

A prospective Nordic study on the use of chromogranin A for the prediction of progression in patients with pancreatic and small intestinal neuroendocrine tumours.

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Background

- Retrospective studies showed that changes in plasma chromogranin A (CgA) may predict change in tumour burden in gastroenteropancreatic neuroendocrine tumour (GEP-NET) patients.
- The aim of this prospective study, was to compare the association between changes in plasma CgA and changes in tumor burden on CT in patients with GEP-NET and unknown primary NET with residual disease

Material & Methods

- In this non-interventional study, patients with GEP-NET and unknown primary NET under treatment with Sandostatin LAR or with non-somatostatin analog anti-tumour treatment, were observed up to a maximum of 24 month.
- 239 patients were included in Denmark, Norway and Sweden from December 2010 to December 2013. A CT was followed by at least one additional CT 1 - 24 months later.
- Matching pairs of CgA and CT assessments were defined for each individual (defined as an event):
 - Primary analysis: CgA within +/- 6 weeks to CT
 - Post-hoc analysis: CgA 3-6 month prior to CT
- Change in tumour size was defined as regression, progression, or stable disease by RECIST1.1. A 25% change in CgA discriminated between increased, unchanged or decreased plasma CgA levels.

Table 1. Patient demographics and history

Category	Statistics	Tumour Location			
		All	Small Intestine	Pancreas	Other/Unknown
Age (years)	n Mean (SD)	239 64 (10)	137 64 (11)	72 63 (10)	30 67 (8)
Sex					
Female	n (%)	100 (42)	61 (45)	27 (38)	12 (40)
Male	n (%)	139 (58)	76 (55)	45 (63)	18 (60)
Number of days from diagnosis to inclusion	n/nmiss Mean (SD)	239/0 1677 (2012)	137/0 1930 (2225)	72/0 1295 (1762)	30/0 1437 (1290)
Octreoscan by Krenning score	n/nmiss	211/28	119/18	64/8	28/2
0	n (%)	0 (0)	0 (0)	0 (0)	0 (0)
1	n (%)	7 (3)	5 (4)	1 (2)	1 (4)
2	n (%)	12 (6)	10 (8)	1 (2)	1 (4)
3	n (%)	64 (30)	39 (33)	20 (31)	5 (18)
4	n (%)	71 (34)	35 (29)	21 (33)	15 (54)
Not done	n (%)	57 (27)	30 (25)	21 (33)	6 (21)
Ki-67 index	n/nmiss Mean (SD)	211/28 6 (6)	121/16 4 (3)	64/8 8 (6)	26/4 10 (11)
Subjects without Baseline CgA	n	13	8	3	2
CgA ≤ULN	n	32	21	7	4
CgA >ULN	n	194	108	62	24

Upper Limit of Normal

Table 2. Correlation analysis between changes in CgA and RECIST1.1, ITT population

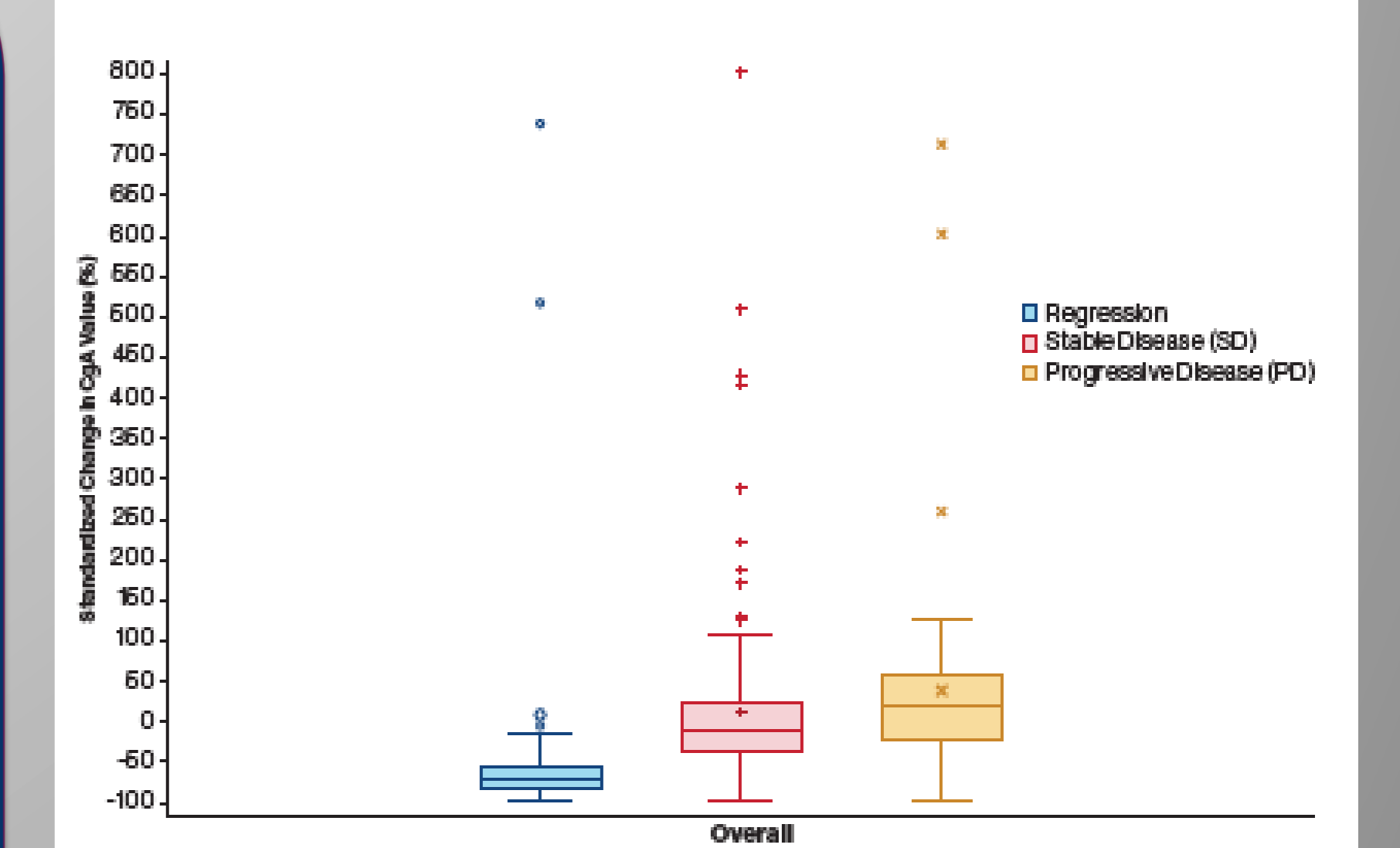
	Small Intestine	Pancreatic	Unknown	Other	Overall
Number of events included in analysis	137	123	40	4	304
Spearman's rank correlation coefficient	0.16	0.18	0.2		0.17
p-value	0.07	0.04	0.21		0.003

Only events with CgA test values that were examined 3 to 6 months prior to the CT-scans were included. The category of 'Other' contain too few numbers to perform correlation.

Results

- Patients demographics and history was similar across subgroups (Table 1)
- Of 304 events in the post-hoc analysis, 58 showed progression, 228 stable disease, and 18 regression (complete and partial response). The median change in plasma CgA was +19 (IQR: -20- (+57))%, -12 (-37- (+23))% and -73 (-83- (-55))%, respectively (Fig. 1)
- The overall Spearman's rank correlation coefficient was 0.17 (p=0.003), and 0.16 (p=0.07), 0.18 (p=0.04) and 0.20 (p=0.21) for small intestinal (137 events), pancreatic (123 events) and unknown GEP primary (40 events), respectively. (Table 2)

Figure 1. Boxplot of standardized change in CgA levels by RECIST1.1, Overall, ITT population



Conclusion

- This prospective observational study of patients with GEP-NET and unknown primary NET showed a positive correlation between CgA change from baseline and RECIST response. The predictive value of CgA change from baseline of RECIST response, however, remains to be established.
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