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Autoantibody against UBE2E3 is a common biomarker for atherosclerotic diseases and digestive tract cancer

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Abstract

Aim: Atherosclerosis and diabetes mellitus (DM) often lead to severe conditions, such as acute ischemic stroke (AIS), cardiovascular disease (CVD), and chronic kidney disease (CKD). Some cancers are also associated with atherosclerosis. Therefore, identifying novel autoantibody biomarkers associated with atherosclerosis-related conditions is crucial for improving early diagnosis and risk assessment.

Methods: We used an array of 9,480 proteins to detect IgG antibodies in the serum of patients with atherosclerosis. Following this screening, we quantified the antibody levels using an amplified luminescent proximity homogeneous assay-linked immunosorbent assay (AlphaLISA) with recombinant antigen proteins.

Results: Ubiquitin conjugating enzyme E2 E3 (UBE2E3) was identified as a candidate antigen recognized by IgG antibodies in the sera of individuals diagnosed with atherosclerosis. Compared with healthy donors,



significantly higher serum antibody levels against UBE2E3 were found in patients with AIS, DM, CVD, CKD, esophageal cancer (EC), and gastric cancer (GC), but not in those with colorectal cancer (CRC). Receiver operating characteristic (ROC) analysis revealed that the higher areas under the ROC curves for anti-UBE2E3 antibodies were observed in DM- or nephrosclerosis-associated CKD than in the others. Spearman's correlation analysis revealed that serum anti-UBE2E3 antibody (s-UBE2E3-Ab) levels were associated with the plaque score, maximum intima-media thickness, and cardio-ankle vascular index, which are typical indices of atherosclerosis and stenosis. In the survival analysis of GC and CRC, patients who were s-UBE2E3-Ab-positive had significantly poorer prognoses than patients who were s-UBE2E3-Ab-negative. The difference became more prominent when s-UBE2E3-Abs were combined with anti-differential screening-selected gene aberrant in neuroblastoma antibody (DAN-Ab) or sclerostin domain-containing protein 1 (SOSTDC1), which are bone morphogenetic protein (BMP) antagonists.

Conclusions: The s-UBE2E3-Ab marker is highly associated with atherosclerosis-related diseases, such as AIS, CVD, DM, CKD, and digestive tract cancers, suggesting the involvement of BMP signals.

Keywords

Autoantibody biomarker, atherosclerosis, chronic kidney disease, diabetes mellitus, acute ischemic stroke, cardiovascular disease, gastrointestinal cancer

Introduction

In contrast to enzymes, antigens, and nucleic acid biomarkers, few antibody biomarkers have been used clinically [1]. The presence of autoantibodies has been noted in individuals with autoimmune diseases and cancer; however, little is known about autoantibodies in other diseases. We have performed a large-scale screening using serological identification of antigens by cDNA expression cloning (SEREX) to select antigenic proteins recognized by serum IgG antibodies in patients with esophageal cancer (EC) from a cDNA library [2, 3]. Some antibodies against the selected antigens showed significantly higher levels in patients than in healthy donors (HDs), suggesting that these antibodies could be tumor markers.

The development of autoantibodies is attributable to the overexpression of antigenic proteins in tumor tissues, followed by tissue damage and leakage of intracellular proteins outside the cells [2–5]. If atherosclerosis is similar and is accompanied by damage to arterial blood vessels, specific antibody markers can be identified. Therefore, we began searching for autoantibody markers of atherosclerosis and related diseases using SEREX and a newly developed protein array method.

Atherosclerosis is caused by various risk factors, such as diabetes mellitus (DM), chronic kidney disease (CKD), hypertension, and dyslipidemia, and is complicated by the onset of acute ischemic stroke (AIS) and cardiovascular disease (CVD) [6]. Autoantibody markers of atherosclerosis, such as anti-ATP2B4 [7], anti-SH3BP5 [8], anti-AP3D1 [9], and anti-KIAA0513 antibodies [10], are commonly associated with AIS, CVD, DM, and CKD. Some of these markers have elevated levels in patients with gastrointestinal cancer. Furthermore, angiogenesis is essential for the development of solid tumors [11], and both DM and atherosclerosis are risk factors for EC and colorectal cancer (CRC) [12].

Autoantibody markers have also been associated with AIS and acute myocardial infarction (AMI). Elevated autoantibodies detected within two weeks of disease onset are thought to have been present before onset, because increases in new antibody levels are not detectable within two weeks. We identified anti-DID01, anti-FOXJ2, anti-CPSF2 [13], and anti-AP3D1 [9] antibodies in patients with AIS within two weeks, and a case-control study nested within the JPHC-based prospective study revealed that patients who were antibody-positive had significantly greater AIS onset frequencies than patients who were antibody-negative [9, 13]. This suggests that autoantibodies may be useful markers for predicting AIS onset.

Each antibody biomarker exhibits different properties. For example, anti-DIDO1-antibodies are closely associated with CKD, whereas anti-FOXJ2 and anti-AP3D1 antibodies reflect arterial stenosis [9, 13]. Anti-CPSF2 antibodies are associated with DM and hypertension [13]. Thus, identifying as many markers as possible allows for a more precise diagnosis.

This study investigates serum antibodies against ubiquitin conjugating enzyme E2 E3 (UBE2E3) in patients with atherosclerosis-related diseases, including AIS, DM, CVD, CKD, and digestive tract cancers.

Materials and methods

Patient and control sera

The local ethical review boards of Chiba University, Graduate School of Medicine, Toho University, Faculty of Medicine, Toho University Omori Medical Center, and Port Square Kashiwado Clinic, and the review boards of the participating hospitals approved this study. All experimental procedures adhered to the guidelines of the Declaration of Helsinki (2013). Serum samples were collected from the donors after obtaining written informed consent.

All serum samples were centrifuged at 3,000 g for 10 min. The supernatants were stored at -80 °C to avoid repeated freezing/thawing.

Serum samples from 127 patients with AIS were obtained from Chiba Prefectural Sawara Hospital within two weeks of disease onset (AIS cohort). The Sawara cohort comprised 665 specimens obtained from Chiba Prefectural Sawara Hospital, including 139 from HDs, 228 from patients with AIS, 44 with transient ischemic attack, 122 with deep and subcortical white matter hyperintensity, 17 with asymptomatic cerebral infarction, 59 with chronic-phase cerebral infarction, and 56 disease controls. Sera of 275 and 100 patients with DM and CVD, respectively, were obtained from Chiba University Hospital (DM and CVD cohorts, respectively); the CVD cases included those with AMI and unstable angina pectoris. Serum samples from 300 patients with CKD were obtained from Kumamoto University (CKD cohort) [14, 15]. Serum samples from 285 patients with EC, gastric cancer (GC), or CRC were collected immediately before surgery, radiotherapy, or chemotherapy at the Department of Surgery, Toho University Hospital (cancer cohort). Serum samples from HDs (HD cohort) were collected from three institutions: Chiba University Hospital, Port Square Kashiwado Clinic, and Chiba Prefectural Sawara Hospital. All HD serum samples were obtained from individuals without any abnormalities on cranial magnetic resonance imaging.

ProtoArray screening

The first screening was performed using the ProtoArray Human Protein Microarrays (version 4.0; Thermo Fisher Scientific, Waltham, MA, USA), which contained 9,480 proteins, as described previously [16–18]. Thirty serum samples (10 from HDs and 10 from patients with atherosclerosis) were analyzed to identify antigens specifically recognized by serum IgG antibodies. The data were processed using Prospector software, which employs M-statistics (Thermo Fisher Scientific). To compare the two groups, a positivity threshold for each protein was determined using M-statistics (Supplementary material), which included background subtraction, normalization of signals, and analysis of differences between patients and HDs. The proportion of subjects in each group showing an immune response above this threshold was scored, and the significance of the difference between the two groups was assessed by calculating the *P*-value, as described previously [19].

Preparation of antigens

Full-length cDNA of human UBE2E3 (accession number: NM_006357) was purchased from Open Biosystems (Huntsville, AL, USA) and recombined into pGEX-6P (Cytiva, Pittsburgh, PA, USA). The cDNA product was expressed by treating *Escherichia coli* (*E. coli*) KRX (Promega, Madison, WI, USA) cells containing the pGEX-6P-UBE2E3 and pMINOR with 0.5 mM isopropyl- β -D-thiogalactoside at 37°C for 3 h [20]. The cells were lysed by sonication in phosphate-buffered saline containing 2 mM dithiothreitol. The proteins were purified using Glutathione-Sepharose 4 Fast Flow medium (Cytiva) and a HiPrep 26/10

desalting column (Cytiva) and concentrated to 1.3 g/mL in phosphate-buffered saline containing 2 mM dithiothreitol. For comparison, antigenic differential screening-selected gene aberrant in neuroblastoma (DAN) protein and sclerostin domain-containing protein 1 (SOSTDC1) peptide were prepared. The region between positions 267 and 731 of the DAN cDNA (accession number: X66872.1) was inserted into the *EcoRI/XhoI* site of pGEX-2T (GE Healthcare Life Sciences, Pittsburgh, PA, USA), which produced a truncated DAN protein (amino acid residues 25–178) lacking its potential signal peptide sequence. The glutathione S-transferase (GST)-fused DAN protein was purified, as described previously [21]. The biotinylated peptide between amino acid residues 156 and 170 of SOSTDC1 (accession number: NM_015464) was synthesized and purified using high-performance liquid chromatography (HPLC), as described previously [16]. The SOSTDC1 peptide structure was biotin-KITVVTACKCKRYTR-COOH, with a purity of > 90%.

Amplified luminescence proximity homogeneous assay-linked immunosorbent assay

Serum antibody levels were examined using amplified luminescence proximity homogeneous assay-linked immunosorbent assay (AlphaLISA) in 384-well microtiter plates (white opaque OptiPlate™, Perkin Elmer, Waltham, MA, USA). AlphaLISA required two types of beads: a donor bead that binds to the antigen via a tag and an acceptor bead that binds the IgG antibody via a secondary antibody. IgG antibody binding to the antigen in solution brings the two beads close together, and excitation at 680 nm results in emission at 618 nm. No plate washing was required. Each well contained 2.5 μ L of 1:100-diluted serum with 2.5 μ L of GST, GST-UBE2E3, or GST-DAN proteins (10 μ g/mL) or biotinylated SOSTDC1 peptide (400 ng/mL) in AlphaLISA buffer. The buffer was composed of 25 mM N-2-hydroxyethylpiperazine-N-2-ethane sulfonic acid (pH 7.4), 0.1% casein, 0.5% Triton X-100, 1 mg/mL dextran-500, and 0.05% ProClin-300, following the manufacturer's instructions (https://www.revvity.co.jp/content/elisa-alphalisa-immunoassay-conversionmade-easy). The reaction mixture was incubated at room temperature for 6-8 h. Anti-human IgGconjugated acceptor beads (2.5 µL at 40 µg/mL) and either glutathione- or streptavidin-conjugated donor beads (2.5 μ L at 40 μ g/mL) were added, and the mixture was incubated at room temperature in the dark for 7-14 days. Chemical emissions at 607-623 nm (alpha photon counts), which indicated the antigenantibody binding level, were measured using an EnSpire Alpha microplate reader (PerkinElmer), as described previously [7–9, 13, 16–18]. Specific reactions were determined by subtracting the alpha emission counts of the GST and buffer control from those of the GST-fused proteins and biotinylated peptides, respectively.

Statistical analysis

We employed the Mann–Whitney *U* test to examine differences between two groups and the Kruskal–Wallis test (Mann–Whitney *U* test with the Bonferroni correction) to evaluate differences among three or more groups. Spearman's correlation analysis, logistic regression analysis, the chi-square test, and univariate and multivariate analyses were used to calculate the correlations. We assessed the predictive values of the putative disease markers using receiver operating characteristic (ROC) curve analysis. The sensitivity and specificity were calculated using the cutoff values of the Youden Index. All statistical analyses were performed using GraphPad Prism (version 5; GraphPad Software, Inc.). Patient survival was evaluated using the Kaplan–Meier method, and the results were compared using the log-rank test. The cutoff values were determined using X-tile software (version 3.6.1; Yale University, New Haven, CT) [22]. All tests were two-tailed, and *P*-values < 0.05 indicated statistically significant differences.

Results

Recognition of UBE2E3 by serum IgG antibodies in patients with atherosclerosis

We used ProtoArray to identify antigens using IgG antibodies from the sera of patients with atherosclerosis. UBE2E3 (accession number: NM_006357) was a candidate antigen because of its high positive reactivity rate (6 of 10 serum samples) in patients with atherosclerosis and low positive reactivity rate (2 of 10 samples) in HDs. All ProtoArray results are available in the public Figshare database (https://figshare.com/articles/dataset/Results_of_protein_array_for_atherosclerosis/25906330).

Elevated anti-UBE1E3 antibody levels in patients with AIS

We examined serum antibody levels in patients with AIS using recombinant UBE2E3 protein. Sera from HDs and patients with AIS (AIS cohort) were obtained from the Chiba Prefectural Sawara Hospital. The sample numbers (total, male, female) and the average age ± standard deviation (SD) are shown in Table 1. The AlphaLISA results revealed that anti-UBE1E3 antibody (UBE2E3-Ab) levels were significantly higher in patients with AIS than in HDs (Figure 1A). Cutoff values were determined using the average plus two SDs of the HD values. The positivity rates of UBE2E3-Abs for the HDs and patients with AIS were 3.9% and 12.6%, respectively (Table 1). ROC analysis revealed that the area under the ROC curve (AUC) for UBE2E3-Abs was 0.6806 [95% confidence interval (CI): 0.6156–0.7456] (Figure 2A). Using the cutoff value based on the Youden index, the sensitivity and specificity for UBE2E3-Abs were 64.6% and 64.8%, respectively.

Sample inform	ample information and alpha analysis			
(A) Sample in	formation	HD	AIS	
Total sample number		128	127	
Male/Female		57/71	71/56	
Age (average ± SD)		47.0 ± 14.5	76.8 ± 11.0	
(B) Alpha analysis (antibody level)		UBE2E3-Ab		
HD	Average	8,025		
	SD	5,811		
	Cutoff value	19,647		
	Positive No.	5		
	Positive rate (%)	3.9%		
AIS	Average	11,960		
	SD	10,035		
	Positive No.	16		
	Positive rate (%)	12.6%		
	<i>P</i> (vs. HD)	1.7E-04		

Table 1. Serum UBE2E3-Ab levels in healthy dono	s (HDs) and patients with acute ischemic stroke (AIS)
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The upper panel (**A**) indicates the information of serum samples, including the number of total samples, male or female, as well as ages [average \pm standard deviation (SD)]. The lower panel (**B**) summarizes the serum antibody levels against the UBE2E3-GST protein examined by the amplified luminescence proximity homogeneous assay-linked immunosorbent assay (AlphaLISA) method. Cutoff values were set at the average HD values plus two SD, and positive samples higher than the cutoff value were scored. *P* values were calculated using the Mann–Whitney *U* test. *P* value < 0.05 and positive rate > 10% are marked in bold text. A scatter dot plot of the same results is shown in Figure 1A

Elevated UBE2E3-Ab levels in patients with DM

Next, we examined the UBE2E3-Ab levels in patients with DM. Serum samples from HDs and patients with DM (DM cohort) were obtained from Chiba University Hospital. The UBE2E3-Ab levels were significantly higher in patients with DM than in HDs (Figure 1B). At the cutoff value (average plus two SDs of the HD values), the positivity rates of UBE2E3-Abs in HDs and patients with DM were 4.9% and 22.2%, respectively (Table 2). ROC analysis was performed to evaluate the diagnostic ability for patients with DM. The AUC for UBE2E3-Abs was 0.6716, yielding a sensitivity and specificity of 50.2% and 79.0%, respectively (Figure 2B).

Table 2. Analysis of the serum anti-UBE2E3 antibody (UBE2E3-Ab) levels among HDs and patients with diabetes mellitus (DM)

Sample information and alpha analysis					
(A) Sample information		HD	DM		
Total sample nu	umber	81	275		
Male/Female		46/35	156/119		
Age (Average ± SD)		45.2 ± 11.0	63.1 ± 12.0		
(B) Alpha anal	ysis (antibody level)	UBE2E3-Ab			
HD Average		4,003			
	SD	2,709			

Table 2. Analysis of the serum anti-UBE2E3 antibody (UBE2E3-Ab) levels among HDs and patients with diabetes mellitus (DM) (continued)

Sample information and alpha analysis			
	Cutoff value	9,421	
	Positive No.	4	
	Positive rate (%)	4.9%	
DM	Average	7,034	
	SD	6,207	
	Positive No.	61	
	Positive rate (%)	22.2%	
	<i>P</i> (vs. HD)	9.8E-10	

The upper panel (**A**) represents the total numbers, sex (male and female), and ages (average \pm SD). The lower panel (**B**) provides a summary of the serum antibody levels, measured as Alpha photon counts using amplified luminescence proximity homogeneous assay-linked immunosorbent assay (AlphaLISA) with UBE2E3-GST protein. Cutoff values were defined as the average healthy donor (HD) values plus two SD, and positive samples higher than the cutoff value were scored. *P* value < 0.05 and positive rate > 10% are marked in bold text. A scatter dot plot of the same results is shown in Figure 1B

Elevated UBE2E3-Ab levels in patients with CVD

We then examined UBE2E3-Ab levels in patients with CVD. Sera from HDs and patients with CVD (CVD cohort) were obtained from Chiba University Hospital. UBE2E3-Ab levels were significantly higher in patients with CVD than in HDs (Figure 1C). Using the cutoff values (average plus two SDs of the HD values), the UBE2E3-Ab positivity rates for HDs and patients with CVD were 5.1% and 8.0%, respectively (Table 3). The AUC value of UBE2E3-Abs was 0.6821 (95% CI: 0.6069–0.7573), with a sensitivity and specificity of 88.4% and 41.7%, respectively (Figure 2C).

Sample information	Sample information and alpha analysis				
(A) Sample info	ormation	HD	CVD		
Total sample nu	mber	78	100		
Male/Female		46/32	84/16		
Age (Average ± SD)		45.3 ± 11.2	66.1 ± 11.3		
(B) Alpha analysis (antibody level)		UBE2E3-Ab			
HD	Average	4,650			
	SD	3,756			
	Cutoff value	12,162			
	Positive No.	4			
	Positive rate (%)	5.1%			
CVD	Average	6,375			
	SD	4,211			
	Positive No.	8			
	Positive rate (%)	8.0%			
	<i>P</i> (vs. HD)	4.4E-03			

Table 3. Contrast of the serum UBE2E3-Ab levels in HDs and patients with cardiovascular disease (CVD)

The upper panel (**A**) displays the total number of samples, along with a breakdown by sex and the age distribution (average \pm SD). The lower panel (**B**) shows the serum antibody levels examined by amplified luminescence proximity homogeneous assaylinked immunosorbent assay (AlphaLISA) using the antigen, purified UBE2E3-GST, as described in the legend of Table 1. *P* value < 0.05 is marked in bold text. A scatter dot plot of the same results is shown in Figure 1C

Elevated UBE2E3-Ab levels in patients with CKD

Next, we analyzed the antibody levels in the sera of patients with CKD (CKD cohort), a condition strongly associated with atherosclerosis. CKD was divided into three subtypes: type 1, diabetic kidney disease; type 2, nephrosclerosis; and type 3, glomerulonephritis. Samples from patients with CKD were obtained from the Kumamoto cohort, and samples from HDs were obtained from Chiba University Hospital. Patients in all three CKD groups had significantly higher serum UBE2E3-Ab levels than the HDs (Figure 1D). The UBE2E3-Ab positivity rates in HDs and patients with types 1, 2, and 3 CKD were 2.4%, 17.2%, 15.6%, and 8.1%,



Figure 1. Comparison of serum anti-UBE2E3 antibody (UBE2E3-Ab) levels between HDs and patients. The UBE2E3-Ab levels of healthy donors (HDs) and patients with acute ischemic stroke (AIS) (**A**), diabetes mellitus (DM) (**B**), cardiovascular disease (CVD) (**C**), chronic kidney disease (CKD) (**D**), and esophageal cancer (EC), gastric cancer (GC), and colorectal cancer (CRC) (**E**) were examined by amplified luminescence proximity homogeneous assay-linked immunosorbent assay (AlphaLISA) using GST-UBE2E3 protein as the antigen, and shown in scatter dot plots. The ordinate represents Alpha emission photon counts, which correspond to the antibody levels. Serum anti-differential screening-selected gene aberrant in neuroblastoma antibody (DAN-Ab) (**F**) and anti-SOSTDC1 antibodies (SOSTDC1-Ab) levels (**G**) between HDs and patients with cancer were also examined. Type-1, type-2, and type-3 CKDs represent diabetic kidney disease, nephrosclerosis, and glomerulonephritis, respectively. The bars represent the average and average \pm SD. *P* values were calculated using the Kruskal–Wallis test. ** *P* < 0.01; *** *P* < 0.001; **** *P* < 0.0001 vs. HD specimens. ns: not significant. The total (male/female) numbers, average ages \pm standard deviations (SDs), average antibody levels \pm SDs, cutoff values, positive numbers, positive rates (%), and *P* values versus HDs are summarized in Tables 1–5

respectively (Table 4). ROC analysis suggested that the AUCs of types 1, 2, and 3 CKD were 0.8531 (95% CI: 0.8005–0.9057) (Figure 2D), 0.8798 (95% CI: 0.8189–0.9408) (Figure 2E), and 0.7941 (95% CI: 0.7300–0.8583) (Figure 2F), respectively. Overall, CKD showed much higher AUC values than AIS and DM, irrespective of the CKD type (Figures 2A-F).

Sample informat	ion and alpha analysis				
(A) Sample inform	mation	HD	Type-1 CKD	Type-2 CKD	Type-3 CKD
Total sample number		82	145	32	123
Male/Female		44/38	106/39	21/11	70/53
Age (average ± S	D)	44.1 ± 11.2	66.0 ± 10.4	76.0 ± 9.8	62.0 ± 11.7
(B) Alpha analys	is (antibody level)	UBE2E3-Ab			
HD	Average	1,625			
	SD	1,403			
	Cutoff value	4,432			
	Positive No.	2			
	Positive rate (%)	2.4%			
Type 1-CKD	Average	3,404			
	SD	2,348			
	Positive No.	25			
	Positive rate (%)	17.2%			
	<i>P</i> (vs. HD)	1.3E-11			
Type 2-CKD	Average	3,177			
	SD	1,409			
	Positive No.	5			
	Positive rate (%)	15.6%			
	<i>P</i> (vs. HD)	2.1E-06			
Type 3-CKD	Average	2,602			
	SD	1,381			
	Positive No.	10			
	Positive rate (%)	8.1%			
	<i>P</i> (vs. HD)	2.1E-06			

Table 4. Analysis of serum UBE2E3-Ab levels of HDs versus those of patients with chronic kidney disease (CKD)

Type-1, -2, and -3 CKDs correspond to diabetic kidney disease, nephrosclerosis, and glomerulonephritis, respectively. The upper panel (**A**) indicates the numbers of all samples and samples from males and females, as well as age (average \pm SD). The lower panel (**B**) summarizes the serum antibody levels examined by AlphaLISA using purified UBE2E3-GST protein as an antigen, as described in the legend of Table 1. *P* values were calculated using the Kruskal–Wallis test (Mann–Whitney *U* test with Bonferroni correction applied). *P* values < 0.05 and positive rates > 10% are marked in bold text. A scatter dot plot of the same results is shown in Figure 2D

UBE2E3-Ab levels in solid cancer

An increasing number of reports have indicated that atherosclerosis is closely associated with cancer [7-9], which is supported by shared biomarkers for both atherosclerosis and cancer. To investigate this further, we analyzed serum samples from patients with EC, GC, or CRC (cancer cohort) obtained from Toho University Hospital. UBE2E3-Ab levels were significantly higher in patients with EC and GC, but not with



Figure 2. Receiver operating characteristic curve (ROC) analysis. The abilities of UBE2E3-Abs to detect acute ischemic stroke (AIS) (**A**), diabetes mellitus (DM) (**B**), cardiovascular disease (CVD) (**C**), type-1 chronic kidney disease (CKD) (**D**), type-2 CKD (**E**), type-3 CKD (**F**), esophageal cancer (EC) (**G**), and gastric cancer (GC) (**H**) and colorectal cancer (CRC) (**I**) were evaluated using ROC analysis. Anti-differential screening-selected gene aberrant in neuroblastoma antibodies (DAN-Abs) to detect EC (**J**) and GC (**K**), and CRC (**L**) and SOSTDC1-Abs to detect EC (**M**), GC (**N**), and CRC (**O**) were also examined. The numbers in the figures represent area under the ROC curve (AUC), cutoff values (Youden index) for antibody levels, sensitivity, specificity, 95% confidence interval (CI), and *P* values. *P* values < 0.05 are marked in bold text

CRC, than in HDs (Figure 1E, Table 5). The AUCs for EC, GC, and CRC were 0.6669, 0.6825, and 0.6071, respectively (Figures 2G–I).

Table 5. Comparison of UBE2E3-Ab levels in HDs an	d patients with cancer
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Sample in	formation and alpha analysis				
(A) Sample information		HD	GC	CRC	EC
Total samp	le number	95	94	96	95
Male/Fema	le	51/44	73/21	66/30	57/38
Age (avera	ge ± SD)	57.9 ± 6.0	66.6 ± 9.1	66.6 ± 9.7	67.2 ± 11.6
(B) Alpha	analysis (antibody level)	UBE2E3-Ab	DAN-Ab	SOSTDC1-Ab	
HD	Average	9,193	218,241	18,075	
	SD	6,267	75,817	13,567	
	Cutoff value	21,728	369,876	45,208	
	Positive No.	5	4	4	
	Positive rate (%)	5.3%	4.2%	4.2%	
EC	Average	12,574	291,060	26,084	
	SD	6,739	118,712	18,331	
	Positive No.	10	24	13	
	Positive rate (%)	10.6%	25.5%	13.8%	
	<i>P</i> (vs. HD)	< 0.001	< 0.001	< 0.05	
GC	Average	13,081	264,614	14,297	
	SD	7,715	97,415	12,101	
	Positive No.	11	14	2	
	Positive rate (%)	11.5%	14.6%	2.1%	
	<i>P</i> (vs. HD)	< 0.001	< 0.01	ns	
CRC	Average	11,197	247,181	19,476	
	SD	6,792	102,736	15,742	
	Positive No.	7	8	5	
	Positive rate (%)	7.4%	8.4%	5.3%	
	<i>P</i> (vs. HD)	ns	ns	ns	

Types of cancer diagnoses included esophageal cancer (EC), gastric cancer (GC), and colorectal cancer (CRC). Purified UBE2E3-GST protein, DAN-GST protein, and antigenic SOSTDC1 peptide were used as antigens. Cutoff values were determined as the average HD values plus two SDs. *P* values were calculated by comparing the results of HDs and patients using the Kruskal–Wallis test. *P* values < 0.05 and positive rates > 10% are marked in bold. A scatter dot plot of the same results is shown in Figure 1E–G. DAN-Ab: anti-differential screening-selected gene aberrant in neuroblastoma antibody

UBE2E3 suppresses the cellular senescence of bone marrow mesenchymal stem cells, leading to osteoporosis [23], possibly via bone morphogenetic proteins (BMPs) [24]. We previously identified serum antibodies against BMP antagonists as atherosclerosis markers, such as anti-DAN antibody (DAN-Ab) [21] and anti-SOSTDC1 antibody (SOSTDC1-Ab) [16]. DAN-Ab levels were elevated in patients with EC and GC, but not with CRC, compared to those in HDs (Figure 1F, Table 5). The AUC values of DAN-Abs for EC, GC, and CRC were 0.6907, 0.6422, and 0.6071, respectively (Figures 2J–L). In contrast, only patients with EC had higher SOSTDC1-Ab levels than those in HDs (Figure 1G, Table 5). The AUC values of SOSTDC1-Abs for EC, GC, and CRC were 0.634, 0.582, and 0.527, respectively (Figures 2M–O).

Prognosis analysis

Next, we tested whether UBE2E3-Ab levels were associated with the postoperative survival of patients with EC, GC, and CRC. We divided the UBE2E3-Ab levels into positive and negative groups using the cutoff values

obtained from the X-tile software [22] to determine the optimal cutoff values for the discrimination of survival rates. Although there were no significant differences in overall survival between the UBE2E3-Abpositive and -negative EC groups (P = 0.1271), the UBE2E3-Abpositive group showed a tendency toward a better prognosis (Figure 3A).

The patients with EC in the DAN-Ab-positive group also showed a more favorable prognosis than the DAN-Ab-negative group, although the difference was not statistically significant (P = 0.1322) (Figure 3B). The prognosis of the UBE2E3-Ab-positive/DAN-Ab-positive group was significantly more favorable than that of the UBE2E3-Ab-negative/DAN-Ab-negative group (P = 0.0179) (Figure 3C). Patients with EC in the SOSTDC1-Ab-positive group showed no difference in prognosis compared with the SOSTDC1-Ab-negative group was more favorable than that of the UBE2E3-Ab-negative/SOSTDC1-Ab-positive group was more favorable than that of the UBE2E3-Ab-negative/SOSTDC1-Ab-negative group was more favorable than that of the UBE2E3-Ab-negative/SOSTDC1-Ab-negative group, but the difference was not significant (P = 0.1545) (Figure 3E). Thus, significant discrimination was observed only for the combination of UBE2E3-Abs and DAN-Abs.

In contrast, the UBE2E3-Ab-positive group had a significantly poorer prognosis than the UBE2E3-Abnegative group in patients with GC (P = 0.0193) (Figure 4A). The DAN-Ab-positive group showed a more unfavorable prognosis than the DAN-Ab-negative group, but the difference was not significant (P = 0.0640) (Figure 4B). The difference in prognosis between the UBE2E3-Ab-positive/DAN-Ab-positive and UBE2E3-Ab-negative/DAN-Ab-negative groups was greater than that of each alone (P = 0.0098) (Figure 4C). Likewise, the SOSTDC1-Ab-positive group showed a somewhat poorer prognosis than the SOSTDC1-Abnegative group (P = 0.1148) (Figure 4D), and the combined UBE2E3-Ab-positive/SOSTDC1-Ab-positive group exhibited a somewhat more unfavorable prognosis than the combined UBE2E3-Abnegative/SOSTDC1-Ab-negative group (P = 0.0063) (Figure 4E). Thus, a more precise prediction of GC prognosis was achieved using a combination of UBE2E3-Abs and DAN-Abs or SOSTDC1-Abs.

The prognostic analysis of CRC showed results similar to those of GC but not of EC. The UBE2E3-Abpositive group had a significantly worse prognosis than the UBE2E3-Ab-negative group in patients with CRC (P = 0.0212) (Figure 5A). The DAN-Ab-positive group tended to have a worse prognosis than the DAN-Ab-negative group (P = 0.1478) (Figure 5B). The prognosis of the combined UBE2E3-Ab-positive/DAN-Abpositive group was worse than that of the combined UBE2E3-Ab-negative/DAN-Ab-negative group (P = 0.0362) (Figure 5C). The SOSTDC1-Ab-positive group had a significantly poorer prognosis than the SOSTDC1-Ab-negative group (P = 0.0290) (Figure 5D), and the combined UBE2E3-Ab-positive/SOSTDC1-Ab-positive group exhibited an even more unfavorable prognosis than the combined UBE2E3-Abnegative/SOSTDC1-Ab-negative group (P = 0.0033) (Figure 5E). Thus, a more precise prediction of CRC prognosis was achieved using a combination of UBE2E3-Abs and DAN-Abs or SOSTDC1-Abs.

Correlation analysis

Spearman's correlation analysis of the CKD cohort of 300 participants revealed significant associations between the plaque score, maximum intima-media thickness (max-IMT) [25, 26], and cardio-ankle vascular index (CAVI) (right and left) [27] (Table 6), which are key indicators of atherosclerosis. Aspartate aminotransferase (AST) levels were weakly associated, and standardized urea clearance (Kt/V), urea nitrogen levels, and high-density lipoprotein cholesterol (HDL-c) levels were inversely correlated with UBE2E3-Ab levels. Other patient data, including age, height, weight, and body mass index (BMI), showed no significant correlation with UBE2E3-Ab levels.

Patient data	s-UBE2E-Ab	
	rho	P value
Age*	0.0726	0.2100
Height	0.0935	0.1065
Weight	0.0211	0.7165
Body mass index (BMI)	-0.0257	0.6584

Table 6. Correlation analysis of serum UBE2E3-Ab levels with the clinical data of CKD cohort

Table 6. Correlation analysis of serum UBE2E3-	Ab levels with the clinical data o	f CKD cohort (continued
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Patient data	s-UBE2E-Ab		
	rho	<i>P</i> value	
Dialysis period	-0.0981	0.0897	
Plaque score	0.1478	0.0109**	
Maximum intima-media thickness (max-IMT)	0.1187	0.0412	
Ankle brachial pressure index (ABI) (right)	0.0253	0.6664	
ABI (left)	-0.0003	0.9952	
Cardio-ankle vascular index (CAVI) (right)	0.1915	0.0013	
CAVI (left)	0.1716	0.0038	
Glycated hemoglobin (HbA1c)	-0.0865	0.2959	
Whole parathyroid hormone (W-PTH)	0.0954	0.0993	
Transferrin saturation ratio	0.0030	0.9584	
Standardized urea clearance (Kt/V)	-0.1515	0.0086	
Red blood cell	-0.0712	0.2187	
Hemoglobin	-0.0354	0.5410	
Hematocrit	-0.0205	0.7238	
Platelet	-0.0681	0.2394	
Total protein	-0.0318	0.5832	
Albumin	-0.0537	0.3544	
Urea nitrogen (UN)	-0.1427	0.0134	
Creatinin	-0.0498	0.3902	
Uric acid	-0.0859	0.1376	
Na	0.0389	0.5018	
К	-0.1030	0.0749	
CI	0.0255	0.6599	
Са	-0.0110	0.8491	
Inorganic phosphate (IP)	-0.0280	0.6287	
Са	0.0186	0.7478	
Mg	0.0899	0.1202	
Fe	-0.0595	0.3046	
Ferritin	0.1319	0.0223	
Aspartate aminotransferase (AST)	0.1212	0.0359	
Alanine amino transferase (ALT)	0.0658	0.2560	
Lactate dehydrogenase (LDH)	0.0701	0.2258	
γ-glutamyl transpeptidase (γ-GTP)	0.0465	0.4223	
Alkaline phosphatase (ALP)	-0.0389	0.5026	
Total bilirubin (tBil)	0.0229	0.6934	
Amylase	-0.0284	0.6246	
Creatin kinase (CK)	-0.0127	0.8273	
Total cholesterol	-0.0540	0.3516	
High-density lipoprotein cholesterol (HDL-c)	-0.1261	0.0290	
Low-density lipoprotein cholesterol (LDL-c)	-0.0399	0.4913	
Triglyceride (TG)	0.1030	0.0747	
C-reactive protein (CRP)	0.1129	0.0507	

Correlation coefficients (*rho*) and *P* values obtained by Spearman's correlation analysis between UBE2E3-Ab levels and the subjects' data are shown. *Subjects' data used were age, height, weight, body mass index (BMI), maximum intima-media thickness (max-IMT), plaque score, cardio-ankle vascular index (CAVI), ankle brachial pressure index (ABI), glycated hemoglobin (HbA1c), whole parathyroid hormone (W-PTH), dialysis period, angiotensin II receptor blocker (ARB), angiotensin converting enzyme (ACE), prothrombin (PTA), iron (Fe), ferritin, Transferrin saturation ratio (TSAT ratio), standardized urea clearance (Kt/V), red blood cell number (RBC), hemoglobin (HGB), hematocrit (HCT), platelet number (PLT), total protein (TP), albumin (ALB), urea nitrogen (UN), creatinine (CRE), uric acid (UA), sodium (Na), potassium (K), chlorine (CI), calcium (Ca), inorganic phosphate (IP), magnesium (Mg), aspartate aminotransferase (AST), alanine amino transferase (ALT), lactate dehydrogenase (LDH), γ -glutamyl transpeptidase (γ -GTP), alkaline phosphatase (ALP), total bilirubin (tBil), amylase (AMY), creatinine kinase (CK), total cholesterol (T-CHO), high-density lipoprotein cholesterol (HDL-c), low-density lipoprotein cholesterol (LDL-c), triglyceride (TG), and C-reactive protein (CRP). **Significant correlations (*P* < 0.05) are marked in bold



Figure 3. Comparison of EC prognosis between the antibody-positive and negative groups. Overall survival of the patients with EC was compared according to UBE2E3-Ab-positive (UBE2E3-Ab+) and negative (UBE2E3-Ab-) groups (**A**). Cutoff values were determined using X-tile software. Statistical analyses were performed using the log-rank test. *P* values and cutoff values are shown in the figures. Similar analyses were performed between anti-differential screening-selected gene aberrant in neuroblastoma antibody (DAN-Ab)-positive and negative groups or SOSTDC1-Ab-positive and negative groups alone (**B** and **D**, respectively) or in combination with UBE2E3-Abs (**C** and **E**, respectively). The numbers of patients at each follow-up period are shown below the figures. The median survival times are also shown



Figure 4. Comparison of GC prognosis between the antibody-positive and negative groups. Overall survival of the patients with GC was compared according to UBE2E3-Ab-positive (UBE2E3-Ab+) and negative (UBE2E3-Ab-) groups (**A**). Cutoff values were determined by X-tile software. Statistical analyses were performed using the log-rank test. *P* values and cutoff values are shown in the figures. Similar analyses were performed between anti-differential screening-selected gene aberrant in neuroblastoma antibody (DAN-Ab)-positive and negative groups or SOSTDC1-Ab-positive and negative groups alone (**B** and **D**, respectively) or in combination with UBE2E3-Abs (**C** and **E**, respectively). The numbers of patients at each follow-up period are shown below the figures





patients with CRC was compared between UBE2E3-Ab-positive (UBE2E3-Ab+) and negative (UBE2E3-Ab-) groups (**A**). Cutoff values, statistical analyses, and *P* values are as described in the legends of Figure 3. Similar analyses were performed between anti-differential screening-selected gene aberrant in neuroblastoma antibody (DAN-Ab)-positive and negative groups or SOSTDC1-Ab-positive and negative groups alone (**B** and **D**, respectively) or in combination with UBE2E3-Abs (**C** and **E**, respectively). The numbers of patients at each follow-up period are shown below the figures

We performed a correlation analysis of UBE2E3-Ab levels in the Sawara cohort. UBE2E3-Ab levels were significantly correlated with carotid artery stenosis, including IMT (right and left) and max-IMT (Table S1), confirming the results from the CKD cohort (Table 6). UBE2E3-Ab levels were strongly associated with blood pressure (P = 0.0015) and smoking duration (P = 0.0020), which are major risk factors for atherosclerosis [28, 29]. Furthermore, UBE2E3 levels were positively correlated with age, alkaline phosphatase levels, thymol turbidity test (TTT) results, and white blood cell numbers and negatively correlated with height, weight, and chlorine levels. The correlation between UBE2E3-Ab levels and the IMT in the Sawara cohort also suggested that UBE2E3-Ab levels were associated with arterial stenosis or atherosclerosis. A chi-square test was conducted to compare the sex differences between the UBE2E3-Ab positive and -negative groups. No significant correlation was observed between sex and UBE2D3-Ab positivity (P = 0.9317) (Table S2). Univariate and multivariate analyses were performed using the AIS cohort. The UBE2E3-Ab level was not a significant independent predictor in the multivariate analysis (P = 0.34) (Table S3).

The correlation of UBE2E3-Ab levels was also examined in 275 patients with DM (DM cohort) at Chiba University Hospital. UBE2E3-Ab levels were correlated with the blood pressure and estimated glomerular filtration rate (eGFR) but were inversely correlated with calcium levels and platelet numbers (Table S4). Thus, the results of the correlation analyses in the CKD, DM, and Sawara cohorts support an association between UBE2E3-Abs levels and the level of atherosclerotic progression.

Discussion

Initial ProtoArray screening identified UBE2E3 as an antigen recognized by serum IgG in patients with atherosclerosis. Serum antibody levels were examined by AlphaLISA using a purified recombinant GST-tagged UBE2E3 protein. The AlphaLISA results indicated significantly higher UBE2E3-Ab levels in patients with AIS, DM, CVD, CKD, EC, and GC than in HDs (Figures 1A–E, Tables 1–5). ROC analysis revealed that the highest AUC values were observed for CKD types 1–3 (Figures 2A–F). Furthermore, Spearman's correlation analysis of the Sawara and CKD cohorts showed significant correlations between UBE2E3-Ab levels and the max-IMT, plaque score, and CAVI (Table 6), which are key indicators of atherosclerosis and arterial stenosis [30–32]. Therefore, UBE2E3-Ab levels may reflect the development of atherosclerosis.

In addition to the UBE2E3-Abs employed in this study, the levels of autoantibodies targeting ATP2B4, BMP-1, KIAA0513, DHPS, LRPAP1, and ASXL2, which are known markers of atherosclerosis, are also elevated in the sera of patients with EC [7, 10, 30–32], suggesting a relationship between atherosclerosis and cancer. Consistently, angiogenesis, the physiological process of the formation of new blood vessels, is crucial for the growth, progression, and metastasis of solid tumors [11], and both diabetes and arteriosclerosis are known risk factors for cancer [12, 33–36]. Therefore, common mechanisms may be involved in the development of atherosclerotic diseases and cancers.

UBE2E3 is a member of the ubiquitin-conjugating enzyme (UBE2) family. Although its biological function remains unclear, another member of the family, UBE2C, is overexpressed in advanced-stage hepatocellular carcinoma tissues [37], head and neck squamous cell carcinoma [38], and uterine corpus endometrial carcinoma [39]. This unfavorable prognosis-associated UBE2C expression is consistent with the results of our survival analyses of UBE2E3-Abs for GC and CRC but not for EC (Figure 3A).

It has been reported that UBE2E3 suppresses cellular senescence in bone marrow mesenchymal stem cells, leading to osteoporosis [23], possibly via BMP signaling [40]. UBE2E3 suppresses cellular senescence in bone marrow mesenchymal stem cells, leading to osteoporosis [23], possibly via BMP signaling [40]. In previous studies, BMP-related proteins have demonstrated causal effects on the development of atherosclerosis [41, 42]. These include BMP-2 [43], BMP-4 [44], the BMP type II receptor BMPRII [45], and the BMP antagonist MGP [46]. Our large-scale screening for atherosclerosis antibody markers also identified BMP-1 [47], BMP antagonists such as SOSTDC1 [16] and DAN [21], and a *BMP* target gene product DIDO1 [13, 48], as antigens recognized by autoantibodies in patients with atherosclerosis. The possible absorption of BMP antagonists by their respective autoantibodies may play a key role in the development of atherosclerosis by increasing active BMP levels.

BMP signaling is also involved in the development of cancer. BMPs play tumor-promoting or tumorsuppressive roles depending on the cancer cell types [49–52]. Thus, we compared the overall survival between the groups that were positive and negative for UBE2F3-Abs, DAN-Abs, and SOSTDC1-Abs, alone or in combination. Among patients with EC, the UBE2E3-Ab-positive group tended to have a more favorable prognosis than the UBE2E3-Ab-negative group, and this prognostic difference became significant in combination with DAN-Abs (Figures 3A and C). In contrast, UBE2E3-Ab positivity was associated with an unfavorable prognosis in GC and CRC, which became significant in combination with positive DAN-Abs or SOSTDC1-Abs. Thus, the tumor-suppressive or -promoting role of BMP signaling may account for the differential prognosis of UBE2E3-Abs between EC and GC or CRC.

Autoantibodies may develop after tissue destruction, followed by leakage of the antigenic proteins that are overexpressed in the lesional tissues, as suggested previously [9, 13]. The repeated leakage of small amounts of antigens results in a marked increase in antibodies to detectable levels. Therefore, antibody markers offer greater sensitivity than antigen markers, and IgG antibodies remain very stable in serum samples. Consequently, serum UBE2E3-Abs are useful for the early diagnosis of AIS, DM, CVD, CKD, and gastrointestinal cancer.

Abbreviations

AIS: acute ischemic stroke

AlphaLISA: amplified luminescence proximity homogeneous assay-linked immunosorbent assay

AMI: acute myocardial infarction

AUC: area under the receiver operating characteristic curve

BMP: bone morphogenetic protein

CAVI: cardio-ankle vascular index

CI: confidence interval

CKD: chronic kidney disease

CRC: colorectal cancer

CVD: cardiovascular disease

DAN: differential screening-selected gene aberrant in neuroblastoma

DAN-Ab: anti-differential screening-selected gene aberrant in neuroblastoma antibody

DM: diabetes mellitus

EC: esophageal cancer

GC: gastric cancer

GST: glutathione S-transferase

HDs: healthy donors

max-IMT: maximum intima-media thickness

ROC: receiver operating characteristic

SD: standard deviation

SEREX: serological identification of antigens by cDNA expression cloning

SOSTDC1: sclerostin domain-containing protein 1

SOSTDC1-Ab: anti-sclerostin domain-containing protein 1 antibody

UBE2E3: ubiquitin conjugating enzyme E2 E3

UBE2E3-Ab: anti-ubiquitin conjugating enzyme E2 E3 antibody

Supplementary materials

The supplementary materials for this article are available at: https://www.explorationpub.com/uploads/ Article/file/101258_sup_1.pdf.

Declarations

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Author contributions

TH: Conceptualization, Investigation, Writing—original draft. YY: Writing—review & editing, Validation. MK: Investigation, Writing—original draft. BSZ: Investigation, Writing—original draft. SYL: Investigation, Writing—original draft. T Matsutani: Resource, Validation, Writing—review & editing. SH: Resource, Validation. MT: Conceptualization, Resource, Validation. KI: Investigation, Validation. SM: Resource, Validation. T Machida: Conceptualization, Validation, Writing—review & editing. YK: Conceptualization, Resource, Validation. HT: Resource, Validation. MI: Resource, Validation. SY: Resource, Validation. HS: Conceptualization, Writing—review & editing, Supervision. KY: Writing—review & editing, Supervision. YH: Writing—review & editing, Supervision. All authors read and approved the submitted version.

Conflicts of interest

The authors declare that they have no conflicts of interest.

Ethical approval

This study was approved by the Local Ethical Review Board of Chiba University, Graduate School of Medicine [approved numbers: 2012-438, 2014-44, 2016-86, 2017-251, 2018-320, 2020-1129, 2022-623, 2023-836], Toho University, Faculty of Medicine [approved numbers: A18103_A17052_A16035_A16001_26095_25024_24038_22047, 25131_23005], Toho University Omori Medical Center [approved number: 26-255], and Port Square Kashiwado Clinic [approved number: 2012-001] as well as by the review boards of the participating hospitals. All experimental procedures were performed in accordance with the Declaration of Helsinki, version 2013.

Consent to participate

Written informed consent was obtained from all participants by following the protocols approved by their institutional ethical committees.

Consent to publication

Not applicable.

Availability of data and materials

All of the results of ProtoArray are available in the public Figshare database (https://figshare.com/articles/ dataset/Results_of_protein_array_for_atherosclerosis/25906330). The other raw data supporting the conclusions of this manuscript will be made available by the authors, without undue reservation, to any qualified researcher.

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References

- Kikuchi Y, Shimada H, Yamasaki F, Yamashita T, Araki K, Horimoto K, et al. Clinical practice guidelines for molecular tumor marker, 2nd edition review part 2. Int J Clin Oncol. 2024;29:512–34. [DOI] [PubMed]
- 2. Nakashima K, Shimada H, Ochiai T, Kuboshima M, Kuroiwa N, Okazumi S, et al. Serological identification of TROP2 by recombinant cDNA expression cloning using sera of patients with esophageal squamous cell carcinoma. Int J Cancer. 2004;112:1029–35. [DOI] [PubMed]
- Kuboshima M, Shimada H, Liu T, Nakashima K, Nomura F, Takiguchi M, et al. Identification of a novel SEREX antigen, SLC2A1/GLUT1, in esophageal squamous cell carcinoma. Int J Oncol. 2006;28:463–8. [PubMed]
- Hou J, Zhao R, Xia W, Chang C, You Y, Hsu J, et al. PD-L1-mediated gasdermin C expression switches apoptosis to pyroptosis in cancer cells and facilitates tumour necrosis. Nat Cell Biol. 2020;22: 1264–75. [DOI] [PubMed] [PMC]
- 5. Routsias JG, Tzioufas AG. Autoimmune response and target autoantigens in Sjogren's syndrome. Eur J Clin Invest. 2010;40:1026–36. [DOI] [PubMed]
- Fularski P, Czarnik W, Dąbek B, Lisińska W, Radzioch E, Witkowska A, et al. Broader Perspective on Atherosclerosis-Selected Risk Factors, Biomarkers, and Therapeutic Approach. Int J Mol Sci. 2024;25: 5212. [DOI] [PubMed] [PMC]
- Hiwasa T, Machida T, Zhang XM, Kimura R, Wang H, Iwase K, et al. Elevated levels of autoantibodies against ATP2B4 and BMP-1 in sera of patients with atherosclerosis-related diseases. Immunome Res. 2015;11:097. [DOI]
- 8. Hiwasa T, Tomiyoshi G, Nakamura R, Shinmen N, Kuroda H, Kunimatsu M, et al. Serum SH3BP5specific antibody level is a biomarker of atherosclerosis. Immunome Res. 2017;13:132. [DOI]
- Li S, Yoshida Y, Kobayashi E, Kubota M, Matsutani T, Mine S, et al. Serum anti-AP3D1 antibodies are risk factors for acute ischemic stroke related with atherosclerosis. Sci Rep. 2021;11:13450. [DOI] [PubMed] [PMC]
- Hiwasa T, Yoshida Y, Kubota M, Li S, Zhang B, Matsutani T, et al. Serum antiKIAA0513 antibody as a common biomarker for mortal atherosclerotic and cancerous diseases. Med Int (Lond). 2024;4:45.
 [DOI] [PubMed] [PMC]
- 11. Makrilia N, Lappa T, Xyla V, Nikolaidis I, Syrigos K. The role of angiogenesis in solid tumours: an overview. Eur J Intern Med. 2009;20:663–71. [DOI] [PubMed]
- 12. Tapia-Vieyra JV, Delgado-Coello B, Mas-Oliva J. Atherosclerosis and Cancer; A Resemblance with Farreaching Implications. Arch Med Res. 2017;48:12–26. [DOI] [PubMed]
- Hiwasa T, Wang H, Goto K, Mine S, Machida T, Kobayashi E, et al. Serum anti-DIDO1, anti-CPSF2, and anti-FOXJ2 antibodies as predictive risk markers for acute ischemic stroke. BMC Med. 2021;19:131.
 [DOI] [PubMed] [PMC]

- 14. Nishiura R, Fujimoto S, Sato Y, Yamada K, Hisanaga S, Hara S, et al. Elevated osteoprotegerin levels predict cardiovascular events in new hemodialysis patients. Am J Nephrol. 2009;29:257–63. [DOI] [PubMed]
- 15. Komatsu H, Fujimoto S, Hara S, Fukuda A, Fukudome K, Yamada K, et al. Recent therapeutic strategies improve renal outcome in patients with IgA nephropathy. Am J Nephrol. 2009;30:19–25. [DOI] [PubMed]
- 16. Goto K, Sugiyama T, Matsumura R, Zhang XM, Kimura R, Taira A, et al. Identification of cerebral infarction-specific antibody markers from autoantibodies detected in patients with systemic lupus erythematosus. J Mol Biomark Diagnos. 2015;6:2. [DOI]
- 17. Hamanaka S, Nakagawa T, Hiwasa T, Ohta Y, Kasamatsu S, Ishigami H, et al. Investigation of novel biomarkers for predicting the clinical course in patients with ulcerative colitis. J Gastroenterol Hepatol. 2018;33:1975–83. [DOI] [PubMed]
- 18. Naito A, Hiwasa T, Tanabe N, Sanada TJ, Sugiura T, Shigeta A, et al. Elevated levels of autoantibodies against EXD2 and PHAX in the sera of patients with chronic thromboembolic pulmonary hypertension. PLoS One. 2019;14:e0211377. [DOI] [PubMed] [PMC]
- Vermeulen N, de Béeck KO, Vermeire S, Steen KV, Michiels G, Ballet V, et al. Identification of a novel autoantigen in inflammatory bowel disease by protein microarray. Inflamm Bowel Dis. 2011;17: 1291–300. [DOI] [PubMed]
- Chumpolkulwong N, Sakamoto K, Hayashi A, Iraha F, Shinya N, Matsuda N, et al. Translation of 'rare' codons in a cell-free protein synthesis system from *Escherichia coli*. J Struct Funct Genomics. 2006;7: 31–6. [DOI] [PubMed]
- 21. Matsumura T, Terada J, Kinoshita T, Sakurai Y, Yahaba M, Tsushima K, et al. Circulating autoantibodies against neuroblastoma suppressor of tumorigenicity 1 (NBL1): A potential biomarker for coronary artery disease in patients with obstructive sleep apnea. PLoS One. 2018;13:e0195015. [DOI] [PubMed] [PMC]
- 22. Camp RL, Dolled-Filhart M, Rimm DL. X-tile: a new bio-informatics tool for biomarker assessment and outcome-based cut-point optimization. Clin Cancer Res. 2004;10:7252–9. [DOI] [PubMed]
- 23. Liu Y, Cai G, Chen P, Jiang T, Xia Z. UBE2E3 regulates cellular senescence and osteogenic differentiation of BMSCs during aging. PeerJ. 2021;9:e12253. [DOI] [PubMed] [PMC]
- 24. Sánchez-Duffhues G, Hiepen C, Knaus P, Dijke PT. Bone morphogenetic protein signaling in bone homeostasis. Bone. 2015;80:43–59. [DOI] [PubMed]
- 25. Tran LTT, Park H, Kim H. Is the carotid intima-media thickness really a good surrogate marker of atherosclerosis? J Atheroscler Thromb. 2012;19:680–90. [DOI] [PubMed]
- 26. Zureik M, Ducimetière P, Touboul PJ, Courbon D, Bonithon-Kopp C, Berr C, et al. Common carotid intima-media thickness predicts occurrence of carotid atherosclerotic plaques: longitudinal results from the Aging Vascular Study (EVA) study. Arterioscler Thromb Vasc Biol. 2000;20:1622–9. [DOI] [PubMed]
- 27. Shirai K, Utino J, Otsuka K, Takata M. A novel blood pressure-independent arterial wall stiffness parameter; cardio-ankle vascular index (CAVI). J Atheroscler Thromb. 2006;13:101–7. [DOI] [PubMed]
- Fan J, Watanabe T. Atherosclerosis: Known and unknown. Pathol Int. 2022;72:151–60. [DOI] [PubMed]
- 29. Libby P, Buring JE, Badimon L, Hansson GK, Deanfield J, Bittencourt MS, et al. Atherosclerosis. Nat Rev Dis Primers. 2019;5:56. [DOI] [PubMed]
- 30. Nakamura R, Tomiyoshi G, Shinmen N, Kuroda H, Kudo T, Doi H, et al. An anti-deoxyhypusine synthase antibody as a marker of atherosclerosis-related cerebral infarction, myocardial infarction, diabetes mellitus, and chronic kidney disease. SM Atheroscler J. 2017;1:1001.

- Sumazaki M, Shimada H, Ito M, Shiratori F, Kobayashi E, Yoshida Y, et al. Serum anti-LRPAP1 is a common biomarker for digestive organ cancers and atherosclerotic diseases. Cancer Sci. 2020;111: 4453–64. [DOI] [PubMed] [PMC]
- 32. Li S, Yoshida Y, Kobayashi E, Adachi A, Hirono S, Matsutani T, et al. Association between serum antiASXL2 antibody levels and acute ischemic stroke, acute myocardial infarction, diabetes mellitus, chronic kidney disease and digestive organ cancer, and their possible association with atherosclerosis and hypertension. Int J Mol Med. 2020;46:1274–88. [DOI] [PubMed] [PMC]
- 33. Jarvandi S, Davidson NO, Schootman M. Increased risk of colorectal cancer in type 2 diabetes is independent of diet quality. PLoS One. 2013;8:e74616. [DOI] [PubMed] [PMC]
- Fujihara S, Kato K, Morishita A, Iwama H, Nishioka T, Chiyo T, et al. Antidiabetic drug metformin inhibits esophageal adenocarcinoma cell proliferation in vitro and in vivo. Int J Oncol. 2015;46: 2172–80. [DOI] [PubMed]
- 35. Gallagher EJ, LeRoith D. Obesity and Diabetes: The Increased Risk of Cancer and Cancer-Related Mortality. Physiol Rev. 2015;95:727–48. [DOI] [PubMed] [PMC]
- 36. Goto A, Yamaji T, Sawada N, Momozawa Y, Kamatani Y, Kubo M, et al. Diabetes and cancer risk: A Mendelian randomization study. Int J Cancer. 2020;146:712–9. [DOI] [PubMed] [PMC]
- 37. Zhang C, Yang M. Functions of three ubiquitin-conjugating enzyme 2 genes in hepatocellular carcinoma diagnosis and prognosis. World J Hepatol. 2022;14:956–71. [DOI] [PubMed] [PMC]
- 38. Yang Y, Chang Y, Tsai K, Hung M, Kang B. UBE2C triggers HIF-1α-glycolytic flux in head and neck squamous cell carcinoma. J Cell Mol Med. 2022;26:3716–25. [DOI] [PubMed] [PMC]
- Ma S, Chen Q, Li X, Fu J, Zhao L. UBE2C serves as a prognosis biomarker of uterine corpus endometrial carcinoma via promoting tumor migration and invasion. Sci Rep. 2023;13:16899. [DOI] [PubMed] [PMC]
- 40. Li S, Xue S, Li Z. Osteoporosis: Emerging targets on the classical signaling pathways of bone formation. Eur J Pharmacol. 2024;973:176574. [DOI] [PubMed]
- 41. Cai J, Pardali E, Sánchez-Duffhues G, Dijke Pt. BMP signaling in vascular diseases. FEBS Lett. 2012;586: 1993–2002. [DOI] [PubMed]
- 42. Dyer LA, Pi X, Patterson C. The role of BMPs in endothelial cell function and dysfunction. Trends Endocrinol Metab. 2014;25:472–80. [DOI] [PubMed] [PMC]
- 43. Zhang M, Sara JD, Wang F, Liu L, Su L, Zhe J, et al. Increased plasma BMP-2 levels are associated with atherosclerosis burden and coronary calcification in type 2 diabetic patients. Cardiovasc Diabetol. 2015;14:64. [DOI] [PubMed] [PMC]
- 44. Sorescu GP, Sykes M, Weiss D, Platt MO, Saha A, Hwang J, et al. Bone morphogenic protein 4 produced in endothelial cells by oscillatory shear stress stimulates an inflammatory response. J Biol Chem. 2003;278:31128–35. [DOI] [PubMed]
- Kim CW, Song H, Kumar S, Nam D, Kwon HS, Chang KH, et al. Anti-inflammatory and antiatherogenic role of BMP receptor II in endothelial cells. Arterioscler Thromb Vasc Biol. 2013;33:1350–9. [DOI] [PubMed] [PMC]
- Yao Y, Bennett BJ, Wang X, Rosenfeld ME, Giachelli C, Lusis AJ, et al. Inhibition of bone morphogenetic proteins protects against atherosclerosis and vascular calcification. Circ Res. 2010;107:485–94. [DOI] [PubMed] [PMC]
- 47. Machida T, Kubota M, Kobayashi E, Iwadate Y, Saeki N, Yamaura A, et al. Identification of strokeassociated-antigens via screening of recombinant proteins from the human expression cDNA library (SEREX). J Transl Med. 2015;13:71. [DOI] [PubMed] [PMC]
- 48. Braig S, Bosserhoff A. Death inducer-obliterator 1 (Dido1) is a BMP target gene and promotes BMPinduced melanoma progression. Oncogene. 2013;32:837–48. [DOI] [PubMed]
- 49. Zhou W, Yan K, Xi Q. BMP signaling in cancer stemness and differentiation. Cell Regen. 2023;12:37. [DOI] [PubMed] [PMC]

- 50. Sharma T, Kapoor A, Mandal CC. Duality of bone morphogenetic proteins in cancer: A comprehensive analysis. J Cell Physiol. 2022;237:3127–63. [DOI] [PubMed]
- 51. Ehata S, Miyazono K. Bone Morphogenetic Protein Signaling in Cancer; Some Topics in the Recent 10 Years. Front Cell Dev Biol. 2022;10:883523. [DOI] [PubMed] [PMC]
- 52. Alkhathami AG, Abdullah MR, Ahmed M, Ahmed HH, Alwash SW, Mahdi ZM, et al. Bone morphogenetic protein (BMP)9 in cancer development: mechanistic, diagnostic, and therapeutic approaches? J Drug Target. 2023;31:714–24. [DOI] [PubMed]