

**The effect of RAS/BRAF mutation status on survival and treatment efficacy in vulnerable older patients with metastatic colorectal cancer a post-hoc exploratory analysis of the randomized NORDIC9-study**

Liposits, Gabor

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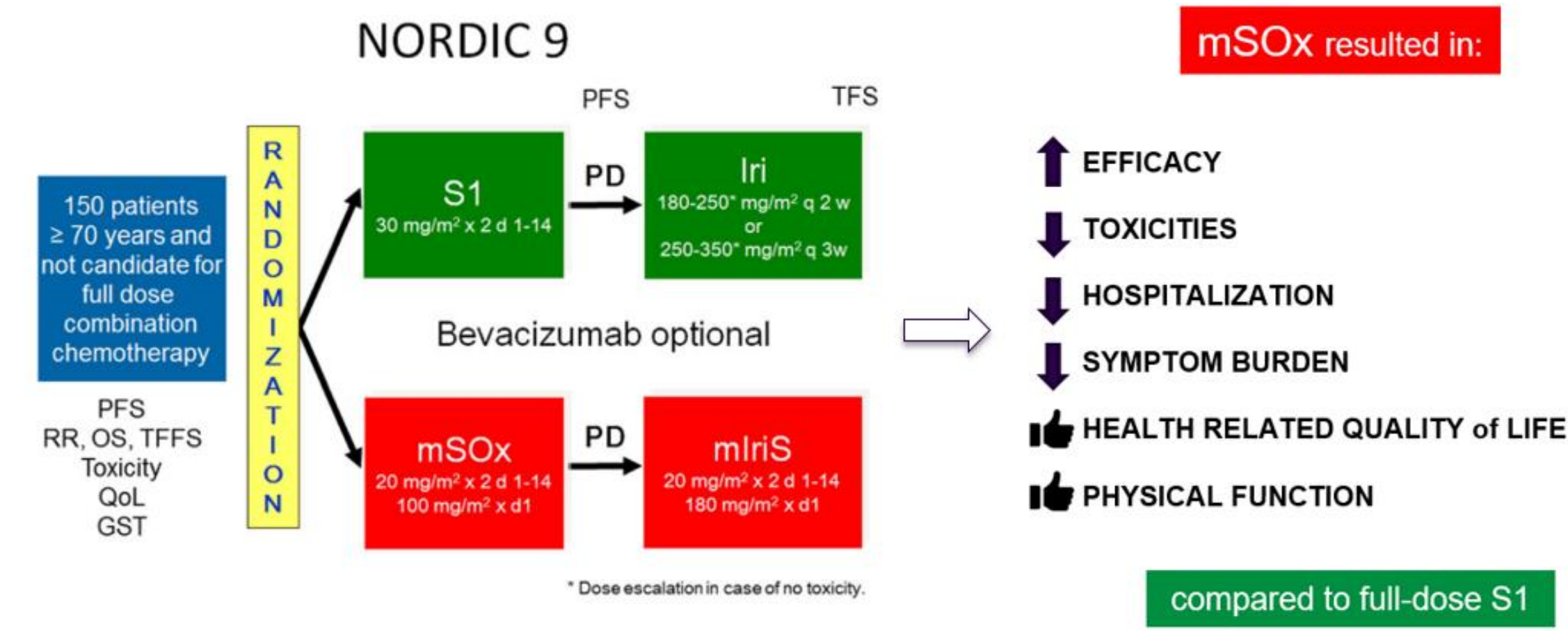
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1. Department of Oncology, Odense University Hospital, Denmark
2. Department of Clinical Research, University of Southern Denmark
3. Academy of Geriatric Cancer Research (AgeCare), Denmark
4. Department of Geriatric Medicine, Odense University Hospital, Denmark
5. Regional Hospital Gødstrup, Denmark
6. Open Patient data Explorative Network, Odense University Hospital, Denmark
7. Trondheim University Hospital, Norway
8. Norwegian University of Science and Technology, Norway
9. Karolinska University Hospital, Sweden
10. Aalborg University Hospital, Denmark
11. Uppsala University, Sweden
12. Tampere University Hospital, Finland
13. Helsinki University Hospital, Finland
14. Haukeland University Hospital, Bergen, Norway
15. University of Bergen, Bergen, Norway



## BACKGROUND



## AIM

Here, we present the post-hoc exploratory analysis of the *RAS/BRAF* mutation status on survival and treatment efficacy in a cohort comprising vulnerable older patients with mCRC. Of the very few randomized trials investigating the efficacy of chemotherapy in older adults with mCRC, none has reported specific outcomes in a *BRAF*<sup>V600E</sup> mutated population.

## METHOD

Overall survival (OS) and progression-free survival (PFS) for subgroups based on *RAS/BRAF* status were compared by log-rank test. Hazard ratios (HR) and 95% confidence intervals (95%CI) were estimated by Cox regression. The prognostic value of *RAS/BRAF* status was evaluated by C-statistics. For multivariate analyses, Cox proportional hazards regression models were applied.

## CONCLUSIONS

We demonstrated a substantial difference in OS favoring patients with *BRAF*<sup>V600E</sup> mutated tumors receiving reduced-dose combination chemotherapy ( $\pm$  bev) compared to those treated with full-dose monotherapy ( $\pm$  bev). The relatively high proportion of patients with *BRAF* mutation is representative for the Nordic population.

## IMPORTANCE

- The largest group of adults with cancer is  $\geq 70$  years.
- *BRAF*<sup>V600E</sup> mCRC is more frequent than seen in clinical trials
- Availability of BEACON doublet and ICIs remains limited in low- and middle-income countries
- Emphasizing the need for effective chemotherapy option for *BRAF*<sup>V600E</sup> mutated patients

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## RESULTS

Between 2015 and 2017, 160 patients were enrolled; the median age was 78 years (IQR): 75-81). *RAS* and *BRAF* status were established in 116 patients with the following distribution: *RAS/BRAF* wild type 31%, *RAS* mutation 51%, and *BRAF*<sup>V600E</sup> mutation in 18% of the patients.

### The prognostic value of *RAS/BRAF* mutation status

mOS for *RAS/BRAF*wt, *RAS*mut, and *BRAF*mut subgroups were 16.0, 14.5, and 10.8 months. In multivariable analysis, *BRAF*mut patients had higher risk for shorter survival.

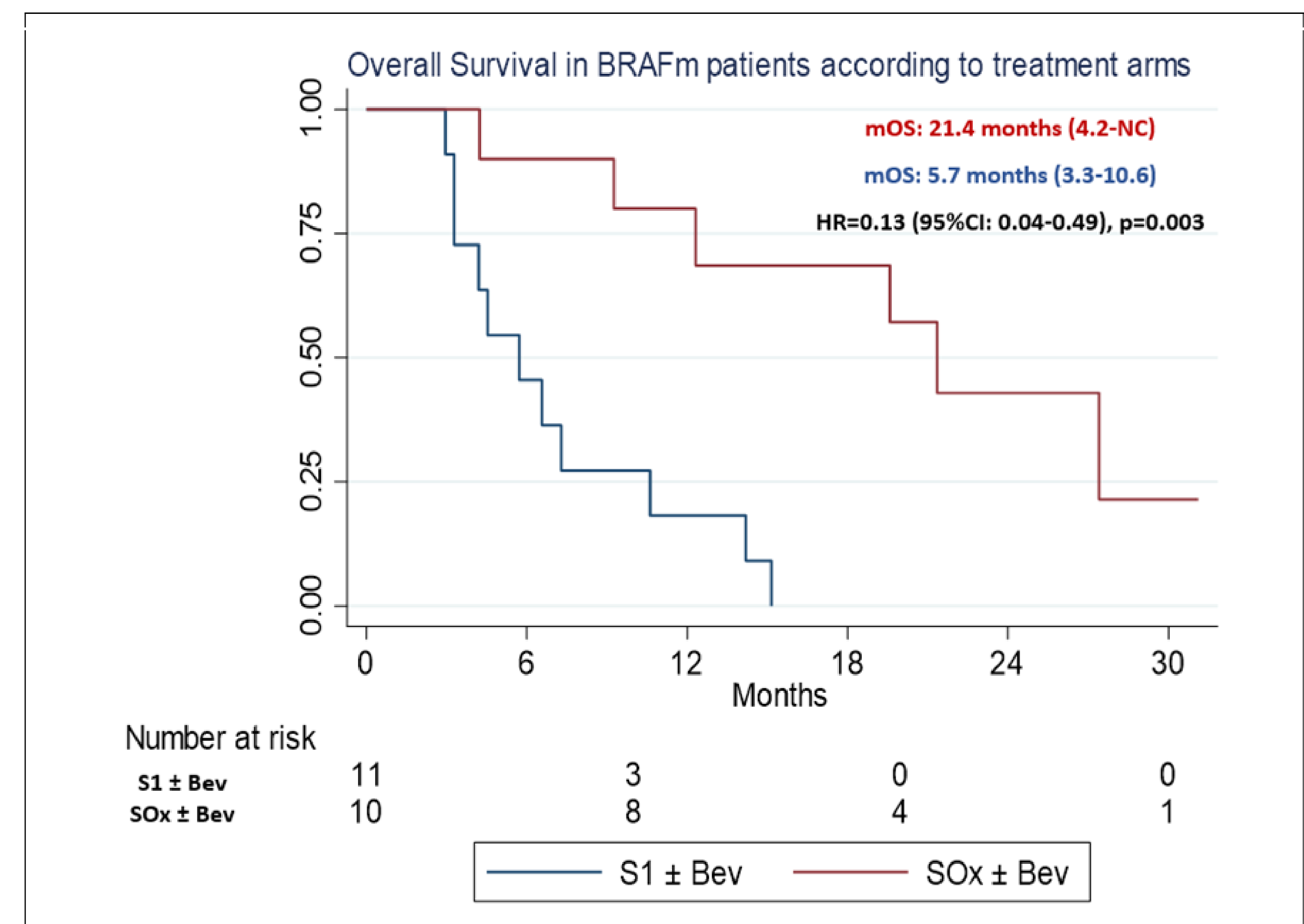
Multivariable analysis adjusted for age, sex, treatment allocation, bevacizumab, ECOG PS, number of metastatic sites, primary tumor in situ, weight-loss, and plasma C-reactive protein

<i>RAS/BRAF</i> status	n	Progression-free survival			Overall survival		
		Hazard ratio (95%CI)	p-value	Harrell's C (95%CI)	Hazard ratio (95%CI)	p-value	Harrell's C (95%CI)
<i>RAS</i> and <i>BRAF</i> wild type	36	1.00	NA	<b>0.69</b> (0.66-0.73)	1.00	NA	<b>0.73</b> (0.69-0.78)
<i>RAS</i> mutated	59	1.19 (0.75-1.88)	0.463		1.13 (0.65-1.95)	0.675	
<i>BRAF</i> mutated	21	<b>1.93</b> (1.04-3.56)	<b>0.037</b>		<b>2.78</b> (1.37-5.64)	<b>0.005</b>	

### The predictive value of treatment approach on survival according to the *RAS/BRAF* status

In the *BRAF*<sup>V600E</sup> subgroup, patients receiving SOx $\pm$ Bev had a significant longer OS compared to those treated with S1 $\pm$ Bev.

The difference remained statistically significant using multivariate analysis: HR=0.20 (95%CI: 0.05-0.84), p=0.028).



Univariate analysis of OS and PFS according to treatment regimens and *RAS/BRAF* status (Abbreviations: bev: bevacizumab, 95%CI: 95% confidence interval, NA: non-applicable, NE: not estimable)

### Progression-free and Overall Survival according to *RAS/BRAF* status and treatment regimens Univariate analyses

Treatment arms	n	Progression-free Survival			Overall Survival		
		Months (95%CI)	Hazard ratio (95%CI)	p-value	Months (95% I)	Hazard ratio (95%CI)	p-value
<b><i>RAS/BRAF</i> wild type</b>	36						
S1 $\pm$ bev	21	6.8 (4.8-9.0)	1.00	0.48	19.5 (10.4-27.3)	1.00	0.602
SOx $\pm$ bev	15	6.7 (4.2-10.2)	0.77 (0.38-1.58)		15.4 (8.8-NE)	1.24 (0.55-2.78)	
<b><i>RAS</i> mutated</b>	59						
S1 $\pm$ bev	30	5.4 (3.1-8.0)	1.00	0.238	10.2 (5.9-20.5)	1.00	0.256
SOx $\pm$ bev	29	8.2 (5.3-9.8)	0.73 (0.43-1.24)		15.5 (12.9-NE)	0.69 (0.36-1.31)	
<b><i>BRAF</i> mutated</b>	21						
S1 $\pm$ bev	11	3.1 (2.0-4.7)	1.00	0.052	5.7 (3.3-10.6)	1.00	<b>0.003</b>
SOx $\pm$ bev	10	6.2 (2.0-11.5)	0.36 (0.13-1.00)		<b>21.4</b> (4.2-NE)	<b>0.13</b> (0.04-0.49)	