ L) at diagnosis, 1 (FU1), 3 (FU3), 6 (FU6) and 12 (FU12) months after surgery.

| | miR 21-5p FU1 | | | miR 23b-3p FU1 | | | miR 34a-5p FU1 | | |
|-------------|------------------|----------------|------------------|-------------------|----------------|------------------|-------------------|----------------|----------|
| | R | R ² | <i>p</i> | R | R ² | <i>p</i> | R | R ² | <i>p</i> |
| FU1 | | | | | | | | | |
| miR 21-5p | | | | 0.987 | 0.974 | <0.001 | 0.310 | 0.096 | 0.242 |
| miR 23b-3p | 0.987 | 0.974 | <0.001 | | | | 0.283 | 0.080 | 0.289 |
| miR 34a-5p | 0.310 | 0.096 | 0.242 | 0.283 | 0.080 | 0.289 | | | |
| FU3 | | | | | | | | | |
| miR 21-5p | 0.115 | 0.013 | 0.829 | 0.154 | 0.024 | 0.770 | -0.343 | 0.118 | 0.505 |
| miR 23b-3p | 0.675 | 0.456 | 0.141 | 0.707 | 0.500 | 0.116 | -0.214 | 0.046 | 0.683 |
| miR 34a-5p | -0.387 | 0.150 | 0.449 | -0.343 | 0.118 | 0.505 | -0.717 | 0.514 | 0.109 |
| FU6 | | | | | | | | | |
| miR 21-5p | -0.032 | 0.001 | 0.946 | 0.063 | 0.004 | 0.894 | -0.039 | 0.002 | 0.934 |
| miR 23b-3p | 0.420 | 0.176 | 0.349 | 0.540 | 0.292 | 0.211 | 0.298 | 0.089 | 0.517 |
| miR 34a-5p | -0.146 | 0.021 | 0.754 | -0.085 | 0.007 | 0.856 | -0.048 | 0.002 | 0.919 |
| FU12 | | | | | | | | | |
| miR 21-5p | 0.431 | 0.186 | 0.569 | -0.063 | 0.004 | 0.937 | -0.108 | 0.012 | 0.892 |
| miR 23b-3p | -0.659 | 0.434 | 0.341 | -0.940 | 0.884 | 0.060 | -0.113 | 0.013 | 0.887 |
| miR 34a-5p | -0.456 | 0.208 | 0.544 | -0.443 | 0.196 | 0.557 | 0.895 | 0.801 | 0.105 |

Table 4 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/ μ L) 1 month (FU1) after surgery and miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/ μ L) 1 (FU1), 3 (FU3), 6 (FU6) and 12 (FU12) months after surgery.

| | miR 21-5p FU3 | | | miR 23b-3p FU3 | | | miR 34a-5p FU3 | | |
|-------------|------------------|----------------|--------------|-------------------|----------------|--------------|-------------------|----------------|--------------|
| | R | R ² | <i>p</i> | R | R ² | <i>p</i> | R | R ² | <i>p</i> |
| FU3 | | | | | | | | | |
| miR 21-5p | | | | 0.760 | 0.578 | 0.028 | 0.689 | 0.475 | 0.059 |
| miR 23b-3p | 0.760 | 0.578 | 0.028 | | | | 0.385 | 0.148 | 0.346 |
| miR 34a-5p | 0.689 | 0.475 | 0.059 | 0.385 | 0.148 | 0.346 | | | |
| FU6 | | | | | | | | | |
| miR 21-5p | 0.585 | 0.342 | 0.300 | 0.717 | 0.134 | 0.849 | 0.940 | 0.884 | 0.018 |
| miR 23b-3p | 0.737 | 0.543 | 0.156 | 0.860 | 0.009 | 0.062 | 0.945 | 0.893 | 0.015 |
| miR 34a-5p | 0.983 | 0.966 | 0.003 | 0.849 | 0.924 | 0.069 | 0.448 | 0.201 | 0.171 |
| FU12 | | | | | | | | | |
| miR 21-5p | - | - | - | - | - | - | - | - | - |
| miR 23b-3p | - | - | - | - | - | - | - | - | - |
| miR 34a-5p | - | - | - | - | - | - | - | - | - |

Table 5 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/ μ L) 3 months (FU3) after surgery and miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/ μ L) 3 (FU3), 6 (FU6) and 12 (FU12) months after surgery.

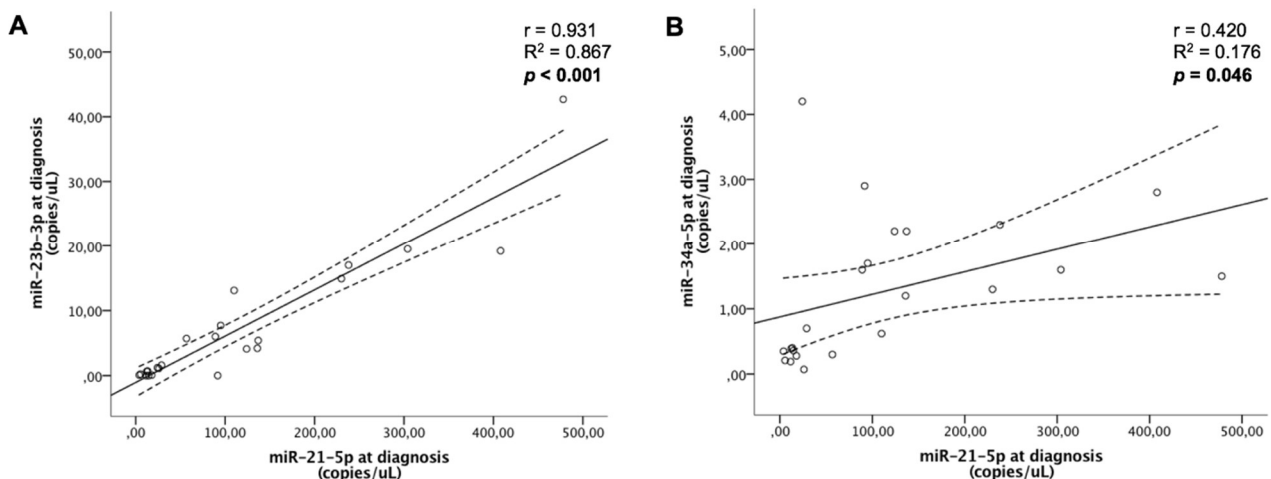
| | miR 21-5p FU6 | | | miR 23b-3p FU6 | | | miR 34a-5p FU6 | | |
|-------------|------------------|----------------|--------------|-------------------|----------------|--------------|-------------------|----------------|----------|
| | R | R ² | <i>p</i> | R | R ² | <i>p</i> | R | R ² | <i>p</i> |
| FU6 | | | | | | | | | |
| miR 21-5p | | | | 0.847 | 0.717 | 0.004 | 0.400 | 0.160 | 0.287 |
| miR 23b-3p | 0.847 | 0.717 | 0.004 | | | | 0.349 | 0.122 | 0.358 |
| miR 34a-5p | 0.400 | 0.160 | 0.287 | 0.349 | 0.122 | 0.358 | | | |
| FU12 | | | | | | | | | |
| miR 21-5p | 0.968 | 0.937 | 0.032 | 0.830 | 0.689 | 0.170 | 0.259 | 0.067 | 0.741 |
| miR 23b-3p | 0.559 | 0.312 | 0.441 | 0.634 | 0.402 | 0.366 | -0.688 | 0.473 | 0.312 |
| miR 34a-5p | 0.200 | 0.040 | 0.800 | -0.001 | 0.000 | 0.999 | -0.729 | 0.531 | 0.271 |

Table 6 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/μL) 6 months (FU6) after surgery and miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/μL) 6 (FU6) and 12 (FU12) months after surgery.

| | miR 21-5p FU12 | | | miR 23b-3p FU12 | | | miR 34a-5p FU12 | | |
|-------------|-------------------|----------------|----------|--------------------|----------------|----------|--------------------|----------------|----------|
| | R | R ² | <i>p</i> | R | R ² | <i>p</i> | R | R ² | <i>p</i> |
| FU12 | | | | | | | | | |
| miR 21-5p | | | | 0.447 | 0.200 | 0.451 | -0.164 | 0.027 | 0.792 |
| miR 23b-3p | 0.447 | 0.200 | 0.451 | | | | -0.027 | 0.001 | 0.965 |
| miR 34a-5p | -0.164 | 0.027 | 0.792 | -0.027 | 0.001 | 0.965 | | | |

Table 7 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations (copies/μL) 12 months (FU12) after surgery.

miR 21-5p levels at diagnosis were significantly correlated to miR 23b-3p ($r=0.931$, $R^2=0.867$; $p<0.001$) and miR 34a-5p levels at diagnosis ($r = 0.420$, $R^2 = 0.176$; $p=0.046$), but also to miR 21-5p levels 1 month after surgery ($r=0.651$, $R^2=0.424$; $p=0.006$) and miR 23b-3p levels 1 month after surgery ($r=0.672$, $R^2=0.452$; $p=0.004$). Similarly, miR 21-5p levels 1 month after surgery were significantly correlated to miR 23b-3p levels 1 month after surgery ($r=0.987$, $R^2=0.974$; $p<0.001$) (**Figure 39**).



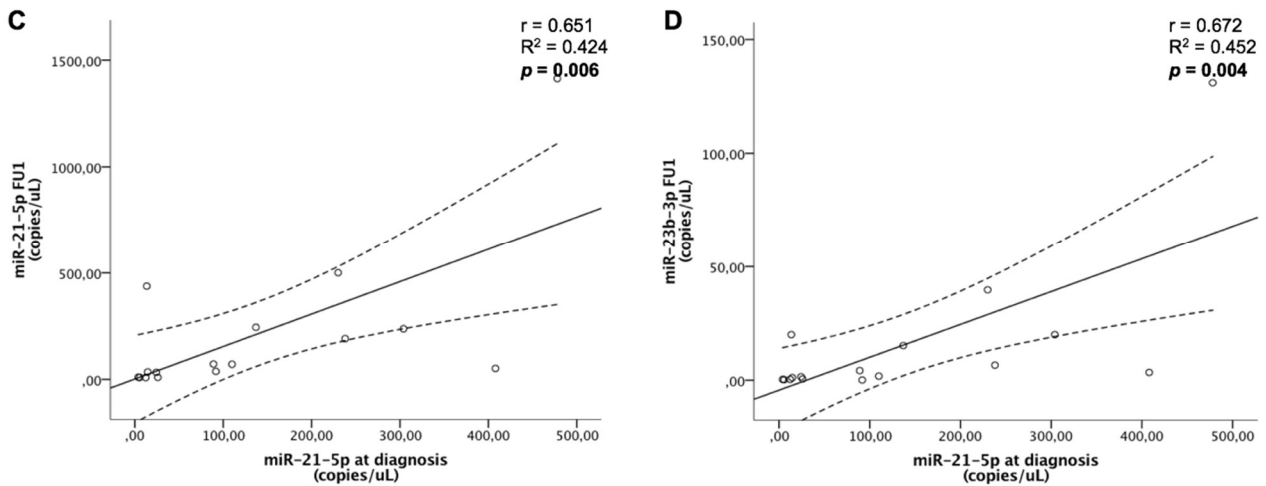
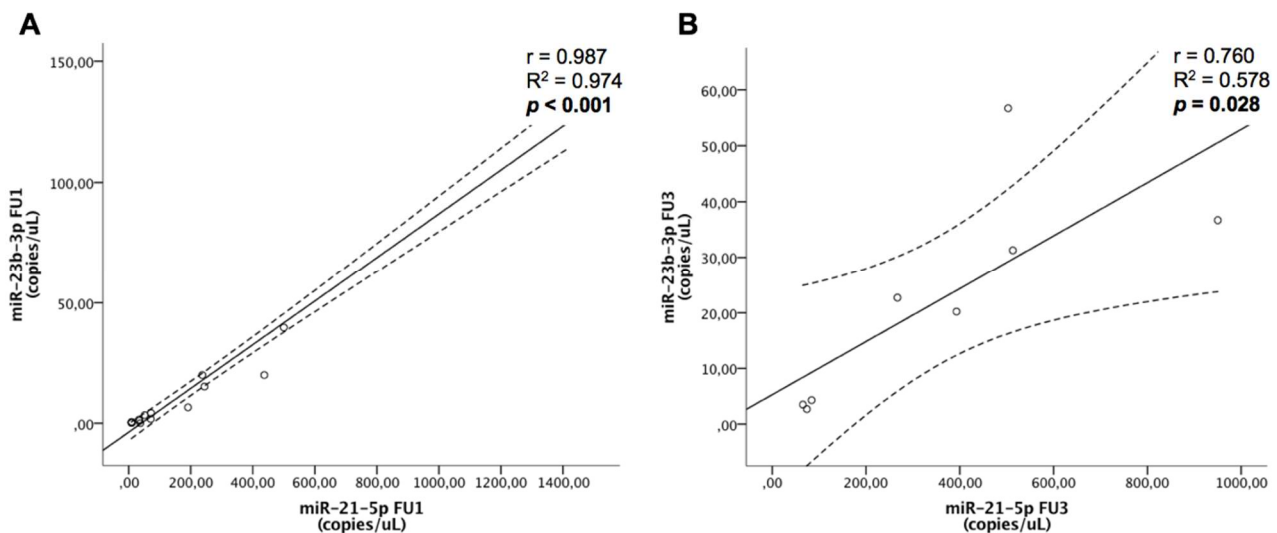


Figure 39 – Scatter plots correlating miR 21-5p plasmatic concentrations at diagnosis with miR 23b-3p plasmatic concentrations at diagnosis (A), miR 34a-5p plasmatic concentrations at diagnosis (B), miR 21-5p plasmatic concentrations 1 month after surgery (C) and miR 23b-3p plasmatic concentrations at 1 month after surgery (D).

miR 21-5p levels 3 months after surgery were significantly correlated to miR 23b-3p levels 3 months after surgery ($r=0.760$, $R^2=0.578$; $p=0.028$) and miR 34a-5p 6 months after surgery ($r=0.983$, $R^2=0.966$; $p=0.003$). miR 21-5p levels 6 months after surgery were related to miR 23b-3p levels 6 months after surgery ($r=0.847$, $R^2=0.717$; $p=0.004$) and to miR 21-5p levels 12 months after surgery ($r=0.968$, $R^2=0.937$; $p=0.032$) (Figure 40).



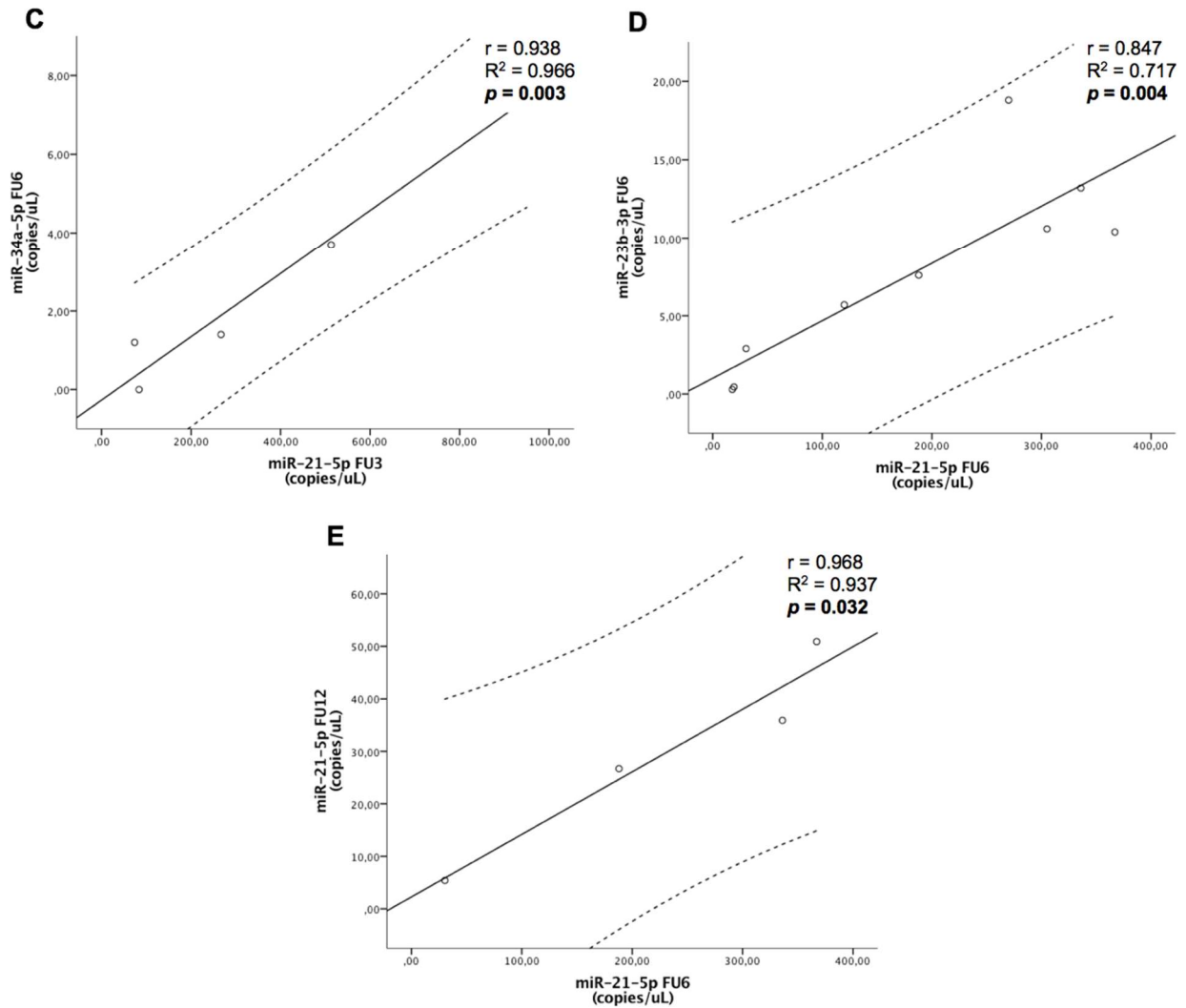


Figure 40 – Scatter plots showing mutual correlation of miR 21-5p plasmatic concentrations at different timing points.

miR 23b-3p levels at diagnosis were significantly correlated to miR 21-5p levels at diagnosis ($r=0.931$, $R^2=0.867$; $p<0.001$), miR 21-5p levels 1 month after surgery ($r=0.796$, $R^2=0.634$; $p<0.001$) and miR 23b-3p 1 month after surgery ($r=0.819$, $R^2=0.671$; $p<0.001$). Similarly, miR 23b-3p levels 1 month after surgery were significantly correlated to miR 21-5p levels 1 month after surgery ($r=0.987$, $R^2=0.974$; $p<0.001$) (**Figure 41**).

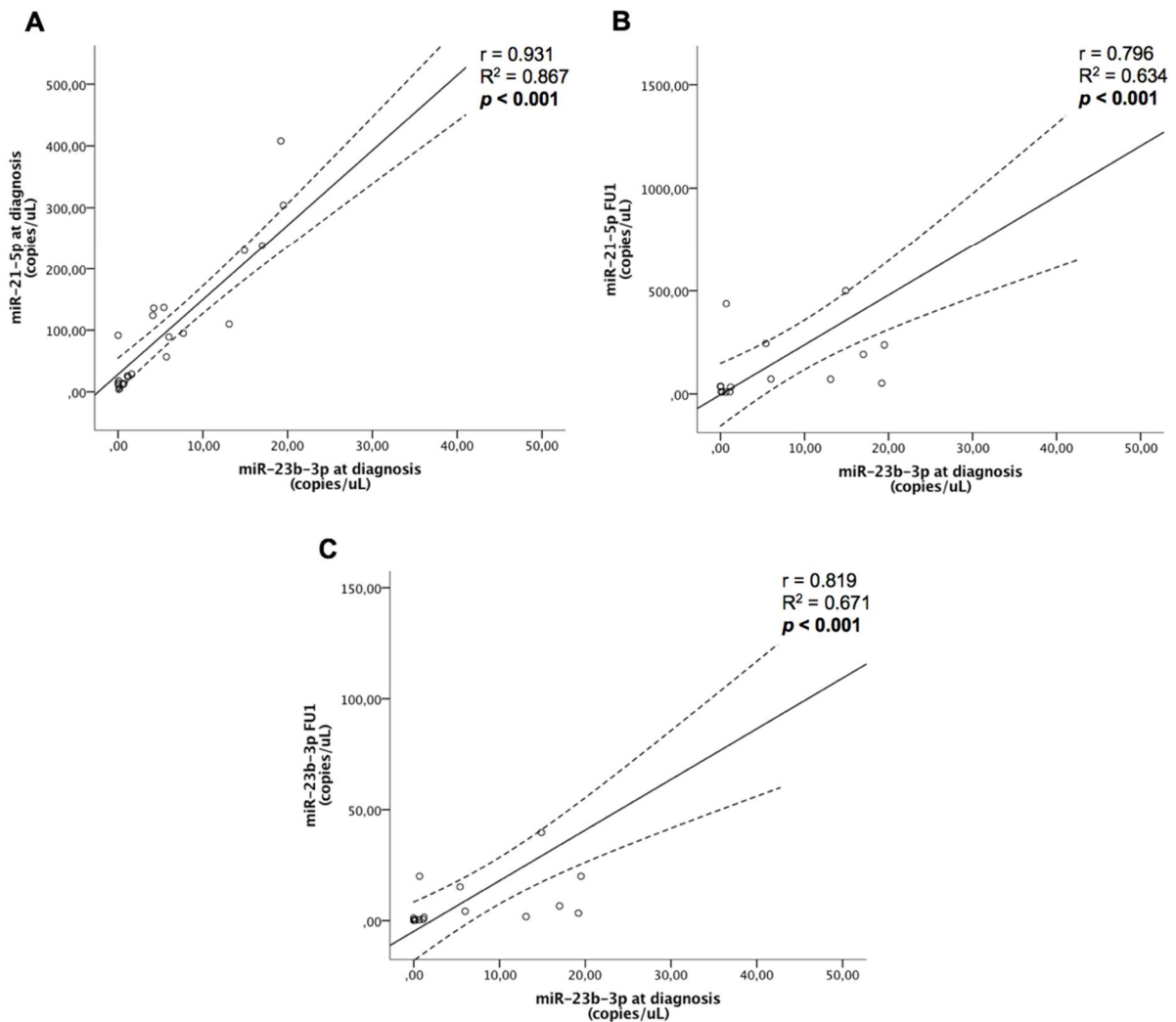


Figure 41 – Scatter plots displaying mutual correlation of miR 23b-3p plasmatic concentrations at different timing points.

miR 34a-5p levels at diagnosis were significantly correlated to miR 21-5p levels at diagnosis ($r=0.420$, $R^2=0.176$; $p=0.046$), miR 34a-5p levels 1 month after surgery ($r=0.522$, $R^2=0.272$; $p=0.038$) and miR 21-5p levels 3 months after surgery ($r=-0.750$, $R^2=0.563$; $p=0.032$). Furthermore, miR 34a-5p levels 3 months after surgery were significantly related to miR 21-5p ($r=0.940$, $R^2=0.884$; $p=0.018$) and miR 23b-3p ($r=0.945$, $R^2=0.893$; $p=0.015$) levels 6 months after surgery (**Figure 42**).

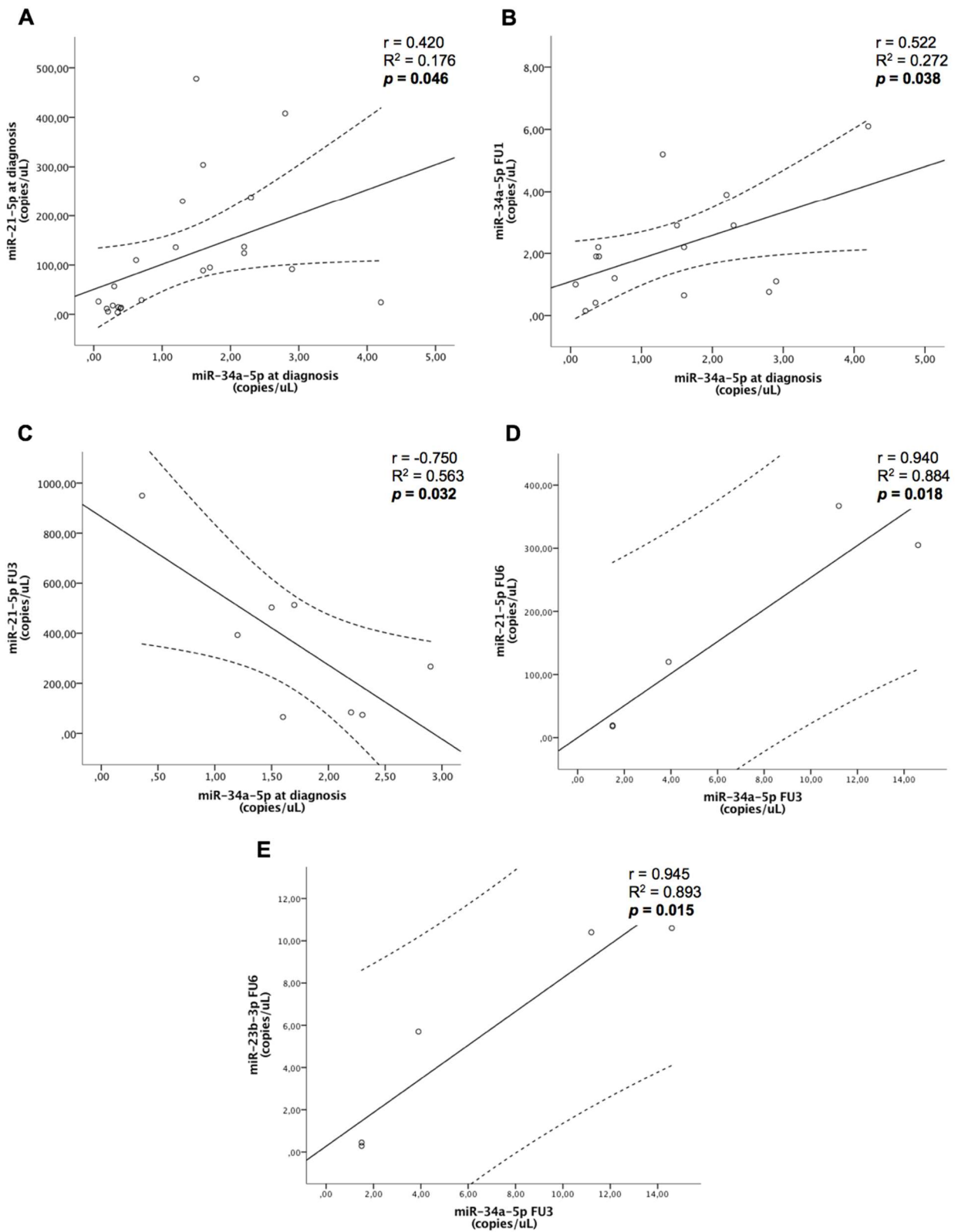


Figure 42 – Scatter plots showing mutual correlation of miR 34a-5p plasmatic concentrations at different timing points.

The multivariate linear regression analyses (see **Appendix – Tables 18-22**) showed that miR 23b-3p (OR 11.50, 95% C.I. 9.53 – 13.47, $p < 0.001$) and miR 34a-5p (OR 23.80, 95% C.I. 5.57 – 42.02, $p = 0.013$) levels at diagnosis were predictors of miR 21-5p levels at diagnosis such as miR 21-5p levels at diagnosis independently predicted miR 23b-3p (OR 0.08, 95% C.I. 0.06 – 0.09, $p < 0.001$) and miR 34a-5p (OR 0.01, 95% C.I. 0.00 – 0.02, $p = 0.013$) levels at diagnosis. Furthermore, miR 23b-3p plasmatic concentrations 1 month after surgery predicted miR 21-5p levels 1 month after surgery (OR 10.83, 95% C.I. 8.34 – 13.32, $p < 0.001$) and vice versa (OR 0.08, 95% C.I. 0.06 – 0.10, $p < 0.001$). miR 21-5p levels 6 months after surgery predicted miR 23b-3p levels 6 months after surgery (OR 0.02, 95% C.I. 0.01 – 0.03, $p = 0.022$) (**Table 8**).

| | OR | 95% CI | <i>p</i> value |
|--------------------------------|--------------|---------------------|------------------|
| miR 21-5p at diagnosis | | | |
| miR 23b-3p at diagnosis | 11.50 | 9.53 – 13.47 | <0.001 |
| miR 34a-5p at diagnosis | 23.80 | 5.57 – 42.02 | 0.013 |
| | | | |
| miR 23b-3p at diagnosis | | | |
| miR 21-5p at diagnosis | 0.08 | 0.06 – 0.09 | <0.001 |
| | | | |
| miR 34a-5p at diagnosis | | | |
| miR 21-5p at diagnosis | 0.01 | 0.00 – 0.02 | 0.013 |
| | | | |
| miR 21-5p at FU1 | | | |
| miR 23b-3p at FU1 | 10.83 | 8.34 – 13.32 | <0.001 |
| | | | |
| miR 23b-3p at FU1 | | | |
| miR 21-5p at FU1 | 0.08 | 0.06 – 0.10 | <0.001 |
| | | | |
| miR 23b-3p at FU6 | | | |
| miR 21-5p at FU6 | 0.02 | 0.01 – 0.03 | 0.022 |

Table 8 – Multivariate linear regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations.

4.9 miRNA network analysis

According to the miRNet analysis (see **Appendix – Table 23**), we identified 5 genes (*E2F1*, *TNPO1*, *MTAP*, *CPEB3*, *NUFIP2*) regulated by miR 21-5p, miR-23b-3p and miR 34a-5p. Among them, *E2F1* was deregulated in gliomas (**Figure 43**).

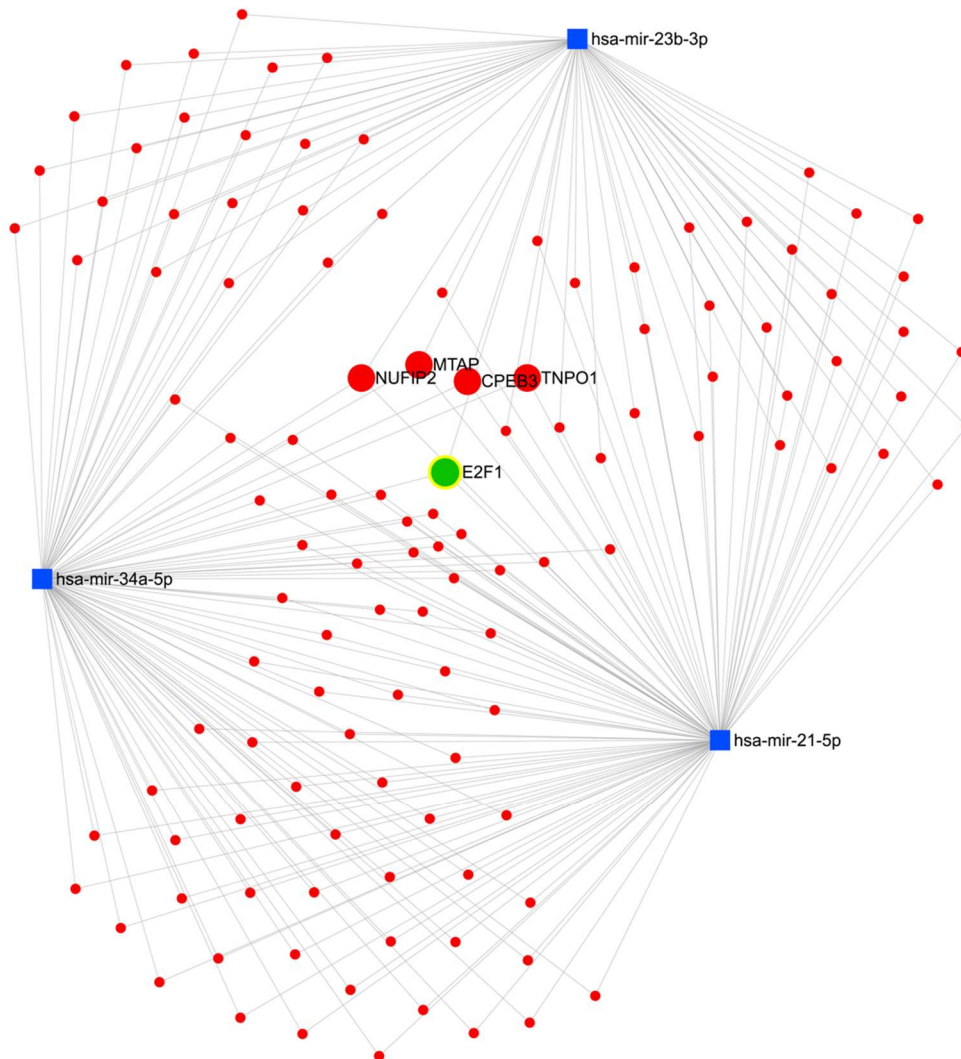


Figure 43 – Network visualization of miR 21-5p, miR-23b-3p and miR 34a-5p (blue squares) and related genes (red dots), highlighting genes deregulated in gliomas (green dots; miRNet version 2.0).

Focusing on miR 21-5p and miR 23b-3p, the network analysis (see **Appendix – Table 24**) identified 36 genes 4 of which (*E2F1*, *PTEN*, *PIK3R1*, *RB1*) were related to gliomas (**Figure 44**).

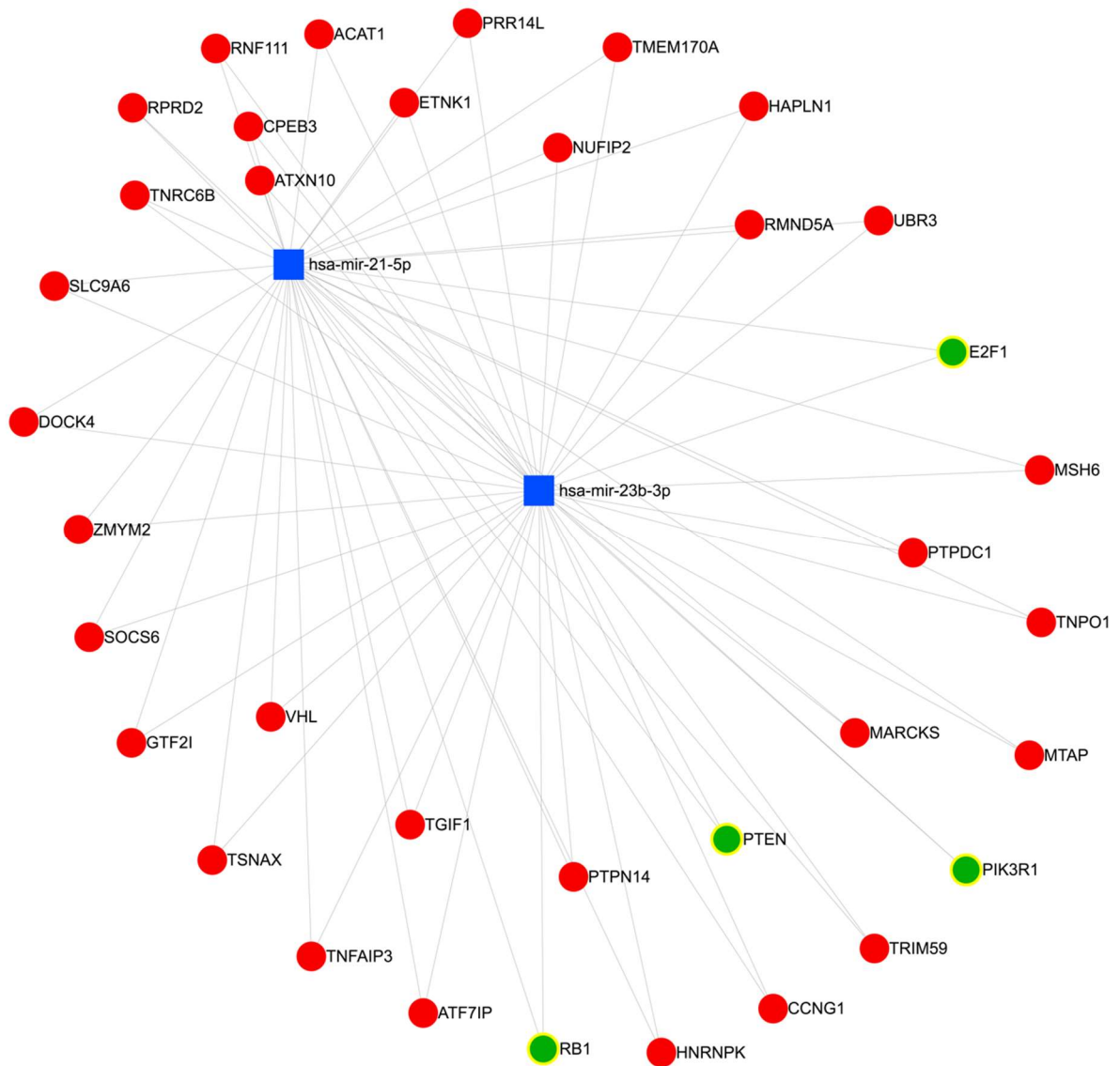


Figure 44 – Network visualization of miR 21-5p and miR 23b-3p (blue squares) and related genes (red dots), highlighting genes deregulated in gliomas (green dots; miRNet version 2.0).

Analysing miR 21-5p and miR 34a-5p, the network analysis (see **Appendix – Table 25**) identified 67 genes 4 of which (*E2F1*, *E2F3*, *CDK6*, *IGF1R*) were involved in the pathogenesis of gliomas (**Figure 45**).

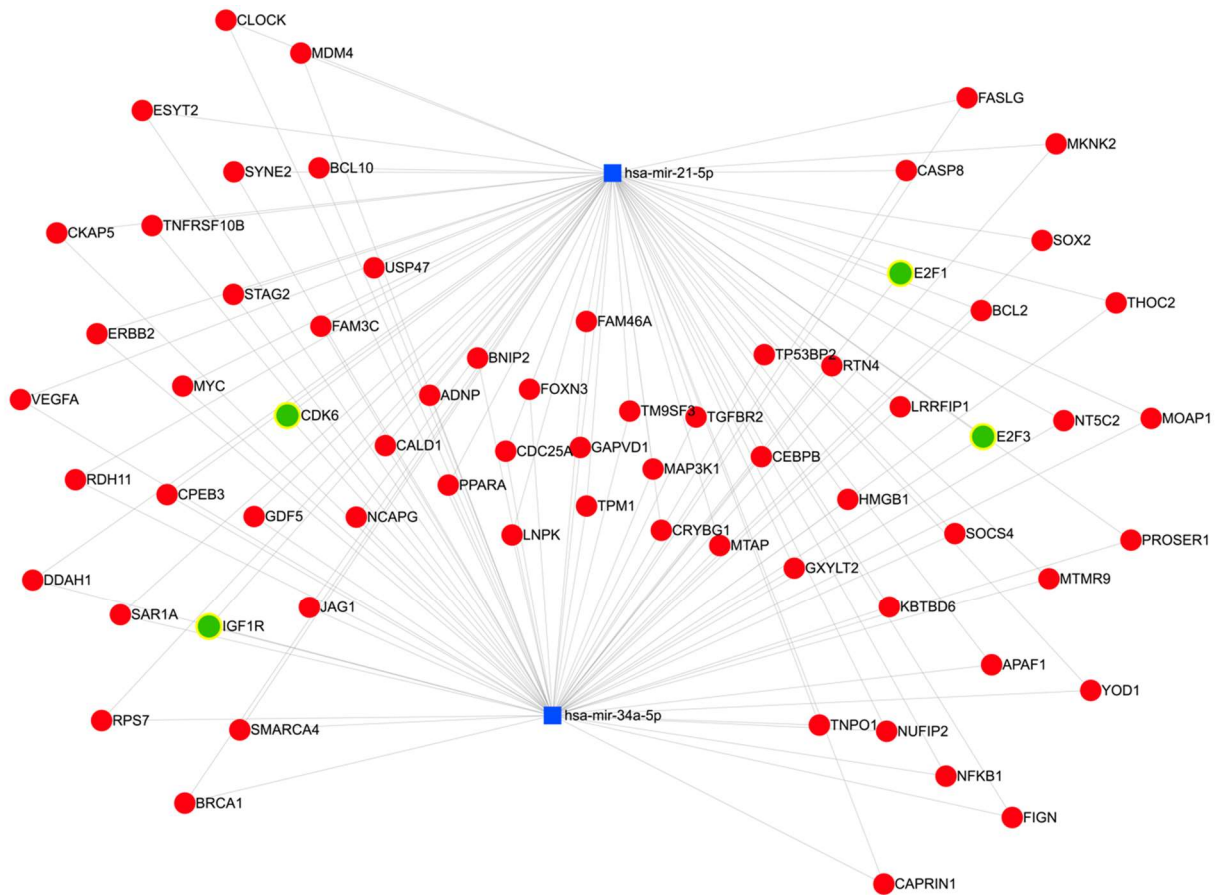


Figure 45 – Network visualization of miR 21-5p and miR 34a-5p (blue squares) and related genes (red dots), highlighting genes deregulated in gliomas (green dots; miRNet version 2.0).

4.10 Overall survival analysis

The mean length of follow-up was 11.9 months (± 9.4 ; median 8.0, IQR: 3.0-17.0). The mean OS was 12.4 months (95% C.I. 8.4 – 16.4) with 39.1% and 21.7% of predicted OS at 1 and 2 years, respectively (**Figure 46**).

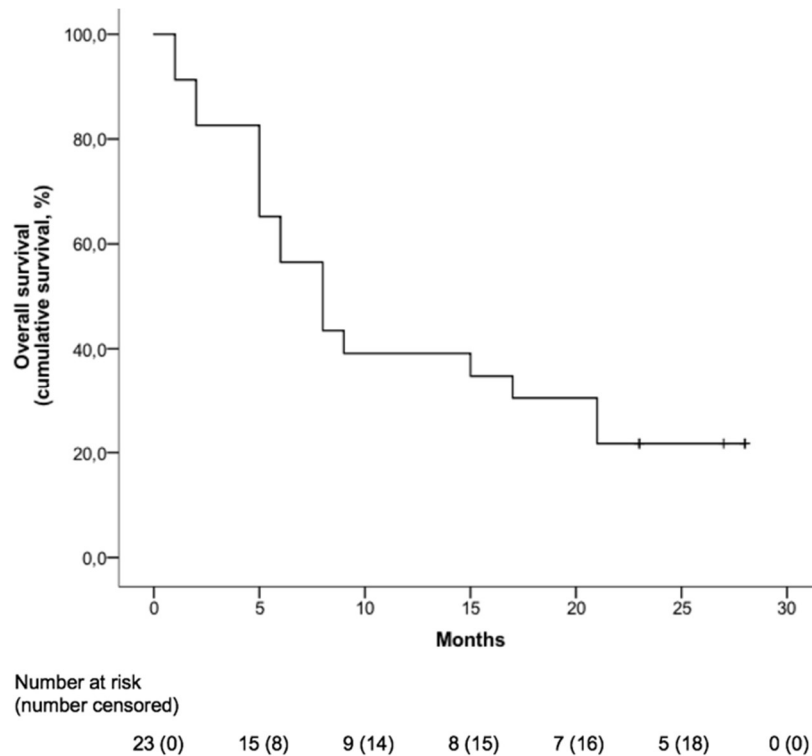


Figure 46 – Overall survival; **B**) Recurrence free survival.

The Cox regression analysis showed no relationships between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at different timing points and OS or RFS (see **Appendix – Tables 26-27**).

4.10.1 Overall survival according to miRNAs' variations at different timing points

After splitting the population in two groups of patients experiencing increase or stability/decrease of miRNAs circulating levels over the follow up, the mean OS were compared (**Table 9**).

| | | Mean OS (95% C.I.) | pOS at 1 year (%) | pOS at 2 years (%) | p value |
|----------------------------|--------------------|---------------------------|-------------------|--------------------|--------------|
| miR 21-5p diagnosis-FU1 | Increase | 14.6 (7.5 – 21.8) | 50.0 | 25.0 | 0.763 |
| | Stability/Decrease | 12.1 (5.3 – 18.9) | 37.5 | 25.0 | |
| miR 23b-3p diagnosis-FU1 | Increase | 15.5 (7.8 – 23.2) | 50.0 | 37.5 | 0.353 |
| | Stability/Decrease | 10.6 (5.5 – 15.8) | 25.0 | 12.5 | |
| miR 34a-5p diagnosis-FU1 | Increase | 11.0 (5.8 – 16.2) | 25.0 | 16.7 | 0.171 |
| | Stability/Decrease | 20.5 (11.3 – 29.7) | 75.0 | 50.0 | |
| miR 21-5p FU1-FU3 | | | | | |
| | Increase | 24.5 (19.7 – 29.4) | 50.0 | 50.0 | 0.049 |
| | Stability/Decrease | 8.5 (4.1 – 12.9) | 25.0 | 0.0 | |
| miR 23b-3p FU1-FU3 | | | | | |
| | Increase | 24.5 (19.7 – 29.4) | 50.0 | 50.0 | 0.049 |
| | Stability/Decrease | 8.5 (4.1 – 12.9) | 25.0 | 0.0 | |
| miR 34a-5p FU1-FU3 | | | | | |
| | Increase | 24.5 (19.7 – 29.4) | 50.0 | 50.0 | 0.049 |
| | Stability/Decrease | 8.5 (4.1 – 12.9) | 25.0 | 0.0 | |
| miR 21-5p FU3-FU6 | | | | | |
| | Increase | - | - | - | 0.156 |
| | Stability/Decrease | - | - | - | |
| miR 23b-3p FU3-FU6 | | | | | |
| | Increase | - | - | - | - |
| | Stability/Decrease | 16.2 (9.6 – 22.8) | 60.0 | 20.0 | |
| miR 34a-5p FU3-FU6 | | | | | |
| | Increase | - | - | - | - |
| | Stability/Decrease | 16.2 (9.6 – 22.8) | 60.0 | 20.0 | |
| miR 21-5p FU6-FU12 | | | | | |
| | Increase | - | - | - | - |
| | Stability/Decrease | 23.5 (18.9 – 28.1) | 100.0 | 50.0 | |
| miR 23b-3p FU6-FU12 | | | | | |
| | Increase | - | - | - | - |
| | Stability/Decrease | 23.5 (18.9 – 28.1) | 100.0 | 50.0 | |
| miR 34a-5p FU6-FU12 | | | | | |
| | Increase | - | - | - | 0.090 |
| | Stability/Decrease | - | - | - | |

Table 9 – Mean and predicted OS according to variations (increase or stability/decrease) of circulating levels of miR 21-5p, miR 23b-3p and miR 34a-5p at different timing points.

pOS: predicted overall survival.

The mean OS was significantly longer in patients presenting increased levels of miR 21-5p, miR 23b-3p and miR 34a-5p (24.5 months, 95% C.I. 19.7 – 29.4 vs. 8.5 months, 95% C.I. 4.1 – 12.9; $p=0.049$) between 1 and 3 months after surgery (**Figures 47-50**).

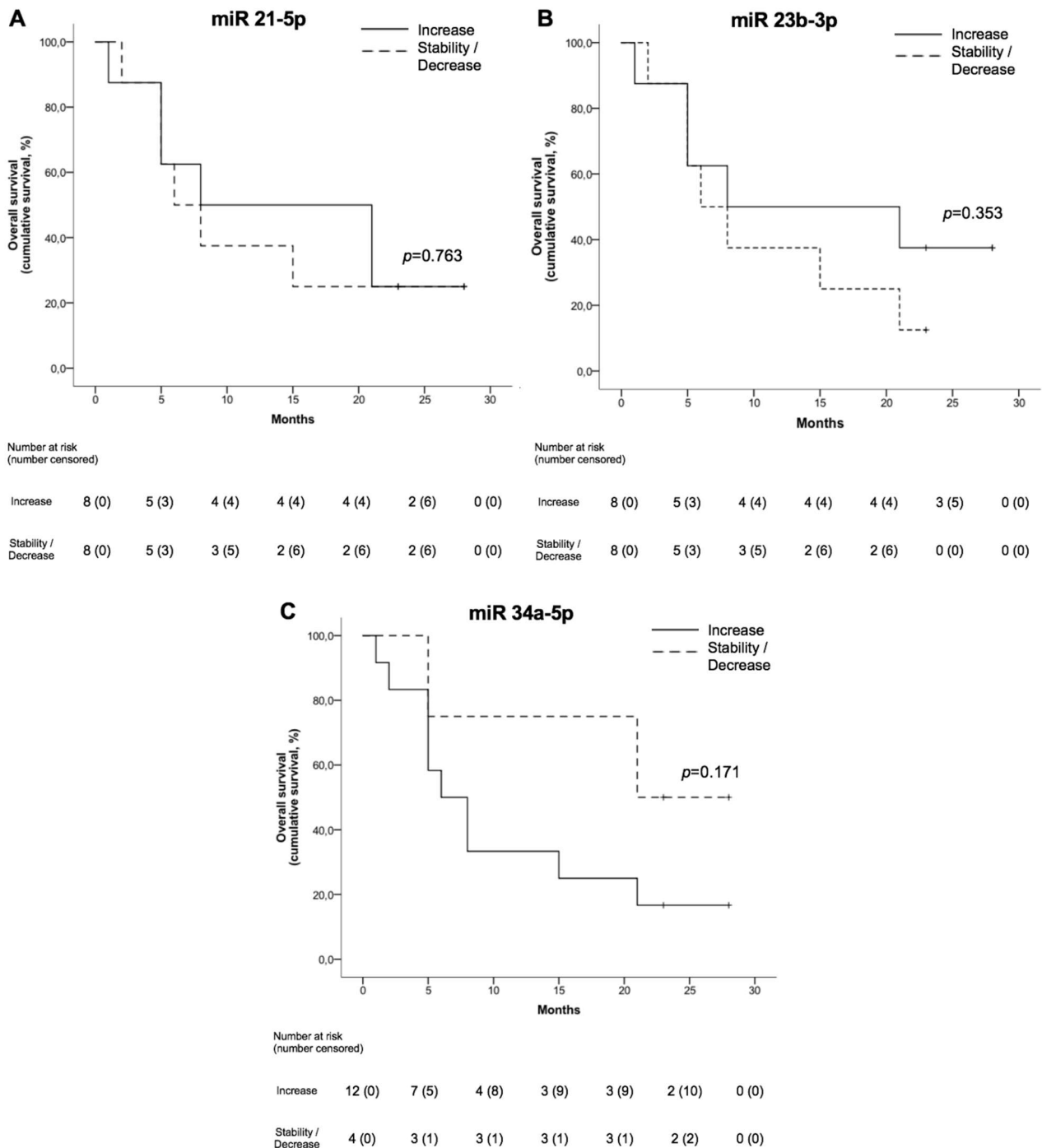


Figure 47 – Overall survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between the diagnosis and 1 month after surgery.

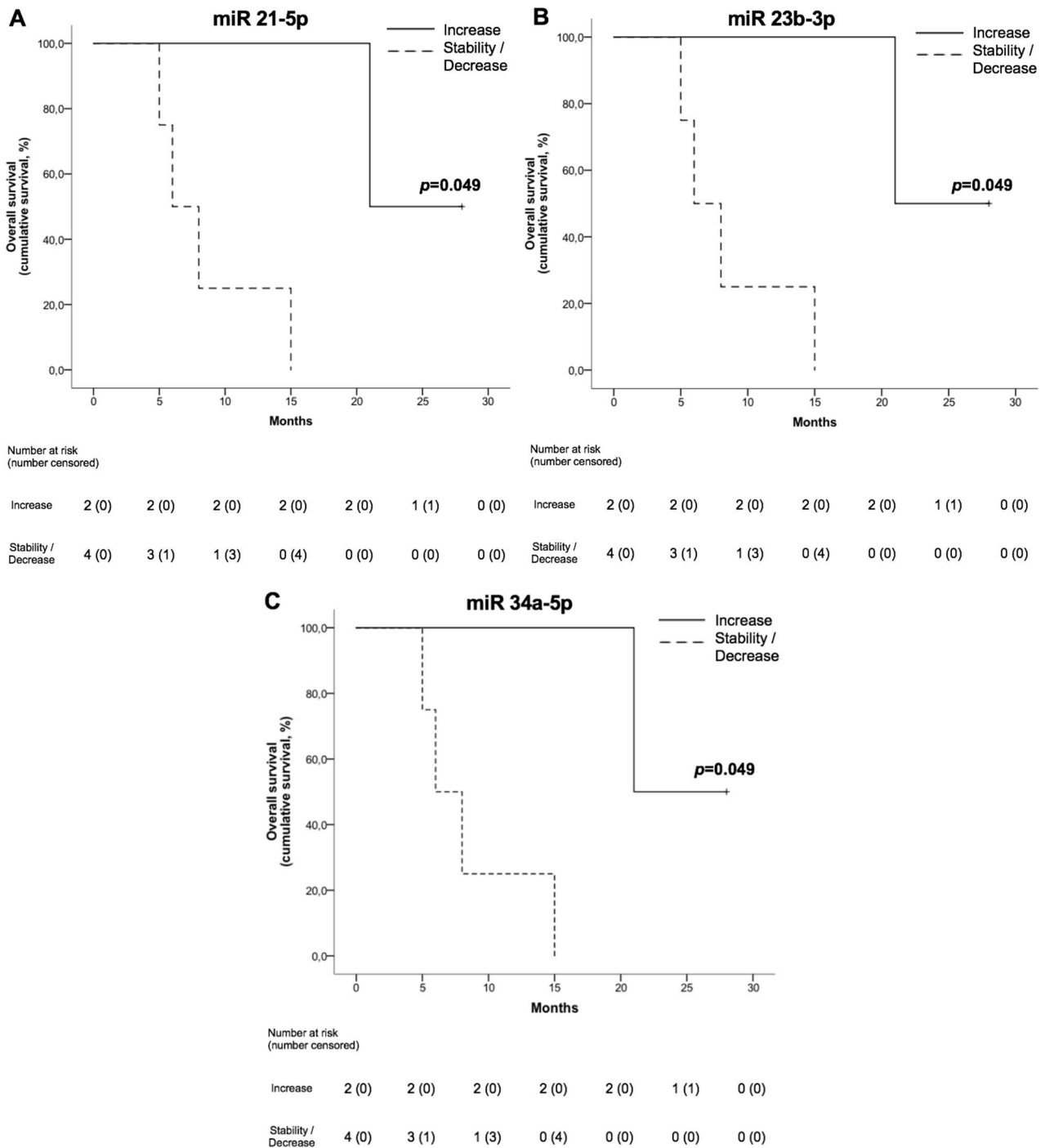


Figure 48 – Overall survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between 1 and 3 months after surgery.

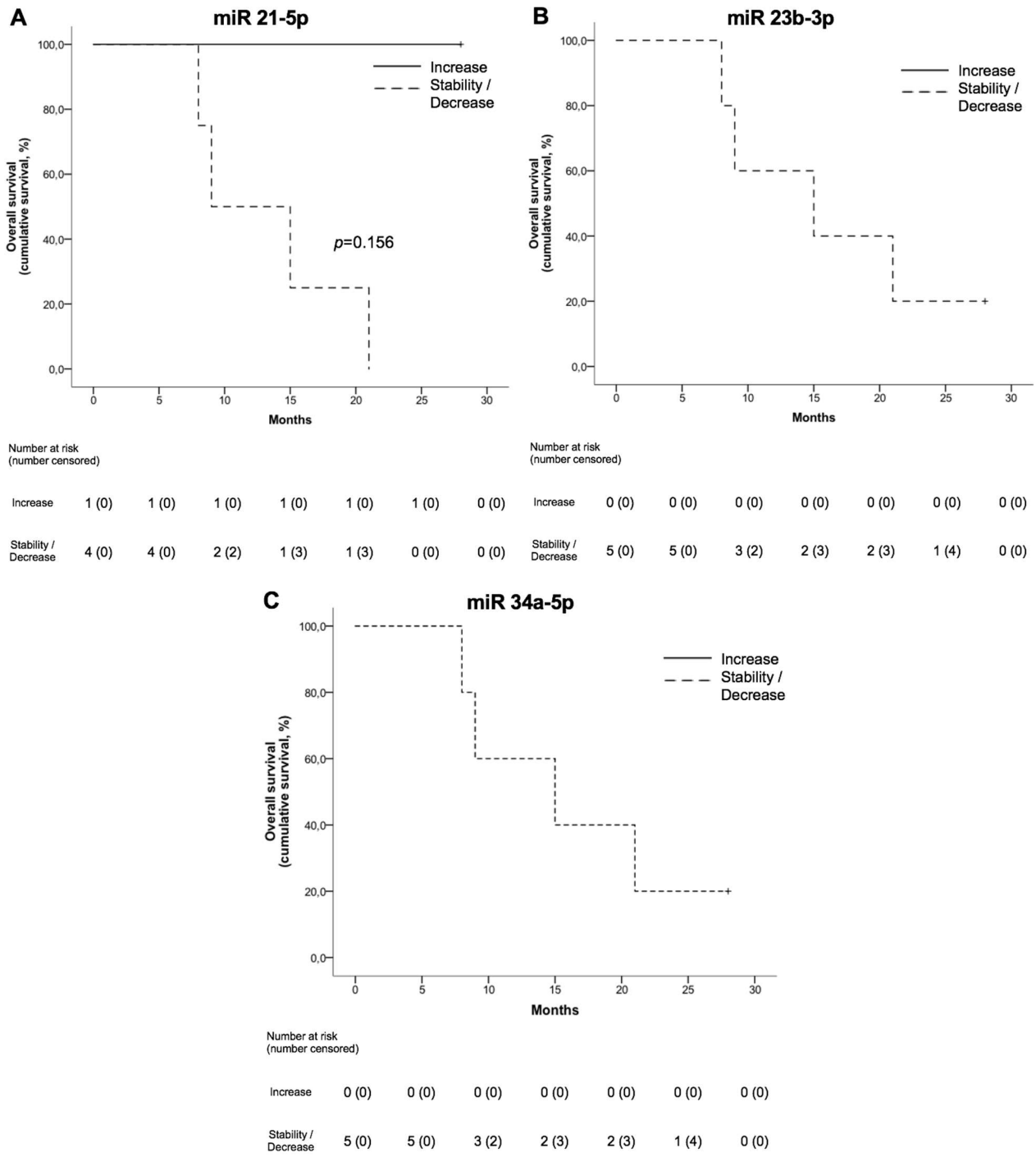


Figure 49 – Overall survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C) between 3 and 6 months after surgery.

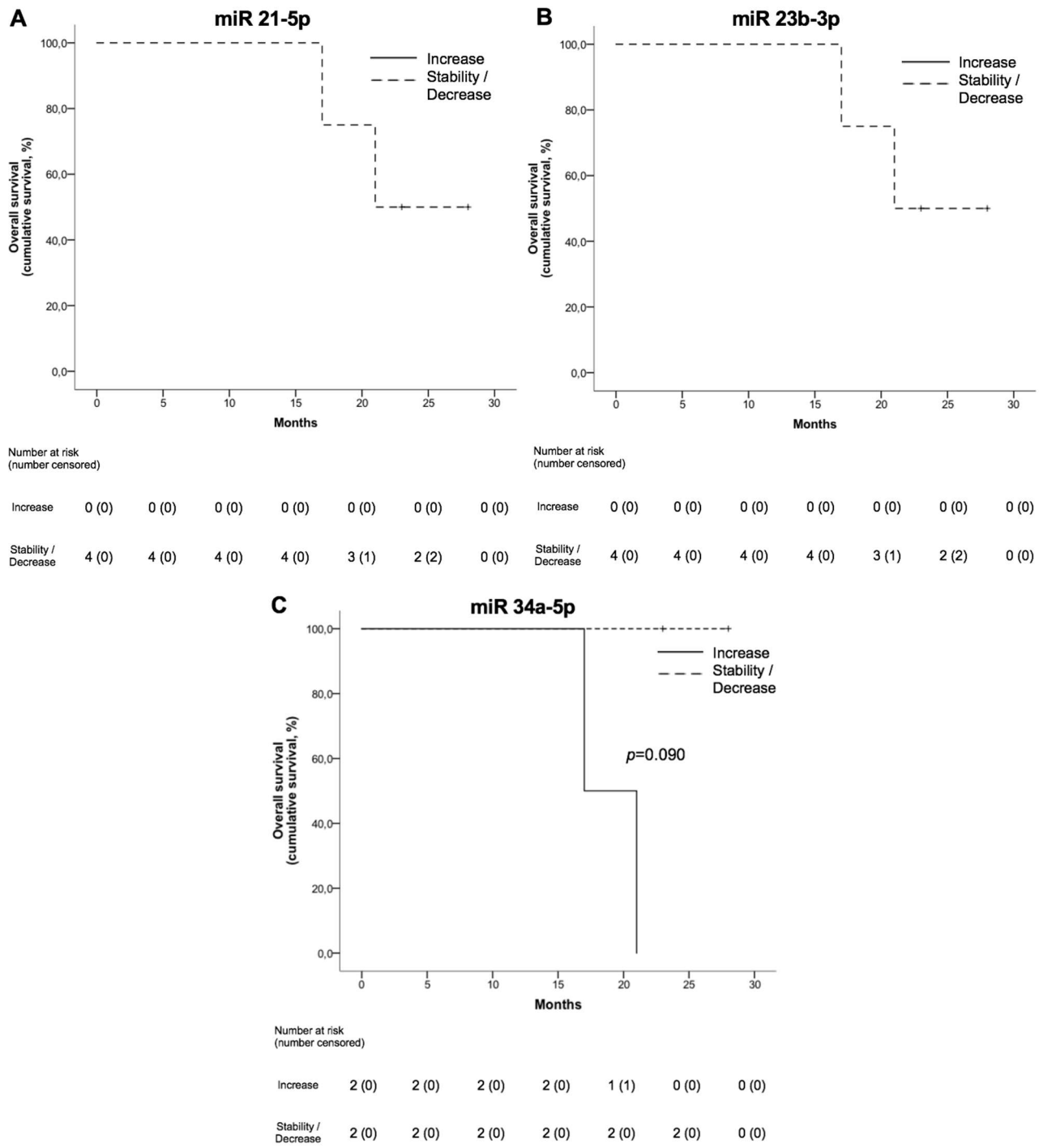


Figure 50 – Overall survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C) between 6 and 12 months after surgery.

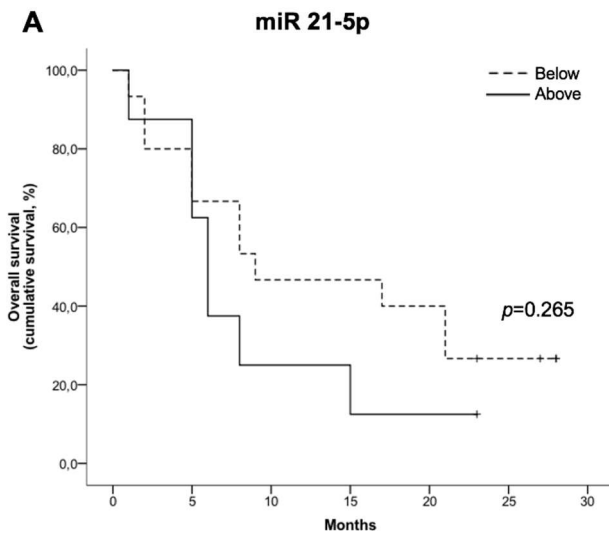
4.10.2 Overall survival according to mean miRNAs' plasmatic concentrations

After splitting the population above and below the mean values of circulating miRNAs at different timing points (**Table 10**). The mean OS was significantly shorter in patients with miR 34a-5p levels below the mean 3 months after surgery (8.2 months, 95% C.I. 5.2 – 11.1 vs 24.5 months, 95% C.I. 19.6 – 29.4; $p=0.032$) and in those with miR 21-5p levels below the mean 6 months after surgery (13.3 months, 95% C.I. 7.4 – 19.2 vs 24.4 months, 95% C.I. 20.4 – 28.4; $p=0.024$) (**Figures 51-55**).

| | | Mean OS (95% C.I.) | pOS at 1 year (%) | pOS at 2 years (%) | p value |
|--------------------------|----------------|---------------------------|-------------------|--------------------|--------------|
| miR 21-5p at diagnosis | Below the mean | 14.1 (8.8 – 19.3) | 46.7 | 26.7 | 0.265 |
| | Above the mean | 8.6 (4.1 – 13.2) | 25.0 | 12.5 | |
| miR 23b-3p at diagnosis | Below the mean | 13.4 (4.3 – 14.3) | 43.8 | 25.0 | 0.456 |
| | Above the mean | 9.3 (8.4 – 16.4) | 28.6 | 14.3 | |
| miR 34a-5p at diagnosis | Below the mean | 10.3 (5.5 – 15.1) | 33.3 | 8.3 | 0.213 |
| | Above the mean | 14.6 (8.4 – 20.9) | 45.5 | 36.4 | |
| miR 21-5p at FU1 | Below the mean | 14.7 (8.6 – 20.9) | 54.5 | 27.3 | 0.618 |
| | Above the mean | 9.4 (3.4 – 15.4) | 20.0 | 20.0 | |
| miR 23b-3p at FU1 | Below the mean | 14.2 (8.4 – 19.9) | 50.0 | 25.0 | 0.734 |
| | Above the mean | 9.8 (2.2 – 17.3) | 25.0 | 25.0 | |
| miR 34a-5p at FU1 | Below the mean | 13.2 (6.4 – 20.1) | 44.4 | 22.2 | 0.808 |
| | Above the mean | 13.6 (6.4 – 20.7) | 42.9 | 28.6 | |
| miR 21-5p at FU3 | Below the mean | 14.3 (5.8 – 22.7) | 50.0 | 25.0 | 0.475 |
| | Above the mean | 10.3 (3.0 – 17.5) | 25.0 | 0 | |
| miR 23b-3p at FU3 | Below the mean | 8.8 (4.6 – 12.9) | 25.0 | 0 | 0.230 |
| | Above the mean | 15.8 (6.7 – 24.8) | 50.0 | 25.0 | |
| miR 34a-5p at FU3 | Below the mean | 8.2 (5.2 – 11.1) | 16.7 | 0 | 0.032 |
| | Above the mean | 24.5 (19.6 – 29.4) | 100.0 | 50.0 | |
| miR 21-5p at FU6 | Below the mean | 13.3 (7.4 – 19.2) | 50.0 | 0 | 0.024 |
| | Above the mean | 24.4 (20.4 – 28.4) | 100.0 | 60.0 | |
| miR 23b-3p at FU6 | Below the mean | 15.2 (9.9 – 20.5) | 60.0 | 20.0 | 0.230 |
| | Above the mean | 23.5 (18.9 – 28.1) | 100.0 | 50.0 | |
| miR 34a-5p at FU6 | Below the mean | 17.8 (12.0 – 23.6) | 80.0 | 20.0 | 0.356 |
| | Above the mean | 19.0 (13.3 – 24.7) | 75.0 | 50.0 | |
| miR 21-5p at FU12 | Below the mean | 25.7 (21.9 – 29.4) | 100.0 | 66.7 | 0.592 |
| | Above the mean | 22.5 (14.9 – 30.1) | 100.0 | 50.0 | |
| miR 23b-3p at FU12 | Below the mean | 25.7 (21.9 – 29.4) | 100.0 | 66.7 | 0.592 |
| | Above the mean | 22.5 (14.9 – 30.1) | 100.0 | 50.0 | |
| miR 34a-5p at FU12 | Below the mean | - | 100.0 | 33.3 | 0.199 |
| | Above the mean | - | 100.0 | 100.0 | |

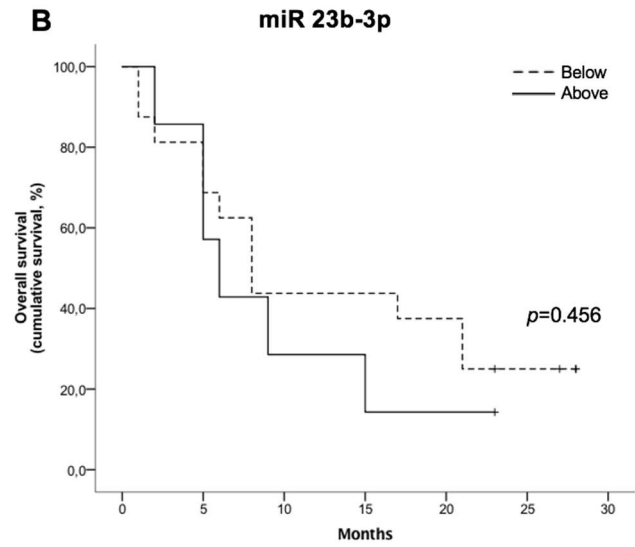
Table 10 – Mean and predicted OS according to mean miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis, 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) after surgery.

pOS: predicted overall survival.



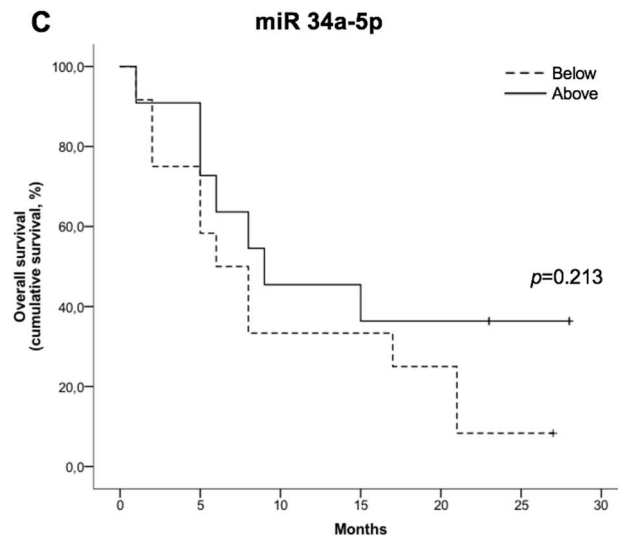
Number at risk
(number censored)

| | | | | | | | |
|-------|--------|--------|-------|-------|-------|--------|-------|
| Below | 15 (0) | 10 (5) | 7 (8) | 7 (8) | 6 (9) | 4 (11) | 0 (0) |
| Above | 8 (0) | 5 (3) | 2 (6) | 1 (7) | 1 (7) | 0 (0) | 0 (0) |



Number at risk
(number censored)

| | | | | | | | |
|-------|--------|--------|-------|-------|--------|--------|-------|
| Below | 16 (0) | 11 (5) | 7 (9) | 7 (9) | 6 (10) | 4 (12) | 0 (0) |
| Above | 7 (0) | 4 (3) | 2 (5) | 1 (6) | 1 (6) | 0 (0) | 0 (0) |



Number at risk
(number censored)

| | | | | | | | |
|-------|--------|-------|-------|-------|-------|--------|-------|
| Below | 12 (0) | 7 (5) | 4 (8) | 4 (8) | 3 (9) | 1 (11) | 0 (0) |
| Above | 11 (0) | 8 (3) | 5 (6) | 4 (7) | 4 (7) | 4 (7) | 0 (0) |

Figure 51 – Overall survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean at diagnosis.

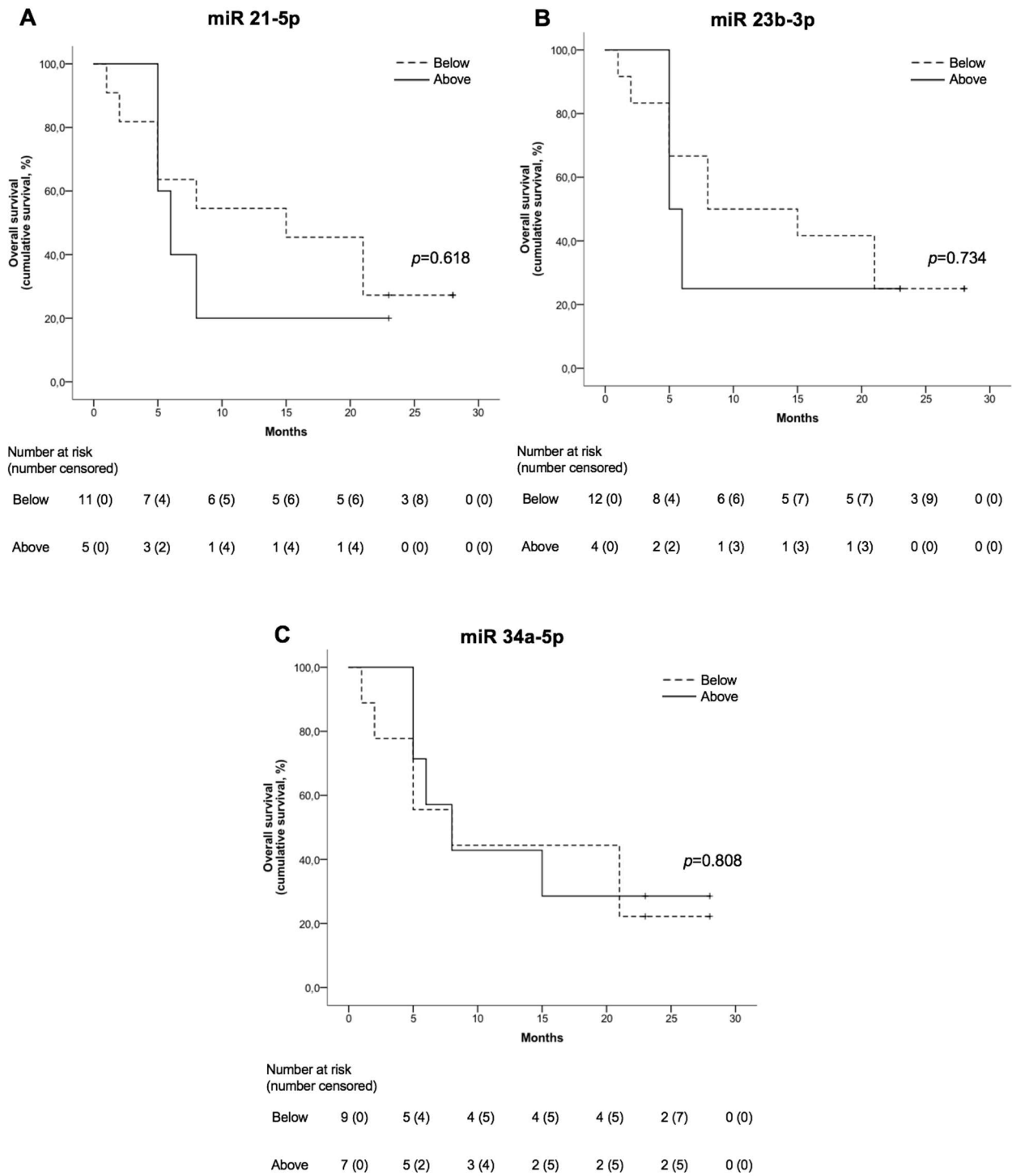


Figure 52 – Overall survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C) below (dashed line) and above (solid line) the mean 1 month after surgery.

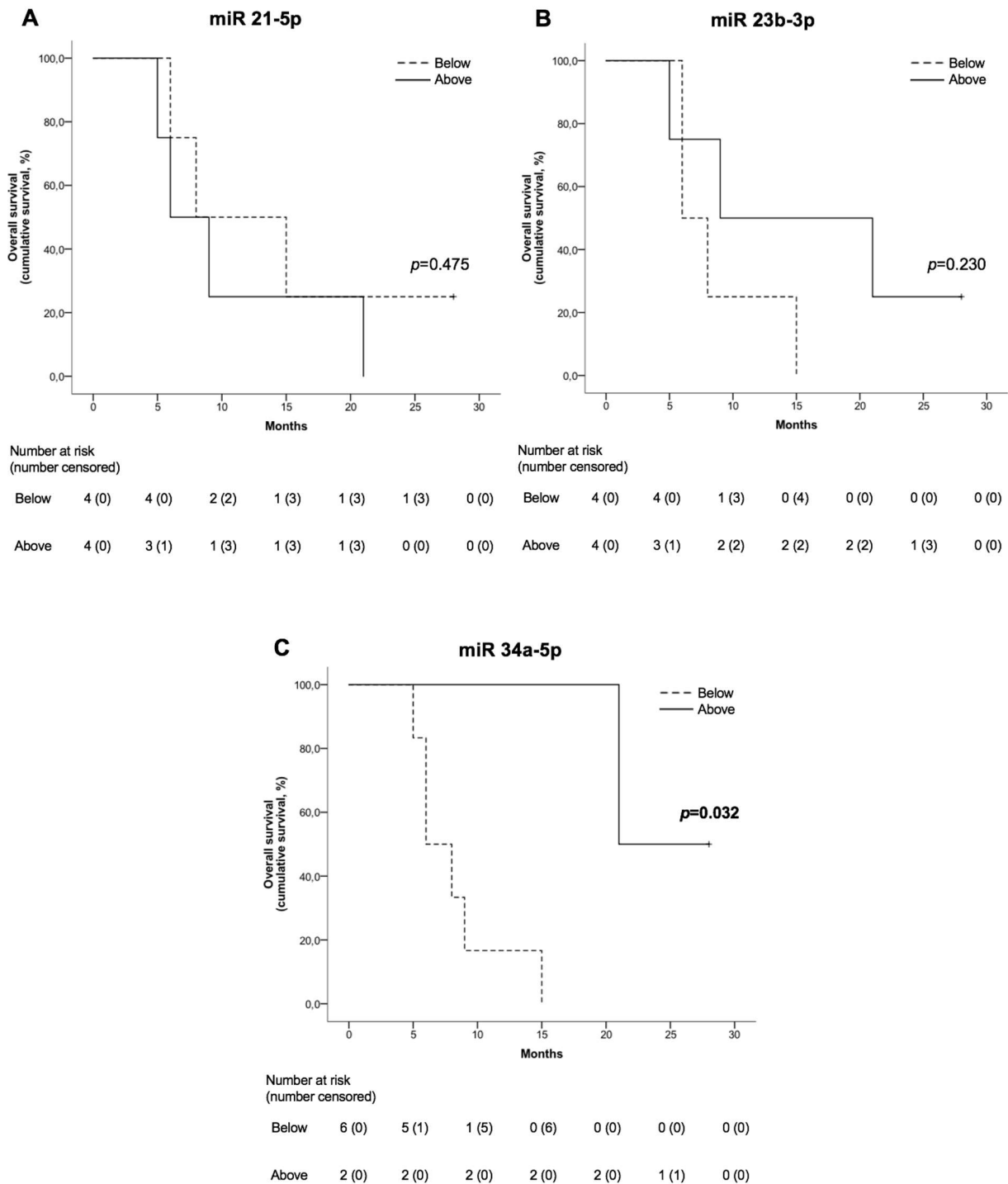


Figure 53 – Overall survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean 3 months after surgery.

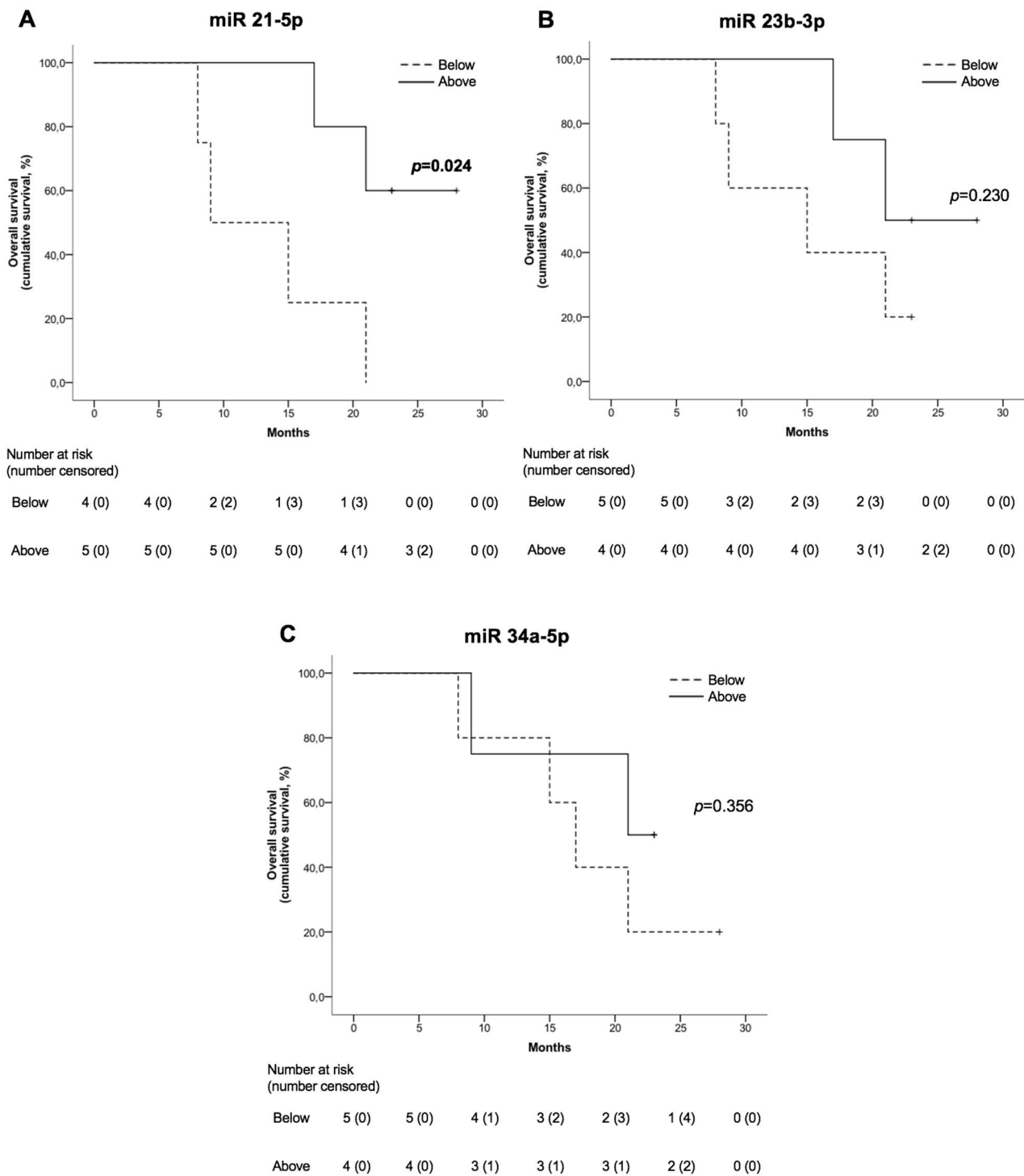


Figure 54 – Overall survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean 6 months after surgery.

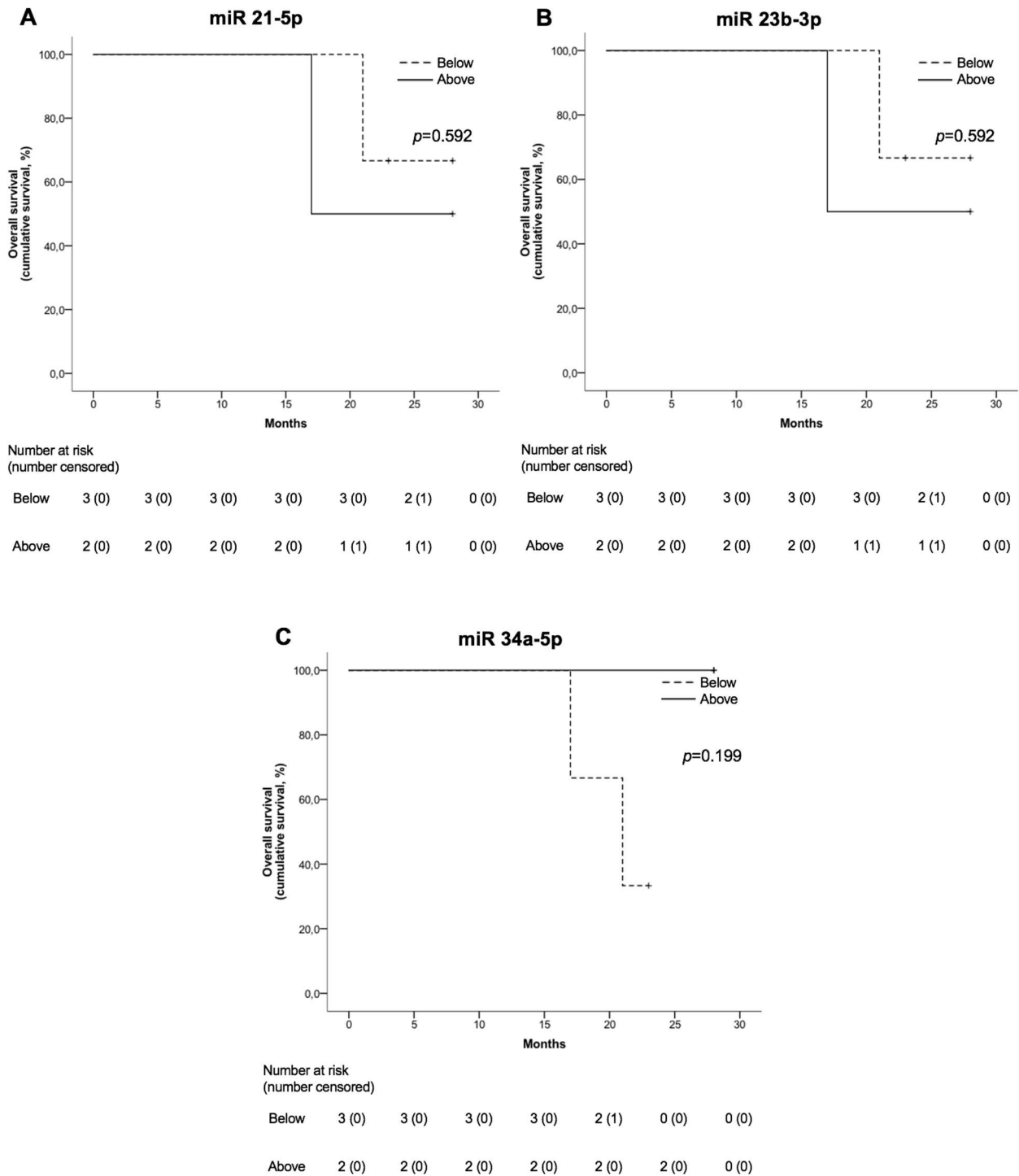


Figure 55 – Overall survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (A), miR 23b-3p (B) and miR 34a-5p (C) below (dashed line) and above (solid line) the mean 12 months after surgery.

4.11 Recurrence free survival analysis

The mean RFS was 11.4 months (95% C.I. 6.8 – 16.0) with 31.9% and 26.6% of predicted RFS at 1 and 2 years, respectively (**Figure 56**).

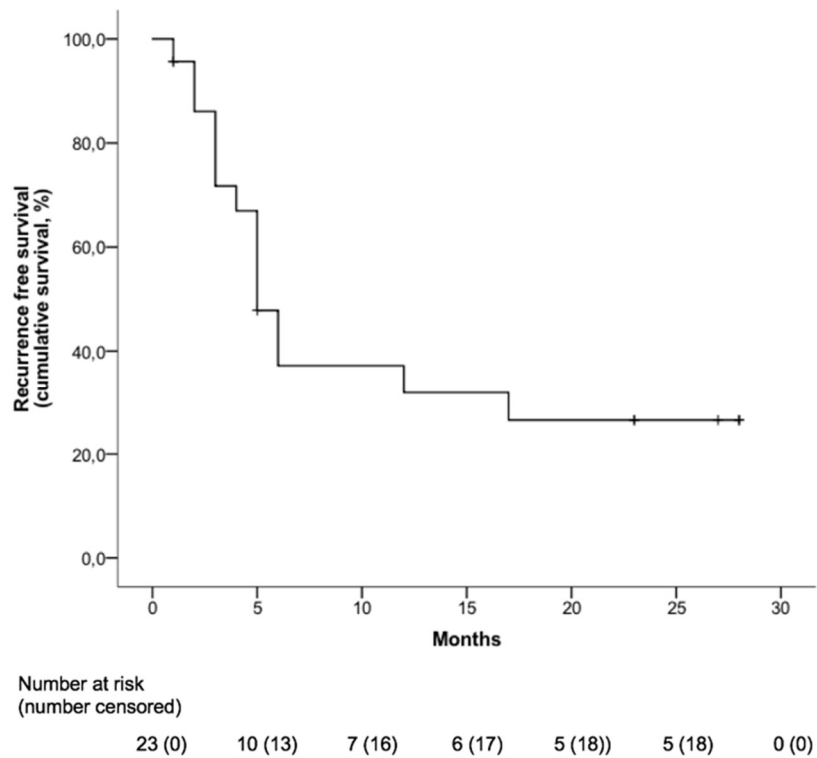


Figure 56 – Recurrence free survival.

4.11.1 Recurrence free survival according to miRNAs' variations at different timing points

After splitting the population in two groups of patients experiencing increase or stability/decrease of miRNAs circulating levels over the follow up, the RFS were compared (**Table 11**) and no significant differences surfaced (**Figures 57-60**).

| | | Mean RFS (95% C.I.) | pRFS at 1 year (%) | pRFS at 2 years (%) | p value |
|--------------------------|--------------------|---------------------|--------------------|---------------------|---------|
| miR 21-5p diagnosis-FU1 | Increase | 15.5 (7.2 – 23.8) | 57.1 | 38.1 | 0.412 |
| | Stability/Decrease | 9.8 (2.4 – 17.1) | 25.0 | 25.0 | |
| miR 23b-3p diagnosis-FU1 | Increase | 17.6 (8.6 – 26.5) | 57.1 | 57.1 | 0.135 |
| | Stability/Decrease | 7.8 (2.6 – 12.9) | 25.0 | 12.5 | |
| miR 34a-5p diagnosis-FU1 | Increase | 9.6 (3.0 – 16.2) | 24.4 | 24.4 | 0.211 |
| | Stability/Decrease | 19.5 (10.2 – 28.8) | 75.0 | 50.0 | |
| | | | | | |
| miR 21-5p FU1-FU3 | Increase | 15.5 (0.0 – 32.8) | 50.0 | 50.0 | 0.535 |
| | Stability/Decrease | 4.5 (2.7 – 6.3) | 0.0 | 0.0 | |
| miR 23b-3p FU1-FU3 | Increase | 15.5 (0.0 – 32.8) | 50.0 | 50.0 | 0.535 |
| | Stability/Decrease | 4.5 (2.7 – 6.3) | 0.0 | 0.0 | |
| miR 34a-5p FU1-FU3 | Increase | 15.5 (0.0 – 32.8) | 50.0 | 50.0 | 0.535 |
| | Stability/Decrease | 4.5 (2.7 – 6.3) | 0.0 | 0.0 | |
| | | | | | |
| miR 21-5p FU3-FU6 | Increase | - | - | - | 0.116 |
| | Stability/Decrease | - | - | - | |
| miR 23b-3p FU3-FU6 | Increase | - | - | - | - |
| | Stability/Decrease | 8.6 (0.0 – 17.2) | 20.0 | 20.0 | |
| miR 34a-5p FU3-FU6 | Increase | - | - | - | - |
| | Stability/Decrease | 8.6 (0.0 – 17.2) | 20.0 | 20.0 | |
| | | | | | |
| miR 21-5p FU6-FU12 | Increase | - | - | - | - |
| | Stability/Decrease | 21.3 (14.4 – 28.1) | 75.0 | 50.0 | |
| miR 23b-3p FU6-FU12 | Increase | - | - | - | - |
| | Stability/Decrease | 21.3 (14.4 – 28.1) | 75.0 | 50.0 | |
| miR 34a-5p FU6-FU12 | Increase | - | - | - | 0.090 |
| | Stability/Decrease | - | - | - | |

Table 11 – Mean and predicted RFS according to variations (increase or stability/decrease) of circulating levels of miR 21-5p, miR 23b-3p and miR 34a-5p at different timing points. pRFS: predicted recurrence free survival.

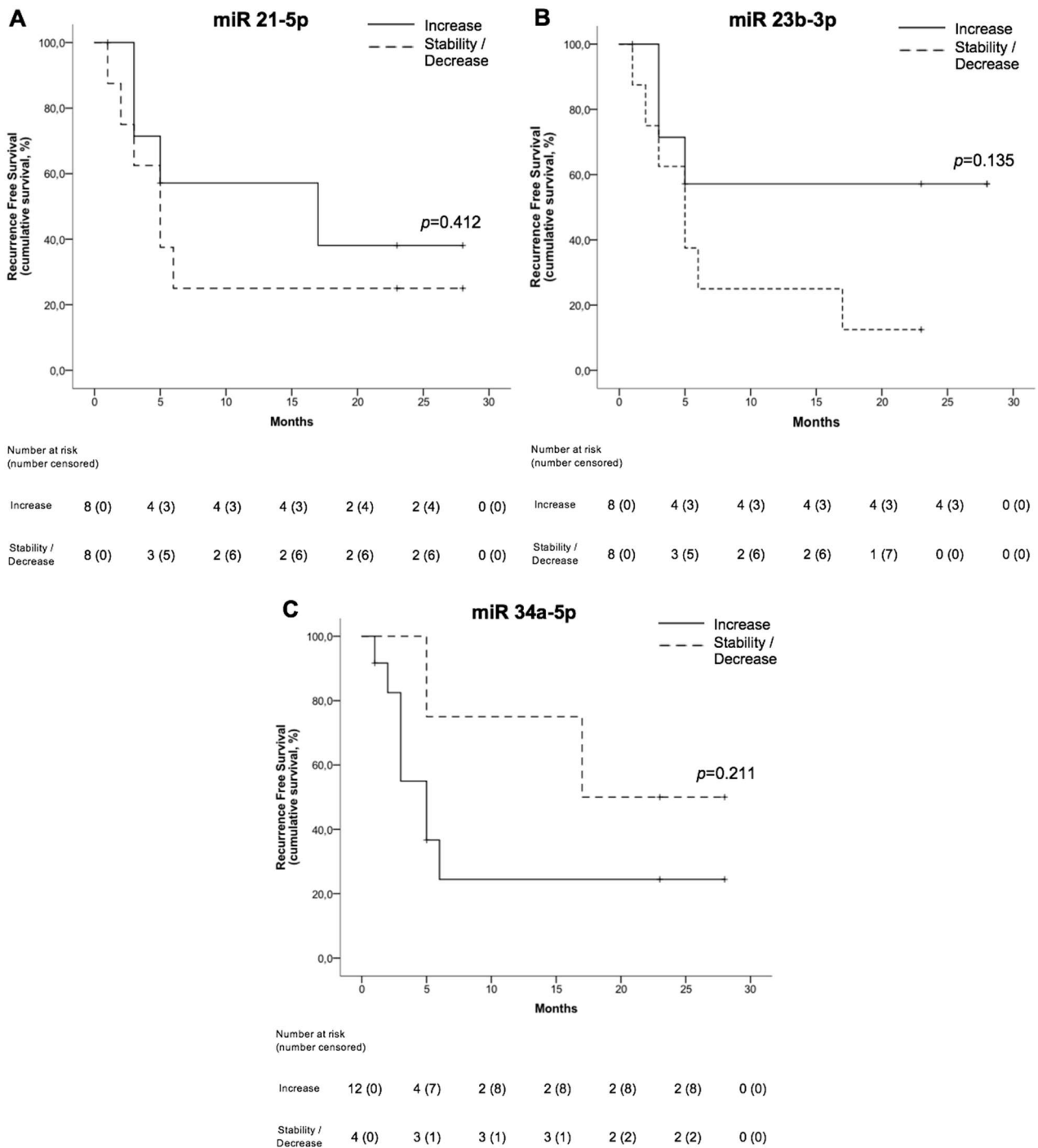


Figure 57 – Recurrence free survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between the diagnosis and 1 month after surgery.

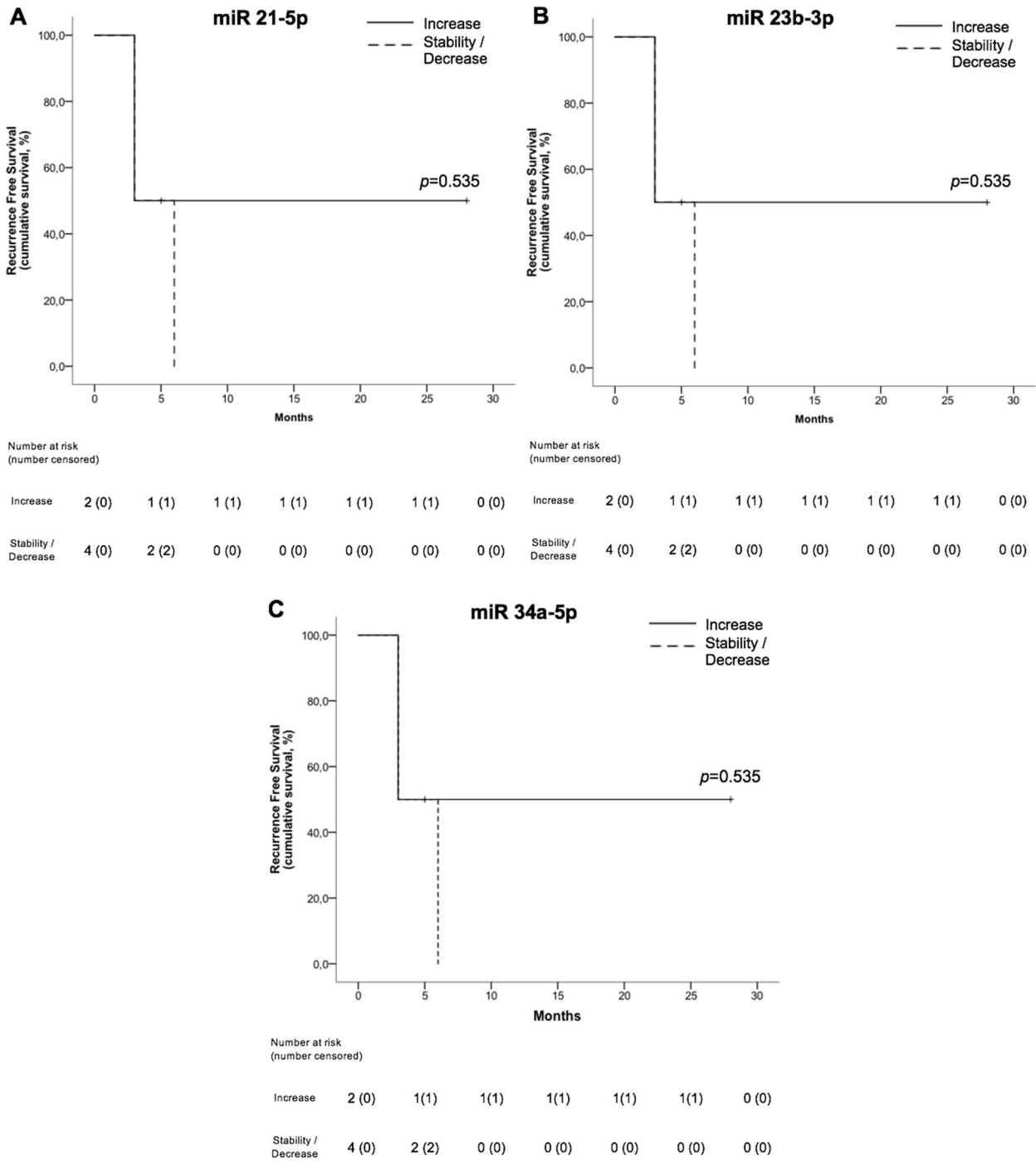


Figure 58 – Recurrence free survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between 1 and 3 months after surgery.

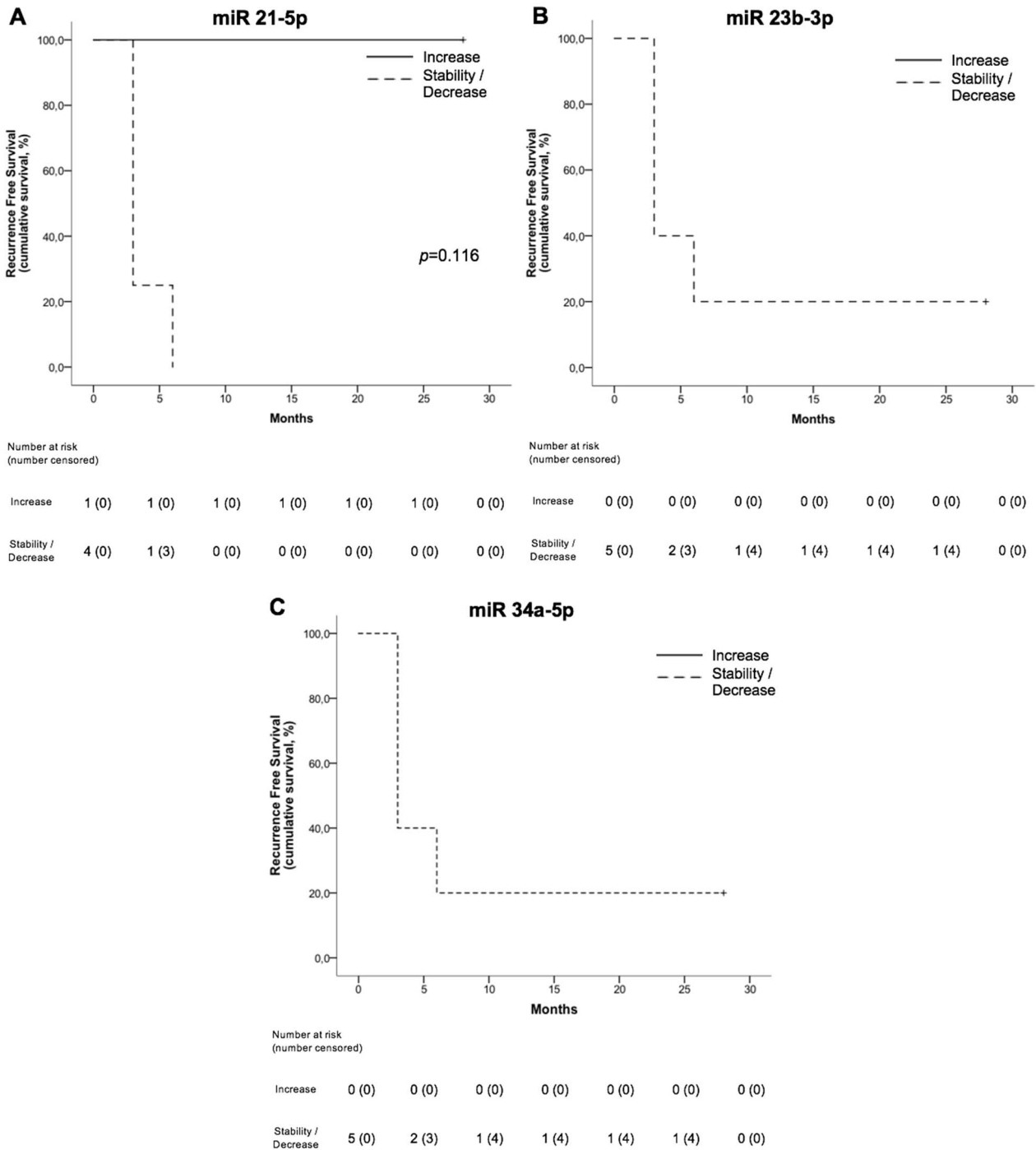


Figure 59 – Recurrence free survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between 3 and 6 months after surgery.

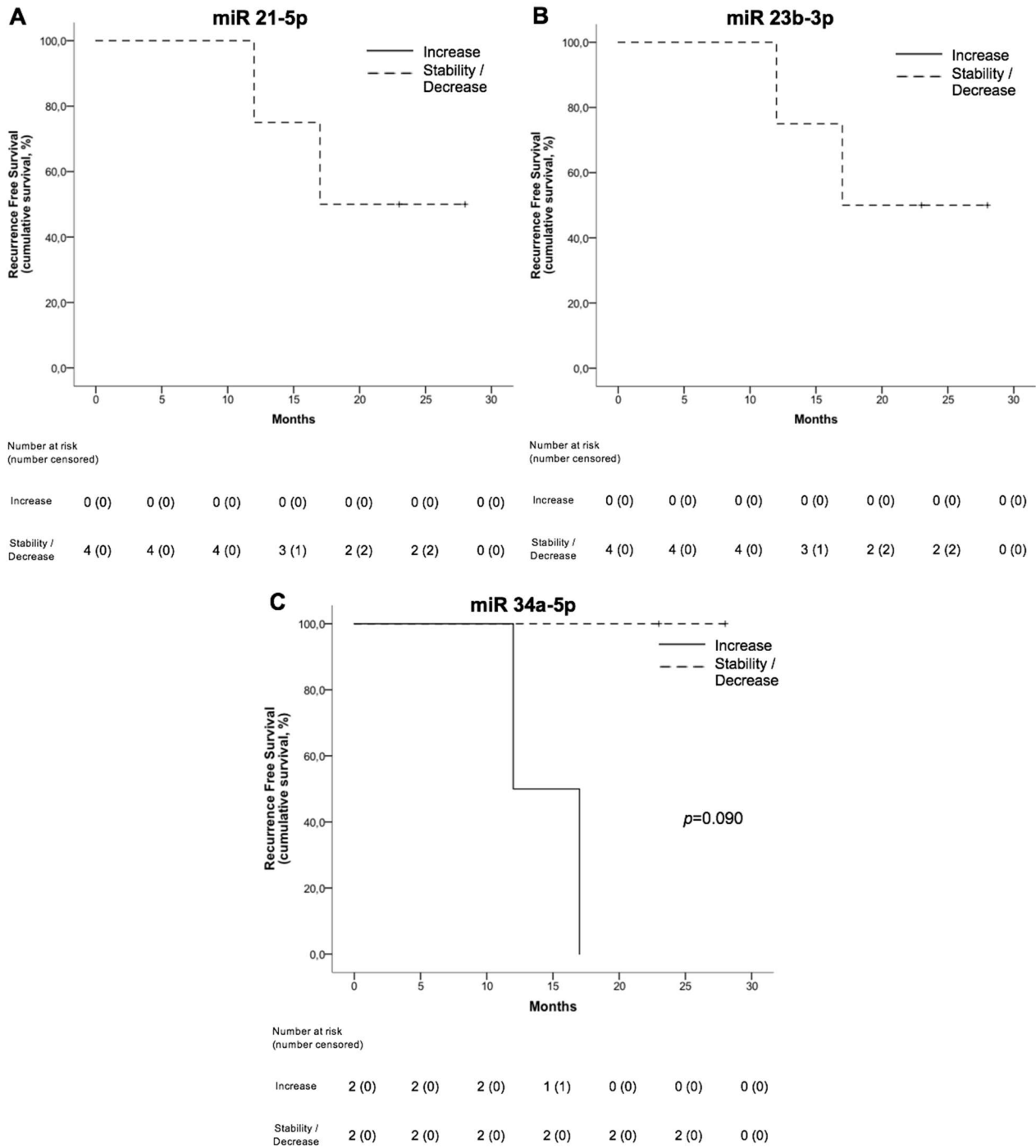


Figure 60 – Recurrence free survival Kaplan–Meier curves comparing patients with increased (solid line) and stable or decreased (dashed line) circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) between 6 and 12 months after surgery.

4.11.2 Recurrence free survival according to mean miRNAs' plasmatic concentrations

After splitting the population above and below the mean values of circulating miRNAs at different timing points (**Table 12**), the mean RFS was significantly shorter in patients with miR 34a-5p levels below the mean at diagnosis (7.6 months, 95% C.I. 3.1 – 12.1 vs 15.6 months, 95% C.I. 8.0 – 23.1, $p=0.049$) (**Figures 61-65**).

| | | Mean RFS (95% C.I.) | pRFS at 1 year (%) | pRFS at 2 years (%) | p value |
|--------------------------------|----------------|--------------------------|--------------------|---------------------|--------------|
| miR 21-5p at diagnosis | Below the mean | 12.1 (6.4 – 17.8) | 35.9 | 28.7 | 0.872 |
| | Above the mean | 8.5 (2.1 – 14.9) | 21.4 | 21.4 | |
| miR 23b-3p at diagnosis | Below the mean | 12.2 (6.5 – 17.9) | 36.1 | 28.8 | 0.878 |
| | Above the mean | 8.1 (2.0 – 14.1) | 19.0 | 19.0 | |
| miR 34a-5p at diagnosis | Below the mean | 7.6 (3.1 – 12.1) | 18.3 | 9.2 | 0.049 |
| | Above the mean | 15.6 (8.0 – 23.1) | 46.7 | 46.7 | |
| | | | | | |
| miR 21-5p at FU1 | Below the mean | 12.1 (5.1 – 19.1) | 40.4 | 30.3 | 0.692 |
| | Above the mean | 10.3 (1.9 – 18.7) | 30.0 | 30.0 | |
| miR 23b-3p at FU1 | Below the mean | 11.3 (4.7 – 17.8) | 36.7 | 27.5 | 0.383 |
| | Above the mean | 12.1 (2.4 – 21.8) | 37.5 | 37.5 | |
| miR 34a-5p at FU1 | Below the mean | 11.6 (4.4 – 18.8) | 37.5 | 25.0 | 0.812 |
| | Above the mean | 12.8 (3.3 – 22.3) | 38.1 | 38.1 | |
| | | | | | |
| miR 21-5p at FU3 | Below the mean | 10.0 (0 – 20.3) | 25.0 | 25.0 | 0.815 |
| | Above the mean | 5.0 (3.5 – 6.5) | 0 | 0 | |
| miR 23b-3p at FU3 | Below the mean | 4.3 (2.8 – 5.7) | 0 | 0 | 0.151 |
| | Above the mean | 13.5 (0.6 – 26.4) | 37.5 | 37.5 | |
| miR 34a-5p at FU3 | Below the mean | 4.8 (3.6 – 6.0) | 0 | 0 | 0.486 |
| | Above the mean | 15.5 (0 – 32.8) | 50.0 | 50.0 | |
| | | | | | |
| miR 21-5p at FU6 | Below the mean | 7.3 (0.7 – 13.8) | 25.0 | 0 | 0.076 |
| | Above the mean | 19.8 (10.7 – 29.0) | 60.0 | 60.0 | |
| miR 23b-3p at FU6 | Below the mean | 10.4 (3.3 – 17.5) | 40.0 | 20.0 | 0.412 |
| | Above the mean | 17.8 (7.2 – 21.2) | 50.0 | 50.0 | |
| miR 34a-5p at FU6 | Below the mean | 12.6 (4.4 – 20.8) | 40.0 | 20.0 | 0.519 |
| | Above the mean | 13.8 (4.6 – 22.9) | 50.0 | 50.0 | |
| | | | | | |
| miR 21-5p at FU12 | Below the mean | 24.3 (18.5 – 30.2) | 100.0 | 66.7 | 0.592 |
| | Above the mean | 20.0 (8.9 – 31.1) | 50.0 | 50.0 | |
| miR 23b-3p at FU12 | Below the mean | 24.3 (18.5 – 30.2) | 100.0 | 66.7 | 0.592 |
| | Above the mean | 20.0 (8.9 – 31.1) | 50.0 | 50.0 | |
| miR 34a-5p at FU12 | Below the mean | - | 66.7 | 33.3 | 0.199 |
| | Above the mean | - | 100.0 | 100.0 | |

Table 12 – Mean and predicted RFS according to mean miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis, 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) of follow up. pRFS: predicted recurrence free survival.

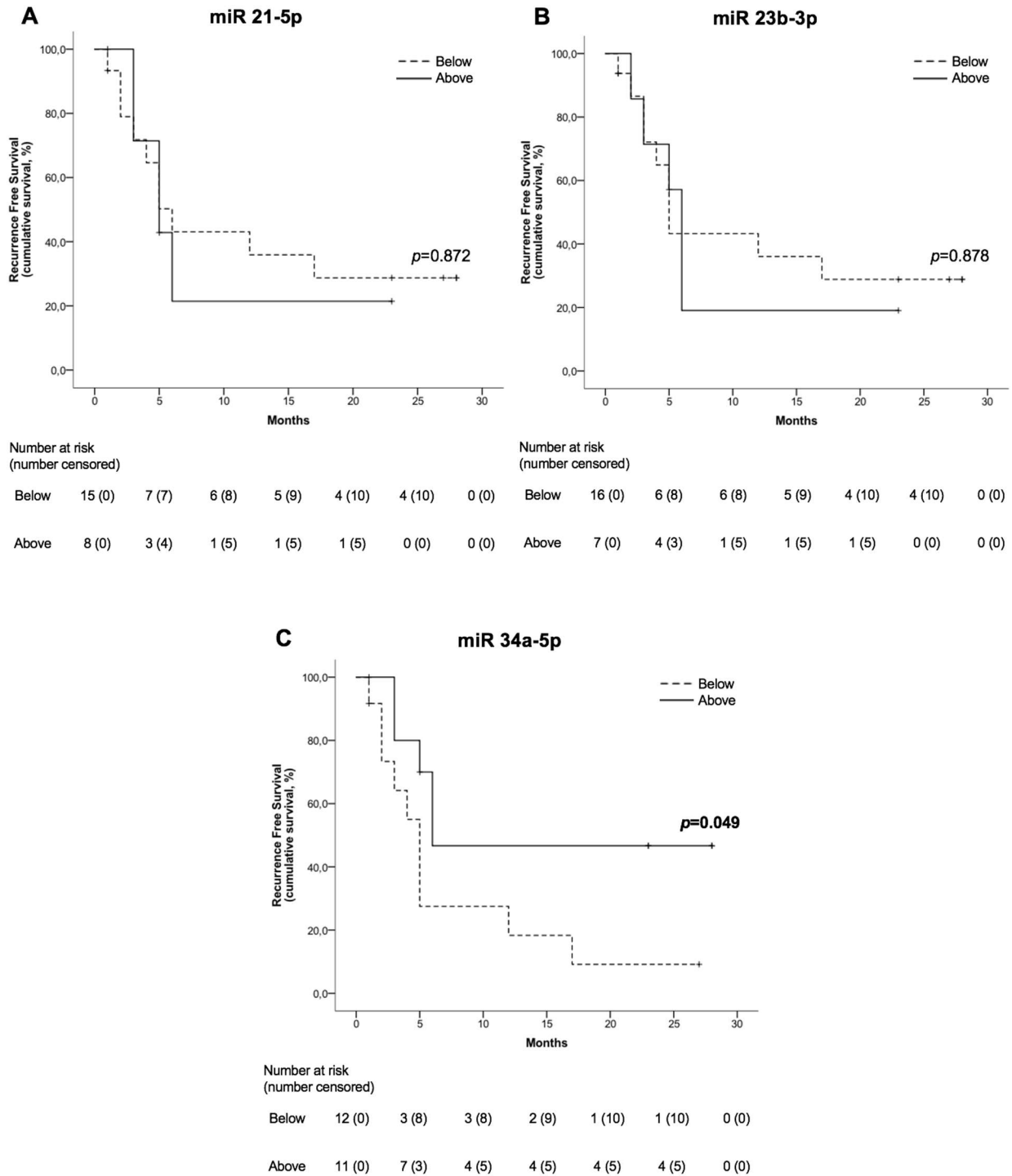


Figure 61 – Recurrence free survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean at diagnosis.

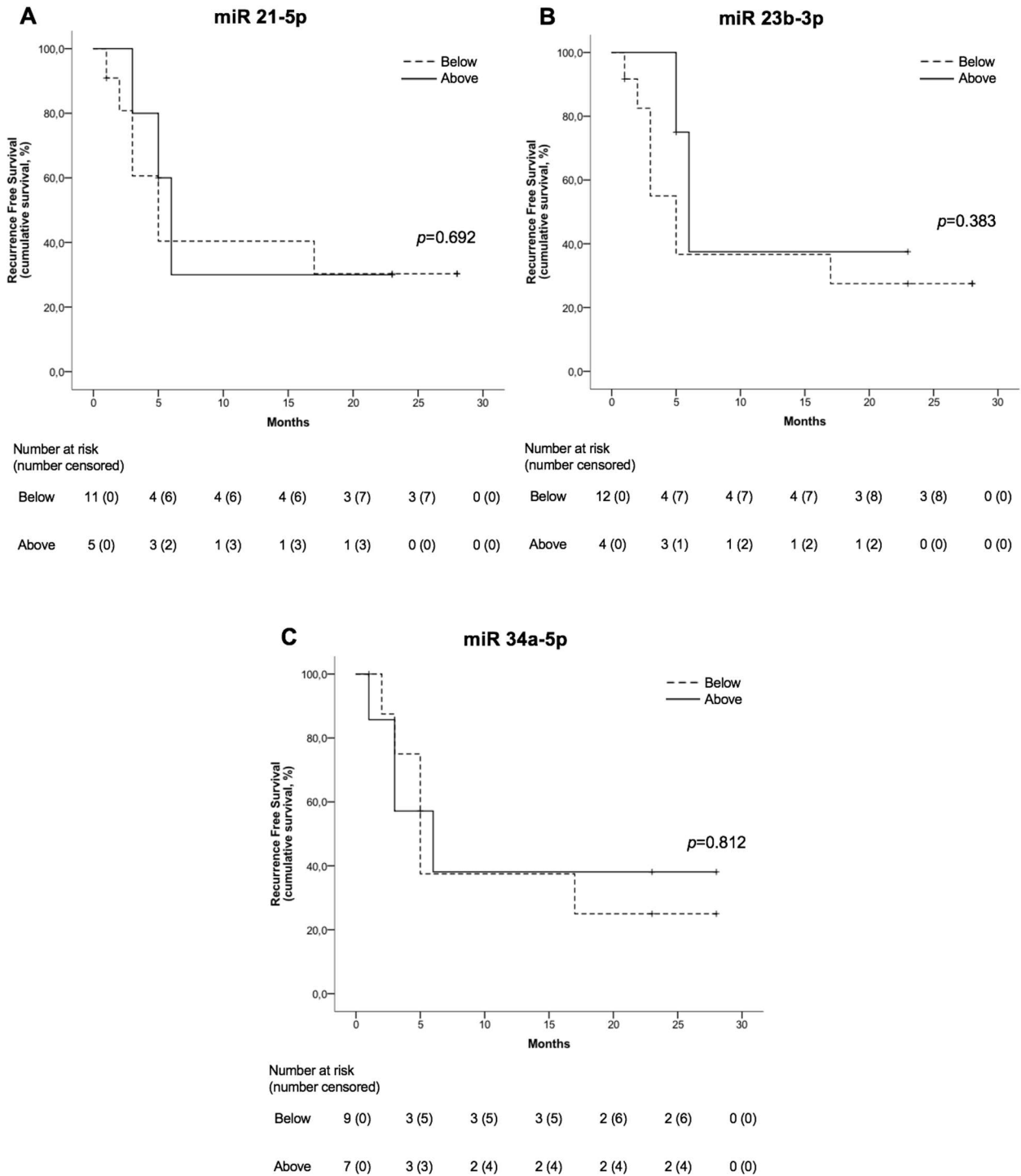


Figure 62 – Recurrence free survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean 1 month after surgery.

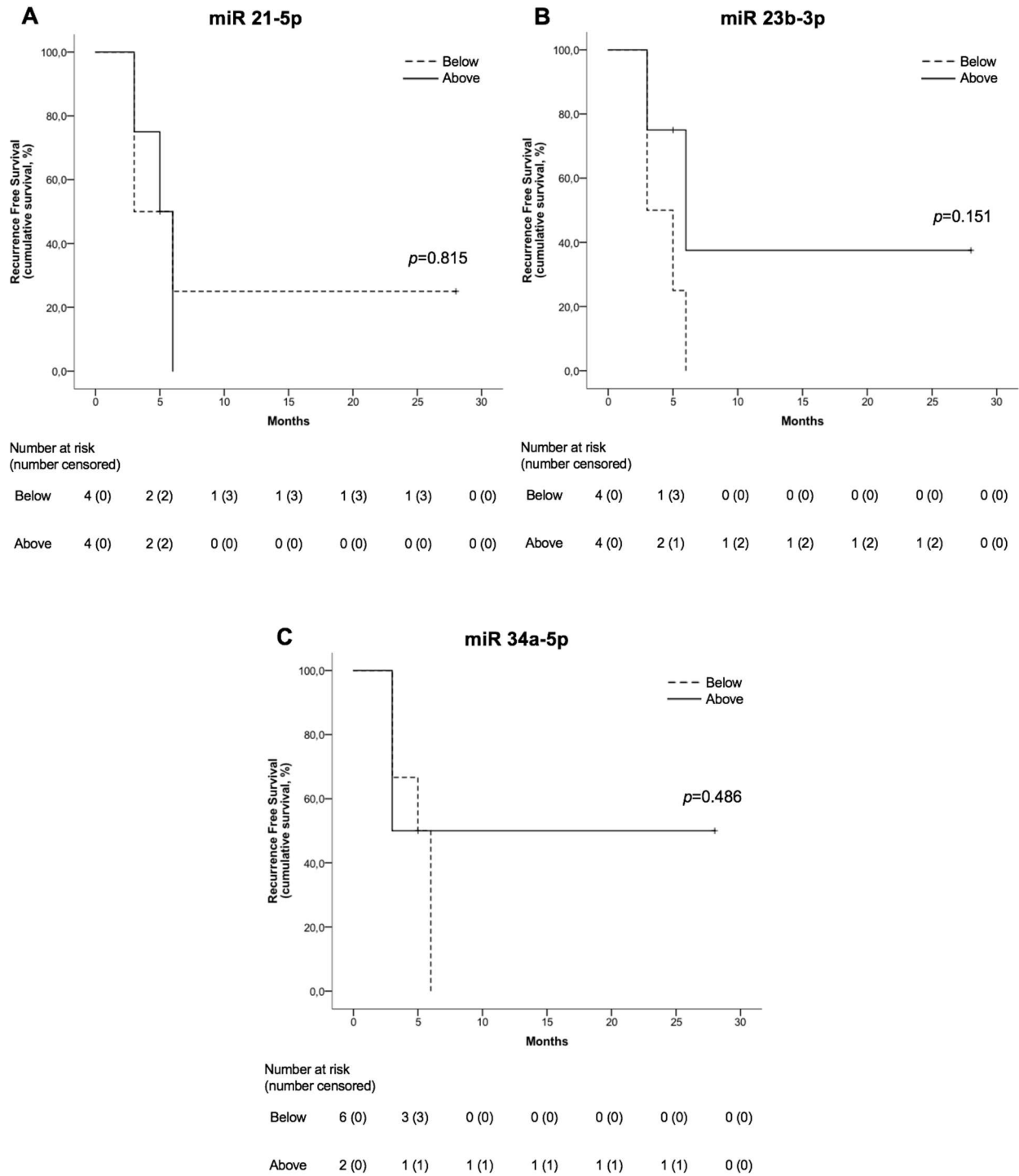


Figure 63 – Recurrence free survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean 3 months after surgery.

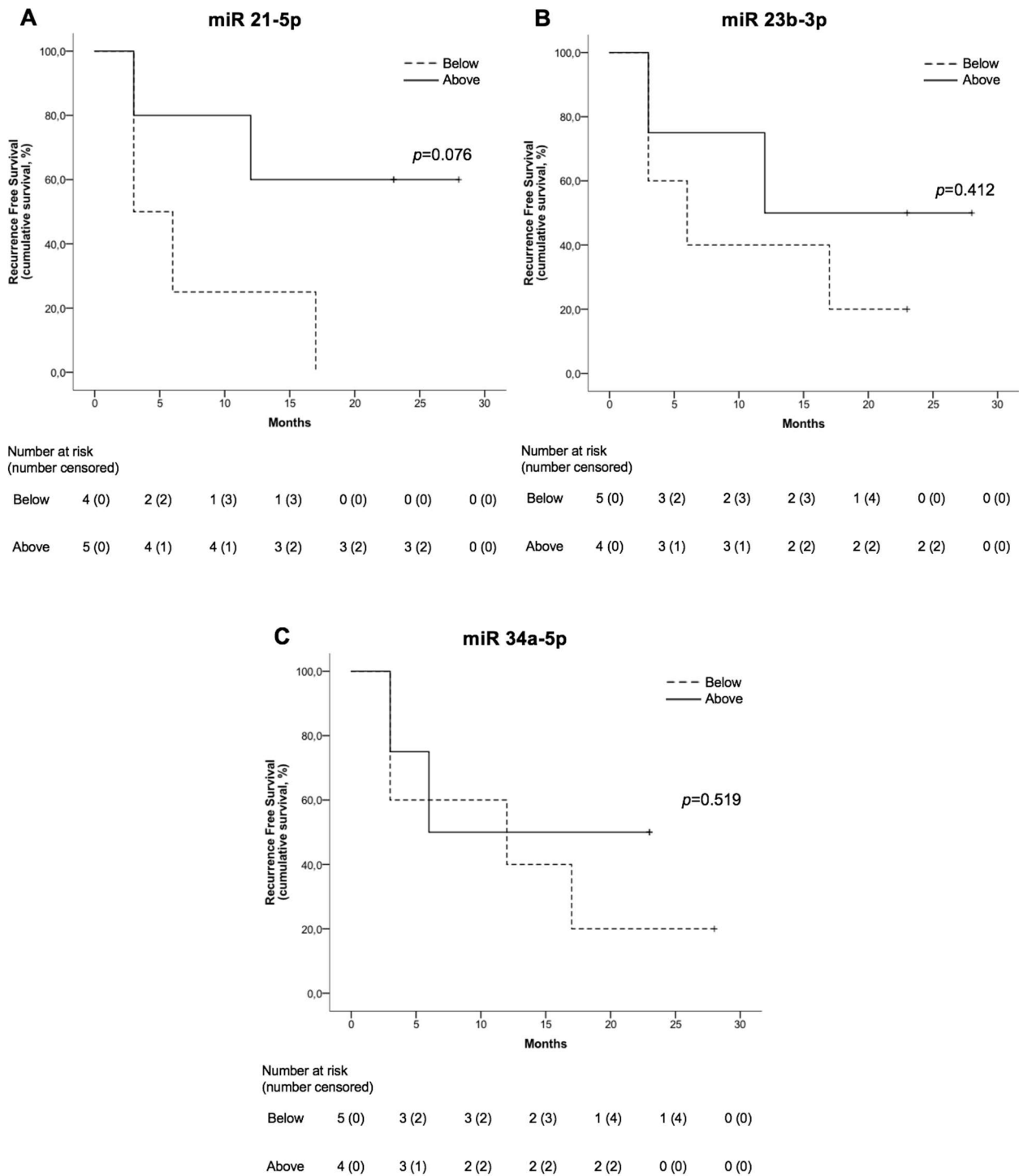


Figure 64 – Recurrence free survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean 6 months after surgery.

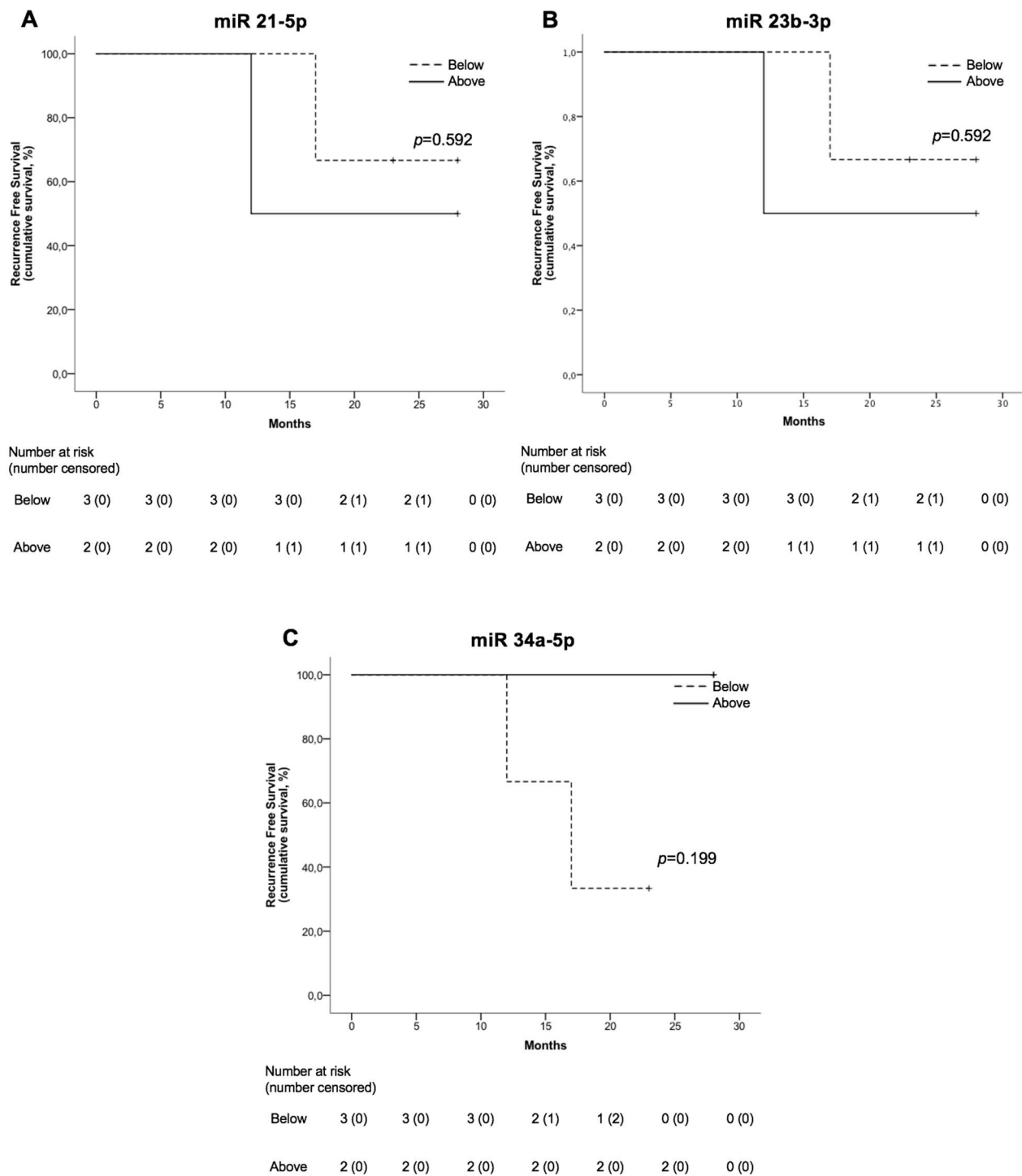


Figure 65 – Recurrence free survival Kaplan–Meier curves comparing patients with circulating levels of miR 21-5p (**A**), miR 23b-3p (**B**) and miR 34a-5p (**C**) below (dashed line) and above (solid line) the mean 12 months after surgery.

5. DISCUSSION

5.1 Liquid biopsy in GBM

In the neuro-oncology field, liquid biopsy has recently surged in popularity as a potential tool for diagnosis, prognosis and treatment monitoring³⁻⁷. GBM is characterized by molecular and genetic heterogeneity, presenting distinct genetic features in different tumour regions⁷¹. Liquid biopsy provides a comprehensive picture of the cancer's landscape more than tumour tissue analysis, capturing genetic materials reversed in the bloodstream by all the tumour components.

In the era of precision medicine, liquid biopsy may offer the potential for early GBM detection, serial monitoring over time for disease progression and response to treatments, allowing the identification of molecular targets for personalized therapies.

However, the identification of GBM-specific circulating biomarkers presents some challenges and limitations. Sensitivity and specificity of the detection techniques might be influenced by the low concentrations of circulating analytes in part due to the presence of the blood brain barrier^{5,6,14-16,44}. Moreover, standardization of methodologies and further validation studies are needed to establish clinical reliability of these biomarkers for GBM.

5.2 Plasmatic miRNAs' assessment by ddPCR

This study was aimed to explore the role of circulating non-coding RNAs as potential GBM-specific longitudinal peripheral biomarkers using ddPCR. In particular, the detection and quantification of selected miRNAs was accomplished by ddPCR on plasma samples, collected at specific timing points.

miRNAs are involved in the tumorigenesis of different types of cancers. Specific miRNA expression profiles are often associated with different tissues and pathological conditions. Changes of their expression can reflect tissue alterations, such as cancerous changes^{15,17,23-25,28,33}. For this reason, circulating miRNAs have been explored as potential tumour-specific biomarkers^{4,17,18,21,23-25,27,28,30}. Using circulating miRNAs for liquid biopsy brings several advantages and we focused our attention on their assessment.

miRNAs are relatively stable in comparison with other circulating nucleic acids^{1,2,17,18,72}. Furthermore, they are protected from degradation thanks to their encapsulation in extracellular vesicles and apoptotic bodies or protein binding, but show also resistance to endogenous RNase activity^{18,31}.

However, clinical application of miRNA as tumour biomarkers has been mainly limited by the lack of standard methods for blood collection, RNA extraction, RNA sequencing and normalization methods based on endogenous controls⁷³⁻⁷⁵.

Although both plasma and serum are suitable for miRNAs' analysis, some studies have reported that the identification of extracellular ("cell-free") miRNAs may be more reliable in plasma samples for different reasons: i) anticoagulation prevents the degranulation of platelets (specifically tumour-educated platelets), a crucial step in the clotting process, that can cause the release of intracellular miRNAs; ii) plasma is not affected by haemolysis that can influence miRNA profiles by leaking intracellular miRNAs; iii) plasma contains higher concentrations of extracellular vesicles (particularly exosomes) that contribute to the global pool of extracellular miRNAs^{7,35,72,76-78}. Accordingly, we decided to quantify miRNAs' plasmatic levels.

We then adopted ddPCR for an absolute quantification of our target molecules, differently from real time PCR that provides a relative quantification, without relying on standard curves based on endogenous controls. Despite different suggestions, there is no universal housekeeping control for miRNAs and the process of choosing a suitable reference is still controversial contributing to the final heterogeneity of data reported in literature for real time PCR^{5,37,79-81}.

The ddPCR partitions the process into thousands of independent reactions (droplets) enabling a highly sensitive detection of low-abundant targets, such as circulating nucleic acids^{32,82}. Moreover, ddPCR is less affected by the presence of PCR inhibitors, an extraordinary advantage when working with complex samples such as blood, and by PCR amplification efficiency variability, that can limit the accuracy of relative quantification. Overall, ddPCR affords for consistent reaction conditions contributing to highly reproducible results^{5,82}.

Circulating miRNAs can also be analysed by microarrays and next generation sequencing. However, the quantitative accuracy of microarrays can be compromised by low concentrations of transcripts. On the other hand, next generation sequencing is significantly less affordable requiring substantial computational resources for data storage and processing compared to PCR³.

5.3 miRNAs' selection: miR 21-5p, miR 23b-3p and miR 34a-5p

According to the literature review, we identified three miRNAs as potential peripheral biomarkers for GBM: miR 21-5p, miR 23b-3p and miR 34a-5p.

miR 21-5p is located on chromosome 17q23.2⁸³. miR 21-5p has been extensively studied in various cancers, including GBM, for its oncogenic properties and implications in promoting cell proliferation, inhibiting apoptosis and contributing to tumour invasion and migration. miR 21-5p has been involved in three different tumour suppressor pathways, namely p53, TGF-beta and the mitochondrial apoptotic pathway^{38,83-86}. It has been found as highly expressed in tissues and serum or plasma of patients affected by GBM compared to healthy controls and/or patients with low grade diffuse astrocytic gliomas, with all limitations of previous studies that did not follow the new WHO Classification of the Tumours of the CNS^{14,15,35-43,87}.

miR 23b-3p is located on chromosome 9p21.3. *miR 23b-3p* has been shown to influence tumour cell proliferation, invasion and migration and some studies reported its overexpression in cell cultures and tumour tissue derived from patients affected by GBM^{45,47,48}.

Eventually, *miR 34a-5p* sited on chromosome 1p36.22⁴⁹ is a direct transcriptional target of p53 and acts as a tumour suppressor miRNA. It promotes cell cycle arrest and apoptosis, suppresses cell invasion and extracellular matrix remodelling and targets key oncogenic signalling pathways. Downregulation of *miR 34a-5p* has been reported in tumour tissues and serum derived from patients affected by GBM^{15,37,39,44,49,50,52,53}.

5.4 Study population features

We analysed a homogenous population of patients presenting with GBM IDH-wildtype WHO grade 4 or astrocytoma IDH-mutant WHO grade 4, according to the fifth edition of the WHO Classification of the Tumours of the CNS^{13,19}.

All patients underwent standardized preoperative and postoperative clinical and neuroradiological evaluations, were treated following the international guidelines^{10-12,59} and were followed at defined timing points: at diagnosis, 1 month after surgery, before starting RT and concomitant chemotherapy, 3 months after surgery, immediately after the end of RT and concomitant chemotherapy, 6 and 12 months after surgery.

5.5 miRNAs plasmatic concentrations in healthy volunteers and patients

miR 21-5p, *miR 23b-3p* and *miR 34a-5p* were all detectable in healthy volunteers. Only *miR 34a-5p* showed significantly higher levels in patients, with a diagnostic cut-off value of 1.25 copies/ μ L (sensitivity: 47.8%; specificity: 87.5%). In our cohort of patients, *miR 21-5p* showed similar plasmatic concentrations in healthy volunteers and patients, differently from data previously reported in literature. A high degree of inter-individual variability in the levels of circulating miRNAs has been reported, even when focusing only on healthy populations^{88,89}. Furthermore, in contrast with literature reports of *miR 21-5p* and *miR 23b-3p* overexpression and *miR 34a-5p* downregulation in GBM tissues, our findings on their circulating levels may be due to the fact that aberrant expression of miRNAs in tissues may not be reflected in the circulation^{40,87}.

According to the absolute concentrations of miRNAs at diagnosis, patients were clustered in two groups presenting low and high levels of circulating miRNAs, respectively. Considering the miRNA expression levels relative to healthy volunteers, the population was grouped in patients with low levels of all miRNAs, patients with low levels of *miR 21-5p* and high levels of *miR 23b-3p*, patients with high levels of all miRNAs and patients with low levels of *miR 21-5p* and *miR 23b-3p* and high levels of *miR 34a-5p*. We can therefore identify at diagnosis different groups of patients.

miR 21-5p, miR 23b-3p and miR 34a-5p showed the same longitudinal course. Their plasmatic levels progressively increased 1 and 3 months after surgery, at the end of RT and concomitant chemotherapy, and then gradually decreased over time, at 6 and 12 months after surgery.

The blood brain barrier may affect the amount of miRNAs' released in the bloodstream of patients affected by GBM⁸⁵. In our study population, miRNAs plasmatic concentrations resulted higher when the damage of the blood brain barrier might be potential maximal, particularly after the combined effect of surgery, RT and chemotherapy (3 months after surgery), to subsequently decrease probably as a consequence of the blood brain barrier healing processes⁴⁴.

5.6 miRNAs' correlations with clinical, neuroradiological and pathological features

Despite we registered higher levels of miR 23b-3p and miR 34a-5p at diagnosis in patients who were not administered steroids and lower levels of miR 21-5p, miR 23b-3p and miR 34a-5p in patients with tumours displaying loss of ATRX expression, these results were not confirmed by multivariate analysis. Furthermore, we did not identify significant correlations between the "tumour load", considered as preoperative tumour volume and postoperative tumour remnant volume, and miRNAs' circulating levels at diagnosis and over the follow up¹⁴.

5.7 Mutual miRNAs' correlations

A direct correlation was recorded for each miRNA between their concentration at diagnosis and 1 month after surgery.

Furthermore, to the best of our knowledge, we firstly reported a significant direct correlation between miR 21-5p and miR 23b-3p circulating levels at each timing point of plasma collection, except for 12 months after surgery. The multivariate analysis confirmed this mutual correlation at diagnosis, 1 and 6 months after surgery. Similarly, miR 21-5p and miR 34a-5p levels at diagnosis were significantly directly correlated.

In keeping with our findings, we performed a network analysis to identify possible common molecular pathways for these miRNAs. Firstly, we identified four genes shared by miR 21-5p and miR 23b-3p in the context of gliomas: *E2F1*, *RB1*, *PTEN* and *PIK3R1*. Furthermore, we found four genes shared between miR 21-5p and miR 34a-5p in the same pathological scenario: *E2F1*, *E2F3*, *CDK6* and *IGF1R*. Eventually, the overall network analysis for miR 21-5p, miR 23b-3p and miR 34a-5p turned towards *E2F1*.

E2F1 and *E2F3* encode for two members of the E2F transcription factor family, key players in the regulation of the cell cycle, promoting the transition from the G1 to the S phase. *E2F1* and *E2F3* aberrant activation or dysregulation can contribute to tumorigenesis. There are growing evidences

of *E2F1* and *E2F3* overexpression in GBM tissue compared to normal brain, associated with tumour aggressiveness and therapy resistance^{37,49,90,91}.

RB1 is a tumour suppressor gene, that inhibits the activity of E2F transcription factors blocking the cell cycle progression. Loss of function of *RB1* has been associated with a more aggressive tumour behaviour, increased cell proliferation and decreased apoptosis in GBM⁹².

PTEN is another tumour suppressor gene that can be dysregulated in GBM, contributing to increased cell survival, proliferation, invasion and resistance to apoptosis. miR 21-5p and miR 23b-3p are known to negatively regulate *PTEN*, leading to the activation of the PI3K/AKT/mTOR signal pathway^{37,38,83,85,86,93}.

PIK3R1 encodes for the regulatory subunit of p85 alpha of PI3K, which increased activity promoting cell survival, proliferation and resistance to apoptosis. miR 21-5p and miR 23b-3p have been associated to this molecular pathway³⁷.

CDK6 encoding for a member of the cyclin-dependent kinase family involved in regulating the cell cycle promotes the transition from the G1 to the S phase. Downregulation of *CDK6* exerted by miR 34a-5p has been described in GBM^{15,37,50,93}.

Finally, *IGF1R* encodes for an insulin-like growth factor receptor, which activation favours cell proliferation and survival in the tumourigenesis including GBM^{94,95}.

5.8 Overall and recurrence free survival

According to previous studies, we documented significant impact of circulating miRNAs levels on OS^{52,83,86,93,96}. Patients with increased levels of miR 21-5p, miR 23b-3p and miR 34a-5p between 1 and 3 months after surgery showed longer OS. Similarly, patients with miR 34a-5p circulating levels above the mean 3 months after surgery and miR 21-5p circulating levels above the mean 6 months after surgery showed longer OS.

By contrast, RFS resulted significantly longer in patients presenting miR 34a-5p circulating levels above the mean at diagnosis.

5.9 Study limitations and future perspectives

The generalization of our findings is limited by limited sizes of the study population. Nevertheless, these represent the preliminary results of an ongoing study, in which a larger number of patients with longer follow up is being enrolled.

This research has many future goals:

- a) investigating possible correlations between plasmatic miRNAs and haematological parameters, such as blood cell counts, the neutrophil-to-lymphocyte ratio, the platelet-to-lymphocyte ratio, the lymphocyte-to-monocyte ratio, the systemic immune-inflammation index, the systemic

inflammation response index, and systemic inflammatory markers, such as fibrinogen, erythrocyte sedimentation rate and C-reactive protein;

- b) early differentiating tumour 'pseudo-progression' from 'tumour progression' after adjuvant treatments during the neuroradiological follow-up according to different plasmatic miRNA expression;
- c) quantifying miRNAs levels in tumour samples by ddPCR, as reported by some other groups⁹⁷⁻¹⁰⁰;
- d) identifying the correlations between circulating miRNAs and the GBM molecular features included in the fifth edition of the WHO Classification of the Tumours of the CNS (homozygous deletion of *CDKN2A/B*, *TERT* promoter mutation, *EGFR* amplification, +7/-10 cytogenetic signature, *BRAF* mutations);
- e) following the longitudinal levels of plasmatic non-coding RNAs in patients who experienced tumour recurrence in order to assess the response to second-line therapies, such as VEGF (Vascular Endothelial Growth Factor) and tyrosine kinase inhibitors.

6. CONCLUSIONS

We reported a prospective case-control longitudinal study on plasmatic miR 21-5p, miR 23b-3p and miR 34a-5p detected by means of ddPCR as peripheral biomarkers of astrocytomas, IDH-mutant WHO grade 4 and GBM, IDH-wildtype WHO grade 4 according to the fifth edition of the WHO Classification of the Tumours of the CNS^{13,19}.

We detected all miRNAs both in healthy volunteers and patients and documented significantly higher levels of circulating miR 34a-5p in patients than healthy volunteers identifying a diagnostic absolute cut-off value of 1.25 copies/ μ L. Furthermore, two groups of patients were recognised showing high and low absolute plasmatic levels of miRNAs at diagnosis, respectively.

miR 21-5p, miR 23b-3p and miR 34a-5p showed the same longitudinal course with a progressive increase 1 and 3 months after surgery, when they all reached their zenith, followed by a gradual decrease 6 and 12 months after surgery.

We identified significant correlations of miRNAs' levels between the diagnosis and 1 month after surgery, and more interestingly we firstly reported a mutual correlation between miR 21-5p and miR 23b-3p at each timing point of plasma collection and between miR 21-5p and miR 34a-5p at diagnosis. A network analysis highlights some possible molecular pathways of interaction in the context of glioma such as cell cycle control (*E2F1*, *E2F3*, *RB1*, *CDK6*) and cell growth (*PTEN*, *PIK3R1*, *IGF1R*).

Eventually, high miRNA circulating levels seem to be good prognosticators in terms of OS and RFS. However, future studies with larger cohorts of patients are needed to support our findings.

7. APPENDIX

Table 1 – Patients' clinical features.

| Variables | | Chi square | p value |
|--------------------------------------|------------------|-------------------|-------------------|
| Gender | | | |
| Males (n, %) | 14 (60.9) | 1.087 | 0.297 |
| Females (n, %) | 9 (39.1) | | |
| mRS | | | |
| 0-2 (n, %) | 21 (91.3) | 15.696 | < 0.001 |
| 3-5 (n, %) | 2 (8.7) | | |
| KPS | | | |
| 70-80 – “Dependent” (n, %) | 5 (21.7) | 7.348 | 0.007 |
| 90-100 – “Independent” (n, %) | 18 (78.3) | | |
| Steroids | | | |
| Yes (n, %) | 21 (91.3) | 15.696 | < 0.001 |
| No (n, %) | 2 (8.7) | | |
| AEDs | | | |
| Yes (n, %) | 10 (43.5) | 0.391 | 0.532 |
| No (n, %) | 13 (56.5) | | |

AEDs: anti-epileptic drugs; KPS: Karnofsky Performance Status; mRS: modified Ranking Score.

Table 2 – Pathological, neuroradiological and surgical tumour's features.

| Variables | | Chi square | p value |
|--|------------------|---------------|-------------------|
| Preoperative MRI | | | |
| Location | | | |
| Right | 9 (39.1) | 4.522 | 0.104 |
| Left | 11 (47.8) | | |
| Midline | 3 (13.1) | | |
| Contrast enhancement | | | |
| Yes (n, %) | 23 (100.0) | - | - |
| No (n, %) | 0 (0.0) | | |
| Necrosis | | | |
| Yes (n, %) | 20 (86.7) | 11.636 | 0.001 |
| No (n, %) | 3 (13.3) | | |
| Primary vs recurrent GBM | | | |
| Primary (n, %) | 19 (82.6) | 9.783 | 0.002 |
| Recurrent (n, %) | 4 (17.4) | | |
| Surgical procedure | | | |
| Craniotomy (n, %) | 21 (91.3) | 15.696 | < 0.001 |
| Frameless neuronavigated biopsy (n, %) | 2 (8.7) | | |
| Intraoperative tumour fluorescence | | | |
| Strong (n, %) | 22 (95.7) | 19.174 | < 0.001 |
| Weak (n, %) | 1 (4.3) | | |
| Absent (n, %) | 0 | | |
| IDH1/IDH2 status | | | |
| Wild type (n, %) | 22 (95.7) | 19.174 | < 0.001 |
| Mutant (n, %) | 1 (4.3) | | |
| MGMT promoter methylation % (mean, SD) | | | |
| | 17.4 (17.3) | | |
| MGMT promoter methylation status (n, %) | | | |
| Un-methylated (< 9 %) | 12 (52.2) | 4.261 | 0.119 |
| Methylated (9-29 %) | 4 (17.4) | | |
| Hyper-methylated (> 29%) | 7 (30.4) | | |
| ATRX expression | | | |
| Lost (n, %) | 3 (13.0) | 13.762 | < 0.001 |
| Retained (n, %) | 20 (87.0) | | |
| p53 expression | | | |
| Absent (n, %) | 2 (8.7) | 7.143 | 0.028 |
| Intermediate (n, %) | 13 (56.5) | | |
| Strong (n, %) | 8 (34.8) | | |
| Ki67 % (mean, SD) | | | |
| | 41.3 (19.8) | | |
| Early postoperative MRI | | | |
| Extent of tumour resection [^] | | | |
| NTR (n, %) | 15 (71.4) | 3.857 | 0.050 |
| STR (n, %) | 6 (28.6) | | |
| Contrast enhancement of the surgical cavity [^] | | | |

| | | | |
|--------------------------------|------------------|---------------|--------------|
| Absent (n, %) | 11 (52.4) | 1.143 | 0.565 |
| Linear (n, %) | 6 (28.6) | | |
| Nodular (n, %) | 4 (19.0) | | |
| | | | |
| RT + concomitant TMZ § | | | |
| Yes (n, %) | 17 (89.5) | 11.842 | 0.001 |
| No (n, %) | 2 (10.5) | | |
| | | | |
| RT protocol | | | |
| Standard RT (n, %) | 15 (88.2) | 9.941 | 0.002 |
| Hypo-fractionated RT (n, %) | 2 (11.8) | | |
| | | | |
| Adjuvant chemotherapy § | | | |
| Yes (n, %) | 12 (52.2) | 0.043 | 0.835 |
| No (n, %) | 11 (47.8) | | |

ATRX: alpha thalassemia/mental retardation syndrome X-linked; IDH: isocitrate dehydrogenase; MGMT: O-6-methylguanine-DNA-methyltransferase; NTR: near total resection; STR: subtotal resection; TMZ: temozolomide.

^ Extent of tumour resection assessed in 21 patients who underwent craniotomy for tumour resection.

§ In 19 patients with primary tumours.

Table 3 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis and clinical, neuroradiological and pathological features.

| | miR 21-5p at diagnosis | | | miR 23b-3p at diagnosis | | | miR 34a-5p at diagnosis | | |
|--|---------------------------|----------------|--------------|----------------------------|----------------|----------|----------------------------|----------------|----------|
| | R | R ² | <i>p</i> | R | R ² | <i>p</i> | R | R ² | <i>p</i> |
| Gender (M vs F) | 0.107 | 0.011 | 0.626 | 0.229 | 0.052 | 0.294 | -0.074 | 0.005 | 0.738 |
| Age (years) | 0.402 | 0.162 | 0.057 | 0.389 | 0.151 | 0.067 | 0.176 | 0.031 | 0.421 |
| mRS (≤ 2 vs > 2) | 0.186 | 0.035 | 0.395 | 0.186 | 0.035 | 0.395 | 0.186 | 0.035 | 0.395 |
| KPS (70-80 vs 90-100) | -0.175 | 0.031 | 0.425 | -0.270 | 0.073 | 0.212 | -0.374 | 0.140 | 0.079 |
| Steroids (Yes vs No) | -0.419 | 0.176 | 0.054 | -0.396 | 0.157 | 0.062 | -0.396 | 0.157 | 0.062 |
| AEDs (Yes vs No) | -0.185 | 0.034 | 0.398 | -0.146 | 0.021 | 0.507 | -0.073 | 0.005 | 0.741 |
| MRI – Location (right, left, midline) | -0.079 | 0.006 | 0.720 | 0.034 | 0.001 | 0.879 | 0.023 | 0.001 | 0.916 |
| MRI – Necrosis (Yes vs No) | 0.010 | 0.000 | 0.963 | 0.052 | 0.003 | 0.817 | -0.125 | 0.016 | 0.578 |
| MRI – Tumour volume (cc) | 0.019 | 0.000 | 0.932 | 0.076 | 0.006 | 0.729 | -0.221 | 0.049 | 0.312 |
| GBM type (primary vs recurrent) | 0.086 | 0.007 | 0.695 | 0.156 | 0.024 | 0.478 | 0.017 | 0.000 | 0.938 |
| MGMT promoter methylation (%) | -0.273 | 0.075 | 0.207 | -0.362 | 0.131 | 0.089 | 0.107 | 0.011 | 0.628 |
| MGMT promoter methylation (unmethylated vs methylated) | -0.223 | 0.050 | 0.306 | -0.315 | 0.099 | 0.143 | 0.072 | 0.005 | 0.743 |
| ATRX status (Lost vs Retained) | 0.0455 | 0.002 | 0.038 | 0.268 | 0.072 | 0.240 | 0.268 | 0.072 | 0.240 |
| p53 status (Lost vs Intermediate vs Diffuse) | -0.072 | 0.005 | 0.756 | -0.156 | 0.024 | 0.499 | 0.039 | 0.002 | 0.867 |
| Ki67 (%) | -0.331 | 0.110 | 0.123 | -0.324 | 0.105 | 0.131 | -0.180 | 0.032 | 0.412 |

AEDs: anti-epileptic drugs; ATRX: alpha thalassemia/mental retardation syndrome X-linked; KPS: Karnofsky Performance Status; mRS: modified Ranking Score; MGMT: O-6-methylguanine-DNA-methyltransferase.

Table 4 – Multivariate linear regression analysis between miR 21-5p plasmatic concentrations at diagnosis and clinical, pathological and neuroradiological features.

ANOVA: R 0.768; R² 0.589; F 0.836; *p* value 0.626.

| | OR | 95% CI | <i>p</i> value |
|--|-----------|------------------|-----------------------|
| Gender (M vs F) | 5.68 | -182.70 – 194.08 | 0.948 |
| Age (years) | 4.19 | -3.72 – 12.10 | 0.265 |
| mRS (≤ 2 vs > 2) | -12.97 | -384.60 – 358.65 | 0.940 |
| KPS (70-80 vs 90-100) | -60.02 | -331.99 – 211.94 | 0.634 |
| Steroids (Yes vs No) | -207.32 | -535.08 – 120.44 | 0.189 |
| AEDs (Yes vs No) | 2.24 | -204.46 – 208.94 | 0.981 |
| MRI Necrosis (Yes vs No) | 52.26 | -166.10 – 270.62 | 0.606 |
| GBM type (Primary vs Recurrent) | -105.38 | -326.77 – 116.00 | 0.314 |
| MGMT promoter methylation (unmethylated vs methylated) | -125.80 | -301.52 – 49.93 | 0.142 |
| ATRX status (Lost vs Retained) | 80.318 | -262.23 – 422.85 | 0.597 |
| Ki67 (%) | -1.24 | -5.22 – 2.75 | 0.505 |

AEDs: anti-epileptic drugs; ATRX: alpha thalassemia/mental retardation syndrome X-linked; KPS: Karnofsky Performance Status; mRS: modified Ranking Score; MGMT: O-6-methylguanine-DNA-methyltransferase; OR: odd ratio.

Table 5 – Multivariate linear regression analysis between miR 23b-3p plasmatic concentrations at diagnosis and clinical, pathological and neuroradiological features.

ANOVA: R 0.724; R² 0.525; adjusted R² -0.290; F 0.644; *p* value 0.760.

| | OR | 95% CI | <i>p</i> value |
|--|-----------|----------------|-----------------------|
| Gender (M vs F) | 2.62 | -17.50 – 22.74 | 0.767 |
| Age (years) | 0.48 | -0.49 – 1.45 | 0.279 |
| mRS (≤ 2 vs > 2) | -0.58 | -64.44 – 64.33 | 0.998 |
| KPS (70-80 vs 90-100) | -4.58 | -38.25 – 29.09 | 0.757 |
| Steroids (Yes vs No) | -8.27 | -55.43 – 38.88 | 0.691 |
| AEDs (Yes vs No) | 3.17 | -23.42 – 29.65 | 0.789 |
| MRI Necrosis (Yes vs No) | 6.75 | -23.58 – 37.08 | 0.615 |
| GBM type (Primary vs Recurrent) | 0.83 | -43.80 – 45.47 | 0.966 |
| MGMT promoter methylation (unmethylated vs methylated) | -12.45 | -33.09 – 8.20 | 0.197 |
| ATRX status (Lost vs Retained) | 2.88 | -25.42 – 31.18 | 0.817 |
| Ki67 (%) | -0.10 | -0.56 – 0.54 | 0.966 |

AEDs: anti-epileptic drugs; ATRX: alpha thalassemia/mental retardation syndrome X-linked; KPS: Karnofsky Performance Status; mRS: modified Ranking Score; MGMT: O-6-methylguanine-DNA-methyltransferase; OR: odd ratio.

Table 6 – Multivariate linear regression analysis between miR 34a-5p plasmatic concentrations at diagnosis and clinical, pathological and neuroradiological features.

ANOVA: R 0.881; R² 0.776; adjusted R² 0.392; F 2.022; p value 0.179.

| | OR | 95% CI | p value |
|--|-----------|---------------|----------------|
| Gender (M vs F) | -0.80 | -2.26 – 0.67 | 0.238 |
| Age (years) | 0.02 | -0.05 – 0.09 | 0.506 |
| mRS (≤ 2 vs > 2) | 2.22 | -2.46 – 6.90 | 0.299 |
| KPS (70-80 vs 90-100) | -1.32 | -3.77 – 1.13 | 0.244 |
| Steroids (Yes vs No) | 0.77 | -2.66 – 4.20 | 0.614 |
| AEDs (Yes vs No) | -0.09 | -2.02 – 1.84 | 0.912 |
| MRI Necrosis (Yes vs No) | -1.58 | -3.79 – 0.62 | 0.134 |
| GBM type (Primary vs Recurrent) | 2.09 | -1.16 – 5.33 | 0.173 |
| MGMT promoter methylation (unmethylated vs methylated) | -0.21 | -1.71 – 1.29 | 0.751 |
| ATR status (Lost vs Retained) | 1.04 | -1.02 – 3.10 | 0.271 |
| Ki67 (%) | 0.02 | -0.02 – 0.06 | 0.274 |

AEDs: anti-epileptic drugs; ATRX: alpha thalassemia/mental retardation syndrome X-linked; KPS: Karnofsky Performance Status; mRS: modified Ranking Score; MGMT: O-6-methylguanine-DNA-methyltransferase; OR: odd ratio.

Table 7 – Comparison of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 1 month after surgery (FU1) according to extent of tumour resection (16 patients).

| | miR 21-5p FU1 (mean, SD) | <i>p</i> value | miR 23b-3p FU1 (mean, SD) | <i>p</i> value | miR 34a-5p FU1 (mean, SD) | <i>p</i> value |
|---|---|----------------|--|----------------|--|----------------|
| Surgical procedure (n, %) | | | | | | |
| Craniotomy (14, 87.5) | 235.69 (375.09) | 0.065 | 17.31 (34.69) | 0.124 | 2.25 (1.79) | 0.465 |
| Neuronavigated biopsy (2, 22.5) | 30.00 (30.41) | | 1.89 (2.14) | | 1.48 (1.02) | |
| Extent of tumour resection (n, %)^ | | | | | | |
| NTR (10, 71.4) | 277.22 (427.14) | 0.410 | 21.96 (40.29) | 0.256 | 2.70 (1.93) | 0.040 |
| STR (4, 28.6) | 131.87 (205.46) | | 5.67 (9.57) | | 1.13 (0.61) | |
| Postoperative contrast enhancement (n, %)^ | | | | | | |
| Absent (6, 42.9) | 179.55 (179.03) | 0.362 | 11.21 (14.94) | 0.388 | 3.31 (2.18) | 0.143 |
| Linear (3, 21.4) | 38.30 (31.10) | | 1.04 (0.80) | | 1.08 (0.88) | |
| Nodular (5, 35.7) | 421.50 (582.75) | | 34.38 (54.89) | | 1.68 (0.99) | |

NTR: near total resection; STR: subtotal resection.

^ Extent of resection data analysed on 14 patients who underwent craniotomy.

Table 8 – Comparison of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 3 months after surgery (FU3) according to extent of tumour resection and adjuvant therapies (8 patients).

| | miR 21-5p FU3 (mean, SD) | <i>p</i> value | miR 23b-3p FU3 (mean, SD) | <i>p</i> value | miR 34a-5p FU3 (mean, SD) | <i>p</i> value |
|---|---|----------------|--|----------------|--|----------------|
| Surgical procedure (n, %) | | | | | | |
| Craniotomy (8, 100.0) | 356.15 (303.89) | - | 22.26 (19.04) | - | 4.79 (5.17) | - |
| Neuronavigated biopsy (0, 0) | - | | - | | - | |
| Extent of tumour resection (n, %) | | | | | | |
| NTR (7, 87.5) | 333.74 (321.02) | 0.620 | 20.97 (20.18) | 0.649 | 4.91 (5.57) | 0.870 |
| STR (1, 22.5) | 513.00 | | 31.30 | | 3.90 | |
| Postoperative contrast enhancement (n, %) | | | | | | |
| Absent (4, 50.0) | 204.45 (153.88) | 0.105 | 12.48 (10.43) | 0.398 | 3.95 (4.83) | 0.393 |
| Linear (2, 25.0) | 731.50 (309.01) | | 34.00 (3.82) | | 9.25 (7.57) | |
| Nodular (2, 25.0) | 284.20 (309.43) | | 30.10 (37.62) | | 2.00 (1.13) | |
| RT + concomitant TMZ (n, %) | | | | | | |
| Yes (6, 75.0) | 377.05 (328.50) | 0.778 | 24.02 (20.27) | 0.718 | 5.48 (5.87) | 0.339 |
| No (2, 25.0) | 293.45 (310.49) | | 17.00 (20.22) | | 2.70 (1.70) | |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 9 – Comparison of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 6 months after surgery (FU6) according to extent of tumour resection and adjuvant therapies (9 patients).

| | miR 21-5p FU6 (mean, SD) | <i>p</i> value | miR 23b-3p FU6 (mean, SD) | <i>p</i> value | miR 34a-5p FU6 (mean, SD) | <i>p</i> value |
|---|---|----------------|--|----------------|--|----------------|
| Surgical procedure (n, %) | | | | | | |
| Craniotomy (9, 100.0) | 183.74 (141.87) | - | 7.77 (6.16) | - | 2.32 (2.43) | - |
| Neuronavigated biopsy (0, 0) | - | | - | | - | |
| Extent of tumour resection (n, %) | | | | | | |
| NTR (8, 88.9) | 191.71 (149.50) | 0.665 | 8.03 (6.53) | 0.747 | 2.15 (2.53) | 0.582 |
| STR (1, 11.1) | 120.00 | | 5.70 | | 3.70 | |
| Postoperative contrast enhancement (n, %) | | | | | | |
| Absent (6, 66.7) | 199.72 (153.10) | 0.664 | 8.46 (7.28) | 0.668 | 1.60 (1.33) | 0.228 |
| Linear (3, 33.3) | 151.80 (140.03) | | 6.40 (3.90) | | 3.77 (3.80) | |
| Nodular (0, 0) | - | | - | | - | |
| RT + concomitant TMZ (n, %) | | | | | | |
| Yes (7, 77.8) | 216.33 (142.89) | 0.122 | 9.11 (6.23) | 0.187 | 2.29 (2.70) | 0.926 |
| No (2, 22.2) | 69.70 (71.13) | | 3.08 (3.71) | | 2.45 (1.77) | |
| Adjuvant chemotherapy (n, %) | | | | | | |
| Yes (8, 88.9) | 164.71 (138.84) | 0.283 | 7.09 (6.21) | 0.385 | 2.53 (2.51) | 0.515 |
| No (1, 11.1) | 336.00 | | 13.20 | | 0.70 | |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 10 – Comparison of miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 12 months after surgery (FU12) according to extent of tumour resection and adjuvant therapies (5 patients).

| | miR 21-5p FU12 (mean, SD) | <i>p</i> value | miR 23b-3p FU12 (mean, SD) | <i>p</i> value | miR 34a-5p FU12 (mean, SD) | <i>p</i> value |
|---|--|----------------|---|----------------|---|----------------|
| Surgical procedure (n, %) | | | | | | |
| Craniotomy (5, 100.0) | 27.42 (17.27) | - | 0.91 (0.42) | - | 1.56 (0.34) | - |
| Neuronavigated biopsy (0, 0) | - | | - | | - | |
| Extent of tumour resection (n, %) | | | | | | |
| NTR (5, 100.0) | 27.42 (17.27) | - | 0.91 (0.42) | - | 1.56 (0.34) | - |
| STR (0, 0) | - | | - | | - | |
| Postoperative contrast enhancement (n, %) | | | | | | |
| Absent (4, 80.0) | 32.93 (13.99) | 0.177 | 0.92 (0.48) | 0.919 | 1.58 (0.39) | 0.873 |
| Linear (1, 20.0) | 5.40 | | 0.86 | | 1.50 | |
| Nodular (0, 0) | - | | - | | - | |
| RT + concomitant TMZ (n, %) | | | | | | |
| Yes (5, 100.0) | 27.42 (17.27) | - | 0.91 (0.42) | - | 1.56 (0.34) | - |
| No (0, 0) | - | | - | | - | |
| Adjuvant chemotherapy (n, %) | | | | | | |
| Yes (4, 80.0) | 25.30 (19.18) | 0.655 | 0.76 (0.29) | 0.113 | 1.60 (0.37) | 0.665 |
| No (1, 20.0) | 35.90 | | 1.50 | | 1.40 | |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 11 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 1 month after surgery (FU1) and extent of tumour resection.

| | miR 21-5p FU1 | | | miR 23b-3p FU1 | | | Mir 34a-5p FU1 | | |
|--|------------------|----------------|---------|-------------------|----------------|---------|-------------------|----------------|---------|
| | R | R ² | p value | R | R ² | p value | R | R ² | p value |
| Surgical procedure (craniotomy vs biopsy) | -0.328 | 0.108 | 0.215 | -0.164 | 0.027 | 0.544 | -0.103 | 0.011 | 0.705 |
| Extent of tumour resection (NTR vs STR) | -0.336 | 0.113 | 0.203 | -0.238 | 0.057 | 0.374 | -0.393 | 0.154 | 0.132 |
| Extent of tumour resection (percentage) | 0.199 | 0.040 | 0.459 | 0.185 | 0.034 | 0.492 | 0.253 | 0.064 | 0.344 |
| Volume of postoperative tumour remnant (cc) | -0.153 | 0.023 | 0.572 | -0.160 | 0.027 | 0.553 | -0.199 | 0.040 | 0.460 |
| Postoperative contrast enhancement (absent vs linear vs nodular) | -0.166 | 0.028 | 0.538 | 0.0 | 0 | 1.0 | -0.325 | 0.106 | 0.219 |

NTR: near total resection; STR: subtotal resection.

Table 12 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 3 months after surgery (FU3), extent of tumour resection and postoperative concomitant RT and chemotherapy.

| | miR 21-5p FU3 | | | miR 23b-3p FU3 | | | Mir 34a-5p FU3 | | |
|---|------------------|----------------|---------|-------------------|----------------|---------|-------------------|----------------|---------|
| | R | R ² | p value | R | R ² | p value | R | R ² | p value |
| Surgical procedure (craniotomy vs biopsy) | - | - | - | - | - | - | - | - | - |
| Extent of tumour resection (NTR vs STR) | 0.412 | 0.167 | 0.310 | 0.247 | 0.061 | 0.555 | 0.249 | 0.062 | 0.552 |
| Extent of tumour resection (percentage) | -0.398 | 0.158 | 0.328 | - 0.495 | 0.245 | 0.212 | 0.016 | 0.000 | 0.970 |
| Volume of postoperative tumour remnant (cc) | 0.324 | 0.105 | 0.434 | 0.615 | 0.378 | 0.105 | -0.144 | 0.021 | 0.735 |
| Postoperative contrast enhancement (absent vs linear vs nodular) | 0.206 | 0.042 | 0.625 | 0.437 | 0.191 | 0.279 | 0.0 | 0 | 1.0 |
| RT + concomitant TMZ (yes vs no) | 0.0 | 0 | 1.0 | 0.252 | 0.064 | 0.547 | 0.063 | 0.004 | 0.881 |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 13 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 6 months after surgery (FU6), extent of tumour resection, postoperative concomitant RT and chemotherapy and adjuvant chemotherapy.

| | miR 21-5p FU6 | | | miR 23b-3p FU6 | | | Mir 34a-5p FU6 | | |
|---|------------------|----------------|---------|-------------------|----------------|---------|-------------------|----------------|---------|
| | R | R ² | p value | R | R ² | p value | R | R ² | p value |
| Surgical procedure (craniotomy vs biopsy) | - | - | - | - | - | - | - | - | - |
| Extent of tumour resection (NTR vs STR) | -0.137 | 0.019 | 0.725 | - 0.137 | 0.019 | 0.725 | 0.413 | 0.171 | 0.270 |
| Extent of tumour resection (percentage) | 0.152 | 0.023 | 0.697 | 0.127 | 0.016 | 0.745 | -0.345 | 0.119 | 0.363 |
| Volume of postoperative tumour remnant (cc) | -0.198 | 0.039 | 0.609 | - 0.156 | 0.024 | 0.689 | 0.259 | 0.067 | 0.500 |
| Postoperative contrast enhancement (absent vs linear vs nodular) | -0.091 | 0.008 | 0.815 | - 0.091 | 0.008 | 0.815 | 0.321 | 0.103 | 0.400 |
| RT + concomitant TMZ (yes vs no) | 0.414 | 0.171 | 0.268 | 0.414 | 0.171 | 0.268 | -0.208 | 0.043 | 0.591 |
| Adjuvant chemotherapy (yes vs no) | -0.411 | 0.169 | 0.272 | - 0.411 | 0.169 | 0.272 | 0.275 | 0.076 | 0.474 |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 14 – Correlation analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 12 months after surgery (FU12), extent of tumour resection, postoperative concomitant RT and chemotherapy and adjuvant chemotherapy.

| | miR 21-5p FU12 | | | miR 23b-3p FU12 | | | Mir 34a-5p FU12 | | |
|--|-------------------|----------------|---------|--------------------|----------------|---------|--------------------|----------------|---------|
| | R | R ² | p value | R | R ² | p value | R | R ² | p value |
| Surgical procedure (craniotomy vs biopsy) | - | - | - | - | - | - | - | - | - |
| Extent of tumour resection (NTR vs STR) | - | - | - | - | - | - | - | - | - |
| Extent of tumour resection (percentage) | 0.713 | 0.508 | 0.177 | 0.064 | 0.004 | 0.919 | 0.100 | 0.010 | 0.873 |
| Volume of postoperative tumour remnant (cc) | -0.713 | 0.508 | 0.177 | -0.064 | 0.004 | 0.919 | -0.100 | 0.010 | 0.873 |
| Postoperative contrast enhancement (absent vs linear vs nodular) | -0.707 | 0.500 | 0.182 | 0 | 0 | 1.0 | 0 | 0 | 1.0 |
| RT + concomitant TMZ (yes vs no) | - | - | - | - | - | - | - | - | - |
| Adjuvant chemotherapy (yes vs no) | -0.354 | 0.125 | 0.559 | -0.707 | 0.500 | 0.182 | 0.354 | 0.125 | 0.559 |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 15 – Multivariate linear regression analysis between miR 21-5p plasmatic concentrations 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) after surgery, extent of tumour resection, postoperative concomitant RT and chemotherapy and adjuvant chemotherapy.

FU1: ANOVA: R 0.600; R² 0.360; adjusted R² 0.040; F 1.125; *p* value 0.407.

FU3: ANOVA: R 0.962; R² 0.925; adjusted R² 0.738; F 4.952; *p* value 0.177.

FU6: ANOVA: R 0.770; R² 0.592; adjusted R² -0.087; F 0.872; *p* value 0.584.

FU12: ANOVA: R 0.719; R² 0.518; adjusted R² 0.035; F 1.073; *p* value 0.482.

| | OR | 95% CI | <i>p</i> value |
|---|----------|--------------------|----------------|
| miR 21-5p FU1 | | | |
| Extent of resection (NTR vs STR) | -976.77 | -2343.43 – 389.89 | 0.142 |
| Extent of resection (percentage) | -29.39 | -104.71 – 45.92 | 0.405 |
| Volume of postoperative tumour remnant (cc) | -0.56 | -65.92 – 64.80 | 0.985 |
| Contrast enhancement (linear vs nodular) | 206.53 | -98.89 – 511.94 | 0.163 |
| miR 21-5p FU3 | | | |
| Extent of resection (NTR vs STR) | -3309.88 | -7096.67 – 476.92 | 0.064 |
| Extent of resection (percentage) | -795.65 | -1570.89 – -20.42 | 0.048 |
| Volume of postoperative tumour remnant (cc) | -1163.49 | -2782.72 – 455.75 | 0.091 |
| Contrast enhancement (linear vs nodular) | -311.35 | -905.22 – 282.52 | 0.153 |
| RT + concomitant TMZ (yes vs no) | 174.07 | -598.14 – 946.27 | 0.434 |
| miR 21-5p FU6 | | | |
| Extent of resection (NTR vs STR) | -2405.07 | -8632.84 – 3822.71 | 0.307 |
| Extent of resection (percentage) | -392.74 | -1344.84 – 559.36 | 0.281 |
| Volume of postoperative tumour remnant (cc) | - | - | - |
| Contrast enhancement (linear vs nodular) | -532.29 | -1786.48 – 721.90 | 0.270 |
| RT + concomitant TMZ (yes vs no) | 191.33 | -334.96 – 717.61 | 0.331 |
| Adjuvant chemotherapy (yes vs no) | -125.28 | -651.56 – 401.01 | 0.504 |
| miR 21-5p FU12 | | | |
| Extent of resection (NTR vs STR) | - | - | - |
| Extent of resection (percentage) | - | - | - |
| Volume of postoperative tumour remnant (cc) | - | - | - |
| Contrast enhancement (linear vs nodular) | -26.53 | -110.83 – 57.76 | 0.308 |
| RT + concomitant TMZ (yes vs no) | - | - | - |
| Adjuvant chemotherapy (yes vs no) | -3.97 | -88.26 – 80.33 | 0.858 |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 16 – Multivariate linear regression analysis between miR 23b-3p plasmatic concentrations 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) after surgery, extent of tumour resection, postoperative concomitant RT and chemotherapy and adjuvant chemotherapy.

FU1: ANOVA: R 0.587; R² 0.345; adjusted R² 0.107; F 1.448; *p* value 0.283.

FU3: ANOVA: R 0.960; R² 0.922; adjusted R² 0.725; F 4.696; *p* value 0.185.

FU6: ANOVA: R 0.650; R² 0.422; adjusted R² -0.542; F 0.438; *p* value 0.804.

FU12: ANOVA: R 0.800; R² 0.640; adjusted R² 0.280; F 1.777; *p* value 0.360.

| | OR | 95% CI | <i>p</i> value |
|---|---------|------------------|----------------|
| miR 23b-3p FU1 | | | |
| Extent of resection (NTR vs STR) | -46.26 | -105.70 – 13.18 | 0.115 |
| Extent of resection (percentage) | 0.23 | -0.91 – 1.38 | 0.665 |
| Volume of postoperative tumour remnant (cc) | 0.83 | -4.35 – 6.00 | 0.732 |
| Contrast enhancement (linear vs nodular) | 24.68 | -0.92 – 50.28 | 0.057 |
| miR 23b-3p FU3 | | | |
| Extent of resection (NTR vs STR) | -161.49 | -404.65 – 81.66 | 0.104 |
| Extent of resection (percentage) | -26.95 | -76.73 – 22.83 | 0.145 |
| Volume of postoperative tumour remnant (cc) | 2.30 | -101.68 – 106.27 | 0.933 |
| Contrast enhancement (linear vs nodular) | -22.54 | -60.67 – 15.59 | 0.126 |
| RT + concomitant TMZ (yes vs no) | 13.03 | -36.55 – 62.62 | 0.375 |
| miR 23b-3p FU6 | | | |
| Extent of resection (NTR vs STR) | -63.70 | -385.76 – 258.37 | 0.574 |
| Extent of resection (percentage) | -11.01 | -60.25 – 38.22 | 0.528 |
| Volume of postoperative tumour remnant (cc) | - | - | - |
| Contrast enhancement (linear vs nodular) | -16.24 | -81.10 – 48.62 | 0.484 |
| RT + concomitant TMZ (yes vs no) | 8.82 | -18.39 – 36.04 | 0.378 |
| Adjuvant chemotherapy (yes vs no) | -3.93 | -31.14 – 23.29 | 0.677 |
| miR 23b-3p FU12 | | | |
| Extent of resection (NTR vs STR) | - | - | - |
| Extent of resection (percentage) | - | - | - |
| Volume of postoperative tumour remnant (cc) | - | - | - |
| Contrast enhancement (linear vs nodular) | 0.133 | -1.64 – 1.90 | 0.777 |
| RT + concomitant TMZ (yes vs no) | - | - | - |
| Adjuvant chemotherapy (yes vs no) | -0.773 | -2.54 – 1.00 | 0.201 |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 17 – Multivariate linear regression analysis between miR 34a-5p plasmatic concentrations 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) after surgery, extent of tumour resection, postoperative concomitant RT and chemotherapy and adjuvant chemotherapy.

FU1: ANOVA: R 0.515; R² 0.266; adjusted R² -0.001; F 0.994; *p* value 0.451.

FU3: ANOVA: R 0.817; R² 0.668; adjusted R² -0.163; F 0.803; *p* value 0.636.

FU6: ANOVA: R 0.918; R² 0.843; adjusted R² 0.581; F 3.222; *p* value 0.182.

FU12: ANOVA: R 0.317; R² 0.100; adjusted R² -0.799; F 0.111; *p* value 0.900.

| | OR | 95% CI | <i>p</i> value |
|---|--------|-----------------|----------------|
| miR 34a-5p FU1 | | | |
| Extent of resection (NTR vs STR) | -1.39 | -4.68 – 1.90 | 0.372 |
| Extent of resection (percentage) | 0.01 | -0.06 – 0.07 | 0.872 |
| Volume of postoperative tumour remnant (cc) | 0.08 | -0.21 – 0.37 | 0.551 |
| Contrast enhancement (linear vs nodular) | -0.57 | -1.98 – 0.85 | 0.398 |
| miR 34a-5p FU3 | | | |
| Extent of resection (NTR vs STR) | -43.87 | -179.62 – 91.88 | 0.299 |
| Extent of resection (percentage) | -11.68 | -39.47 – 16.11 | 0.212 |
| Volume of postoperative tumour remnant (cc) | -22.32 | -80.36 – 35.73 | 0.240 |
| Contrast enhancement (linear vs nodular) | -3.89 | -25.18 – 17.40 | 0.514 |
| RT + concomitant TMZ (yes vs no) | 3.27 | -24.42 – 30.95 | 0.662 |
| miR 34a-5p FU6 | | | |
| Extent of resection (NTR vs STR) | -69.91 | -135.99 – -3.84 | 0.043 |
| Extent of resection (percentage) | -10.87 | -20.97 – -0.77 | 0.042 |
| Volume of postoperative tumour remnant (cc) | - | - | - |
| Contrast enhancement (linear vs nodular) | -11.67 | -24.97 – 1.64 | 0.068 |
| RT + concomitant TMZ (yes vs no) | 0.73 | -4.86 – 6.31 | 0.707 |
| Adjuvant chemotherapy (yes vs no) | 1.23 | -4.36 – 6.81 | 0.535 |
| miR 34a-5p FU12 | | | |
| Extent of resection (NTR vs STR) | - | - | - |
| Extent of resection (percentage) | - | - | - |
| Volume of postoperative tumour remnant (cc) | - | - | - |
| Contrast enhancement (linear vs nodular) | -0.133 | -2.37 – 2.11 | 0.822 |
| RT + concomitant TMZ (yes vs no) | - | - | - |
| Adjuvant chemotherapy (yes vs no) | 0.233 | -2.01 – 2.47 | 0.698 |

NTR: near total resection; RT: radiotherapy; STR: subtotal resection; TMZ: temozolomide.

Table 18 – Multivariate linear regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis.

miR 21-5p: ANOVA: R 0.950; R² 0.902; adjusted R² 0.893; F 92.341; p value <0.001.

miR 23b-3p: ANOVA: R 0.943; R² 0.889; adjusted R² 0.878; F 80.079; p value <0.001.

miR 34a-5p: ANOVA: R 0.563; R² 0.317; adjusted R² 0.249; F 4.649; p value 0.022.

| | OR | 95% CI | p value |
|--------------------------------|--------------|---------------------|------------------|
| miR 21-5p at diagnosis | | | |
| miR 23b-3p at diagnosis | 11.50 | 9.53 – 13.47 | <0.001 |
| miR 34a-5p at diagnosis | 23.80 | 5.57 – 42.02 | 0.013 |
| | | | |
| miR 23b-3p at diagnosis | | | |
| miR 21-5p at diagnosis | 0.08 | 0.06 – 0.09 | <0.001 |
| miR 34a-5p at diagnosis | -1.55 | -3.13 – 0.40 | 0.056 |
| | | | |
| miR 34a-5p at diagnosis | | | |
| miR 21-5p at diagnosis | 0.01 | 0.00 – 0.02 | 0.013 |
| miR 23b-3p at diagnosis | -0.11 | -0.23 – 0.00 | 0.056 |

Table 19 – Multivariate linear regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 1 month (FU1) after surgery with miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis and 1 month (FU1) after surgery.

miR 21-5p: ANOVA: R 0.988; R² 0.976; adjusted R² 0.964; F 80.547; p value <0.001.

miR 23b-3p: ANOVA: R 0.989; R² 0.978; adjusted R² 0.967; F 88.313; p value <0.001.

miR 34a-5p: ANOVA: R 0.696; R² 0.485; adjusted R² 0.227; F 1.882; p value 0.185.

| | OR | 95% CI | p value |
|--------------------------|--------------|---------------------|------------------|
| miR 21-5p at FU1 | | | |
| miR 21-5p at diagnosis | 0.132 | -0.82 – 1.09 | 0.763 |
| miR 23b-3p at diagnosis | -2.22 | -16.67 – 12.22 | 0.739 |
| miR 34a-5p at diagnosis | -11.53 | -60.99 – 37.94 | 0.615 |
| miR 23b-3p at FU1 | 10.83 | 8.34 – 13.32 | <0.001 |
| miR 34a-5p at FU1 | 11.21 | -19.67 – 42.09 | 0.437 |
| miR 23b-3p at FU1 | | | |
| miR 21-5p at diagnosis | -0.02 | -0.11 – 0.06 | 0.571 |
| miR 23b-3p at diagnosis | 0.53 | -0.69 – 1.75 | 0.353 |
| miR 34a-5p at diagnosis | 0.55 | -3.84 – 4.93 | 0.788 |
| miR 21-5p at FU1 | 0.08 | 0.06 – 0.10 | <0.001 |
| miR 34a-5p at FU1 | -0.54 | -3.32 – 2.23 | 0.672 |
| miR 34a-5p at FU1 | | | |
| miR 21-5p at diagnosis | -0.01 | -0.03 – 0.01 | 0.404 |
| miR 23b-3p at diagnosis | 0.05 | -0.27 – 0.37 | 0.737 |
| miR 34a-5p at diagnosis | 1.00 | 0.15 – 1.86 | 0.025 |
| miR 21-5p at FU1 | 0.01 | -0.01 – 0.02 | 0.437 |
| miR 23b-3p at FU1 | -0.03 | -0.21 – 0.14 | 0.672 |

Table 20 – Multivariate linear regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 3 months (FU3) after surgery and miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis, 1 (FU1) and 3 months (FU3) after surgery.

miR 21-5p: ANOVA: R 0.999; R² 0.999; adjusted R² 0.994; F 214.546; *p* value 0.051.

miR 23b-3p: ANOVA: R 0.999; R² 0.997; adjusted R² 0.986; F 88.945; *p* value 0.079.

miR 34a-5p: ANOVA: R 0.992; R² 0.984; adjusted R² 0.918; F 14.950; *p* value 0.191.

| | OR | 95% CI | <i>p</i> value |
|--------------------------|---------|-----------------|----------------|
| miR 21-5p at FU3 | | | |
| miR 21-5p at diagnosis | -0.33 | -3.13 – 2.46 | 0.372 |
| miR 21-5p at FU1 | -4.62 | -9.68 – 0.44 | 0.055 |
| miR 23b-3p at FU3 | 121.16 | 1.55 – 240.77 | 0.049 |
| miR 34a-5p at FU3 | -310.05 | -666.09 – 47.99 | 0.058 |
| miR 23b-3p at FU3 | | | |
| miR 23b-3p at diagnosis | 0.02 | -3.06 – 3.10 | 0.954 |
| miR 23b-3p at FU1 | 0.39 | -0.46 – 1.24 | 0.109 |
| miR 21-5p at FU3 | 0.01 | -0.09 – 0.10 | 0.457 |
| miR 34a-5p at FU3 | 2.35 | -4.63 – 9.33 | 0.146 |
| miR 34a-5p at FU3 | | | |
| miR 34a-5p at diagnosis | 4.23 | -16.71 – 25.17 | 0.237 |
| miR 34a-5p at FU1 | -1.82 | -13.93 – 10.29 | 0.307 |
| miR 21-5p at FU3 | 0.03 | -0.05 – 0.10 | 0.138 |
| miR 23b-3p at FU3 | -0.15 | -0.86 – 0.57 | 0.237 |

Table 21 – Multivariate linear regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 6 months (FU6) after surgery and miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 3 (FU3) and 6 months (FU6) after surgery.

miR 21-5p: ANOVA: R 0.992; R² 0.983; adjusted R² 0.934; F 19.735; *p* value 0.164.

miR 23b-3p: ANOVA: **R 1.000; R² 1.000; adjusted R² 0.999; F 1098.073; *p* value 0.022.**

miR 34a-5p: ANOVA: R 0.992; R² 0.984; adjusted R² 0.936; F 20.662; *p* value 0.160.

| | OR | 95% CI | <i>p</i> value |
|--------------------------|-------------|--------------------|----------------|
| miR 21-5p at FU6 | | | |
| miR 21-5p at FU3 | -0.22 | -5.18 – 4.74 | 0.677 |
| miR 23b-3p at FU6 | 38.66 | -49.83 – 127.15 | 0.113 |
| miR 34a-5p at FU6 | 10.24 | -535.82 – 556.31 | 0.851 |
| | | | |
| miR 23b-3p at FU6 | | | |
| miR 23b-3p at FU3 | 0.10 | -0.08 – 0.28 | 0.087 |
| miR 21-5p at FU6 | 0.02 | 0.01 – 0.03 | 0.022 |
| miR 34a-5p at FU6 | 0.05 | -0.69 – 0.78 | 0.581 |
| | | | |
| miR 34a-5p at FU6 | | | |
| miR 34a-5p at FU3 | 0.72 | -1.83 – 3.27 | 0.172 |
| miR 21-5p at FU6 | -0.06 | -0.19 – 0.08 | 0.116 |
| miR 23b-3p at FU6 | 1.33 | -3.09 – 5.75 | 0.163 |

Table 22 – Multivariate linear regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations 12 months (FU12) after surgery.

miR 21-5p: ANOVA: R 0.472; R² 0.223; adjusted R² -0.554; F 0.287; *p* value 0.777.

miR 23b-3p: ANOVA: R 0.449; R² 0.202; adjusted R² -0.596; F 0.253; *p* value 0.798.

miR 34a-5p: ANOVA: R 0.172; R² 0.030; adjusted R² -0.941; F 0.031; *p* value 0.970.

| | OR | 95% CI | <i>p</i> value |
|---------------------------|-------|------------------|----------------|
| miR 21-5p at FU12 | | | |
| miR 23b-3p at FU12 | 18.21 | -92.15 – 128.57 | 0.551 |
| miR 34a-5p at FU12 | -7.82 | -145.70 – 130.06 | 0.830 |
| | | | |
| miR 23b-3p at FU12 | | | |
| miR 21-5p at FU12 | 0.01 | -0.06 – 0.08 | 0.551 |
| miR 34a-5p at FU12 | 0.06 | -3.38 – 3.50 | 0.948 |
| | | | |
| miR 34a-5p at FU12 | | | |
| miR 21-5p at FU12 | -0.01 | -0.07 – 0.06 | 0.830 |
| miR 23b-3p at FU12 | 0.05 | -2.64 – 2.73 | 0.948 |

Table 23 – Node explorer table for miR 21-5p, miR 23b-3p and miR 34a-5p network analysis (miRNet version 2.0).

| ID | Degree | Betweenness |
|----------------|---------------|--------------------|
| hsa-mir-21-5p | 98 | 3393.358 |
| hsa-mir-34a-5p | 89 | 2679.215 |
| hsa-mir-23b-3p | 58 | 1182.427 |
| E2F1 | 3 | 3.134753 |
| TNPO1 | 3 | 3.134753 |
| MTAP | 3 | 3.134753 |
| CPEB3 | 3 | 3.134753 |
| NUFIP2 | 3 | 3.134753 |

Networks: a) *queries*, miRNA: 3; b) *nodes*, gene: 120, miRNA: 3; c) *edges*, 245.

Table 24 – Node explorer table for miR 21-5p and miR 23b-3p network analysis (miRNet version 2.0).

| ID | Degree | Betweenness |
|----------------|---------------|--------------------|
| hsa-mir-21-5p | 36 | 315 |
| hsa-mir-23b-3p | 36 | 315 |
| ACAT1 | 2 | 0.027778 |
| CCNG1 | 2 | 0.027778 |
| HAPLN1 | 2 | 0.027778 |
| E2F1 | 2 | 0.027778 |
| MSH6 | 2 | 0.027778 |
| GTF2I | 2 | 0.027778 |
| HNRNPK | 2 | 0.027778 |
| TNPO1 | 2 | 0.027778 |
| MARCKS | 2 | 0.027778 |
| MTAP | 2 | 0.027778 |
| PIK3R1 | 2 | 0.027778 |
| PTEN | 2 | 0.027778 |
| PTPN14 | 2 | 0.027778 |
| RB1 | 2 | 0.027778 |
| TGIF1 | 2 | 0.027778 |
| TNFAIP3 | 2 | 0.027778 |
| TSNAX | 2 | 0.027778 |
| VHL | 2 | 0.027778 |
| ZMYM2 | 2 | 0.027778 |
| SOCS6 | 2 | 0.027778 |
| DOCK4 | 2 | 0.027778 |
| SLC9A6 | 2 | 0.027778 |
| CPEB3 | 2 | 0.027778 |
| TNRC6B | 2 | 0.027778 |
| RPRD2 | 2 | 0.027778 |
| ATXN10 | 2 | 0.027778 |
| RNF111 | 2 | 0.027778 |
| ETNK1 | 2 | 0.027778 |
| ATF7IP | 2 | 0.027778 |
| NUFIP2 | 2 | 0.027778 |
| RMND5A | 2 | 0.027778 |
| TMEM170A | 2 | 0.027778 |
| UBR3 | 2 | 0.027778 |
| PTPDC1 | 2 | 0.027778 |
| PRR14L | 2 | 0.027778 |
| TRIM59 | 2 | 0.027778 |

Networks: a) *queries*, miRNA: 2; b) *nodes*, gene: 36, miRNA: 2; c) *edges*, 72.

Table 25 – Node explorer table for miR 21-5p and miR 34a-5p network analysis (miRNet version 2.0).

| ID | Degree | Betweenness |
|----------------|---------------|--------------------|
| hsa-mir-21-5p | 67 | 1105.5 |
| hsa-mir-34a-5p | 67 | 1105.5 |
| JAG1 | 2 | 0.014925 |
| CRYBG1 | 2 | 0.014925 |
| APAF1 | 2 | 0.014925 |
| FASLG | 2 | 0.014925 |
| BCL2 | 2 | 0.014925 |
| BNIP2 | 2 | 0.014925 |
| BRCA1 | 2 | 0.014925 |
| CALD1 | 2 | 0.014925 |
| CASP8 | 2 | 0.014925 |
| CDC25A | 2 | 0.014925 |
| CDK6 | 2 | 0.014925 |
| CEBPB | 2 | 0.014925 |
| FOXN3 | 2 | 0.014925 |
| E2F1 | 2 | 0.014925 |
| E2F3 | 2 | 0.014925 |
| ERBB2 | 2 | 0.014925 |
| MKMK2 | 2 | 0.014925 |
| HMGB1 | 2 | 0.014925 |
| IGF1R | 2 | 0.014925 |
| TNPO1 | 2 | 0.014925 |
| CAPRIN1 | 2 | 0.014925 |
| MDM4 | 2 | 0.014925 |
| MAP3K1 | 2 | 0.014925 |
| MTAP | 2 | 0.014925 |
| MYC | 2 | 0.014925 |
| NFKB1 | 2 | 0.014925 |
| PPARA | 2 | 0.014925 |
| RPS7 | 2 | 0.014925 |
| SMARCA4 | 2 | 0.014925 |
| SOX2 | 2 | 0.014925 |
| TGFBR2 | 2 | 0.014925 |
| TP53BP2 | 2 | 0.014925 |
| TPM1 | 2 | 0.014925 |
| VEGFA | 2 | 0.014925 |
| GDF5 | 2 | 0.014925 |
| TNFRSF10B | 2 | 0.014925 |
| BCL10 | 2 | 0.014925 |
| LRRFIP1 | 2 | 0.014925 |
| CLOCK | 2 | 0.014925 |
| CKAP5 | 2 | 0.014925 |
| FAM3C | 2 | 0.014925 |
| STAG2 | 2 | 0.014925 |
| CPEB3 | 2 | 0.014925 |
| NT5C2 | 2 | 0.014925 |
| SYNE2 | 2 | 0.014925 |
| ADNP | 2 | 0.014925 |

| | | |
|---------|---|----------|
| DDAH1 | 2 | 0.014925 |
| GAPVD1 | 2 | 0.014925 |
| RDH11 | 2 | 0.014925 |
| USP47 | 2 | 0.014925 |
| FIGN | 2 | 0.014925 |
| YOD1 | 2 | 0.014925 |
| FAM46A | 2 | 0.014925 |
| SAR1A | 2 | 0.014925 |
| TM9SF3 | 2 | 0.014925 |
| RTN4 | 2 | 0.014925 |
| THOC2 | 2 | 0.014925 |
| ESYT2 | 2 | 0.014925 |
| NUFIP2 | 2 | 0.014925 |
| MOAP1 | 2 | 0.014925 |
| NCAPG | 2 | 0.014925 |
| MTMR9 | 2 | 0.014925 |
| PROSER1 | 2 | 0.014925 |
| LNPK | 2 | 0.014925 |
| KBTBD6 | 2 | 0.014925 |
| SOCS4 | 2 | 0.014925 |
| GXYLT2 | 2 | 0.014925 |

Networks: a) *queries*, miRNA: 2; b) *nodes*, gene: 67, miRNA: 2; c) *edges*, 134.

Table 26 – Cox regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis, 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) after surgery and overall survival.

At diagnosis: chi square 3.044; dof 3; *p* value 0.385

At FU1: chi square 2.637; dof 3; *p* value 0.451

At FU3: chi square 7.841; dof 3; *p* value 0.049

At FU6: chi square 3.304; dof 3; *p* value 0.347

At FU12: chi square 4.886; dof 3; *p* value 0.180

| | HR | 95% CI | <i>p</i> value |
|-------------------------|-----------|---------------|----------------|
| miR 21-5p at diagnosis | 1.019 | 0.991 – 1.019 | 0.503 |
| miR 23b-3p at diagnosis | 0.982 | 0.834 – 1.157 | 0.829 |
| miR 34a-5p at diagnosis | 0.612 | 0.301 – 1.242 | 0.174 |
| miR 21-5p at FU1 | 1.004 | 0.994 – 1.015 | 0.405 |
| miR 23b-3p at FU1 | 0.965 | 0.865 – 1.076 | 0.521 |
| miR 34a-5p at FU1 | 0.715 | 0.453 – 1.127 | 0.148 |
| miR 21-5p at FU3 | 1.005 | 0.995 – 1.015 | 0.336 |
| miR 23b-3p at FU3 | 1.051 | 0.943 – 1.170 | 0.370 |
| miR 34a-5p at FU3 | 0.343 | 0.064 – 1.834 | 0.211 |
| miR 21-5p at FU6 | 1.001 | 0.984 – 1.018 | 0.921 |
| miR 23b-3p at FU6 | 0.809 | 0.478 – 1.368 | 0.428 |
| miR 34a-5p at FU6 | 1.081 | 0.708 – 1.650 | 0.719 |
| miR 21-5p at FU12 | 0.883 | 0.606 – 1.286 | 0.517 |
| miR 23b-3p at FU12 | 27497.812 | 0.000 – 3.499 | 0.472 |
| miR 34a-5p at FU12 | 0.008 | 0.000 – 8.192 | 0.869 |

Table 27 – Cox regression analysis between miR 21-5p, miR 23b-3p and miR 34a-5p plasmatic concentrations at diagnosis, 1 (FU1), 3 (FU3), 6 (FU6) and 12 months (FU12) after surgery and recurrence free survival.

At diagnosis: chi square 2.239; dof 3; *p* value 0.524

At FU1: chi square 1.987; dof 3; *p* value 0.575

At FU3: chi square 3.295; dof 3; *p* value 0.348

At FU6: chi square 2.701; dof 3; *p* value 0.440

At FU12: chi square 4.886; dof 3; *p* value 0.180

| | HR | 95% CI | <i>p</i> value |
|-------------------------|-----------|---------------|----------------|
| miR 21-5p at diagnosis | 1.005 | 0.990 – 1.020 | 0.523 |
| miR 23b-3p at diagnosis | 0.945 | 0.786 – 1.136 | 0.546 |
| miR 34a-5p at diagnosis | 0.606 | 0.288 – 1.272 | 0.185 |
| miR 21-5p at FU1 | 1.006 | 0.995 – 1.017 | 0.321 |
| miR 23b-3p at FU1 | 0.911 | 0.765 – 1.086 | 0.298 |
| miR 34a-5p at FU1 | 0.920 | 0.619 – 1.367 | 0.679 |
| miR 21-5p at FU3 | 1.013 | 0.993 – 1.034 | 0.204 |
| miR 23b-3p at FU3 | 0.801 | 0.578 – 1.109 | 0.181 |
| miR 34a-5p at FU3 | 0.780 | 0.433 – 1.406 | 0.409 |
| miR 21-5p at FU6 | 1.002 | 0.984 – 1.020 | 0.823 |
| miR 23b-3p at FU6 | 0.777 | 0.434 – 1.392 | 0.397 |
| miR 34a-5p at FU6 | 1.268 | 0.774 – 2.077 | 0.345 |
| miR 21-5p at FU12 | 0.883 | 0.606 – 1.286 | 0.517 |
| miR 23b-3p at FU12 | 27497.812 | 0.000 – 3.499 | 0.472 |
| miR 34a-5p at FU12 | 0.008 | 0.000 – 8.192 | 0.869 |

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List of Publications between November 2021 and January 2024

1. “Anatomical study of the mastoid foramina and mastoid emissary veins: classification and application to localizing the sigmoid sinus” – Chaiyamoong A., Schneider K., Iwanaga J., **Donofrio C.A.**, Badaloni F., Fioravanti A., Tubbs R.S. – *Neurosurg Rev.* 2023 Dec 19;47(1):16. doi: 10.1007/s10143-023-02229-4.
2. “An anatomical study of the sigmoid sinus artery: Application to the transmastoid approach” – Iwanaga J., Jackson N., Komune N., Johnson K., **Donofrio C.A.**, Badaloni F., Fioravanti A., Dumont A.S., Tubbs R.S. – *Neurosurg Rev.* 2023 Dec 7;47(1):4. doi: 10.1007/s10143-023-02245-4.
3. “Omics sciences and precision medicine in breast and ovarian cancer” – Bonetti G., Madeo G., Michelini S., Ricci M., Cestari M., Michelini S., Gadler M., Benedetti S., Guerri G., Cristofoli F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Bernini A., Fulcheri E., Stuppia L., Gatta V., Cecchin S., Marceddu G., Bertelli M. – *Clin Ter.* 2023 Nov-Dec;174(Suppl 2(6)):104-118. doi: 10.7417/CT.2023.2477.
4. “Omics sciences and precision medicine in prostate cancer” – Medori M.C., Micheletti C., Gadler M., Benedetti S., Guerri G., Cristofoli F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Bernini A., Fulcheri E., Calogero A.E., Cannarella R., Stuppia L., Gatta V., Cecchin S., Marceddu G., Bertelli M. – *Clin Ter.* 2023 Nov-Dec;174(Suppl 2(6)):95-103. doi: 10.7417/CT.2023.2476.
5. “Omics sciences and precision medicine in pancreas cancer” – Donato K., Micheletti C., Medori M.C., Maltese P.E., Tanzi B., Tezzele S., Mareso C., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Beccari T., Ceccarini M.R., Iaconelli A., Aquilanti B., Matera G., Ahmed R., Stuppia L., Gatta V., Cecchin S., Marceddu G., Bertelli M. – *Clin Ter.* 2023 Nov-Dec;174(Suppl 2(6)):85-94. doi: 10.7417/CT.2023.2475.
6. “Omics sciences and precision medicine in glioblastoma” – Micheletti C., Bonetti G., Madeo G., Gadler M., Benedetti S., Guerri G., Cristofoli F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Manganotti P., Caruso P., Bernini A., Fulcheri E., Stuppia L., Gatta V., Cecchin S., Marceddu G., Bertelli M. – *Clin Ter.* 2023 Nov-Dec;174(Suppl 2(6)):77-84. doi: 10.7417/CT.2023.2474.
7. “Omics sciences and precision medicine in sarcoma” – Bonetti G., Donato K., Dhuli K., Gadler M., Benedetti S., Guerri G., Cristofoli F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Bernini A., Fulcheri E., Cavalca D., Stuppia L., Gatta V., Cristoni S., Cecchin S., Marceddu G., Bertelli M. – *Clin Ter.* 2023 Nov-Dec;174(Suppl 2(6)):68-76. doi: 10.7417/CT.2023.2473.
8. “Omics sciences and precision medicine in colon cancer” – Madeo G., Bonetti G., Gadler M., Benedetti S., Guerri G., Cristofoli F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio

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9. “Omics sciences and precision medicine in kidney cancer” – Dhuli K., Micheletti C., Medori M.C., Maltese P.E., Tanzi B., Tezzele S., Mareso C., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Beccari T., Ceccarini M.R., Stuppia L., Gatta V., Cristoni S., Cecchin S., Marceddu G., Bertelli M. – Clin Ter. 2023 Nov-Dec;174(Suppl 2(6)):46-54. doi: 10.7417/CT.2023.2471.
10. “Omics sciences and precision medicine in lung cancer” – Micheletti C., Dhuli K., Donato K., Gadler M., Benedetti S., Guerri G., Cristofoli F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Bernini A., Fulcheri E., Stuppia L., Gatta V., Cristoni S., Cecchin S., Marceddu G., Bertelli M. – Clin Ter. 2023 Nov-Dec;174(Suppl 2(6)):37-45. doi: 10.7417/CT.2023.2470.
11. “Omics sciences and precision medicine in melanoma” – Medori M.C., Donato K., Dhuli K., Maltese P.E., Tanzi B., Tezzele S., Mareso C., Miertus J., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Beccari T., Ceccarini M.R., Gisondi P., Bellinato F., Stuppia L., Gatta V., Cecchin S., Marceddu G., Bertelli M. – Clin Ter. 2023 Nov-Dec;174(Suppl 2(6)):29-36. doi: 10.7417/CT.2023.2469.
12. “Omics sciences and precision medicine in testicular cancer” – Madeo G., Bonetti G., Maltese P.E., Tanzi B., Tezzele S., Mareso C., Agostini F., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Beccari T., Ceccarini M.R., Calogero A.E., Cannarella R., Stuppia L., Gatta V., Nughman M., Cecchin S., Marceddu G., Bertelli M. – Clin Ter. 2023 Nov-Dec;174(Suppl 2(6)):21-28. doi: 10.7417/CT.2023.2468.
13. “Omics sciences and precision medicine in thyroid cancer” – Dhuli K., Medori M.C., Donato K., Maltese P.E., Tanzi B., Tezzele S., Mareso C., Miertus J., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Beccari T., Ceccarini M.R., Stuppia L., Gatta V., Cristoni S., Cecchin S., Marceddu G., Bertelli M. – Clin Ter. 2023 Nov-Dec;174(Suppl 2(6)):11-20. doi: 10.7417/CT.2023.2467.
14. “Omics sciences and precision medicine in Urothelial Carcinoma” – Medori M.C., Micheletti C., Madeo G., Maltese P.E., Tanzi B., Tezzele S., Mareso C., Generali D., **Donofrio C.A.**, Cominetti M., Fioravanti A., Riccio L., Beccari T., Ceccarini M.R., Stuppia L., Gatta V., Cristoni S., Ahmed R., Cecchin S., Marceddu G., Bertelli M. – Clin Ter. 2023 Nov-Dec;174(Suppl 2(6)):1-10. doi: 10.7417/CT.2023.2466.
15. “Intrasellar dermoid cyst: case report of a rare lesion and systematic literature review comparing intrasellar, suprasellar and parasellar locations” – **Donofrio C.A.**, Bertazzoni G., Riccio L., Pinacoli A., Pianta L., Generali D., Ungari M., Servadei F., Roncaroli F., Fioravanti A. – World Neurosurg. 2023 Nov 21:S1878-8750(23)01619-4. doi: 10.1016/j.wneu.2023.11.057. Online ahead of print.

16. "Endoscope-Assisted Microvascular Decompression of Cochleo-Vestibular Nerve for Treating Unilateral Pulsatile Tinnitus, Sensorineural Hearing Loss, and Paroxysmal Vertigo: 2-Dimensional Operative Video" – **Donofrio C.A.**, Riccio L., Badaloni F., Rosellini E., Servadei F., Fioravanti A. – *Oper Neurosurg* (Hagerstown). 2023 Nov 1;25(5):e277-e278. doi: 10.1227/ons.0000000000000830. Epub 2023 Jul 11.
17. "Colorectal carcinoma to pituitary tumour: tumour to tumour metastasis" – **Donofrio C.A.***, Pizzimenti C.*, Djoukhadar I., Kearney T., Gnanalingham K., Roncaroli F. – *Br J Neurosurg*. 2023 Oct;37(5):1367-1370. doi: 10.1080/02688697.2020.1823937. Epub 2020 Sep 21.
18. "How I do it – The "drum skin" duraplasty technique after foramen magnum decompression for Chiari malformations" – Fioravanti A., Badaloni F., Tubbs R.S., **Donofrio C.A.** – *Acta Neurochir* (Wien). 2023 Oct;165(10):3045-3050. doi: 10.1007/s00701-023-05576-9. Epub 2023 Apr 15.
19. "The Occipitalis Muscle as an Adjunct Superficial Landmark for the Transverse Sinus and Transverse-Sigmoid Junction: An Anatomical Study With Application to Posterior Cranial Fossa Surgery" – Gilkes A., Rajaram-Gilkes M., Cardona J.J., Reina F., Carrera A., Iwanaga J., Dumont A.S., **Donofrio C.A.**, Badaloni F., Fioravanti A., Tubbs R.S. – *Cureus*. 2023 May 30;15(5):e39723. doi: 10.7759/cureus.39723. eCollection 2023 May.
20. "A Newly Discovered Dural Venous Sinus of the Skull Base: The Anterior Petroclinoid Sinus" – Spencer P.S., Cardona J.J., Reina F., Carrera A., Iwanaga J., Dumont A.S., **Donofrio C.A.**, Badaloni F., Fioravanti A., Tubbs R.S. – *World Neurosurg*. 2023 Apr;172:e581-e584. doi: 10.1016/j.wneu.2023.01.087. Epub 2023 Jan 27.
21. "Transcortical endoportalsubchoroidal endoscope-assisted approach to the third ventricle: from virtual reality to anatomical laboratory" – **Donofrio C.A.**, Riccio L., Capitanio J.F., Herur-Raman A., Panni P., Gagliardi F., Caputy A.J., Mortini P. – *J Neurosurg Sci*. 2023 Apr;67(2):175-184. doi: 10.23736/S0390-5616.20.05122-X. Epub 2020 Sep 28
22. "Posterior petrous meningiomas: surgical classification and postoperative outcomes in a case series of 130 patients operated via the retrosigmoid approach" – **Donofrio C.A.**, Badaloni F., Riccio L., Morandini A., Bertuccio A., Generali D., Calbucci F., Servadei F., Fioravanti A. – *World Neurosurg*. 2023 Mar;171:e301-e308. doi: 10.1016/j.wneu.2022.12.022. Epub 2022 Dec 10.
23. "Analysis of PD-L1 and CD3 Expression in Glioblastoma Patients and Correlation with Outcome: A Single Center Report" – Sobhani N., Bouchè V., Aldegheri G., Rocca A., D'Angelo A., Giudici F., Bottin C., **Donofrio C.A.**, Pinamonti M., Ferrari B., Panni S., Cominetti M., Aliaga J., Ungari M., Fioravanti A., Zanconati F., Generali D. – *Biomedicines*. 2023 Jan 22;11(2):311. doi: 10.3390/biomedicines11020311.
24. "A challenging case of sporadic melanocytoma of the jugular foramen." – **Donofrio C.A.**, Roncaroli F., Riccio L., Pereira M., O'Sullivan J., Mayers H., Potter G.M., Djoukhadar I., Rutherford S.A. – *Neurochirurgie* 2022 Jul;68(4):453-457. doi: 10.1016/j.neuchi.2021.06.001. Epub 2021 Jun 19.

25. "Classificazione e nomenclatura dei tumori neuroendocrini dell'ipofisi anteriore" – Roncaroli F., **Donofrio C.A.** – L'Endocrinologo 2022. <https://doi.org/10.1007/s40619-022-01039-y>
26. "Endoscopic sublabial transmaxillary approach to the inferior orbit: pearls and pitfalls – A comparative anatomical study" – **Donofrio C.A.**, Riccio L., Pathmanaban O.N., Fioravanti A., Caputy A.J., Mortini P. – Neurosurg Rev 2021 Dec;44(6):3297-3307. doi: 10.1007/s10143-021-01494-5. Epub 2021 Feb 10.
27. "BRAF Signaling Inhibition in Glioblastoma: Which Clinical Perspectives?" – Bouchè V., Aldegheri G., **Donofrio C.A.**, Fioravanti A., Roberts-Thomson S., Fox S.B., Schettini F., Generali D. – Front Oncol. 03 Nov 2021. doi: 10.3389/fonc.2021.772052. eCollection 2021.
28. "Microsurgical endoportals MRI/US-navigated approach for the resection of large intraventricular tumours: a 20-consecutive patients case series." – Capitanio J.F., **Donofrio C.A.**, Panni P., Barzaghi L.R., Bailo M., Gagliardi F., Mortini P. – Br J Neurosurg. 2021 Oct;35(5):570-577. doi: 10.1080/02688697.2021.1918632. Epub 2021 Aug 5
29. "Malignant behaviour of primary intracranial Rosai Dorfman disease: A rare presentation of a benign disease" – Riccio L.*, **Donofrio C.A.***, Serio G., Melatini A. – Neurochirurgie 2021 Apr;67(2):205-209. doi: 10.1016/j.neuchi.2020.11.006. Epub 2020 Dec 1. 2.

List of Abstracts, Posters and Oral Presentations between November 2021 and January 2024

1. XXVI Congresso Nazionale e Corso Residenziale dell'Associazione Italiana di Neuro-Oncologia – 23-25 November 2023 – Messina, Italy – "Determination of plasmatic microRNA levels by ddPCR as peripheral glioblastoma biomarkers" – **Donofrio C.A.**, Fioravanti A., Riccio L., Cominetti M., Cappelletti M.R., Ungari M., Generali D., Grossi I., Pelisenco I., De Petro G., Salvi A.
2. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – "Combined endoscopic transorbital and transmaxillary approach to the infratemporal fossa: anatomical feasibility study" – Corrivetti F., **Donofrio C.A.**, Riccio L., Fioravanti A.
3. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – "Transcranial approaches to tuberculum sellae meningiomas: the 'dark side of the moon' in the endoscopic endonasal era" – **Donofrio C.A.**, Riccio L., Badaloni F., Bresciani E., Muzzi M., Grappa E., Morandini A., Fioravanti A.
4. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – "Unilateral pulsatile tinnitus, sensorineural hearing loss and paroxysmal vertigo: endoscope-assisted microvascular decompression of cochleo-vestibular nerve" – **Donofrio C.A.**, Riccio L., Badaloni F., Rosellini E., Servadei F., Fioravanti A.
5. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – "Endoscope-assisted microvascular decompression for trigeminal neuralgia: preliminary results in a cohort of 30 consecutive patients" – **Donofrio C.A.**, Riccio L., Bianchini S., Fioravanti A.

6. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Five-aminolevulinic acid-guided glioblastoma surgery: can surgeons rely only on fluorescence? a comparative surgical and histopathological study on 97 patients” – Fioravanti A., Riccio L., Cominetti M., Cappelletti M.R., D’Auria P., Paioli G., Farinaro G., Generali D., Grappa E., Ungari M., **Donofrio C.A.**
7. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “The immune microenvironment in glioblastoma: PD-L1 expression and tumour infiltrating lymphocytes – new potential therapeutic targets” – **Donofrio C.A.**, Sobhani N., Bouché V., Aldegheri G., Rocca A., D’Angelo A., Giudici F., Bottin C., Pinamonti M., Cominetti M., Beretta E., Riccio L., Ungari M., Zanconati F., Generali D., Fioravanti A.
8. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Combined endoscopic endonasal transclival and contralateral transmaxillary approach to the petrous apex and the petroclival fissure – a reverse translational anatomical study” – **Donofrio C.A.**, Corrivetti F., Riccio L., Fioravanti A., De Notaris M.
9. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Postoperative outcomes in a series of 130 posterior petrous meningiomas operated via the retrosigmoid approach” – **Donofrio C.A.**, Badaloni F., Riccio L., Morandini A., Bertuccio A., Generali D., Calbucci F., Servadei F., Fioravanti A.
10. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Vertigini come sintomo d’esordio della sindrome da ipotensione liquorale” – Bianchini S.C.P.G., **Donofrio C.A.**, Ambrosi C., Beretta E., D’Auria P., Morandini A., Muzzi M., Paioli G., Riccio L., Fioravanti A.
11. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Cisti sinoviali della colonna lombare: diagnosi e trattamento. nostra esperienza” – Muzzi M., Morandini A., Paioli G., Bianchini S., **Donofrio C.A.**, D’Auria P., Riccio L., Fioravanti A.
12. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Gestione perioperatoria multidisciplinare e setting intraoperatorio nella “chirurgia da sveglio” per lesioni cerebrali in aree eloquenti” – D’Auria P., **Donofrio C.A.**, Grappa E., Subacchi S., Maccabelli G., Bianchini S., Morandini A., Muzzi M., Paioli G., Riccio L., Fioravanti A.
13. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – ““Hybrid” surgical treatment (open and endovascular) of a vertebral artery thrombosed giant aneurysm: case report and operative video” – Rosellini E., Raspagliesi L., Farinaro G., **Donofrio C.A.**, Paioli G., Muzzi M., Fioravanti A.
14. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Exeresi microchirurgica per via suboccipitale retrosigmoidea di un meningioma premeatale della faccia posteriore della rocca petrosa” – Bresciani E., **Donofrio C.A.**, Farinaro G., Rosellini E., Morandini A., Bianchini S., Beretta E., Fioravanti A.

15. 72° Congresso Nazionale SINch 2023 – 12-14 October 2023 – Padua, Italy – “Surgical technique resection of a bleeding brainstem cavernoma: case report and operative video” – Farinaro G., Rosellini E., **Donofrio C.A.**, Morandini A., Riccio L., Fioravanti A.
16. 2023 CNS (Congress of Neurological Surgeons) Annual Meeting – 9-13 September 2023 – Washington D.C., U.S.A. – “The Posterior Auricular Muscle as a Superficial Landmark for the Sigmoid Sinus and Transverse-Sigmoid Sinus Junction: An Anatomical Study” – **Donofrio C.A.**, Cardona J.J., Riccio L., Chaiyamon A., Shekhawat D., Iwanaga J., Dumont A.S., Badaloni F., Jackson N., Tubbs R.S., Fioravanti A.
17. 71° Congresso Nazionale SINch 2022 – 13-15 October 2022 – Naples, Italy – “Contralateral transmaxillary approaches: the Caldwell-Luc approach revival for ventral skull base surgery” – **Donofrio C.A.**, de Notaris M., Corrivetti F., Morandini A., Riccio L., Fioravanti A.
18. 71° Congresso Nazionale SINch 2022 – 13-15 October 2022 – Naples, Italy – “Endoscopic transorbital approaches: pearls of orbital anatomy for neurosurgeons” – **Donofrio C.A.**, Riccio L., Badaloni F., Fioravanti A.
19. EACR (European Association for Cancer Research) Liquid Biopsies – 24-26 May 2022 – Bergamo, Italy – “Determination of plasmatic microRNA levels by ddPCR as peripheral biomarkers for IDH-wild type glioblastomas: a pilot study” – **Donofrio C.A.**, Fioravanti A., Riccio L., Cominetti M., Cappelletti M.R., Generali D., Grossi I., De Petro G., Salvi A.
20. 70° Congresso Nazionale SINch 2021 – 14-16 October 2021 – Milan, Italy – “Covid-19 pandemic affected surgical treatment of elderly patients with glioblastoma: the Cremona Hub experience in the ‘Hub-Spokes’ model” – **Donofrio C.A.**, Nodari G., Bianchini S., d’Auria P., Muzzi M., Aliaga Arias J.M., Riccio L., Giudici S., Valtulina C., Morandini A., Fioravanti A.
21. 70° Congresso Nazionale SINch 2021 – 14-16 October 2021 – Milan, Italy – “‘Combined cranio-nasal approach’: transcranial and endoscopic endonasal combined approaches to treat invasive tumours of the anterior cranial fossa” – **Donofrio C.A.**, Riccio L., Nodari G., Paioli G., Grappa E., Morandini A., Fioravanti A.
22. 70° Congresso Nazionale SINch 2021 – 14-16 October 2021 – Milan, Italy – “Surgical treatment of supratentorial meningiomas in elderly patients: safety and efficacy” – **Donofrio C.A.**, d’Auria P., Bianchini S., Nodari G., Ambrosi C., Besana M., Varotti E., Ungari M., Valtulina C., Fioravanti A.
23. 70° Congresso Nazionale SINch 2021 – 14-16 October 2021 – Milan, Italy – “Surgical treatment of glioblastoma in elderly patients: a monocentric and retrospective study” – **Donofrio C.A.**, Cappelletti M., Catenacci E., Aliaga Arias J.M., Subacchi S., Generali D., Maccabelli G., Varotti E., Ungari M., Riccio L., Giudici S., Morandini A., Fioravanti A.
24. 70° Congresso Nazionale SINch 2021 – 14-16 October 2021 – Milan, Italy – “Functional performances in elderly neuro-oncological patients” – Aliaga Arias J.M., Subacchi S., **Donofrio C.A.**, Catenacci E., Paioli G., Grappa E., Muzzi M., Morandini A., Fioravanti A.

25. eEANS 2021: Neurosurgery in translation – 03-07 October 2021 – Virtual Congress – “Meningiomas of the posterior petrous bone: surgical anatomy and postoperative results of 130 consecutive patients” – **Donofrio C.A.**, Badaloni F., Nodari G., Paioli G., d’Auria P., Muzzi M., Calbucci F., Fioravanti A.
26. eEANS 2021: Neurosurgery in translation – 03-07 October 2021 – Virtual Congress – “Haemorrhagic brainstem cavernous malformation: early and long term functional outcomes after microsurgical resection” – **Donofrio C.A.**, Badaloni F., Aliaga Arias J.M., Valtulina C., Bianchini S., Catenacci E., Morandini A., Calbucci F., Fioravanti A.
27. eEANS 2021: Neurosurgery in translation – 03-07 October 2021 – Virtual Congress – “Degenerative cervical myelopathy: the preoperative Nurick score as predictor of surgical outcomes” – **Donofrio C.A.**, Muzzi M., Catenacci E., Paioli G., Nodari G., Valtulina C., Morandini A., Fioravanti A.