ORIGINAL ARTICLE

Circulating tumour-derived microvesicles in plasma of gastric cancer patients

Jaroslaw Baran · Monika Baj-Krzyworzeka · Kazimierz Weglarczyk · Rafal Szatanek · Maria Zembala · Jakub Barbasz · Antoni Czupryna · Antoni Szczepanik · Marek Zembala

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Abstract Cell membrane microfragments called microvesicles (MV) originating from different cells are circulating in the blood of healthy subjects and their elevated numbers are found in different diseases, including cancer. This study was designed to characterise MV present in plasma of gastric cancer patients. Since majority of MV in blood are platelets-derived (PMV), plasma samples deprived of PMV were used. In comparison to control, the number of MV in patients was significantly elevated in all stages, higher in more advanced disease. Patients' MV showed an increased membrane expression of CCR6 and HER-2/neu. The proportion of MV carrying some leucocyte determinants was low and similar in patients and control. Transmission electron microscopy showed their substantial heterogeneity in size and shape. The size

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J. Baran · M. Baj-Krzyworzeka · K. Weglarczyk · R. Szatanek · M. Zembala (⊠) Department of Clinical Immunology, Polish-American Institute of Paediatrics, Jagiellonian University Medical College, Wielicka Str. 265, 30-663 Cracow, Poland e-mail: mizembal@cyf-kr.edu.pl

M. Zembala · J. Barbasz Institute of Catalysis and Surface Chemistry, Polish Academy of Sciences, Niezapominajek Str. 8, 30-239 Cracow, Poland

A. Czupryna · A. Szczepanik

First Department of General and Gastrointestinal Surgery, Jagiellonian University Medical College, Kopernika Str. 40, 31-501 Cracow, Poland

J. Barbasz

Institute of Physics, Jagiellonian University, Reymonta Str. 4, 30-059 Cracow, Poland determined by dynamic light scattering analysis confirmed this heterogeneity. The MV size distribution in patients was broader within the range of 10-800 nm, while in control MV showed 3-mode distribution within the range of 10-400 nm. Atomic force microscopy confirmed MV size heterogeneity with implication that larger objects represented aggregates of smaller microparticles. Patients' MV exhibited increased absolute values of zeta potential, indicating a higher surface charge. Tumour markers HER-2/neu, MAGE-1, c-MET and EMMPRIN were detected both in control and patients' samples with stronger expression in the latter. Significantly higher expression of MAGE-1 and HER-2/neu mRNA was observed in individual patients. All together, it suggests that at least some MV in plasma of gastric cancer patients are tumourderived. However, their role in cancer requires further studies.

Keywords Microvesicles · Gastric cancer · Chemokine receptors · Tumour markers

Abbreviations

- AFM Atomic force microscopy
- MV Microvesicles
- PFP Platelet-free plasma
- PMV Platelet-derived microvesicles
- TEM Transmission electron microscopy
- TMV Tumour-derived microvesicles

Introduction

Microvesicles (MV) are membrane microfragments secreted from cytoplasmic membrane compartments by normal and malignant cells in vitro and in vivo. MV are defined as spherical proteolipids and divided by size into two groups: smaller (30–100 nm), more homogeneous in size, released by endosomal compartment (called exosomes) and larger (0.1–1 μ m), released from the surface membranes during membrane blebbing through calcium-dependent mode [1]. Although biological activity of these two types of MV appears to be similar [2], exosomes have a more homogeneous protein composition than surface membrane MV and are viewed as antigen carrying vehicles [1, 3, 4]. Surface membrane MV are characterised by the presence of procoagulant phospholipids, which confer possible prothrombotic potential, and antigens similar to those present in the membranes of the cells from which MV originate [5, 6].

Tumour cells release MV (tumour-derived microvesicles, TMV) continuously, and the rate of their shedding is increased in more malignant tumours [7, 8]. Our and others results showed that TMV released by different cell lines in vitro expressed CD44, β 1 integrins, EMMPRIN, CEA, tetraspanin, HLA class I and II, TRAIL, FasL, ADAM 10, L1 and CX3CR1 [3, 9, 10] and carry mRNA for some of them [5, 9].

In in vitro cultures TMV stimulate secretion of cytokines [11] and chemokines by monocytes (Baj-Krzyworzeka, unpublished). TMV also induce secretion of several proangiogenic factors by stromal fibroblasts, chemoattract endothelial cells and enhance their proliferation [12, 13]. An increasing attention has recently been focused on MV as they are present in plasma and other body fluids of cancer patients [14–16].

Plasma level of MV is independent of donor gender or age [17], but depends on the clinical status of the patient [14, 18, 19]. In plasma of normal subjects, platelet-derived microvesicles (PMV) constitute the largest proportion of MV $(\sim 80\%)$ [20]. The remaining 20% is composed of leucocyte, erythrocyte and endothelial cell-origin MV [13, 20]. The number of circulating MV changes during inflammation or malignant processes. The elevated counts of MV, mainly PMV, were observed in plasma of gastric cancer patients, and their level may be a predictive marker for metastasis formation [14]. An increasing number of PMV and monocyte-derived MV was found in patients with lung cancer and may be associated with vascular complications [21]. The number of endothelial and hepatic origin MV, but not total MV, correlates with the tumour size in hepatocellular carcinoma [22]. FasL-bearing MV, described in sera of patients with oral squamous [12], ovarian [2], colorectal [23] carcinomas and melanoma [24] induce apoptosis of activated T cells. TMV may also impair monocyte differentiation to dendritic cells in vitro [25].

The TMV may be involved in tumour escape (relocation of determinants) [26], induction of immunotolerance or

immunosuppression [25, 27], chemoresistance of tumours [28] and promotion of angiogenesis [12]. On the other hand, MV isolated from plasma and dendritic cells from cancer patients may carry tumour-associated antigens (TAA) and induce antitumour response [29]. It may suggest that circulating MV/TMV are important factors in affecting immune cells at a distance from the tumour microenvironment [30]. To the best of our knowledge, there is no complete characterisation of circulating TMV in patients with cancer.

Here, we present, for the first time, the broad characterisation of MV found in PMV-depleted plasma of patients with gastric cancer. The analysis comprises their quantity, immunophenotype, size and physical characteristics at single particle level.

Materials and methods

Patients

Thirty-seven patients (median age 66 years, range 40–76) with biopsy proven gastric cancer at different clinical stages were studied. The postsurgical clinicopathological staging system of the UICC TNM Classification of Malignant Tumours (sixth edition) was used [31]. The control group consisted of ten healthy subjects (median age 41 years, range 33–61). All study participants signed informed consent form, and the Bioethical Committee of the Jagiellonian University approved the protocol.

Blood samples and CD61⁻ plasma from donors and patients with gastric cancer

Peripheral blood was drawn into polypropylene tubes containing EDTA (Vacutainer System, BD Biosciences, San Jose, CA, USA). The patients' blood was taken during routine laboratory tests. The whole blood samples were centrifuged for 15 min at $1,500 \times g$; within 60 min of drawing plasma was collected and spun down for 30 min at $3,000 \times g$ to remove platelets. Next, plasma was carefully aspirated from the pellet, centrifuged at $15,000 \times g$ for 30 min, aliquoted and kept frozen at -80° C until use. In some analysis plasma samples were additionally centrifuged at $50,000 \times g$ for 60 min to obtain purified MV (called isolated MV).

Immunophenotyping of plasma MV

The following monoclonal antibodies (mAbs) were used to characterise the phenotype of MV: FITC-conjugated against CD3, CD11a, CD61, CD64 (all from BD Pharmingen, San Diego, CA, USA); PE-conjugated against CD29, CD34, CD41, HLA-class I, HLA-DR, CCR6, CCR7 (all from BD Pharmingen), CCR1, CCR2, CCR3, CXCR1, CXCR4 (all from R&D Systems Minneapolis, MN, USA), and APCconjugated anti-CD14 and anti-HER-2/neu (both from BD Biosciences). Plasma samples (20 μ l) were incubated with mAbs or appropriate isotype control for 30 min at 4°C, and after washing resuspended in 0.5 ml of PBS. Samples were analysed in FACSCanto flow cytometer (BD Biosciences, Immunocytometry Systems, San Jose, CA, USA) after gating on a forward (FSC) and side scatter (SSC) parameters set in log scale. Data from 500,000 events were acquired and analysed using FACS DiVa software. To create histogram overlays, data were saved in FCS 2.0 format and exported to CellQuest software (BD Biosciences).

Determination of absolute number of MV

To determine absolute counts of MV, flow cytometry with the use of BD TruCOUNTTM Tubes (BD Biosciences) was used. MV gating was accomplished by preliminary standardisation experiments using beads of different size. Samples (0.5 ml) of plasma (diluted 1:100) in TruCOUNTTM Tubes containing known quantity of 4.2-µm beads were analysed in FACSCanto flow cytometer on FSC, and SSC set in log scale. By this approach, we were able to use FSC as a trigger and analyse beads and MV on the same cytogram without using fluorochrome conjugated mAb to label MV. Samples were run at the "low flow" mode, and acquisition was stopped after 1,000 events were counted in the beads gate. Absolute numbers of MV were calculated as previously described [32], according to the formula:

number of events in region containing MV	
number of events in absolute count bead region	
number of beads per test $-$ shealute count of MN	,
test volume = absolute could of M	V

Transmission electron microscopy (TEM) of MV

Purified MV obtained from pooled plasma samples of ten stage IV gastric cancer patients were mixed with 2% of uranyl acetate and applied to carbon coated copper grid. The microphotographs were taken with Philips EM 300 electron microscope (Philips, Eindhoven, The Netherlands) operating at 80 kV.

Determination of MV size distribution

For particle size determination a dynamic light scattering method was used. Samples were diluted with 0.15 M NaCl to the level giving an appropriate optical signal (usually 1:100) and measured in Zetasizer Nano ZS apparatus (Malvern Instruments, Malvern, UK) equipped with a laser of $\lambda = 633$ nm. Particle size distributions were obtained from measured diffusion coefficients assuming spherical shape of particles. The obtained values represent the radius

of spherical particles which move in viscous media with the same velocity as studied particles.

Zeta potential determination

Particle electrophoretic mobility was measured in samples fourfold diluted with distilled water to keep sample conductivity at the level of about 4 mS/cm assuring measurable signal. Electrophoretic mobility was recalculated to zeta potential using Smoluchowski equation valid under conditions of thin electric double layer, in comparison to particle size. Measurements were performed using Zeta Pals apparatus (Brookhaven Instruments Corporation, New York, NY, USA).

Atomic force microscopy (AFM)

Samples for AFM were prepared from isolated MV diluted 1:10,000 to 1:100,000 with 0.15 M NaCl by adsorbing onto HOPG (ZYH-mosaic spread 3.5-5.0°) highly ordered pyrolytic graphite supports (NT-MDT Co., Zelenograd, Russia). 5 µl of diluted sample was placed onto freshly exposed HOPG surface and kept there for the time necessary to get an appropriate coverage (usually from 0.5 to 5 min). After adsorption, the sample was carefully washed with distilled water to remove any traces of NaCl which could crystallise at the support surface. Next, the dry sample was placed under 7-10 nm AFM tip. The measurements were performed using Ntegra Vita and Solver P47-PRO microscopes (NT-MDT). The AFM images were recorded in semicontact mode. Experiments were performed using a commercial silicon tip with polysilicon lever (NT-MDT) with resonance frequency within a range of 140-240 kHz and spring constants k = 3.4 N/m or k = 5.8 N/m.

Western blot analysis

Plasma samples were treated for 4 min at 95°C in a sodium dodecyl sulphate (SDS) reducing buffer containing β -mercaptoethanol, electrophoresed on SDS-polyacrylamide gel (12% polyacrylamide) and transferred onto polyvinylidene difluoride membrane (BioRad, Hercules, CA, USA). The membranes were blocked for 1 h at room temperature in Tris buffered saline buffer (TBS) with 0.1% Tween and 1% bovine serum albumin (BSA, Sigma, St. Louis, MO, USA). The membranes were then incubated overnight at 4°C in TBS-Tween-BSA with various antibodies: goat anti-EMMPRIN (N-19), mouse anti-HER-2/ neu (F-11), rabbit anti-MAGE-1 (FL-309) and rabbit antic-Met (C-12). The membranes were washed in TBS-Tween-BSA and incubated with donkey anti-goat, goat anti-mouse or goat anti-rabbit secondary antibody conjugated with horseradish peroxidase (dilution 1:2,000). All antibodies were purchased from Santa Cruz Biotech (Santa Cruz, CA, USA). The protein bands were visualised with the SuperSignal West Pico Chemiluminescence Substrate kit as recommended by the manufacturer (Pierce, Rockforld, IL, USA) and analysed with KODAK GEL LOGIC 1500 Digital Imaging System (Kodak, Rochester, NY, USA).

Expression of MAGE-1 and HER-2/neu mRNA

Total RNA from isolated MV of individual plasma samples was obtained using the P.A.L.M. RNA Isolation Kit (PALM, Berniard, Germany) according to the manufacturer's protocol. Reverse transcription (RT) of the obtained RNA as well as MAGE-1 "real-time" PCR were performed as previously described [33]. HER-2/neu "realtime" PCR was performed in the LightCycler (Roche, Mannheim, Germany) using the protocol and set of primers described previously [34]. The quantification of relative mRNA gene expression (fold increase), normalised to an endogenous control β -actin mRNA, was analysed using the $2^{-\Delta\Delta C_t}$ method, where $\Delta\Delta C_t = (C_t \text{ target} - C_t \text{ actin})_{\text{patient}} - (C_t \text{ target} - C_t \text{ actin})_{\text{donor}}$. To verify the amplified product, melting curve analysis using the LightCycler software was performed for each sample.

Statistical analysis

Statistical analysis was performed by nonparametric Mann–Whitney test in all experiments using GraphPad Prism 4 Software. Differences were considered significant at P < 0.05.

Results

The number of MV present in platelet-free plasma of gastric cancer patients

In the initial observations, the number of MV in individual plasma of patients and healthy subjects was determined by flow cytometry. Significantly elevated level of MV was observed in patients' plasma (Fig. 1a). However, FACS analysis revealed that these MV consist of two different populations, which also differ in the expression of CD61 (Fig. 2a). Over 90% of MV located in R2 were CD61⁺ indicating their platelet origin (Fig. 2a, right). As in the present study we wished to analyse MV of none platelet origin, plasma samples were centrifuged for 30 min at $15,000 \times g$ [35]. The occurrence of PMV was determined by staining with anti-CD61 mAb. Figure 2b shows data from a representative experiment, which indicates that following centrifugation at $15,000 \times g$ CD61⁺ MV in plasma were substantially decreased and the number of MV in plasma dropped by 4 logs (Fig. 1b). The majority of CD61⁺ MV was seen in the pellet (Fig. 2b, right dot plot). For simplicity such samples, platelet and PMV depleted will be called thereafter platelet-free plasma (PFP). All data presented below is based on PFP samples. Next, we determined the number of MV in PFP of patients with different stages of gastric cancer (Fig. 1b). In comparison to control, the number of CD61⁻ MV was significantly increased in the whole group of patients, as well as in stage II-IV disease. There were no differences between stages, but the highest level was observed in the most advanced disease.

Phenotype of MV

Determination of the expression of various leucocyte surface determinants by flow cytometry revealed low level of MV expressing CD45 (mean 0.8%; range 0.2–1.2%), CD64 (1.7%; range 1–2.2%), CD11a (1.9%; range 0.9–2.6%) and CD18 (1.4%; range 0.8–2.0%) both in normal and patients' plasma and a lack of expression of CD3, CD19, CD1a, CD14, CD34 and HLA-DR. Approximately up to 2% MV were HLA-class I positive (range 1.5–2.6%). Interestingly enough, expression of several chemokine receptors (CR) was observed (Table 1). Patients' MV were characterised by a higher expression of CCR6 and lower expression of CXCR4. CCR1 was increased only in patients with stage I cancer.

Fig. 1 The number of MV (mean \pm SD) in **a** whole plasma of control and patients' samples. **b** PFP of control and patients with different stages of gastric cancer. Significance in comparison to control; **P* < 0.001

Then, we looked for the presence of HER-2/neu on MV from both patients (stage IV) and control (Fig. 3).





Fig. 2 FACS analysis of morphology and CD61 expression in different preparations of plasma. Acquisition parameters were setup to exclude remaining platelets. **a** Events from plasma samples were gated into two populations according to FSC and SSC parameters (R1 and R2) (*left dot plot*) and analysed for CD61

expression. Data from R1 (*middle*) and R2 (*right*) are shown. **b** Plasma samples were centrifuged $(15,000 \times g)$ and both supernatants and pellets were stained for CD61 expression. *Dot-plots* show CD61 staining in whole plasma (*left dot plot*), supernatant (*middle*) and pellet (*right*). Data from representative experiment are shown

Table 1 Expression of chemokine receptors (%) on MV from gastric cancer patients (different stage) or healthy donors

Marker	Control	Patients	Stage I	Stage II	Stage III	Stage IV
CCR1	1.3 ± 0.9	0.7 ± 0.8	$0.3 \pm 0.3*$	0.8 ± 0.7	0.6 ± 0.7	0.9 ± 0.9
CCR2	3.5 ± 2.0	2.5 ± 2.6	2.9 ± 3.2	3.7 ± 3.8	2.1 ± 1.9	2.1 ± 1.6
CCR3	4.0 ± 3.0	4.5 ± 2.7	6.1 ± 0.7	6.6 ± 1.6	1.9 ± 0.8	4.2 ± 1.0
CCR6	0.8 ± 1.2	$4.5 \pm 2.7*$	$6.4 \pm 0.6^{***}$	$4.2 \pm 0.3*$	2.8 ± 0.5	$4.5 \pm 0.5^{**}$
CCR7	2.4 ± 2.8	2.7 ± 2.4	2.8 ± 2.0	2.0 ± 1.7	1.1 ± 0.7	3.6 ± 2.7
CXCR1	0.8 ± 0.8	0.7 ± 0.5	0.8 ± 2.3	0.4 ± 3.3	0.7 ± 2.1	0.7 ± 2.6
CXCR4	4.8 ± 4.3	$1.1 \pm 1.0^{*}$	$0.8\pm2.7^*$	1.6 ± 2.0	$0.9 \pm 1.4^{*}$	$1.2 \pm 2.4^{**}$

Data are presented as mean \pm SD from eight different experiments

* P < 0.05 comparing to control

** P < 0.005 comparing to control

*** P < 0.0001 comparing to control

The patients showed significantly higher proportion of HER-2/neu⁺ MV when compared to control. Expression of MUC-1 was observed on approximately 1-2% MV (data not shown).

Characteristics of MV

This has been performed on isolated MV from two different pools of PFP from both patients with stage IV of



Fig. 3 Membrane expression of HER-2/neu on MV of stage IV gastric cancer patients (n = 13) and control (n = 10); *P < 0.05

gastric cancer and control, each containing ten individual samples.

Morphology and the size

Transmission electron microscopy revealed substantial heterogeneity of isolated MV both in size and shape (Fig. 4a), seen in patients' and control plasma samples. The dynamic light scattering analysis was used to determine the size distribution of MV present in PFP plasma of donors and patients. The size range of MV in control was approximately 10–400 nm and showed 3-mode distribution with sizes of approximately 10, 30–50 and 140–200 nm (Fig. 4b). The most frequent events in patients and control were around 10 nm (data not shown). In plasma from patients' MV size distribution was very dispersed showing the whole range of sizes from around 10 to 800 nm, the most common being also around 10 nm (Fig. 4c). This indicated that population of MV in patients is more heterogeneous in size.

Zeta potential analysis

Zeta potential characterises the effective charge of any objects, including particles, membranes, etc. Using PALS Zeta Potential Analyzer it was found that control samples had -11.9 ± 1.8 mV and patients' -16.5 ± 1.13 mV (significantly different at P < 0.05), indicating a higher electrokinetic charge of patients' MV.

Atomic force microscopy

Purified MV isolated from control samples were analysed in atomic force microscopy to identify precisely the shape of the objects seen. Single particles were detected only when adsorption was performed on samples 100,000 times diluted. Pictures taken by AFM characterise well object



Fig. 4 Morphology and size of isolated MV from PFP. **a** Transmission electron microscopy of purified MV from plasma of gastric cancer patients (magnification $60,000\times$) and size distribution of MV in PFP from **b** control and **c** patients analysed by dynamic light scattering

shapes and sizes in two dimensions (X and Y) whereas Z value cannot be directly correlated with the height of the real object. Figure 5a shows that majority of MV formed aggregates of the sizes corresponded to those found by light scattering analysis (Fig. 5b). AFM pictures of single particles indicated that they are spheroid in shape with diameters of about: 8.0-9.5 nm × 13.0-16.5 nm. These values corresponded well with the most frequent fraction of objects measured by light scattering (first peak of maximum size of about 10 nm) (Fig. 5c). Such small objects tend to form larger aggregates, and the first step of this process can be already noticed when analysing the cross-section of them (Fig. 5d). The object shown is formed from a few subunits.

Expression of tumour-associated proteins

Western blot was used for detection of HER-2/neu, c-MET, MAGE-1 and EMMPRIN, which are known to be overexpressed by cancer cells. In patients' samples, the



1000



b

12 10%

Fig. 5 AFM analysis of MV. a Scan of MV taken at scan area 500 nm \times 500 nm. b Marked three representative MV sizes detected by dynamic light scattering. c Scan of MV with quantified cross-section. d Complex structure of MV after cross-section

presence of all of them was found (supplementary data). However, weaker bands of these markers were also found in control samples (data not shown). These proteins were not detected in plasma samples after centrifugation at $50,000 \times g$, i.e. following removal of MV.

Expression of MAGE-1 and HER-2/neu mRNA

Isolated MV from individual plasma samples of five different patients with stage IV gastric cancer and healthy donors were analysed for MAGE-1 and HER-2/neu mRNA expression. The mRNA for each of the two genes was found both in control and patients' MV; however, it was substantially higher in the latter. The MAGE-1 mRNA expression was about 2¹⁰ higher in the patients' samples than in control. HER-2/neu-mRNA expression was 2³ higher in the patients' samples (Fig. 6).

Discussion

In this study, the attempts were made to characterise MV present in plasma of gastric cancer patients. Since it is known that the vast majority (over 80%) of MV present in plasma are of platelet origin [20], initial centrifugation at

 $3,000 \times g$ to remove platelets was followed by centrifugation at 15,000 $\times g$. This was effective in depletion of CD61⁺ PMV, which were detected in the pellet (up to 91% PMV). By this approach, the number of MV in PFP dropped substantially. Although the number of MV in the initial plasma of patients was higher than in control, the use of PFP revealed much more pronounced differences. The levels of MV did not correlate significantly with different stages of cancer, but they were more numerous in more advanced cancer. However, as AFM revealed that MV form aggregates, it may be that the numbers detected are underestimated. Determination of immunophenotype by flow cytometry indicated a very small proportion of MV expressing CD45, CD64, CD11a, CD18 and HLA-class I, but no other T and B lymphocytes, monocytes and probably no endothelial/stem cells determinants, as judged by the absence of CD34. This may indicate that the vast majority of MV is not of leucocyte origin. However, it is also possible that mAbs may not be able to bind to very small particles, which are the most frequent. On the other hand, expression of surface determinants on MV often does not reflect that on the membrane of cells they are originating from [5, 6]. However, when the small percentages of MV with leucocyte determinants are recalculated to absolute numbers, e.g. CD45⁺ cells in normal plasma samples,

Fig. 6 Fold increase in **a** MAGE-1 and **b** HER-2/neu mRNA expression in isolated MV of patients with stage IV gastric cancer (n = 5) as compared to control subjects (n = 5); *P < 0.05



they are similar to those found in the other studies [36]. MV expressed several chemokine receptors, but only CCR6 was elevated on MV from the patients. We have previously shown that micrometastatic tumour cells which have a high expression of CCR6 are present in the blood, bone marrow and lymph nodes from patients with gastric cancer [37]. Also, several cell lines established from gastric cancer and MV released by them expressed CCR6 [our unpublished data]. It may suggest that tumour cells are a source of CCR6⁺ MV. CCR6 have recently been shown to be involved in prostate cancer progression [38] and metastasis formation in colorectal cancer [39]. Another possible source of CCR6⁺ MV might be CD3⁺ T cells infiltrating gastric mucosa as they have a significantly increased CCR6 expression. However, it is not the case here as no CD3 on MV was detected. We also observed the elevated level of CCR1 positive MV in plasma of stage I patients, which is in keeping with an increased expression of CCR1 on gastric cancer tissue described by Sugasawa [40]. Surprisingly, no CCR7 which is associated with gastric cancer metastasis to lymph nodes [41] was detected, but there are no data on other metastatic lesions. Furthermore, none of gastric cancer cell lines established by us expressed CCR7 (data not shown). CXCR4 and its ligands influence the dissemination, immune rejection and neoangiogenesis of human gastrointestinal cancers [42]. Strong expression of CXCR4 on gastric cancer was described, which correlated with lymph node metastases [43]. We observed lower level of CXCR4 positive MV in all patients, and the reason for this downregulation is unclear. Finally, MV showed membrane expression of HER-2/neu. Although HER-2/neu⁺ MV were seen in both control and patients, their proportion was substantially higher in the latter. The presence of this antigen in normal plasma is not surprising as it is present on some epithelial cells but overexpressed on cancer cells, which may explain the increased HER-2/neu⁺ MV in patients samples. This suggests that at least some HER-2/neu⁺ MV in patients are of tumour cells origin. To our knowledge, this is the first demonstration of the expression of this TAA on plasma MV.

TEM pictures showed heterogeneity and large differences both in size and shape with the most common round shaped particles. Differences in size were confirmed by dynamic light scattering technique which measures temporary fluctuation in intensity of light dispersed by particles and provides precise measure of particle size, even as small as 10 nm. Although flow cytometry is recommended as a reference method for MV analysis [36, 44], it should be noted that this method is unable to determine particles smaller than 200 nm. There were some differences in MV sizes between patients and control, and their distribution was also different. Size distribution was more dispersed in patients, while three clear cut peaks were seen in control. However, in both cases the most frequent MV size was around 10 nm. This corresponded well to AFM scans of control sample in which MV of three sizes were identified. The larger size MV represented aggregates of small particles. However, AFM pictures indicated that even small size particles are composed of a few subunits. It should be stressed that in contrast to TEM objects observed in AMF are much closer to those existing in solutions.

Patients' MV were characterised by significantly higher absolute values of zeta potential, which provides information about electrokinetic charge of the particle. Increased values of zeta potential may indicate higher stability of suspension containing similar type particles [45]. The small particles adsorbed by local interactions at various surfaces produce or modify their heterogeneity [46]. It may be suggested that higher zeta potential facilitate known role of small particles in adherence to other particles, surfaces, phospholipid membranes, etc., as shown for MV in patients with gastrointestinal diseases [47].

Western blot analysis showed the presence of HER-2/ neu, c-MET, MAGE-1, EMMPRIN, seen both in patients' and control samples, though were less expressed in the latter (supplementary data). However, it should be treated with caution as we have no direct proof that they are MVassociated. However, since no such proteins were detected in plasma samples following centrifugation at $50,000 \times g$, typically used to obtain MV in the pellet, it may suggest indirectly that these found in PFP are MV-associated. Expression of MAGE-1 and HER-2/neu mRNA was found in isolated MV of individual plasma samples both from patients and control; however, the mRNA expression was substantially higher in the former. This corresponded well to these proteins detected by Western blot. The indirect evidence that these mRNAs were MV-associated was obtained by finding that they were also present in the pellet of plasma samples obtained by centrifugation at $50,000 \times g$, i.e. containing purified MV. This is also in keeping with our previous observations that MV from cancer cell lines carry mRNA for IL-8, VEGF and HGF mRNA [5]. Therefore, it could be stated that circulating MV present in plasma also carry mRNA for some TAA.

In summary, this study shows that plasma level of MV of non-platelet origin is elevated in gastric cancer patients and describes their characteristics, including immunophenotype, morphology, size, structure at the single particle level, expression of some TAA proteins and their mRNA. However, further studies are required on their biological role in cancer as TMV may be involved in induction of immunosuppression, dysfunction or death of immune cells and metastasis formation [30]. On the other hand, they may play a role as a cell-free antigen source for anticancer vaccines [26].

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