Prospective Evaluation of Neuromediator Dynamics in Castration-Resistant Prostate Cancer Patients During Docetaxel

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Abstract. Aim: Aim of the study was to detect small cell/neuroendocrine (SCNC) transformation in metastatic castration-resistant prostate cancer (mCRPC) that is a challenging procedure. We investigated the role of neuromediator dynamics as potential evidence of SCNC in patients undergoing docetaxel therapy. Patients and Methods: A multi-institutional, prospective observational study was conducted. Patients undergoing docetaxel treatment were included. Chromogranin A (CGA), neuron-specific enolase (NSE), and pro-gastrin releasing peptide (Pro-GRP) were sequentially evaluated at predefined time points. Outcome measures were overall survival (OS), progression-free survival (PFS) and PSA nadir. Results: Fifty-two patients were included. A general rise in CGA levels was observed. Patients with a high CGA rise (100%ULN: CGA≥98.lng/ml) between

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the 1st and 3rd cycle trended towards a decreased OS (p=0.0649) and showed a decreased PFS (p=0.0369). In multivariate analysis, continuous CGA rise correlated with PFS (p=0.0553; HR 1.136), but was not an independent predictor of OS. Conclusion: Patients with an early high CGA rise may demonstrate a subgroup with poor outcome due to underlying SCNC transformation. Monitoring of CGA appears to be an option worth considering.

The wide range of survival of patients with mCRPC is, at least partly, attributed to tumor heterogeneity. Its identification, prognostic impact, and consequent influence in treatment remain a leading challenge in mCRPC.

The development of CRPC is caused by androgendeprivation therapy (ADT) (1). Even though the first therapeutic hit with ADT is highly efficient, therapy resistance develops. This generates molecular changes of the androgen receptor axis (ARA) (1). An interference with the ARA also increases the emergence of SCNC during ADT (2-4). ARA-targeted agents abiraterone acetate (AA) and enzalutamide (ENZ) were granted approval. An ongoing prospective multicenter trial is evaluating the emergence of SCNC by biopsies in mCRPC that have progressed under treatment with AA or ENZ. First results show a SCNC or an intermediate atypical carcinoma in 40% of samples being significantly correlated with a decreased overall survival and the elevation of CGA and NSE (5). It has been shown that already CGA-positive cell clusters represent an early sign of invasion (6). The prognosis of SCNC is poor due to late recognition and heterogeneous clinical features (7).

The microtubule-stabilizing, semi-synthetic taxane DOC is the standard first-line chemotherapy for CRPC (8, 9). Earlier, researchers suggested that treatment with DOC significantly impairs the ARA by interfering with AR nuclear transport in CRPC (10, 11). *In vitro* findings showed that the acquisition of a neuroendocrine phenotype by prostate cancer cells leads to resistance of DOC (12).

Neuroendocrine cells synthesize several neuromediators. To detect neuroendocrine areas in prostate cancer, the most recent methodological directive suggests that any evaluation of the three common markers (CGA, CD56 or SYN) is sufficient. Although the diagnosis of SCNC is *a priori* a pathological diagnosis (13), there is good evidence in support of a significant relationship between neuromediator levels, tissue expression, and neuroendocrine activity in prostate cancer patients (5, 14, 15).

With growing emergence of SCNC, and possible treatment resistance to conventional therapies, it is important to identify patient cohorts that might benefit from biopsy sampling of metastases to detect SCNC transformation. Early detection of SCNC would result in change of therapy.

Here, we hypothesized that pretreatment or treatment with DOC may have influenced neuromediators, which, in turn, may impact patient outcome. To test our hypothesis, we conducted a prospective, multi-institutional observational study to sequentially evaluate serum neuromediators during DOC in mCRPC.

Patients and Methods

Study design and patients. Our investigator-initiated study was designed and coordinated by JvH, EH and CW. We conducted a prospective, observational ethics-committee-approved study at six institutions (two tertiary centers, one teaching hospital, and three outpatient practices) with the primary objective of evaluating dynamics from baseline of neuromediators in serum of patients with mCRPC undergoing DOC chemotherapy. Secondary objectives included to detect correlations of changes in neuromediators for OS, PFS and PSA nadir. The study is registered with Germanctr.de, number DRKS00004797.

Eligibility criteria were histologically-confirmed adenocarcinoma, mCRPC (newly diagnosed or with progression after regimens with AA and/or ENZ and/or DOC). Exclusion criteria were any history of cancer apart from low grade urothelial cancer of the bladder within the last 5 years; neuroendocrine cancer in the past medical history. Potential clinical manifestations associated with SCNC were scored as described previously (13). All study participants provided written informed consent after full explanation of the purpose and nature of all procedures.

Patients received DOC 75 mg/m² intravenously 3-weekly. The dose of DOC was reduced if toxic effects deemed unacceptable were noted. The decision to end treatment due to toxic effects, progression, or other reason was in hands of the treating physician.

Serum collection, measurement and storage. Blood was collected in serum-gel tubes during routine blood withdrawal within seven days

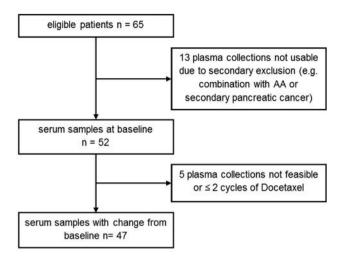


Figure 1. CONSORT diagram on enrollment and attrition.

before application of the 1st, the 3rd, and the 6th cycle of systemic chemotherapy with DOC. In case of dose modifications, the blood withdrawal (bw) scheme was adapted. All serum samples were sent by medical express service to a reference laboratory. For quantitative determination of CGA the Time-Resolved Amplified Cryptate Emission (TRACE) method (Cisbio Bioassays, France), of NSE the chemoluminescence immunoassay (CLIA) LIAISON® NSE (DiaSorin, Italy) and of Pro-GRP the chemiluminescent microparticle immunoassay (CMIA) (Abbot, Germany) were used.

Statistical analysis. First, we calculated the change from baseline of neuromediators under DOC. From clinical experience and reported case series, we assumed that at least a total increase of 100% of the upper limit of normal (ULN) of the widely used serum markers CGA and NSE would be clinically relevant (high rise). Second, we therefore investigated the correlation of an early high rise and an early change (1st – 3rd cycle) from baseline for OS, PFS and PSA nadir. For OS, patients were followed until the date of death from any cause. Patients who did not experience the event by the end of the study were censored at the time of last available follow-up. We analyzed PFS, and PSA nadir according to recent recommendations (16).

All statistical calculations were conducted on SAS® release 9.3 (SAS Institute Inc., NC, USA). For categorical factors, absolute frequencies and relative proportions are given; approximately normally distributed variables (*i.e.* age) are presented by mean value and standard deviation; for skewed variables, median and range are given. In order to compare measurements over time, repeated measures ANOVA was performed using the SAS procedure PROC MIXED. If the ANOVA yielded a statistically significant result Dunnett's test was used as a *post hoc* test for pairwise comparisons. The test was conducted as two-sided tests. For these techniques, the logarithm of marker values were used in order to get approximately normally distributed data.

Furthermore, Kaplan-Meier curves were created to illustrate event rates. The Gehan-Wilcoxon test was used to investigate the influence of an early high rise of CGA and NSE for OS and PFS. As the beginning of the observation is crucial for the patient's course of the

Table I. Characterization of the study cohort.

Patient characteristics (N=52)	N (%)
Age (years)	
Mean±SD	71.3±7.2 (55-88)
ECOG performance status	
0	34 (65.4%)
1	19 (34.6%)
Charlson comorbidity index	
<2	42 (90.8%)
≥2	10 (19.2%)
First diagnosis prostate cancer –	
study entry (months)	
median (range)	34 (6-220)
Gleason score in biopsy/	
surgical specimen (missing n=5)	
6	2 (4.3%)
7	12 (26.7%)
8	12 (26.7%)
9	20 (42.6%)
10	1 (2.2 %)
Site of metastatic disease*	
Bone	42 (80.8%)
Visceral metastasis	14 (26.9%)
Lymph node metastasis	35 (67.3%)
Clinical manifestation of NED	45 (86.5%)
Prior systemic treatment*	
Androgen deprivation therapy	52 (100%)
Months, median (range)	24 (3-139)
Docetaxel	4 (7.7%)
Abiraterone Acetate	24 (46.2%)
Enzalutamide	7 (13.5%)
Blood based marker	· · ·
Hemoglobin (Hb), g/dl	12.1 (7.0-15.1)
Alkaline phosphatase (AP), U/I	139 (46-2,226)
Lactate dehydrogenase (LDH), U/I	276 (128-1,586)
Intake of proton pump inhibitors	16 (30.8%)

Data are number (%) unless stated otherwise. ECOG=Eastern Cooperative Oncology Group; *patients could have more than one.

disease, the Gehan-Wilcoxon test seems to be appropriate because it gives greater weight to the earlier differences at the beginning of the observation than the log rank or the cox regression analysis (17). Influences for OS and PFS were evaluated by Cox regression models for hazard ratios (HR) and 95% confidence interval (CI). All variables in the univariate analysis with a significant test result were included in the multivariable model (using the "selection=forward" method with alpha=0.10). To test for influence of the variables on PSA nadir, the Mann-Whitney *U*-test was used.

A test result was considered statistically significant for p < 0.05. A test result with 0.05 has been regarded as trending towards significance.

Results

Study population. Between February 2013 and July 2015, 65 patients met the eligibility criteria and were included in the

Table II. Marker distribution (bw: blood withdrawal).

Marker	1st bw	2nd bw	3rd bw			
_	Median (range)					
CGA (ng/ml)	132 (22-1023)	146 (12-2027)	160 (39-961)			
	(n=49)	(n=48)	(n=39)			
NSE (ng/ml)	17 (10-60)	19 (6-117)	15 (9-68)			
	(n=49)	(n=45)	(n=37)			
Pro-GRP ($\mu g/ml$)	19 (6-122)	15 (3-77)	15 (5-36)			
	(n=49)	(n=47)	(n=36)			
PSA (ng/ml)	143(0-2525)	62.4 (0-2164)	22 (0-1765)			
	(n=52)	(n=46)	(n=37)			

study. Thirteen patients had to be excluded secondarily due to *e.g.* combination with AA or primary hormone chemotherapy (Figure 1). The clinical and pathological data of all patients are listed in Table I. Thirty-seven (71.1%) patients had died at the time of analysis. For 19 (47.5%), PFS could be provided, whereas 21 (52.5%) patients were censored.

Expression and dynamics of neuromediators. Serum concentrations of CGA, NSE, Pro-GRP, and PSA at all time points of bw are given in Table II. The distribution of CGA, NSE and Pro-GRP is shown in boxplots in Figure 2. Levels of CGA increased significantly from baseline compared to the 2nd and the 3rd blood withdrawal (p=0.0146 and p=0.0330). Levels of NSE did not change during therapy with DOC (p=n.s.). Levels of Pro-GRP significantly decreased during therapy compared to baseline (p=0.0424 and p=0.0114). Waterfall plots depicting changes of the markers are presented in Figure 3. A total increase $\geq 100\%$ of the upper limit of normal (ULN) of CGA was observed in 11 patients and of NSE in 5 patients. The total change was shown for Pro-GRP, as no reference range exists.

Associations of neuromediators dynamics with clinical outcome. A high rise from baseline of CGA trended towards significance with OS (p=0.0649). A high change of NSE was not associated with OS (p=0.7744). A high rise from baseline of CGA was associated with PFS (p=0.0369). A high change of NSE not associated with PFS (p=0.2895). Kaplan-Meier curves for association of early high rise of CGA with OS and PFS are shown in Figure 4. Cox regression analysis for associations of change of neuromediators and other variables with OS and PFS are presented in Table III and Table IV. PSA-nadir was associated with previous therapy with AA (p=0.0285), ENZ (p=0.0593), and AP-levels at baseline (0.0386).

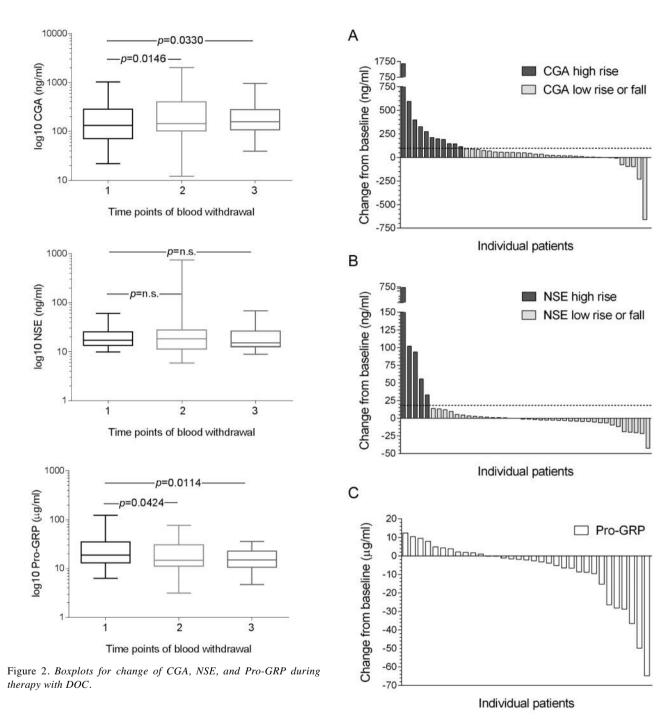


Figure 3. Waterfall diagrams for change from baseline during therapy with DOC (1st to 3rd cycle): (A) CGA with dashed line signifies total rise of ≥ 98.1 ng/ml, (B) NSE with dashed line signifies total rise of ≥ 18.3 ng/ml, (C) Pro-GRP.

Discussion

We present here the first prospective study evaluating dynamics in serum neuromediators in men with mCRPC undergoing DOC. We found that CGA levels rise during therapy with DOC, but only an early high rise in CGA levels

trended towards significance with shorter OS and was associated with PFS indicating potential underlying SCNC transformation in a small subgroup. We observed that dynamics of NSE vary, whereas those of Pro-GRP decrease.

Table III. Cox regression analysis for associations of change (1st to 3rd cycle) of CGA, NSE, Pro-GRP, and further clinical and laboratory variables with OS.

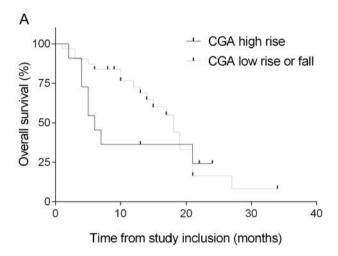
Variable	Categories	Univariate analysis			Multivariate analysis		
		N	HR (95 % CI)	<i>p</i> -Value	N	HR (95 % CI)	p-Value
CGA change; 100 units	Continuous	43	1.069 (0.926-1.235)	0.3633			
NSE change; 20 units	Continuous	42	0.956 (0.827-1.105)	0.5419			
Pro-GRP change; 10 units	Continuous	43	0.986 (0.764-1.272)	0.9110			
PSA; 100 units	Continuous	50	1.075 (1.019-1.133)	0.0076	46	1.085 (1.020-1.153)	0.0155
AP; 100 units	Continuous	48	1.121 (1.046-1.200)	0.0011	46	1.136 (1.020-1.153)	0.0002
Hb	Continuous	51	0.830 (0.695-0.992)	0.0403			
LDH; 100 units	Continuous	46	1.117 (1.005-1.242)	0.0402			
Age	Continuous	52	1.047 (0.994-1.104)	0.0856	46	1.052 (0.997-1.109)	0.0609
ECOG	Binary 0 vs. ≥1	34 vs. 19	1.322 (0.660-2.645)	0.4310			
AA therapy	Binary		1.805 (0.921-3.536)	0.0855			

Table IV. Cox regression analysis for associations of change (1st to 3rd cycle) of CGA, NSE, Pro-GRP, and further clinical and laboratory variables with PFS.

Variable	Categories -	Univariate analysis			Multivariate analysis		
		N	HR (95 % CI)	p-Value	N	HR (95 % CI)	p-Value
CGA change; 100 units	Continuous	37	1.136 (0.999-1.291)	0.0520	34	1.136 (0.999-1.291)	0.0520
NSE change; 20 units	Continuous	33	0.960 (0.804-1.146)	0.6516			
Pro-GRP change; 10 units	Continuous	34	1.121 (0.736-1.708	0.5946			
PSA; 100 units	Continuous	39	1.031 (0.956-1.112)	0.4303			
AP; 100 units	Continuous	39	1.042 (0.895-1.214)	0.5952			
Hb	Continuous	39	0.806 (0.606-1.072)	0.1384			
LDH; 100 units	Continuous	36	1.070 (0.920-1.245)	0.3782			
Age	Continuous	39	1.013 (0.943-1-089)	0.7174			
ECOG	Binary 0 $vs. \ge 1$	39	2.027 (0.727-5.652)	0.1767			
AA therapy	Binary	39	1.673 (0.632-4.428)	0.3003			

Aparicio et al. suggested potential clinical manifestations associated with SCNC (18). We applied these criteria and discovered that a majority of patients (86.5%) have these features. The role of DOC as a putative driver of transformation to SCNC was not studied, although about one-third of DOC-pretreated men were found to have increased neuroendocrine marker levels (19). Based on our analysis DOC drives CGA levels, generally, during the course of therapy. During DOC-rechallenge combined with carboplatin, 33 patients were followed for CGA and NSE change, but the change was not correlated with time to progression or survival (20). In a small pilot study investigating Octreotide acetate in CRPC, CGA decrease during therapy could not identify clinical responders, but the study is biased by the small sample size (n=17) (21). In gastroenteropancreatic neuroendocrine tumors change of CGA predicted treatment response (22). As the general increase of CGA in our study was not correlated to disease outcome, we believe that it may show a neuroendocrine stress response of neuroendocrine cells within the tumor or in general, but it should not be interpreted that DOC is a general driver of SCNC transformation.

Few patients showed a high rise of CGA during DOC. SCNC transformation in mCRPC is a histological diagnosis (13). To date, circulating tumor cells as an alternative are still highly experimental biomarkers and not used in the clinic routinely (23). Therefore, to identify SCNC transformation, sequential biopsies of metastases would be needed prior to several lines of therapy. Single core biopsy sampling may not be an accurate representation of the tumor as a whole due to intratumor heterogeneity shown in advanced prostate cancer (24). Small *et al.* reported that both visceral metastases and lymph node metastases can harvest SCNC (5). Due to the invasive and sometime cost-intensive (if CT-guided) procedure,



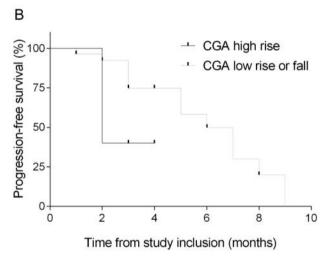


Figure 4. Kaplan-Meier curves for the association of CGA with OS (A) and PFS (B).

a preselection of patients at risk for SCNC transformation using circulating neuromediators could be useful.

From the immunohistochemical markers listed above, only CGA can be used as an established serum biomarker. Eleven patients experienced an early high rise of CGA, which was correlated with PFS and OS. Circulating CGA was used as objective tumor biology assessment tool in five trials including >500 patients with mCRPC in total according to a recent review (25). A high rise has not been studied in these trials. In the literature, a predominance of arbitrary thresholds were chosen after one-time analysis before therapy. Two recently published trials analyzed CGA only before AA therapy and found predictive and prognostic significance (26, 27). In our study some patients in the subgroup of patients with early high CGA rise still lived long. This could be in line with a review of Wang *et al.* that

analyzed the clinical course of patients after diagnosis of neuroendocrine prostate cancer (7). Approximately 20% of the patients lived longer than 12 months, making the statistical analysis with our sample size for OS difficult. In the multivariate analysis, markers for the general volume of the disease were prognostic and CGA failed to show independent significance. Dynamics might have fewer biases as they are compared to baseline levels. Second dynamics can reflect the activity of the compartment under therapy. An early high rise of CGA should be further tested as a preselection tool for multi-core biopsy-sampling of metastases.

NSE and Pro-GRP are not recommended in immunohistochemistry, but they are serum-based and provide information about dynamics of other SCNC cell types. We were unable to detect any relevant associations of NSE with disease outcome. NSE in comparison to CGA showed to be a superior marker in small cell neuroendocrine cancers, whereas CGA was more sensitive in large cell neuroendocrine cancers (28). An explanation could, therefore, be that NSE is expressed by an aggressive small cell subtype of neuroendocrine cells and develops late in CRPC. The GRP/Bombesin receptor was shown to be over expressed in malignant prostate cells, and secretion of GRP by neuroendocrine cells in a paracrine manner might be one escape mechanism for prostate cancer when in an androgen independent stage (29). The growing field of GRP/Bombesin-based imaging for prostate cancer raises the need for more detailed knowledge concerning the role of Pro-GRP/GRP in the development and progression of prostate cancer (30). Yashi et al. found similarly to our results, increased Pro-GRP values in CRPC patient and observed also a decrease of Pro-GRP during ADT (31).

This analysis has limitations. No tandem evaluation of neuroendocrine serum markers and biopsy immunostaining in parallel was performed. Tandem evaluation was suggested to be the most reliable method to assess transformation to SCNC in mCRPC (15). Secondly, the patients did not receive a standardized staging protocol. Therefore radiographic progression-free survival could not be assessed. Thirdly, due to the small sample size it was difficult to gain significant results. Further larger studies are required to confirm our results.

Our prospective study demonstrated that dynamics of neuromediators vary, but early high CGA rise independently from PSA may indicate patients with poor outcome. Through the established low-cost serum marker CGA, the early reanalysis to assess dynamics could easily be used in daily practice. The high rise is possibly due to expression of a subclone of prostate cancer cells with SCNC transformation. Therefore, the cohort might benefit from multi-core biopsy sampling to detect SCNC transformation, if our results have been validated in larger cohorts. The ability to detect and monitor tumor heterogeneity like SCNC transformation remains a leading challenge for patients suffering from mCRPC.

Conflicts of Interest

EH and JvH have received speaker honoraria from Sanofi-Aventis. All remaining authors have declared no conflicts of interest.

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