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Increased primary breast tumor expression of CD73 is associated with development of bone metastases and is a potential biomarker for adjuvant bisphosphonate use

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Purpose. Increased CD73 expression has been associated with progression in various cancer types. Results of the AZURE and other trials suggest that, in postmenopausal breast cancer patients, adjuvant bisphosphonates inhibit bone relapses and prolong overall survival. Based on these findings, adjuvant bisphosphonates (typically zoledronic acid) are standard-of-care in postmenopausal patients with high-risk early breast cancer. However, biomarkers are needed for improved patient selection. The aim of this study was to investigate the association of primary tumor CD73 expression with later development of bone metastases. Methods. To determine whether CD73 levels correlated with tumor parameters (hormone receptor status, tumor stage and grade), patient outcomes (bone metastases and survival) or other patient characteristics (menopausal status, chemotherapy or statin use), we analyzed primary breast tumor CD73 expression immunohistochemically in tumor microarray samples from the AZURE (BIG01/04) trial. Results. In the AZURE control arm, high CD73 score are significantly prognostic for overall survival (p-value = 0.03, HR = 1.87, 95% CI = 1.06-3.29), disease-free survival (p-value = 0.06, HR = 1.66, 95% CI = 0.982-2.8) and time to first metastasis to bone (p-value = 0.04, HR = 2.23, 95% CI = 1.04–4.81), as compared with low CD73 scores. However, high CD73 score did not display an association with time to non-bone metastasis or first recurrence to a non-skeletal site. In the zoledronate arm, high CD73 score did not have association with patient outcomes, first metastasis to bone, nor with bone recurrence at any time (distant recurrence, including skeletal) or first non-skeletal recurrence. In multivariate testing, CD73 had no significant association with age, ER status, tumor stage, histological grade, menopausal status, chemotherapy or statin use in either arm. Conclusions. High CD73 expression is associated with development of bone metastases. Zoledronate counteracts this effect. These results suggest that CD73 expression might serve as a biomarker for adjuvant zoledronic acid use.

Keywords CD73, Breast cancer, TMA, Bone metastases

Disease recurrence remains a significant problem among patients with early breast cancer that have been treated with curative intent. Approximately 70–80% of patients who do relapse, develop bone metastases¹. Bone metastases may linger in a senescent stage for years, possibly decades, before they grow into symptomatic macroscopic metastases but, once manifest, they may cause pain, pathological fractures, hypercalcemia and

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spinal cord compression. However, time to relapse depends greatly on breast cancer subtype. Patients with triple-negative breast cancer (TNBC) typically relapse within the first five years after treatment, whereas those with hormone receptor-positive subtype may relapse many years later^{2,3}.

Dormant cells play a critical role in the recurrence of cancer, particularly in breast cancer. These cells, known as disseminated tumor cells, can remain in a dormant state within the bone marrow for years, after the primary tumor has been surgically removed. Dormant disseminated tumor cells in the bones depend on pro-survival signals from the microenvironment and develop complex immune evasion mechanisms, leading to disease recurrence⁴. In ER-positive breast cancer patients and those with larger tumor sizes, pre-operative dormant tumor cells were a significant predictor of late systemic recurrences⁴. For cancers that commonly spread to bones, such as breast cancer, treatment plans could be based on prognostic markers to improve long-term outcomes and prevent late recurrences. Currently, bisphosphonates are the only standard treatment that may prevent the formation of bone metastases. When given in the adjuvant setting to post-menopausal patients with early breast cancer with high risk of relapse, they may prevent bone metastases, but this benefit does not extend to pre-menopausal patients. The reason for this is currently unknown, but it may be immunologically regulated⁵. However, although a number of potential biomarkers to assess the risk of recurrence and predict response to bisphosphonates have been identified⁶⁻⁹, none are currently in routine clinical use for identification of risk and response, and the number needed to be treated remains substantially higher than the numbers who will benefit. Although bisphosphonate drugs are relatively safe and well understood, it would be beneficial to have biomarkers to guide their clinical use.

The large (N=3300), academic, multi-center, randomized phase III AZURE (BIG01/04) study evaluated the addition of the bisphosphonate zoledronic acid (4 mg) to standard therapy (neo/adjuvant chemotherapy and/or endocrine therapy) for 5 years 10,11 . In this and other adjuvant bisphosphonate studies, analyzed in a large (N=18 700) meta-analysis, the benefits of treatment were limited to post-menopausal women and the benefit remained rather low at 3.3% for breast cancer mortality and 2.2% for recurrence at 10 years (EBCTCG). Currently, the most promising predictive biomarker for adjuvant bisphosphonate use is the MAF (mesenchymal aponeurotic fibrosarcoma) gene. Specifically, patients with MAF-negative breast cancers appear to benefit from adjuvant bisphosphonates. The predictive ability of c-MAF amplification to determine the outcome of treatment with bisphosphonates (either zoledronic acid or clodronate) was recently outlined in two key studies 9,12 . Despite these promising studies, further validation of potential biomarkers prognostic for breast cancer spread to bone and metastasis treatment are required before these markers are considered for routine clinical use.

CD73 is a membrane-bound receptor, the main function of which is to hydrolyze ribonucleoside 5′-monophosphates (AMP, uridine monophosphate, inosine monophosphate and cytidine monophosphate) to extracellular adenosine and inorganic phosphate¹³. Enzymatic activity of CD73, due to adenosine production, is associated with immunosuppressive events in cells¹⁴. As a regulator of tumor immunity, CD73 is also a novel target for immunotherapies. There are several ongoing phase I (NCT03875573, NCT03616886, NCT05431270) or phase II¹⁵ trials to investigate the efficacy of CD73 inhibition in breast cancer, as well as in other types of cancer¹⁶. We and others have demonstrated previously that CD73 expression is associated with metastases^{17,18}, especially among TNBC¹⁹. The aim of this study was to investigate whether primary tumor CD73 expression is prognostic specifically for bone metastases, using tissue samples and the associated clinical database from the AZURE trial.

Materials and methods Sample information

AZURE study

AZURE (BIG01/04) was an academic, prospective, randomized controlled phase III, open-label multi-national and multi-center trial (ISRCTN79831382). Patients with histologically confirmed invasive breast cancer of any biological subtype, with either pathologically confirmed axillary lymph node metastases or a T3/T4 primary tumor were enrolled. Patients were not eligible for this study if there was clinical or imaging evidence of distant metastases prior to study entry or current or recent (previous year) use of bisphosphonates or pre-existing bone disease likely to require bone-targeted treatment 11,20. All patients gave written informed consent for this study. Between September 2003 and February 2006, 3360 patients from 174 centers were randomized on a 1:1 basis, using a central automated 24-hour computer-generated telephone randomization system to receive (neo)adjuvant systemic therapy with zoledronic acid or without (Control) zoledronic acid. A minimization method ensured balance in key prognostic and treatment variables across the two groups 11. AZURE patients were reviewed annually and followed for 10 years. All analyses on patient samples were ethically approved (Ethics reference number: 55/03/182). Analyses for correlation of CD73 levels with risk of bone metastasis were performed on tissue microarrays (TMAs) constructed from primary tumors from a subset of patients within the AZURE trial (n = 689). Due to the relatively high prevalence of bone metastatic outcome within this group, and the long follow-up (median 117 months [interquartile range 70.4–120.4]), these TMAs provide an excellent resource for validation of protein biomarkers emerging from our histological analysis.

Immunohistochemistry (IHC)

Protein expression was assessed on TMAs using immunohistochemistry. Briefly, 5 µm serial tissue sections were dewaxed in xylene and rehydrated through graded alcohols. Endogenous peroxidases were blocked (3% H2O2 for 10 min), and antigens retrieved by microwaving the slides in sodium citrate buffer, pH 6 (800 W for 7 min). After cooling and washing, slides were blocked with normal goat serum for 1 h at room temperature, after which primary antibodies against CD73 (CST, 13160, 1:1000) were applied (overnight, 4 °C). The slides were incubated with biotinylated secondary rabbit antibodies (BA-2000 Vector, UK) at room temperature for 1 h, followed by incubation with avidin/biotinylated enzyme complex (ABC) for 1 h, 3,3′-diaminobenzidine (DAB)

for 1 min and counterstaining with hematoxylin for 10 s (both Vector laboratories, UK). Slides were scanned using Pannoramic 250 slide scanner (3DHISTECH Ltd, Hungary).

IHC scoring

A standard IHC approach of semi-quantitative scoring based on operator assessment of weak to intense staining was performed using a simple scoring system of 0 (no expression), 1 (weak expression), 2 (intermediate expression) and 3 (strong expression). For antibody optimization, IHC staining was performed using a range of primary antibody dilutions. The dilution that gave the widest range of cytoplasmic staining intensities across different TMA samples was chosen.

This optimized approach was then applied to TMAs from the AZURE study. IHC staining was scored as either level 0, 1, 2 or 3 as described above. Scoring was performed blindly by two trained operators. Scoring was based on staining intensity only, and not on percentage of positive cells. This was appropriate for CD73, since when staining was present, it was of a similar intensity throughout the tumor cells, rather than being patchy or focal in nature, and therefore the number of stained cells was not an issue. Kappa coefficients were calculated for the two independent scores demonstrating very significant agreement (Kappa coefficient > 0.85). Discrepant scores were assessed by an independent adjudicator and values assigned. CD73 IHC scores were submitted via patient identification numbers to the clinical database of the AZURE study for alignment to the AZURE clinical data and statistical analyses (Walter Gregory).

Statistical analysis

CD73 scoring data was analyzed within Excel to determine the frequency of CD73 scores within defined patient subgroups (e.g. tumor stage) for each arm of the trial. The distribution of CD73 scores between different subgroups of each such variable was then tested for association by use of a heterogeneity Chi-squared test, with degrees of freedom equal to the number of categories for that variable minus 1. Immunohistochemical analyses followed REMARK guidelines²¹. Statistical analyses evaluated the associations between CD73 protein expression and relevant clinical and pathological variables (e.g. ER/PR/HER2 status) using Fisher's exact test (categorical variables) and the Kruskal–Wallis test (continuous variables), before assessing prognostic and predictive associations with time-to-event data (time to first distant recurrence, time to first skeletal recurrence, time to first non-skeletal recurrence) using Cox proportional hazards regression, the Kaplan–Meier estimate of the survival function and the log-rank test. Time to first distant recurrence was defined as the time from the date of randomization to the date of the distant recurrence. In analyses, other types of events were censored. For example, if a local recurrence occurred prior to any distant recurrence, the patient would be censored at the date of the local recurrence. Time to first skeletal recurrence and first non-skeletal recurrence were defined similarly. Time to first skeletal recurrence irrespective of all other previous recurrences was also investigated. Time to event analysis was first performed within treatment arms to identify prognostic associations with CD73.

Multivariate analysis looking for potential correlation of CD73 levels with patient variables (including menopausal status, ER, HER2 and PR status) were performed using Mann-Whitney U-test. The predictive heterogeneity of effect between treatment arms for time to distant events was then assessed in multivariable analysis by including an interaction term in the Cox proportional hazard regressions for treatment arm and biomarker (while adjusting for systemic therapy plan, ER status and lymph node involvement). All significance tests were two-sided and were designated significant at the 5% level.

E+hice

The study was performed in accordance with ethical principles originating in the Declaration of Helsinki. This research study at the University of Turku was approved by the Finnish National Board on Research Integrity TENK (ISBN 978-952-5995-88-6). Data were anonymized for statistical analyses and handled in a manner that meets the EU General Data Protection Regulation 2016/679 (GDPR) on data protection.

Results

Clinical association of CD73 expression with patient characteristics in breast cancer patients in the AZURE trial

In the study, CD73 staining intensities were analyzed from a TMA cohort from the AZURE (BIG/01/04) trial. All analyses comparing CD73 scores with patient variables, or with time-to-event data, employed data from IHC staining of tumor cores, where CD73 staining intensity was scored as either 0, 1, 2 or 3 (Fig. 1). Dichotomized staining scores were used to compare high CD73 expression (Score = 3) with low CD73 expression (score = 1 or 2). We observed that most cores exhibited low CD73 expression, with negligible instances of high expression. Cores which were not of sufficient quality to score accurately were assigned as non-scorable (NS). The possible association of CD73 with outcome was tested for 422 patients (204 in the Control Arm and 218 in the Zoledronate Arm). The demographic data for the patients with assessable cores and the overall AZURE population were similar (Table 1).

In multivariate testing, CD73 had no significant associations with age, ER status, tumor stage, histological grade, menopausal status, chemotherapy or statin use (Table 2). The measurement of HER2 status was not mandated within the AZURE trial and HER2 measurements are available for 307 participants in total, of which 168 of these patients had assessable cores for CD73 staining. Within this analysis, CD73 did not associate significantly with HER2 status. In addition to these analyses within the control arm of AZURE, no significant associations of CD73 level with clinical variables were observed within the zoledronic acid arm (Table 2).

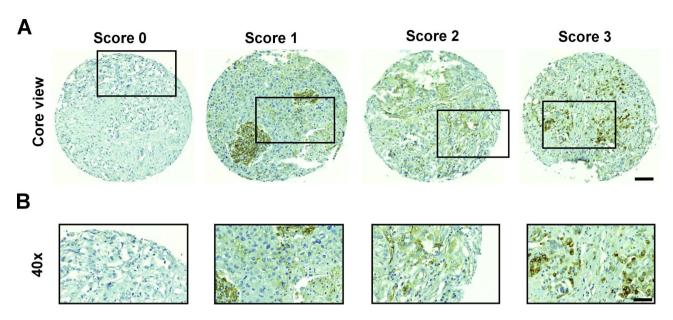


Fig. 1. Representative images of immunostaining for CD73 in TMA cores from patients in the AZURE study. (A) Images of protein expression obtained using the anti-CD73 antibody and visualized at a magnification of 20x. Score 0–3 categories were determined by the intensity of staining in the cytoplasmic compartment of tumor cells only. Scale bar = 200 μ m. (B) 40x magnification view of from the boxed regions depicted in panel A. Scale bar of core view = 100 μ m. Scale bar of $40x = 50 \mu$ m.

Association of CD73 expression with patient outcome

Kaplan-Meier analysis of overall survival showed significant association of high CD73 expression with worse prognosis compared with those for low CD73 expression (p-value=0.03, HR=1.87, 95% CI=1.06–3.29) in control arm but not in the zoledronic acid arm (Fig. 2A, B; Table 3). Association between high CD73 expression and worse disease-free survival (DFS) almost reached significance compared with those with low CD73 expression (p-value=0.06, HR=1.66, 95% CI=0.982–2.8) in the control arm but not in the zoledronic acid arm (Fig. 2C, D; Table 3).

CD73 expression is associated with distant recurrence

Kaplan-Meier estimates for the survival function for the time to distant recurrence confirmed that high CD73 expression is a significantly prognostic factor for time to first metastasis to bone (p-value=0.04, HR=2.23, 95% CI=1.04–4.81) within the control arm (Fig. 3A; Table 3). Expression of CD73 did not correlate with any first recurrence (including skeletal) (p < 0.06, HR=1.77, 95% CI=0.97–3.21) or to a non-skeletal site (p < 0.17, HR=1.74, 95% CI=0.791–3.82) within control arm (Fig. 3C, E; Table 3). We did not observe an association between CD73 level and first skeletal event whether other distant events had occurred first (p < 0.06, HR=1.86, 95% CI 0.97-3.55) within control arm (Table 3). The inclusion of metastasis to another site thus would appear to reduce the association seen during the bone only analysis.

In the zoledronic acid arm, we did not observe association between high CD73 levels and time to first metastasis to bone (p-value=0.92, HR=0.945, 95% CI 0.316–2.83), bone recurrence at any time (distant recurrence including skeletal) (p-value=0.50, HR=0.717, 95% CI 0.275–1.87) or first non-skeletal recurrence (p-value=0.70, HR=0.85, 95% CI 0.374–1.93) (Fig. 3B, D, F; Table 3). These results suggest that the increased risk for first metastasis to bone patients with high tumor CD73 expression is counteracted by treatment with zoledronate.

Discussion

Breast cancer metastasis to bone is associated with increased patient morbidity with development of bone-specific effects, including bone fractures, spinal cord compression and hypercalcemia. In the current study, we have demonstrated that high levels of CD73 expression within primary breast tumors are associated with a greater risk of cancer spread first to bone.

CD73 is a cell surface protein found on normal epithelial and stromal cells^{22–26} and tumor cells^{27–32}. CD73 has prognostic value in TNBC³³. Specifically, low tumor cell expression of CD73 is associated with prolonged disease-free survival in TNBC patients. Furthermore, high CD73 expression on epithelial cells was negatively associated with tumor infiltrating lymphocytes, suggesting an immunoregulatory mechanism behind the survival difference³³. In the present study, we found a correlation between high CD73 expression and OS or DFS in breast cancer patients from control arm of the AZURE trial. However, we did not observe such correlation in patients treated with adjuvant zoledronate.

In normal tissues, CD73-derived adenosine exerts tissue protective functions, but in cancer, it may help tumors escape immune surveillance⁵. Because of this aspect in tumor immunology, CD73 has emerged as a

	CD73 dataset		Full AZURE Trial p	opulation		
Characteristic	Control (n = 204)	Zoledronate (n = 218)	Control (n = 1678)	Zoledronate (n = 1681) 51 (21-89)		
Age (years) Median (Range)	53 (30-79)	52 (26–77)	51 (20-87)	51 (21–89)		
Tumour stage						
T1	66 (32.35)	77 (34.84)	523 (31.2)	542 (32.2)		
T2	104 (50.98)	117 (52.94)	867 (51.7)	851 (50.6)		
T3	32 (15.69)	22 (9.95)	228 (13.6)	227 (13.5)		
T4	2 (0.98)	5 (2.26)	59 (3.5)	58 (3.5)		
Neo-adjuvant therapy						
Yes	2 (0.98)	7 (3.17)	1564 (97.60)	1598 (93 – 50)		
No	202 (99.02)	214 (96.83)	38 (2.40)	104 (6.50)		
Systemic therapy plan	•					
Endocrine Therapy Alone	13 (6.37)	18 (8.14)	74 (4.5)	76 (4.5)		
Chemotherapy Alone	50 (24.51)	41 (18.55)	360 (21.5)	362 (21.5)		
Endocrine and Chemotherapy	141 (69.12)	162 (73.30)	1243 (74.1)	1243 (73.9)		
Anthracyclines	I	1	I.	I		
Yes	186 (91.18)	207 (90)	1564 (97.6)	1564 (97.6)		
No	18 (8.82)	23 (11.11)	38 (2.4)	38 (2.4)		
Taxanes			1	1		
Yes	31 (15.20)	30 (13.57)	385 (24.0)	390 (24.3)		
No	173 (84.80)	191 (86.43)	1219 (76.0)	1215 (75.7)		
Use of statins		()	. ()	(, , , ,		
Yes	12 (5.88)	12 (5.43)	101 (60)	97 (5.8)		
No	192 (94.12)	209 (94.57)	67 (40)	1575 (94.2)		
Menopausal status	,			,		
Pre-menopausal	92 (45.10)	94 (43.53)	752 (44.8)	751 (44.7)		
≤5 years since menopause	38 (18.63)	35 (15.84)	244 (14.5)	247 (14.7)		
> 5 years since menopause	58 (28.43)	69 (31.22)	522 (30.9)	519 (30.9)		
Menstrual status unknown	16 (7.84)	23 (10.40)	160 (9.5)	164 (9.8)		
ER status	10 (7.01)	20 (10:10)	100 (3.0)	101 (310)		
ER Positive	155 (76.35)	177 (80.09)	1316 (78.4)	1319 (78.5)		
ER Negative	48 (23.65)	44 (19.90)	355 (21.2)	349 (20.8)		
Status Unknown	0	0	7 (0.4)	13 (0.8)		
PR status	0	0	7 (0.1)	13 (0.0)		
PR Positive	70 (34.31)	76 (34.90)	698 (41.6)	725 (43.1)		
	44 (21.57)	40 (18.10)	424 (25.3)	382 (22.7)		
PR Negative	ļ	104 (47.06)				
Status Unknown Missing	88 (43.14)	, ,	548 (32.7)	571 (34.0)		
	2 (0.98)	0	1 (0.45)	0		
Lymph node involvement	122 (65 02)	144 (65 45)	1022 (61.5)	1041 (61.0)		
1–3 Nodes involved	132 (65.02)	144 (65.45)	1032 (61.5)	1041 (61.9)		
≥4 Nodes involved	70 (34.48)	76 (34.55)	608 (36.2)	604 (35.9)		
Unknown Involvement	1 (0.49)	0				
Missing Data	0	0				
HER2 status	20 (4 (20)	24 (0.50)	200 (40.0)	1.00 (4.4.1)		
HER2 Positive	29 (14.29)	21 (9.50)	223 (13.3)	192 (11.4)		
HER2 Negative	52 (25.62)	66 (29.86)	603 (35.9)	648 (38.5)		
Unknown/ Not Measured	121 (59.61)	132 (59.73)	843 (50.1)	831 (49.5)		
Missing Data	1 (0.49)	2 (0.91)				
Histological grade	1	T /:	T	T		
Grade 1	13 (6.34)	13 (5.88)	140 (8.3)	145 (8.6)		
Grade 2	82 (40)	92 (41.63)	708 (42.2)	731 (43.5)		
Grade 3	109 (53.17)	114 (51.58)	787 (46.9)	765 (45.4)		
Grade 4	0	1 (0.45)				
Missing Data	1 (0.49)	1 (0.45)				

Table 1. Characteristics of the patients whose tissue was assessed on TMAs in this study (as at baseline in the AZURE study).

	Control	-									Zolodronoto	onoto						
	Collin	5									rollen	Ollate						
	Score 0	0	Score 1		Score 2	2	Score 3				Score 0	0	Score 1		Score 2	Score 3	3	
Media (range)	50.5 (33-79) $n = 172$	(3-79)	$52 (32-73) \\ n = 33$	-73)	49		19		Mec	Media (Range)	51 (29–77) n=175	-77)	50.5 (33-79) n=45		61.5 $(58-65)$ $n=2$			
	Total	%	Total	%	Total	%	Total	% p-value	lue		Total	%	Total	%	Total %	Total	%	p-value
Tumour stage									Tun	Tumour stage								
T1	99	84.84	6	13.64	1	1.52	0	NS 0	T1		64	84.21	12	15.79	0 0	0	0	SN
T2	84	80.77	17	16.35	3	2.88	0	0	T2		68	76.07	28	23.93	0 0	0	0	
Т3	25	78.13	7	21.88	0	0	0	0	T3		18	81.82	4	18.18	0 0	0	0	
T4	2	100	0	0	0	0	0	0	T4		4	80.0	1	20.0	0 0	0	0	
Neo-adjuvant therapy									Neo	Neo-adjuvant therapy								
Yes	2	100	0	0	0	0	0	SN 0	Yes		9	85.71	0	0	0 0		14.29	NS
No	165	81.68	33	16.34	4	1.98	0	0	%		169	78.97	45	21.03	0 0	0	0	
Systemic therapy plan									Syst	Systemic therapy plan								
Endocrine Therapy Alone	12	92	1	8	0	0	0	O NS	End	Endocrine Therapy Alone	14	77.78	3	16.67	0 0	-	5.56	SN
Chemotherapy Alone	37	74	10	20	3	9	0	0	Che	Chemotherapy Alone	30	73.17	11	26.83	0 0	0	0	
Endocrine and Chemotherapy	118	83.69	22	15.60	1	0.71	0	0	End	Endocrine and Chemotherapy	131	78.26	31	18.56	0 0	0	0	
Anthracyclines									Ant	Anthracyclines								
Yes	153	82.26	31	16.67	2	1.08	0	O NS	Yes		157	75.85	41	18.61	0 0	6	4.35	NS
No	14	78	2	11.11	2	11.11	0	0	Š		18	78.26	4	17.39	0 0	_	4.35	
Taxanes									Tax	Taxanes								
Yes	25	80.65	5	16.13	1	3.23	0	SN 0	Yes		22	73.33	8	26.67	0 0	0	0	SN
No	142	82.08	28	16.18	3	1.73	0	0	No		153	80.10	37	19.37	0 0	1	0.53	
Use of statins									Use	Use of statins								
Yes	6	75	2	16.67	1	8.33	0	SN 0	Yes		6	75.00	2	16.67	0 0	1	8.33	SN
No	158	82.29	31	16.15	3	1.56	0	0	Š		166	79.43	43	20.57	0 0	0	0	
Menopausal status									Mer	Menopausal status								
Pre-menopausal	9/	82.61	15	16.30	1	1.09	0	O NS	Pre-	Pre-menopausal	75	79.79	19	20.12	0 0	0	0	NS
≤5 years since menopause	28	73.68	7	18.42	3	7.90	0	0	≤5)	≤5 years since menopause	27	77.14	7	20.0	0 0	-	2.86	
> 5 years since menopause	48	82.76	10	17.24	0	0	0	0	>5)	>5 years since menopause	55	70.71	14	20.29	0 0	0	0	
Menstrual status unknown	15	93.75	1	6.25	0	0	0	0	Мет	Menstrual status unknown	18	78.26	5	21.74	0 0	0	0	
ER status									ER	ER status								
ER Positive	130	83.87	23	14.84	2	1.29	0	O NS	ER]	ER Positive	143	80.79	33	18.64	0 0	-	0.57	SN
ER Negative	36	75.00	10	20.83	2	4.17	0	0	ER 1	ER Negative	32	72.73	12	27.27	0 0	0	0	
Status Unknown	0		0		0		0	0	Stat	Status Unknown	0		0		0 0	0		
PR status									PR	PR status								
PR Positive	58	82.86	11	15.71	1	1.43	0	0 NS	PR 1	PR Positive	99	86.84	10	13.16	0 0	0	0	NS
PR Negative	36	81.82	8	18.18	0	0	0	0	PR 1	PR Negative	25	62.50	14	35.00	0 0	1	2.55	
Status Unknown	71	89.08	14	15.91	3	3.41	0	0	Stat	Status Unknown	83	79.81	21	20.19	0 0	0	0	
Missing	2	100	0	0	0	0	0	0	Mis.	Missing	0		0	0	0 0	0	0	
Lymph node involvement									Lyn	Lymph node involvement								
1–3 Nodes involved	113	85.61	16	12.12	3	2.27	0	O NS	1–3	1-3 Nodes involved	117	81.25	27	18.75	0 0	0	0	SN
Continued																		

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	Control	10									Zoledronate	onate							
	Score 0		Score 1		Score 2		Score 3				Score 0		Score 1		Score 2		Score 3		
Media (range)	50.5 (33-79) n=172	(3-79)	52 (32–73) n=33	-73)	69		61			Media (Range)	51 (29-77) n=175	-77)	50.5 (33-79) $n = 45$		61.5 $(58-65)$ $n=2$				
	Total	%	Total	%	Total	%	Total	%	p-value		Total	%	Total	%	Total	L %	Total %		p-value
≥4 Nodes involved	53	75.71	17	24.29	0	0	0	0		≥4 Nodes involved	57	75.00	18	23.68	0	0		1.32	
Unknown Involvement	0	0	0	0	1	100	0	0		Unknown Involvement	0		0		0	0 0			
Sentinel Node Positive	0		0		0		0	0		Sentinel Node Positive	0		0		0	0 0	_		
HER2 status										HER2 status									
HER2 Positive	24	82.76	5	17.24	0	0	0	0	NS	HER2 Positive	14	29.99	9	28.57	0	0 1		4.76	NS
HER2 Negative	41	78.85	6	17.31	2	3.86	0	0		HER2 Negative	99	84.85	10	15.15	0	0 0	0		
Unknown	4	57.14	3	42.86	0	0	0	0		Unknown	3	75.00	1	25.00	0	0 0	0		
Not Measured	26	85.09	15	13.16	2	1.75	0	0		Not Measured	100	78.13	28	21.88	0	0 0	0		
Missing Data	0	0	1	100	0	0	0	0		Missing Data	2	100.00	0	0	0	0 0	0		
Histological grade										Histological grade									
Grade 1	11	84.62	2	15.38	0	0	0	0	NS	Grade 1	13	100.00	0	0	0	0 0	0	Z	NS
Grade 2	70	85.37	6	10.98	3	3.66	0	0		Grade 2	70	76.09	21	22.82	0	0		1.09	
Grade 3	98	78.90	22	20.18	1	0.92	0	0		Grade 3	06	78.95	24	21.05	0	0 0	0		
Grade 4	0		0		0		0	0		Grade 4	1	100.00	0	0	0	0 0	0		
Missing Data	0	0	1	100	0	0	0	0		Missing Data	2	100.00	0	0	0	0 0	0		

Table 2. Complete multi-variate analysis of CD73 association with patient.

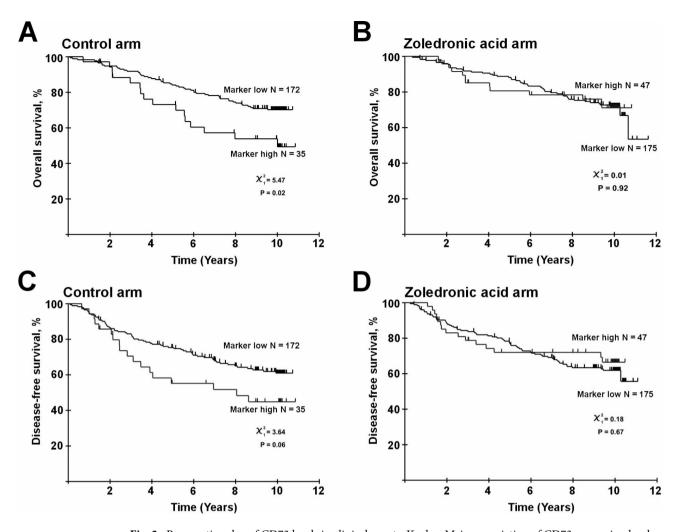


Fig. 2. Prognostic value of CD73 levels in clinical events. Kaplan-Meier association of CD73 expression level in Overall Survival (OS) in **(A)** control and **(B)** zoledronic acid arms. Relationship between level of expression of CD73 in Disease-Free-Survival (DFS) in **(C)** control and **(D)** zoledronic acid arms. P-value is obtained from the log-rank test for testing quality of survival Q15 functions.

		Standard	treatment				Standard	treatment	+ zoledı	ronate	
End point		# at risk	# Events	HR	HR: 95% CI	P-value	# at risk	# Events	HR	HR: 95% CI	P-value
OS	Unadjusted	207	66	1.87	1.06-3.29	0.03	222	61	1.06	0.572 - 1.96	0.86
	Adjusted	207	66	1.53	0.85-2.77	0.15	222	61	0.833	0.444 - 1.56	0.57
DFS	Unadjusted	207	81	1.66	0.982 - 2.80	0.06	222	79	0.885	0.504 – 1.55	0.67
	Adjusted	207	81	1.35	0.78-2.34	0.28	222	79	1.35	0.76-2.41	0.30
Skeletal-first distant recurrence#	Unadjusted	207	33	2.23	1.04-4.81	0.04	222	20	0.945	0.316 - 2.83	0.92
	Adjusted	207	33	2.44	1.08-5.49	0.03	222	20	1.05	0.34-3.29	0.93
Bone metastasis at any time	Unadjusted	207	51	1.86	0.97-3.55	0.062	222	31	0.717	0.275 - 1.87	0.50
	Adjusted	207	51	1.91	0.97-3.75	0.06	222	31	0.694	0.260 - 1.85	0.46
Distant recurrence including skeletal	Unadjusted	207	60	1.77	0.97-3.21	0.06	222	52	0.872	0.438 - 1.74	0.70
	Adjusted	207	60	1.56	0.84-2.90	0.16	222	52	0.694	0.344 - 1.41	0.31
Non-skeletal first distant recurrence##	Unadjusted	207	36	1.74	0.791 - 3.82	0.17	222	38	0.850	0.374 - 1.93	0.70
	Adjusted	207	36	1.25	0.55-2.84	0.60	222	38	0.606	0.264 - 1.39	0.24

Table 3. Associations between CD73 expression and clinical and pathological variables in each arm of the AZURE trial. # (Can include concomitant non-skeletal first distant recurrence). ## (Can include concomitant skeletal first distant recurrence).

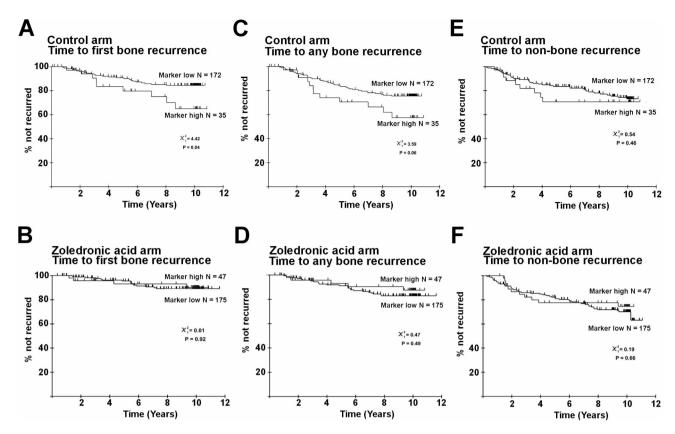


Fig. 3. Kaplan-Meier association of CD73 levels with clinical metastatic events. Relationship between CD73 expression to time to first bone recurrence, – where cancer spread to bone is the first recorded event in **(A)** control and **(B)** zoledronic acid arm. Relationship between level of CD73 expression to spread to bone at any time (for which patients may have had spread to non-bone sites first) in **(C)** control and **(D)** zoledronic acid arm. Relationship between CD73 expression to time to spread to non-bone sites in **(E)** control and **(F)** zoledronic acid arm. P-value is obtained from the log-rank test for testing quality of survival Q15 function.

possible target for cancer therapy. High tumor CD73 expression has also been associated with poor response to anthracyclines³⁴ and NACT²⁷. The poor response may be also associated with tumor immunity, as various chemotherapeutic agents were shown to induce upregulation of PD-L1 and other checkpoint inhibitors³⁵. In our recent study, silencing of CD73 expression (using shRNA) within breast cancer cells was demonstrated to increase the sensitivity of these cells to the growth inhibitory actions of nitrogen-containing bisphosphonates in vivo⁵.

CD73 has been so have several roles within the bone microenvironment. High CD73 expression level was demonstrated to promote osteoblastic differentiation in mice³⁶. Osteoblasts play a key role in the maintenance of breast cancer cell dormancy within the bone marrow, enabling cancer cells to evade cytotoxic drugs that target rapidly proliferation cells. Later, these dormant cells may emerge as macro-metastases³⁷. The importance of CD73 in breast cancer spread to bone was also demonstrated in a study targeting both CD73 and transforming-growth-factor- β (TGF- β) in mouse models of TNBC³⁸. Xing and coworkers demonstrated that high expression of CD73 was associated with epithelial-to-mesenchymal transition, as well as the development of a fibrotic stroma and an immune-tolerant tumor micro-environment. Dual blockade of CD73 and TGF β promoted a multifaceted inflammatory tumor microenvironment with diminished levels of myeloid-derived suppressor cells (MDSCs) and M2-macrophages, as well as increased levels of activated dendritic cells, cytotoxic T cells, and B cells. Their study suggested that dual targeting of CD73 and TGF- β may serve as a potential therapeutic option within TNBC³⁸.

Our study has several limitations. The low frequency of high CD73 expression in the cohort may limit the generalizability of our findings. Additionally, the incomplete data on HER2 status, due to its non-mandatory measurement in the AZURE trial, is another limitation that may impact the comprehensiveness of our analysis. We demonstrated that high CD73 expression did not show a significant association with patient outcomes in the zoledronic acid arm. However, we did not investigate whether CD73 expression is affected by zoledronate treatment in patient samples but it was not regulated in mouse TNBC cells upon zoledronate in vitro, according to our previous study⁵. In tumor microenvironment, CD73 expression is regulated by various factors, such as hypoxia¹⁹, signaling pathways PI3K, MAPK³⁹, transcription factors⁴⁰ or miRNAs⁴¹. However, studies, whether CD73 protein expression is affected by anti-cancer treatment when it is not specific CD73 inhibitor^{42,43} are limited.

CD73 showed association with metastases in different cancers. High tumor CD73 expression significantly correlated with worse outcome of patients with colorectal cancer liver metastases or metastatic melanoma 44. A comprehensive meta-analysis of 2951 cancer cases revealed that high CD73 expression is significantly associated with shorter overall survival in various cancers, including breast cancer and is strongly associated with lymph node metastases 45. We show here that high CD73 expression in the primary tumors is significantly associated with the development of bone metastases in the control arm of the study. Furthermore, Kaplan-Meier analysis revealed a significant prognostic value of high CD73 expression for time to first metastasis to bone in the control arm, which was not evident in the zoledronic acid arm. These findings suggest a potential role for CD73 as a prognostic marker on bone metastasis. However, additional investigation is necessary to uncover the underlying mechanisms and assess the clinical implications of targeting CD73 in breast cancer.

Data availability

Detailed data analysis and statistical analysis are provided within the manuscript. The raw data includes sensitive patient information that cannot be fully anonymized, and we are bound by confidentiality agreements and institutional regulations that restrict data sharing.

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Author contributions

Design – K.S. Materials – J.B. Analysis – N.P, A.L. and M.O. Statistical analysis – W.G, S.W. Data curation – S.W, J.B. Methodology – J.S and A.J. Writing the original draft – N.P, S.W, J.B and K.S. Writing and editing the manuscript – N.P, S.W, J.S, A.J, J.B and K.S. All authors reviewed and approved the final manuscript.

Declarations

Competing interests

The authors declare no competing interests.

Additional information

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