

Human osteopontin: Potential clinical applications in cancer (Review)

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Abstract. Human osteopontin (OPN) is a glycosylated phosphoprotein which is expressed in a variety of tissues in the body. In recent years, accumulating evidence has indicated that the aberrant expression of OPN is closely associated with tumorigenesis, progression and most prominently with metastasis in several tumour types. In this review, we present the current knowledge on the expression profiles of OPN and its main splice variants in human cancers, as well as the potential implications in patient outcome. We also discuss its putative clinical application as a cancer biomarker and as a therapeutic target.

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1. Introduction

Osteopontin (OPN) is a bone associated, extracellular matrix glycosylated phosphoprotein which is produced by several cell types, including osteoblasts, osteoclasts, immune cells, endothelial cells, epithelial cells and extra-osseous cells (skin, kidney and lung) (1-3). Due to differences in post-translational modification (PTM) (phosphorylation, glycosylation, sulfation and proteolysis) from different cellular sources, OPN has a molecular weight ranging from 41 to 75 kDa, which may have a cell type-specific structure and function (4-7). OPN plays a major role in various normal physiological processes, including

bone remodelling, immune-regulation, inflammation and vascularisation (8,9). In addition, OPN has also been shown to be involved in carcinogenesis with multi-functional activities (10-12).

OPN is involved in a series of biological functions through interactions with different integrins and CD44. Therefore, OPN is classified as a member of the 'small integrin-binding ligand N-linked glycoproteins' (SIBLINGs) together with other molecules, including bone sialoprotein (BSP), dentin matrix protein 1 (DMP1), dentin sialophosphoprotein (DSPP) and matrix extracellular phosphoglycoprotein (MEPE) (13). Two critical integrin binding sequences of OPN have been identified: arginine-glycine-aspartic acid (RGD) and serine-valine-valine-tyrosine-glutamate-leucine-arginine (SVVYGLR). OPN interacts mainly with various α v (particularly α v β 1, α v β 3, α v β 5) integrin receptors via the classical RGD sequence, and interacts with α 9 β 1, α 4 β 1, α 4 β 7 via SVVYGLR (14-16). In addition, it also interacts with the CD44 splice variants, CD44v3, CD44v6 and CD44v7, via the C-terminal fragment calcium binding site (17-20). These properties of OPN induce the activation of signal transduction pathways, leading to cell proliferation, adhesion, invasion and migration, which have been demonstrated by both *in vitro* and *in vivo* models (21-23). The binding of OPN to integrins and CD44 initiates a downstream signalling cascade via the PI3K/AKT signalling pathway leading to NF- κ B mediated cell proliferation and survival (24-26). In addition, through the Ras/Raf/MEK/ERK signalling pathway, an OPN-integrin complex and subsequent induction of AP-1-dependent gene expression, urokinase-type plasminogen activator (uPA) and matrix metalloproteinases (MMPs) confer a metastatic phenotype on some cancer cell types (27-29). Induced by vascular endothelial growth factor (VEGF), OPN and certain integrins are able to promote angiogenesis through enhanced endothelial cell migration, proliferation and the subsequent formation of capillaries, which are all essential requirements for the process of angiogenesis (30,31).

2. Expression of OPN in human cancer

OPN has been shown to correlate with tumorigenesis, as well as with the progression and metastasis of different malignancies in both experimental and clinical studies (Table I). The upregulation

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Table I. Summary of OPN functions and clinical significance in human cancers.

| Cancer type | Biological functions | Clinical significance | Refs. |
|------------------|--|--|-------|
| NSCLC | There was a significant correlation between OPN and VEGF expression in NSCLC patients. OPN(+)/VEGF(+) are identified as independent prognostic factors of worse overall survival. OPN and other bone remodelling markers, such as OPG demonstrated independent positive correlations between clinical and tumour parameters. | OPN and VEGF could serve as prognostic factors and treatment targets for NSCLC. | (32) |
| Prostate cancer | Long-term treatment of cytotoxic drug could further upregulate OPN secretion from tumour cells. <i>In vivo</i> animal studies showed that knockdown of OPN enhanced the cytotoxicity of a chemotherapeutic drug. | Serum OPN combined with other markers of bone turnover may be able to determine the time-to-tumour progression, metastatic potential and overall survival of NSCLC patients. | (33) |
| LABC | Patients with above median baseline OPN levels were significantly more likely to die of their disease than those with below median baseline OPN levels, and overall baseline OPN level was significantly associated with survival. | OPN could be a potential drug target for reducing drug resistance in prostate cancer therapy. | (34) |
| Glioblastoma | OPN was an important player in dedifferentiation of neural cells during tumour formation. | Baseline plasma OPN level was a prognostic biomarker in the group of LABC patients, and could also be helpful in identifying LABC patients who will respond to neoadjuvant chemotherapy. | (35) |
| Malignant glioma | High OPN plasma levels were shown to be associated with a more aggressive phenotype, integrating known factors such as grade, tumour volume and extent of necrosis after radiotherapy. | OPN can be a therapeutic target for glioblastoma | (36) |
| SCHNC | OPN expression was associated with an increase in local recurrence in patients who were treated with primary radiotherapy for locally advanced SCHN. | High OPN plasma levels at the end of radiotherapy are associated with poor survival. | (37) |
| Gastric cancer | Strong OPN expression was significantly associated with a low apoptotic index, a high proliferative index, depth of invasion, lymphatic invasion, and venous invasion. OPN mRNA was upregulated in 83% of the tumours; OPN positivity was significantly associated with a shorter survival time | Tirapazamine and/or gene therapy or small molecule inhibitors targeting OPN, should be considered to be included in the treatment during radiotherapy in SCHNC. | (38) |
| CRC | High post-operative OPN levels correlated with post-operative distant metastasis. OPN expression was higher in CRC patients who were resistant to oxaliplatin-involved chemotherapy treatment. | OPN may play an important role in the invasiveness and the progressive nature of gastric cancer. | (39) |
| | | OPN positivity may be useful for predicting poor prognosis in gastric cancer patients. | (40) |
| | | Post-operative plasma OPN level is a potential non-invasive biomarker for monitoring CRC patients after curative resection of their primary tumour. | (41) |
| | | OPN inhibition is a potential therapeutic approach to combat CRC progression and chemoresistance. | (42) |

Table I. Continued.

| Cancer type | Biological functions | Clinical significance | Refs. |
|----------------|--|--|-------|
| PDAC | Serum levels of both OPN and TIMP-1 were markedly upregulated in PDAC; high serum levels of OPN were significantly correlated with reduced patient survival. Strong OPN mRNA signal was found in tumour-infiltrating macrophages; serum OPN levels were significantly elevated in PDAC patients with moderate sensitivity and specificity. | Combining utilisation of OPN, TIMP-1 and CA 19-9 in a panel could improve diagnostic accuracy in PDAC. | (43) |
| HCC | Secreted OPN induced autophagy via binding with its receptor integrin $\alpha\beta3$ and sustaining FoxO3a stability, which further promoted stem-like phenotype of HCCs, chemoresistance and tumour growth. Serum OPN levels were significantly elevated in patients with early and advanced HCC; there was no correlation between serum OPN and AFP levels. Plasma OPN levels were significantly elevated in patients with HCC and directly correlated with the tumour number; there was significant correlation between OPN and AFP levels. | Serum OPN may have utility as a diagnostic marker in patients with PDAC. | (44) |
| Bladder cancer | RNAi-targeting OPN inhibited the proliferation, invasion and tumorigenicity of T24 bladder cancer cells <i>in vitro</i> . | Blockade OPN and its receptors by using specific antibodies or small-molecule inhibitors combined with autophagy inhibitors might provide effective avenues for the treatment of HCC. | (45) |
| Ovarian cancer | The diagnostic performance of OPN assessment is similar to that of subjective ultrasonographic, morphological assessment for the differential diagnosis of ovarian tumours. Pre-operative plasma OPN levels were significantly higher in patients with ovarian cancer; the sensitivity and specificity of pre-operative plasma OPN in detecting ovarian cancer almost reached that of CA125. | Serum OPN is a useful diagnostic and prognostic marker for HCC. The combined use of serum OPN and AFP improved the diagnosis of early HCC. Plasma OPN levels appear to be an additional biomarker for HCC detection. | (46) |
| OSCC | The expression of OPN was elevated in 95% of tumours and 55% of histologically tumour free margin samples. | OPN may serve as a potential therapeutic target for human bladder cancer. | (47) |
| Melanoma | OPN mRNA expression was significantly increased in thicker melanomas, and high OPN mRNA and protein expression were associated with tumour progression and metastasis formation. | OPN may be useful in differential diagnosis for less experienced ultrasonographers and is especially valuable for differential diagnosis of endometriotic cysts. Pre-operative OPN is a useful biomarker for predicting ovarian cancer. It is especially useful when used alongside CA125. | (48) |
| | | OPN can be used as a diagnostic marker in OSCC. In the tumour free surgical margins, elevated levels of OPN may predict a significantly increased risk of recurrence. | (49) |
| | | OPN is a potential target for primary melanoma diagnosis and therapy. | (50) |
| | | | (51) |
| | | | (52) |

OPN, osteopontin; NSCLC, non-small cell lung cancer; VEGF, vascular endothelial growth factor; LABC, locally advanced breast cancer; SCHNC, squamous cell head and neck cancer; CRC, colorectal cancer; PDAC, pancreatic ductal adenocarcinoma; HCC, hepatocellular carcinoma; OSCC, oral squamous cell carcinoma.

of OPN expression has been identified in a variety of human cancers, including breast, prostate, lung, stomach, pancreatic and colorectal cancer, glioma and melanoma (Table I).

In these studies, *in vitro* experiments have also established a causal link between OPN expression and cell functions. *In vitro* transfection experiments have demonstrated that downregulating OPN gene expression suppressed the progression of breast cancer cells, which have invasive potential and metastatic competence (21). Experimental evidence has further suggested that enhanced OPN expression renders a highly metastatic phenotype of cancer cells. For example, the enforced expression of OPN in non-metastatic rat mammary tumour cells has been shown to result in lung metastasis occurring in half of the animals that developed primary tumours (23). In another study, in glioma cells, elevated levels of OPN and VEGF synergistically enhanced the angiogenic properties, via integrin $\alpha\beta3$, on the endothelial cell surface (25). Another study demonstrated that the knockdown of OPN in prostate cancer cell lines enhanced the cytotoxicity of the chemotherapeutic drug, daunomycin (DUN), through integrin-mediated FAK/P-GP signalling (34). The ectopic expression of OPN in DLD1 colon cancer cells has been shown to stimulate EMT activation and subsequent migration (41).

In clinical investigations, elevated OPN levels have been shown to correlate with increased tumour burden (stage, grade and tumour size), a poor prognosis and reduced survival, although discrepancies remain. For example, the increased expression of OPN in plasma and tumour tissues has been identified in breast cancer patients and has been shown to be associated with metastatic disease and decreased survival (35). Similarly, in advanced gastric cancer, patients with OPN-positive cancer (as determined by immunohistochemistry) have a decrease in their 5-year survival, when compared with those with OPN-negative cancer (40). In colorectal cancer (CRC) patients, increased OPN mRNA levels have been shown to significantly correlate with stage, lymph node metastasis and lymphatic or venous invasion, as well as with lower disease-free and overall survival rates (41,42). A study on patients with pancreatic ductal adenocarcinoma (PDAC) demonstrated that serum levels of OPN are elevated with advanced tumour grades (44). Elevated OPN levels have also been strongly associated with increased stage, grade and tumour size in melanoma patients (52). In early stage non-small cell lung cancer (NSCLC) patients high OPN plasma levels were observed which were reduced after surgical tumour resection. However, in those patients which showed recurrence after surgery OPN plasma levels re-elevated, indicating that OPN is not only a potential diagnostic marker in NSCLC, but also has potential as a tool for detecting tumour recurrence after treatment (53). Thus, OPN may be a useful biomarker to monitor cancer progression and a significant predictor of poor prognosis and survival rates.

3. Expression of OPN splice variants in human cancers

The alternative splice-generation of multiple mRNA products from a single gene is a critical mechanism for generating proteomic diversity. The OPN precursor-mRNA (pre-mRNA) is subject to alternative splicing, which generates three splice variants, OPN-a (consists of all exons), OPN-b (which lacks exon 5) and OPN-c (which lacks exon 4) (Fig. 1).

Recent studies have demonstrated that the expression of OPN splice variants in malignancies is cell-type/tissue specific and may have functional heterogeneity (Table II). For example, Pang *et al* reported that OPN-c is selectively expressed in breast cancer tissue (57). In their study on 170 breast cancer patients, OPN-c protein staining was positive in 70% of all of the samples and was significantly higher in tumour tissues compared to normal tissues. The study indicated an inverse correlation between OPN-c and E-cadherin, an established tumour suppressor. However, the observations by the same authors that OPN-c also inversely correlated with β -catenin and that E-cadherin positively correlated β -catenin are intriguing. Despite β -catenin working with E-cadherin to form cell-cell adhesion complexes, β -catenin has been classically regarded as an oncogenic protein. Thus, the true link between OPN-c and the cell adhesion complex requires further study and one has to delineate the cellular location of β -catenin (namely membrane-associated, cytoplasmic and nucleus) in this context. However, the study by Pang has clearly demonstrated that high levels of OPN-c staining in breast tumours are associated with TNM staging, nodal involvement, recurrence and metastasis, and interestingly with the triple negativity of the tumours and, thus is an independent prognostic indicator for these patients (57). Another study by Sun *et al* demonstrated that each of the three OPN splice variants was able to increase the growth of breast tumours, *in vivo* (56). It is noteworthy that OPN-a appears more effective in promoting tumour growth than the other two splice forms.

Conflicting evidence also exists regarding the function of OPN isoforms in hepatocellular carcinoma (HCC). It has been previously demonstrated that tumour tissue predominantly expressed OPN-a and OPN-b, and the ratio of OPN-a and OPN-b to OPN-c increased substantially as the tumours progressed in SK-Hep1 cells and Hep3B cells (67). Thus, it is possible that OPN-a and OPN-b may be associated with a poor prognosis, and OPN-c may prevent both cell migration and invasion in more migratory and invasive cells. By contrast, in another study, Takafuji *et al* (68) reported that the increased expression of OPN-c in Hep3B cells appeared to enhance cellular invasion and metastatic potential due to the formation of OPN-c fragment by MMP-9.

A number of studies have also explored OPN splice variants in lung cancer. Sun *et al* (59) found that, compared with normal lung tissue, the majority of NSCLC samples predominantly expressed OPN-a. A similar observation was made by Blasberg *et al* (61), who reported that OPN-a was the dominant isoform, whereas little OPN-b and no OPN-c expression was detected in lung cancer cell lines that endogenously expressed OPN (A549, H460, H157, H1299 and Calu-3). Functionally, OPN-a promotes angiogenesis in lung cancer by stimulating endothelial cells, likely by binding to the $\alpha\beta3$ integrin and increasing VEGF expression and secretion. In their study, OPN-b appeared to increase tubule formation merely by activating endothelial cells, and did so without a concomitant enhancement of VEGF secretion or expression. OPN-c had a predominantly negative effect significantly inhibiting angiogenesis and VEGF secretion. However, Zhao *et al* (60) suggested that OPN-b strongly affected cell proliferation, while OPN-c was closely related to the invasive behaviour of the NSCLC cell line, A549.

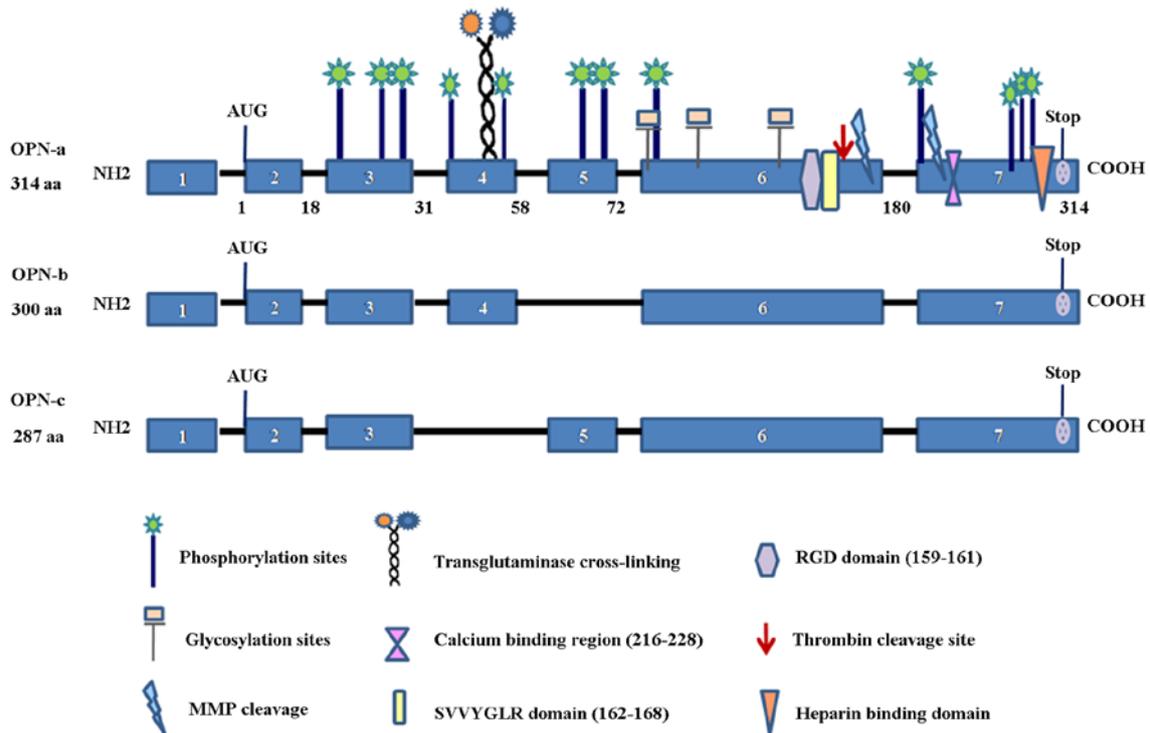


Figure 1. Structural features of the three osteopontin (OPN) splice variants. OPN-a is the full-length isoform with 6 translated exons (314 aa), OPN-b lacks exon 5 (300 aa), and OPN-c lacks exon 4 (287 aa). All OPN variants possess functional domains: arginine-glycine-aspartic acid (RGD) domain (159-161), serine-valine-valine-tyrosine-glutamate-leucine-arginine (SVVYGLR) domain (162-168), thrombin cleavage site, calcium binding domain (216-228) and heparin binding domain. Integrin binding occurs at RGD and SVVYGLR sequences, whilst the CD44 variant receptor binding occurs near the C-terminus. Phosphorylation sites are throughout the whole precursor-mRNA (pre-mRNA). Thrombin cleavage and matrix metalloproteinase (MMP) cleavage sites are present in all three splice variants. Compared with the full-length isoform (OPN-a), OPN-b misses certain phosphorylation sites, while OPN-c is unable to undergo transglutaminase cross-linking. To some extent, the structural differences that exist between the three OPN isoforms may contribute to their clinical and functional differences.

These studies emphasise that OPN splice variants may have diverse expression patterns and different functional roles, which may be cancer type-specific.

4. Polymorphism of OPN and OPN splice variants in cancers

Secreted OPN (sOPN) and intracellular OPN (iOPN). OPN has mainly been studied as a secreted protein, but recent studies have shown that, in some cases, OPN is not secreted and instead will be found in the cytoplasm and nucleus. Mouse OPN mRNA has the canonical AUG translation initiation site and an alternative translation initiation site. When translation initiates from the canonical site, the peptide produced is targeted to secretory vesicles. On the other hand, when translation starts from the alternative translation initiation site, the peptide produced is accompanied by deletion of a signal sequence and localises in the cytoplasm (76,77). Although sOPN and iOPN are generated from the same OPN mRNA species, the biological outcomes mediated by the two isoforms may differ (78). The role of human sOPN in cell physiology has been extensively studied in different cellular contexts. Compared with sOPN, the biological roles of iOPN have only begun to emerge over the past several years. Thus far, iOPN has been found in calvarial cells, dendritic cells, macrophages and nerve cells (77,79,80). It is involved in cell motility, cytoskeletal rearrangement and mitosis by physical association with polo-like kinase-1 (Plk-1) (81-83). Similarly, these two isoforms

may play distinct roles in cancer progression. Indeed, in patients with various solid tumours, sOPN has been proposed as a diagnostic marker to distinguish either resectable cases or predict survival rates (43,84,85). However, cytoplasmic OPN expression has appeared not to correlate with average tumour size, tumour stage and nodal status (10,86). The same has been observed for OPN splice variants. For example, in a study on expression patterns of OPN variants and its functions on cell apoptosis and invasion in glioma cells, Yan *et al* presumed that the secretory nature of OPN splice variants may be induced by the absence of exon 5 or exon 4. As a consequence, OPN-b without out exon 5 may aggregate within the cytoplasm and exert a significant anti-apoptotic effect, while OPN-c without exon 4 may be easily secreted to culture supernatants and has a remarkable effect on cellular invasion (74).

Therefore, it would be necessary to delineate which isoform of OPN is responsible for pathophysiological events, and furthermore, sOPN and iOPN should be independently targeted in any potential therapies.

Host-derived OPN and tumour-derived OPN. Tumour cells and their microenvironment mutually influence tumour formation, progression and metastasis. The tumour microenvironment includes a variety of non-tumour cell types, such as fibroblasts, immune cells, vascular and smooth muscle cells. The host's reaction to neoplastic cells and the ability of environmental modification by tumour cells themselves are both involved in

Table II. Summary of OPN splice variants (OPN-a, OPN-b and OPN-c) expression and function in human cancer tissues and cells.

| Cancer type | Sample size | Methods | Expression pattern of OPN splice variants | Biological function | Refs. |
|-----------------|-------------|--|---|--|-------|
| Breast cancer | 671 | IHC | Most specimens displayed, OPN-a and OPN-b, selectively in the cytoplasm, while OPN-c predominantly in the nuclei | High staining intensity of nuclear OPN-c was strongly associated with mortality in patients with early breast cancer. Cytosolic staining of OPN-a and OPN-b also predicted poor outcome | (54) |
| | 309 | IHC, qPCR | OPN isoforms expression pattern not analysed | Increased OPN-c took a higher risk of recurrence among immunophenotypes, especially in triple-negative/basal-like subtype | (55) |
| | - | RT-PCR | OPN isoforms expression pattern not analysed | Overexpression of OPN transcripts promoted local tumour formation, but there was no significant difference among OPN-a, OPN-b and OPN-c when tumour formation <i>in vivo</i> | (56) |
| Lung cancer | 170 | IHC | OPN-c is positive expression in breast cancer tissue, OPN-a and OPN-b not detected | OPN-c could serve as a prognostic factor of breast cancer | (57) |
| | 127 | qPCR | Three OPN isoforms were upregulated in breast cancer specimens, especially OPN-b and OPN-c | Increased OPN-b and OPN-c expression were significantly associated with adverse pathological and clinical outcomes in breast cancer, especially OPN-c; OPN-a had the inverse effect | (58) |
| | - | RT-PCR | Among the three OPN isoforms, OPN-a is the most highly expressed in lung cancer cell lines | OPN-a can inhibit growth of cells with high integrin $\beta 3$ levels and increase growth via activation of the CD44/NF- κ B pathway in cells with low integrin $\beta 3$ levels. Functional studies of OPN-b and OPN-c not performed | (59) |
| Prostate cancer | - | - | OPN isoforms expression pattern not analysed | OPN-b affected cell proliferation and OPN-c played a role in invasive behavior | (60) |
| | - | RT-PCR | OPN-a and OPN-b were dominantly expressed in OPN secreting cell lines (OPN-a > OPN-b). OPN-c was not endogenously expressed in these cell lines | OPN-a was significantly associated with tubule formation; OPN-b also significantly increased tubule length, but to a lesser extent than OPN-a; OPN-c resulted in a significant decrease in tubule formation | (61) |
| | - | qPCR | OPN isoforms expression pattern not analysed | Compared to OPN-a, PC3 cells overexpressing OPN-b and OPN-c are more resistant to DXT-induced cell death | (62) |
| | - | qPCR | OPN isoforms expression pattern not analysed | OPN-b and OPN-c, but not the OPN-a, exerted pro-tumourigenic roles in prostate cancer cells | (63) |
| - | RT-PCR | OPN isoforms expression pattern not analysed | OPN-c activated capabilities of pro-oncogenicity in OvCar-3 and PC-3 cancer cell lines | (64) | |

Table II. Continued.

| Cancer type | Sample size | Methods | Expression pattern of OPN splice variants | Biological function | Refs. |
|-------------------|-------------|--------------|---|---|-------|
| Esophageal cancer | - | qPCR | All OPN isoforms were highly overexpressed in primary adenocarcinoma EACs | OPN-b significantly increased cell adhesion, OPN-c enhanced cell detachment | (65) |
| | 63 | IHC, qPCR | Expression of OPN-c is significantly elevated in squamous cell carcinoma ESCC tissue, OPN-a and OPN-b not detected | OPN-c was closely related to invasion and stage of cancer, its upregulation could be a potential diagnostic marker | (66) |
| HCC | - | RT-PCR | Tumour tissues and migratory cell lines mainly expressed both OPN-a and OPN-b; while normal liver tissues and non-migratory cell lines mainly expressed OPN-c. OPN isoforms expression pattern not analysed | OPN-a and OPN-b could induce migration of HCC cells, while OPN-c had no significant effects | (67) |
| Gastric cancer | 101 | qPCR | OPN-a is the major isoform in healthy and OPN-b was the dominant isoform in six GC cell lines; three OPN isoforms are all markedly increased in GC tissues | Increased expression of OPN-c was associated with clinical metastatic HCC | (68) |
| PDAC | 40 | RT-PCR | OPN-a and OPN-b were expressed in almost all PDAC samples. OPN-c was present in 80% of patients with metastatic disease | Elevated OPN-b or OPN-c expression could correlate with clinicopathological features; OPN-b exerts the strongest anti-apoptotic effect and OPN-c most effectively stimulated GC metastatic activity | (69) |
| | - | RT-PCR | OPN isoforms expression pattern not analysed | OPN-c was found to significantly correlate with presence of metastasis, and OPN-b with poor survival | (70) |
| MTC | 6 | qPCR | OPN-a expression levels are higher than OPN-b and OPN-c expression levels (OPN-a > OPN-b > OPN-c) both in tissues and cells | Overexpression of OPN-b and OPN-c promoted colony formation, cell migration, and enhance an inflammatory gene profile in PDAC cells | (71) |
| TC | 109 | qPCR | Compared to OPN-b and OPN-c, OPN-a is the prevalent variant in TC tissues and cell lines | TT cells overexpressing OPN-a, OPN-b and OPN-c significantly reduced growth when compared with control cells | (72) |
| Malignant glioma | - | RT-PCR | OPN-b is markedly upregulated in U251MG cells | OPN-a can potentially mediate invasive and metastatic potential in TC cell lines | (73) |
| Gliomas cancer | 45 | qPCR, RT-PCR | The mRNA levels of three OPN isoforms were markedly increased in gliomas tissues, and all OPN splice variants were also found in U251 and U87 cells | Therapeutic targeting of OPN must address all splice variants, in particular OPN-b | (37) |
| STS | 124 | qPCR | OPN-a and OPN-b are expressed at a distinctly higher level than OPN-c in STS tissues | Both OPN-a and OPN-c promoted glioma cell invasion (OPN-c > OPN-a), while OPN-b showed no effect on the invasion of U251 and U87 cells | (74) |
| | | | | The mRNA expression levels of OPN-b and OPN-c were significantly correlated with the clinical outcome of STS patients | (75) |

OPN, osteopontin; HCC, hepatocellular carcinoma; PDAC, pancreatic ductal adenocarcinoma; MTC, medullary thyroid carcinoma; TC, thyroid carcinoma; STS, soft tissue sarcoma.

tumour formation. A number of studies have demonstrated that OPN can be synthesised by tumour and other cells types in the tumour microenvironment, such as macrophages and stromal cells (87,88). However, whether tumour-derived OPN differs from host-derived OPN, either structurally or functionally, is largely unknown. There is evidence indicating that the role of OPN in metastasis is dependent on the site of production. For example, historically, stroma-derived OPN has been considered to be intrinsically involved in the defensive mechanism against tumour development by acting as a macrophage chemoattractant, yet stroma-derived OPN can effectively regulate melanoma growth, angiogenesis and metastasis with the host OPN-tumour interaction (89-91). In addition, tumour-derived OPN may promote the metastatic process by inhibiting macrophage cytotoxicity against tumours (92). By contrast, some evidence suggests that at least in some instances, the presence of OPN in macrophages promote the development and activity of type I natural killer (NK) T cells (NK T cells) and that OPN may inhibit tumour activity via these NK cells (93).

Thus, host-derived OPN and tumour-derived OPN may mutually affect each other and the interaction between the tumour and tumour microenvironment may determine whether the overall effect of OPN will be tumour promoting or tumour inhibiting.

OPN splice variants polymorphism in cancers. The similar heterogeneity of OPN splice variants exists in cancers. The differential effects of the three splice variants may be due to the distinctive functions of the spliced exons that individual OPN isoforms contain. For example, the sequences encoded by exon 5, absent in OPN-b, contain one of several clusters of phosphorylated serine/threonine residues. Exon 4 however encodes two glutamine residues essential for transglutaminase cross-linking. As exon 4 is absent in OPN-c, but present in OPN-a and OPN-b, OPN-c has no specific functions exerted by exon 4 (94). In breast cancer cells, it has been reported that OPN-a, not OPN-c may have a tendency to aggregate in response to physical concentrations of calcium, and thus have a greater capacity to enhance cell adhesion (95). By contrast, OPN-c may be more soluble and able to support anchorage-independent growth of breast cancer cells (96,97). OPN-c possesses a more exposed Arg-Ser-Lys (RSK) motif compared to OPN-a (98). Using the RSK motif as a cleavage site, systemically circulating thrombin can cleave OPN-c easily into two fragments, the N-terminal and C-terminal, of approximately equivalent size (98). The N-terminal GRGDS-containing fragment produced by thrombin cleavage has the potential to enhance tumour cell migration (98). The thrombin-cleaved C-terminal fragment of OPN has also been reported to influence breast cancer cell migration and invasion *in vitro* (99). The absence of the transglutaminase acting site in OPN-c may explain why OPN-c cannot be cross-linked with the extracellular matrix (94).

On the other hand, some investigations suggest that the possible reasons for OPN isoforms cell-type specific patterns and their roles are closely related to these differing PTMs of the three OPN splice variants. Gimba *et al* demonstrated that the potential of OPN-a, OPN-b and OPN-c for specific phosphorylation patterns, and the deletion of exon 4 or 5 altered the pattern of PTMs, ultimately resulting in functional modifications (46).

5. Conclusion and future perspectives

OPN was initially identified in bone and later characterised based on its splice variants and their structures and functions. It is now well established that the isoforms of OPN are differentially expressed in many tumour types and play critical roles at different stages of cancer development and progression. OPN may be a useful biomarker to monitor cancer progression and one of the significant predictors of poor prognosis and survival rates, in certain cancers for example in breast cancer and lung cancer. Thus, OPN may have several potential clinical implications: firstly, as a convenient tool for clinical test. As a secreted protein, OPN can easily detected in body fluids, such as blood, urine and body cavity fluids (namely pleural and peritoneal ascites). This gives rise to convenient sampling, taking advantages of sample quantity, safe detection, procedure speed and minimal invasiveness, collectively providing a convenient approach for clinical testing. Secondly, the power of combining OPN with other traditional biomarkers, such as VEGF, MMPs and E-cadherin, for example, provide more accurate predictions for prognosis and potential response to therapies (43,50,100).

Yet, changes remain, especially given the complexity of the distribution patterns and presence of the variant forms for OPN. Methods of OPN detection are currently very limited. Traditional protein detection methods such as ELISA have shown their limit due to a lack of sensitivity and the nature of samples required for the testings. Several studies have focused on the cost-effectiveness and ease of implementing immunosensors for OPN detection demonstrating better sensitivity than ELISA assays, and providing greater potential to develop simple test kits for OPN combined with other protein biomarkers (101-104).

Due to the pre- and post-translational regulation, the expression and function patterns of OPN isoforms usually exhibit tissue specificity. It has been suggested that this provides the possibility to develop cancer therapy strategies to target those OPN isoforms which are specific to tumour cells or play a key role in tumour progression. But the pre- and post-translational regulation are complex and ubiquitous, developing drugs that target only cancer cells with minimal impact on healthy cells is extremely difficult. Thus the expression patterns and activities of tumour-specific OPN isoforms should be further defined with multidisciplinary and large scale clinical study. The development of specific OPN-based approaches for individual cancer types is equally important as well.

Collectively, OPN and its variants have been shown to play an important role in regulating the aggressive nature of cancer cells and promote the growth of tumours. Although there is more to learn with regards to the mechanisms of action of the specific OPN variants, it is clear that there is clinical prospect(s) for this protein and its variants. OPN appears to have good clinical value in evaluating disease progression and cancer patient outcome. There is a good prospect for developing a OPN based clinical test for cancer patients in order to evaluate their prognosis and response to therapies. What is most exciting is the prospect of developing tools to target the protein and its variants in novel ways. It is anticipated that in the coming years we will see some significant progress along these fronts.

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