

1 *Article*

2 **Predictive potential of tumour-stroma ratio on benefit from adjuvant bevacizumab in**  
3 **high-risk stage II and stage III colon cancer.**

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26 **Running title:** TSR as predictive biomarker in stage II/III CC

27 **Keywords:** Colon cancer, Stroma, Predictive, Bevacizumab, AVANT

28 **Introduction**

29 In Europe colorectal cancer (CRC) is the second most common cause of cancer related death  
30 in both men and women.(1) The 5-year survival is strongly dependent on disease stage and  
31 rapidly decreases in individuals with lymph node or distant metastasis. Current guidelines for  
32 high-risk stage II and stage III patients, advice adjuvant fluoropyrimidine-based  
33 chemotherapy with addition of oxaliplatin as standard therapy. This combination has shown  
34 to significantly improve disease-free survival (DFS) and overall survival (OS).(2, 3)  
35 Adjuvant therapy with bevacizumab, a humanized anti-VEGF monoclonal antibody, has only  
36 demonstrated to improve outcome in patients with metastatic stage IV disease and is  
37 therefore currently not recommended in other stages. (3-8) However, due to heterogeneity of  
38 colon cancer, one could argue that some subpopulations could possibly benefit from targeted  
39 therapy in an adjuvant setting. To identify such potential groups, predictive parameters are  
40 necessary. Currently most biomarkers focus on tumour cells. However, recently the “seed-  
41 and-soil” principle has been revisited, focusing on the tumour –microenvironment as a major  
42 factor responsible for metastasis.(9, 10) Studies have shown that during cancer progression,  
43 the normal stromal host compartments transform, due to complex intercellular  
44 communication between surrounding stromal host cells and cancer cells, in which a cross-talk  
45 of signalling molecules between these compartments, leads to an activated state with  
46 production of various cytokines and growth factors creating an area favouring cancer  
47 progression and invasion. Thus, illustrating the importance of intratumoural stroma. (11-14)  
48 Consistent with this principle, it has been proven that in colon cancer, high amounts of  
49 intratumoural stroma are associated with poor survival compared to tumours with low  
50 amounts of stroma. (15-18) This prognostic parameter is also known as the tumour-stroma  
51 ratio (TSR), and entails a simple microscopic quantification of the amount of intratumoural  
52 stroma on a tumour tissue slide, which is derived after surgical resection. It has been

53 validated in multiple studies, thereby demonstrating the robustness and potential of this fairly  
54 simple, quick and cost effective pathological technique. (15, 17, 18)

55 Since the prognostic quality of the TSR is clear, it is interesting to evaluate whether this  
56 parameter could also serve as a predictive marker to improve risk stratification of patients  
57 with high-risk stage II and III colon cancer, in order to determine if subpopulations  
58 could benefit from the VEGF antibody bevacizumab in an adjuvant setting. Our hypothesis  
59 was that patients with high stromal tumours would benefit from adjuvant bevacizumab,  
60 considering these tumours hold features promoting cancer progression and metastasis, hence  
61 possessing a more aggressive phenotype. (11, 12, 14)

62 To study this concept, we used the study population from the AVANT trial (BO17920), a  
63 prospective randomized trial studying the addition of bevacizumab to oxaliplatin-based  
64 chemotherapy in an adjuvant setting. This was a negative study, showing no prolongation of  
65 DFS and for OS even suggesting a potential detrimental effect when adding bevacizumab to  
66 the chemotherapy regime. We considered that if the TSR is able to identify patients that do  
67 benefit from bevacizumab in an adjuvant setting, it could serve as a selection tool to optimize  
68 adjuvant treatment outcomes in colon cancer.

69 Therefore, the aim of this study was to investigate the predictive potential of TSR, by  
70 determining the effects on DFS and OS in patients with high-risk stage II and stage III colon  
71 cancer who received standard oxaliplatin-based chemotherapy with or without addition of  
72 bevacizumab.

73

## 74 **Patients and Methods**

### 75 **Study design**

76 Available Haematoxylin and Eosin (H&E) stained tumour slides from patients randomized in  
77 the AVANT trial were included in our analysis. Patients entering the AVANT trial had

78 undergone potential curative treatment, including surgery (before randomization) followed by  
79 adjuvant chemotherapy.

80 Inclusion criteria were histologically confirmed high-risk stage II or stage III colon  
81 carcinoma. The study had an open label design, in which patients were randomly assigned  
82 1:1:1 to one of the three treatment regimens; FOLFOX-4 for 24 weeks followed by  
83 observation for 24 weeks, bevacizumab–FOLFOX-4 or bevacizumab–XELOX for 24 weeks  
84 followed by bevacizumab monotherapy for 24 weeks. Patients were recruited in 330 centres  
85 in 34 countries. For detailed trial design, see de Gramont et al. (5)

86 For our study, archival material was used in an anonymized matter, therefore no additional  
87 informed consent was needed.

88

### 89 **Histopathologic scoring**

90 The TSR was determined in all patients from whom a H&E stained formalin-fixed paraffin-  
91 embedded tissue slide from the primary tumour was available.

92 Pathological examination was performed as described by Mesker et.al 2007 (For detailed  
93 description see Appendix 1). Two investigators (SZ, GvP) scored stromal percentage in a  
94 blinded manner. Scoring percentages were given per 10-fold (10%, 20% etc.) per image field.  
95 For statistical analysis, we defined two groups; stroma-high (> 50%) and stroma-low ( $\leq$ 50%)  
96 as determined a priori to have maximum discriminative power (Figure S1). (17, 18)

97

### 98 **Statistical analysis**

99 Statistical analysis was performed using SPSS software version 23.0. The primary endpoint,  
100 DFS, was defined as the time between randomization and recurrence, new occurrence of  
101 colon cancer, or death from any cause. Alive and event-free patients at the clinical cut-off  
102 date were censored at the last date at which they were known to be disease-free and/or alive.

103 The secondary endpoint, OS, was defined as time from randomization to death. Patients who  
104 were still alive at the clinical cut-off date were censored at the date at which they were last  
105 confirmed to be alive.

106 Kaplan-Meier method and log rank test were used to analyse time-to-event endpoints. Intra-  
107 observer variability was tested using Cohen's kappa coefficient.

108 Univariate and multivariable analyses were performed using Cox-regression analysis. For  
109 predictive analysis, a Cox proportional hazard model including an interaction term between  
110 treatment arms and TSR was used. The interaction test was used to test the null hypothesis  
111 that TSR is not predictive for response to bevacizumab.

112 Parameters with a *p*-value less than 0.10 in the univariate analysis, were included in  
113 multivariable analyses.

114

## 115 **Results**

### 116 **Study population**

117 In the AVANT trial, a total of 3451 patients were recruited between 2004 and 2007. We  
118 received a total of 1213 histological samples. After scoring all samples, baseline clinical  
119 patient information was used for analysis. Upon this, one patient was excluded due to the  
120 presence of stage IV disease at time of randomization.

121 The final study population comprised 1212 patients, with respectively 405 (33.4%) patients in  
122 the FOLFOX-4 arm, 401 (33.1%) in the bevacizumab – FOLFOX-4 arm and 406 (33.5%) in  
123 the bevacizumab - XELOX arm.

124 Patient characteristics were reasonably balanced between the different groups (Table 1).

125 Considering our study population comprised only a selection of the total AVANT  
126 population, we compared our study population to the total AVANT population. There were  
127 no apparent differences in distribution between treatment arms, stage, gender and age.

128 Noteworthy to mention, in the AVANT trial high-risk stage II patients were recruited solely  
129 for exploratory analysis. Efficacy (intention-to-treat (ITT)) analysis was only performed on  
130 stage III disease. Our study population consists of 205 (16.9%) high-risk stage II and 1007  
131 (83.1%) stage III cases, which were both used in the analysis because both groups are  
132 considered as candidates for adjuvant chemotherapy according to current European  
133 guidelines.(22)

134

### 135 **Scoring tumour stroma-ratio**

136 Of 1212 evaluated patients, 339 (28.0%) were scored as stroma-high and 824 (68.0%) as  
137 stroma-low. Forty-nine (4.0%) samples could not be scored for TSR due to poor histological  
138 quality and were therefore excluded. These samples consisted either of too little tissue  
139 material to score (i.e. biopsies), exclusively muscle tissue and/or lymph node tissue.  
140 Cohen's kappa coefficient revealed a good level of agreement in the classification.  
141 Cox regression interaction term for TSR and treatment arms showed a significant value for  
142 DFS ( $p = 0.005$ ) and OS ( $p=0.007$ ) (Table S2).

### 143 **Disease-free survival**

144 DFS was significantly shorter in patients with stroma-high tumours compared to patients with  
145 stroma-low tumours, HR 1.75 (95% CI 1.32-2.33;  $p < 0.001$ ) (Figure 1).

146 In the total BEP study population the addition of bevacizumab did not prolong the DFS ( $p =$   
147  $0.23$ ) compared to FOLFOX-4 monotherapy and suggests a potential detrimental effect on  
148 DFS (Figure S2). In the Cox-regression analysis, TSR had a HR of 2.92 (95% CI 1.78 – 4.79;  
149  $p < 0.001$ ) for the low versus high stromal tumours. The interaction model for treatment arms  
150 and TSR, showed a significant predictive value ( $p = 0.005$ ) for treatment effect in the two  
151 TSR-groups for DFS (Table S2). In the stroma-low group this effect was significant, with a  
152 HR of 1.94 (95% CI 1.24 – 3.04;  $p = 0.004$ ) for bevacizumab –FOLFOX-4 versus FOLFOX-

153 4. For bevacizumab – XELOX this was not seen, with a HR of 1.07 (95% CI 0.64 – 1.77;  $p=$   
154  $0.80$ ). In the stroma-high tumours a trend for better DFS outcome was seen in the  
155 bevacizumab – FOLFOX-4 group versus FOLFOX-4 (HR 0.61 (95% CI 0.35-1.07;  $p= 0.08$ ).  
156 For bevacizumab- XELOX versus FOLFOX-4 this was not seen (HR 0.78 (95% CI 0.47-  
157 1.30;  $p= 0.35$ )) (Table S2, Figure 2).

158 The univariate Cox regression analysis revealed TSR ( $p < 0.001$ ), gender ( $p= 0.05$ ), disease  
159 stage ( $p= 0.002$ ) and MMR status ( $p= 0.04$ ) as statistically significant prognosticators for  
160 DFS. In the multivariable analysis TSR ( $p= 0.003$ ), gender ( $p= 0.013$ ) and disease stage ( $p=$   
161  $0.004$ ) maintained significance (Table S1).

162

### 163 **Overall survival**

164 As shown in Figure 1, patients with stroma-high tumours had a significant shorter OS  
165 compared to patients with stroma-low tumours (HR 1.54 (95% CI 1.04-2.29;  $p= 0.03$ )). In the  
166 total BEP study population, the addition of bevacizumab did not prolong the OS ( $p = 0.17$ )  
167 compared to FOLFOX-4 monotherapy (Figure S2).

168 Cox-regression analysis for OS showed a HR of 3.14 (95%CI 1.57 – 6.26;  $p= 0.001$ ) for TSR  
169 with regard to high versus low stromal tumours. The interaction model showed a similar  
170 pattern as for DFS, with a significant interaction term between treatment and TSR-group ( $p=$   
171  $0.007$ ) (Table S2). Stroma-low tumours in the bevacizumab – FOLFOX-4 arm versus  
172 FOLFOX-4 arm had a significant worse OS, HR of 2.53 (95%CI 1.36-4.71;  $p= 0.003$ ). For  
173 stroma-high tumours this was not significant, with a HR of 0.50 (95%CI 0.22-1.14;  $p= 0.10$ ).  
174 For bevacizumab – XELOX versus FOLFOX-4 the HR was 1.13 (95% CI 0.55-2.31;  $p=$   
175  $0.74$ ) for stroma-low tumours and HR 0.74 (95% CI 0.37-1.51;  $p= 0.41$ ) for stroma-high  
176 tumours (Table S2, Figure 3).

177 The univariate analysis for OS showed TSR ( $p= 0.03$ ), gender ( $p= 0.006$ ), disease stage ( $p=$

178 0.04) and BRAF status ( $p= 0.10$ ) as statistically significant prognosticators. In the  
179 multivariable analysis TSR ( $p= 0.05$ ), gender ( $p= 0.002$ ) and disease stage ( $p= 0.05$ )  
180 maintained significance (Table S1).

181 No additional exploratory analyses were performed on patients from whom molecular  
182 variables were available (i.e. MMR status, KRAS and BRAF), due to non-significance in the  
183 Cox-regression analysis.

## 184 **Discussion**

185 In our study, we evaluated the predictive potential of TSR in hopes of being able to select  
186 subpopulations with high-risk stage II and III colon cancer that could benefit from adjuvant  
187 bevacizumab. Prior research failed to show benefit from addition of bevacizumab to standard  
188 chemotherapy regimens in these patients and is therefore currently only recommended in  
189 metastatic disease.(4-8, 23) Our hypothesis was that high-risk stage II and III patients with  
190 high stromal tumours would benefit from adjuvant bevacizumab, considering the pro-  
191 carcinogenic features these tumours possess and association with a worse survival.(15-18, 24)

192 In our study the TSR validated as a predictive parameter, however without clinical  
193 implications. As assumed, the stroma-low group had no benefit whatsoever from addition of  
194 bevacizumab and even showed a significantly detrimental effect on survival, most  
195 pronounced in the bevacizumab- FOLFOX-4 group. This was in accordance with the  
196 AVANT ITT- analysis and supports current guidelines which discommend adjuvant anti-  
197 VEGF in stage II/III disease. It is not completely understood why this was so evident in this  
198 group and not as pronounced in the XELOX-group. Considering capecitabine is  
199 biotransformed into active metabolites that mimic 5-FU infusion, one could consider these  
200 biologically equivalent and of similarly efficacy when administrated correctly.(25) Previous  
201 studies investigating non-inferiority of capecitabine in combination with oxaliplatin versus  
202 5-FU with oxaliplatin, correspondingly showed either similar efficacy or inconclusive results

203 regarding non-inferiority. (26-30) The NO16966 accordingly showed similar performance of  
204 XELOX and FOLFOX in terms of OS, when adding bevacizumab. (31) Taking this into  
205 account, it would be less likely to regard the observed results as due to an interaction of  
206 FOLFOX with bevacizumab. The AVANT ITT-analysis does show considerably less adverse  
207 events, doses reductions, -delays or interruptions in the XELOX-group compared to the other  
208 groups, suggesting less toxicity and perhaps therefore better survival outcomes (for details,  
209 see de Gramont et al).(5) However, since the ITT-analysis only entails stage III patients,  
210 these results have to be adjusted for stage before correlation to our cohort is possible.

211 In contrast with low stromal tumours, in patients with stroma-high tumours we did observe a  
212 beneficial trend with addition of bevacizumab. Although not significant, this was an  
213 anticipated effect when regarding high stromal tumours as more aggressive due to the cross-  
214 talk between their local microenvironment and tumour cells. This finding, in combination  
215 with previous research validating the TSR as an independent prognostic parameter, does  
216 suggest that there could be potential in the TSR as a predictive tool with clinical  
217 implications.(15, 17, 18) Perhaps not solely with TSR, but in combination with additional  
218 markers.(32) However, that would compromise the simplicity and costs effectiveness of the  
219 current technique, which could be easily incorporated in routine diagnostics. Currently  
220 extensive research is being performed regarding the tumour-microenvironment and response  
221 to anti-angiogenic therapy. It has become increasingly clear that stromal cells not only  
222 provide a target for cancer therapy, but also have an essential role in anti-angiogenic  
223 resistance. (33) An issue, which is already relevant to patient groups receiving these agents in  
224 routine clinical practice, since benefit on overall survival with addition of bevacizumab is  
225 often borderline significant or lacking depending on the chemotherapy regimen.(34-36)  
226 Better understanding of these mechanisms will make it possible to identify sensitive targets  
227 and/or phenotypes to overcome these tumour escape mechanisms. For instance, Smith et.al

228 reported two stromal phenotypes (i.e. tumour-vessel and stromal-vessel) based on CD31 and  
229  $\alpha$ -smooth muscle actin ( $\alpha$ -SMA) staining. In mCRC, tumour-vessel phenotype tumours  
230 appeared to be more sensitive to combination oxaliplatin-based chemotherapy with  
231 bevacizumab compared to the stromal-vessel phenotype.(37) It would be interesting to  
232 correlate these phenotypes to the TSR, to possibly improve the predictive performance, but  
233 also to determine whether there is any prognostic relevance in metastatic disease.  
234 A possible limitation of this study is the fact we only investigated a selection of the total  
235 AVANT study population, though evenly balanced, making it possible that the study is  
236 underpowered.  
237 Nevertheless, despite the fact the findings were non-significant, we do find the potential  
238 beneficial survival trend that was observed in the stroma-high tumours with addition of  
239 bevacizumab, is worthwhile for further investigation with or without additional markers.  
240 Since this is one of the first studies evaluating this principle, we feel that we should not  
241 abandon this principle right away and validation of the findings would be necessary, to  
242 definitely rule out a coincidental finding. Considering very limited new targeted therapies  
243 have come available for treatment of colorectal cancer after the introduction of bevacizumab  
244 over a decade ago, maximum efficient utilization of this drug would be desirable.

245 **Ethics approval and consent to participate**

246 Current study was performed by using archival material in an anonymized matter, therefore  
247 no additional informed consent was needed. Archival materials were derived from the  
248 AVANT trial (BO17920), that study was done in accordance with the declaration of Helsinki.  
249 Protocol approval was obtained from the ethics review committees or institutional review  
250 boards at participating sites. Patients provided written informed consent before study  
251 participation. For more details, see de Gramont et al. (5)

252 **Disclosures**

253 C. Mancao is a fulltime employee of Genentech Roche and holds stock/options in Genentech  
254 Roche.

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258 **Contributions**

259 SZ performed TSR scoring, statistical analyses and wrote the first draft of the manuscript.  
260 GvP performed TSR scoring and helped to write and review the manuscript. WE initiated the  
261 study with Roche, wrote the study proposal, delivered clinical input and helped to write the  
262 manuscript. HG and RT delivered clinical input and helped to write the manuscript. CM  
263 arranged material and data transfer. HP helped with the statistical analysis.  
264 All authors approved the final version of the manuscript.

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413 **Titles and legends to figures**

414 **Figure 1. Kaplan-Meier survival curves of DFS (A) and OS (B) of stroma-low versus**  
415 **stroma-high in the total patient population [DFS HR 1.75 (95% CI 1.32-2.33;  $p <$**   
416  **$0.001$ ) | OS HR 1.54 (95% CI 1.04-2.29;  $p = 0.03$ )]**

417 — Tumour stroma-low

418 — Tumour stroma-high

419 **Figure 2. Disease-free survival: (A) Stroma-low, (B) Stroma-high**

420 — 1: FOLFOX-4

421 — 2: FOLFOX-4 + bevacizumab

422 — 3:XELOX + bevacizumab

423 **Figure 3. Overall survival: (A) Stroma-low, (B) Stroma-high**

424 — 1: FOLFOX-4

425 — 2: FOLFOX-4 + bevacizumab

426 — 3: XELOX + bevacizumab

**Table 1. Patient characteristics**

		Total study population	Tumour - stroma ratio				<i>p</i> - value
		N (%)	stroma -low		stroma-high		
			N =	(%)	N =	(%)	
Treatment	FOLFOX-4	405 (33,4%)	267	68%	123	32%	0.32
	FOLFOX-4 +bevacizumab	401 (33,1%)	284	73%	103	27%	
	XELOX +bevacizumab	406 (33,5%)	273	71%	113	29%	
Gender	Male	673 (55,5%)	453	70%	195	30%	0.43
	Female	539 (44,5%)	371	72%	144	28%	
Age (years)	<= 50	278 (22,9%)	189	72%	72	28%	0.75
	51 - 64	556 (45,9%)	379	71%	152	29%	
	65 - 70	247 (20,4%)	166	69%	75	31%	
	71 - 80	129 (10,6%)	88	69%	40	31%	
	> 80	2 (0,2%)	2	100%	0	0%	
Disease stage	stage II (high-risk)	205 (16,9%)	136	69%	61	31%	0.54
	stage III	1007 (83,1%)	688	71%	278	29%	
Previous hypertension	No	786 (64,9%)	545	72%	208	28%	0.12
	Yes	426 (35,1%)	279	68%	131	32%	
KRAS mutation*	Positive	445 (36,7%)	296	68%	139	32%	0.04
	Negative	328 (27,1%)	226	70%	95	30%	
BRAF mutation*	Mutation	78 (6,4%)	56	72%	22	28%	0.84
	Wildtype	994 (82,0%)	688	71%	285	29%	
MMR status*	MSS	930 (76,7%)	631	69%	281	31%	0.01
	MSI	121 (10,0%)	97	80%	24	20%	
CEA (ng/L)	<=5.0	1171 (96,6%)	799	71%	325	29%	0.08
	>5.0	28 (2,3%)	15	56%	12	44%	

Abbreviations: *MMR status* Mismatch Repair status, *MSI* Microsatellite instability, *MSS* Microsatellite stable, *CEA* Carcinoembryonic antigen

\* Data not available from all patients

**Figure 1. Kaplan-Meier survival curves of DFS (A) and OS (B) of stroma-low versus stroma-high in the total patient population [DFS HR 1.75 (95% CI 1.32-2.33;  $p < 0.001$ ) | OS HR 1.54 (95% CI 1.04-2.29;  $p = 0.03$ )]**

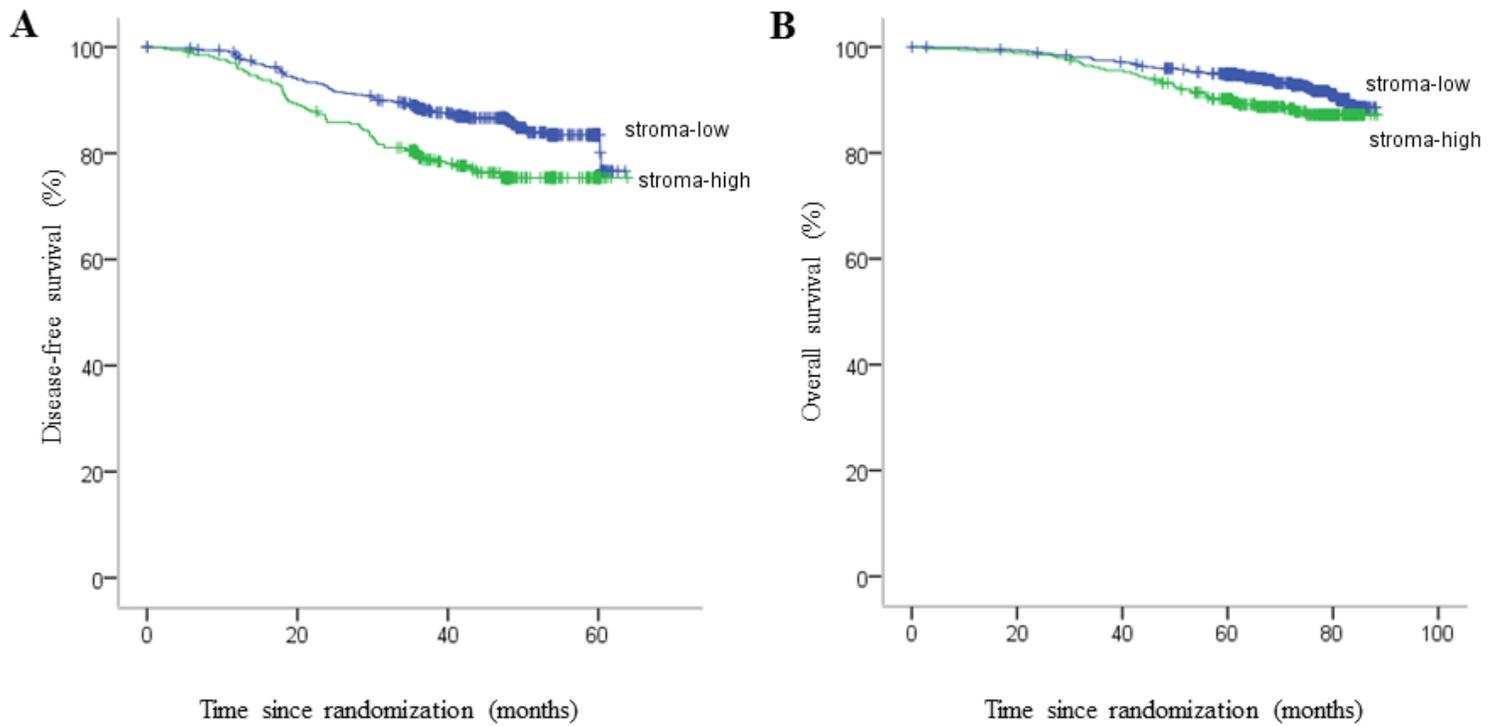


Figure 2. Disease-free survival: (A) Stroma-low, (B) Stroma-high

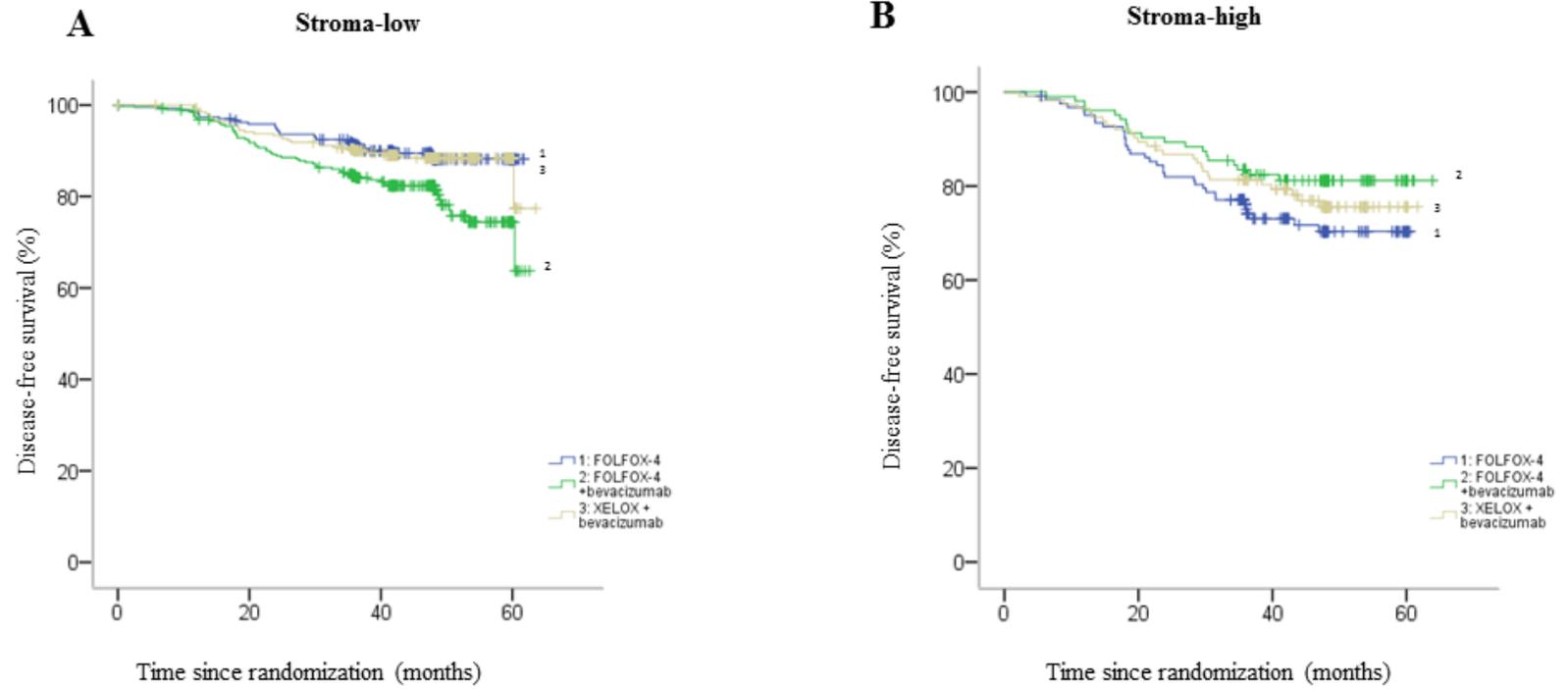


Figure 3. Overall survival: (A) Stroma-low, (B) Stroma-high

