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The Wnt signalling pathway: a tailored target in cancer

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Abstract: Cancer is one of the greatest public health challenges. According to the World Health Organization (WHO) 9.6 million cancer deaths have been reported in 2018. The most common cancers include lung, breast, colorectal, prostate, skin cancer (non-melanoma) and stomach. The unbalance of physiological signalling pathways due to the acquisition of mutations in tumour cells is considered the most common cancer driver. The Wingless-related integration site (Wnt)/β-catenin pathway is crucial for tissue development and homeostasis in all animal species and its dysregulation is one of the most relevant events linked to cancer development and dissemination.

The canonical and the non-canonical Wnt/β-catenin pathways are known to control both physiological and pathological processes including cancer. Herein the impact of the Wnt/β-catenin cascade in driving cancers from different origin has been examined. Finally, based on the impact of Extracellular Vesicles (EVs) on tumour growth, invasion and chemoresistance, and their role as tumour diagnostic and prognostic tools, an overview of the current knowledge linking EVs to the Wnt/β-catenin pathway is also discussed.

Keywords: Wnt/β-catenin dependent pathway; Wnt/β-catenin independent pathway; colorectal cancer; breast cancer; ovarian cancer; extracellular vesicles

Introduction

The human wingless-related integration site (Wnt) genes encode 19 evolutionarily conserved glycoproteins with 22-24 Cys residues. In the endoplasmic reticulum (ER), the Wnt ligands are post-translationally acetylated by porcupine, a membrane associated O-acyl transferase. Acetylation leads to palmitoylation which is required for the release and binding of Wnt to the frizzled (FZD) receptors. This, on turn, drives the biological response[1].

The Wnt signalling pathway regulates crucial cellular processes including cell fate determination, organogenesis during embryonic development, normal adult homeostasis, motility, polarity and stem cell renewal[2]. Moreover, its contribute in cancer has been extensively investigated[3].

The Wnt pathway has been widely studied and reviewed, and a general understanding of the transduction cascade has been clarified. The Wnt cascade has been subdivided into different branches due to its complexity[4,5]. They include the canonical Wnt/β-catenin (Wnt/β-catenin dependent pathway) and the non-canonical Wnt/β-catenin pathway (β-catenin-independent pathway). The latter was further allocated into two additional branches, the planar cell polarity (PCP) and the Wnt/calcium pathways[2]. Both of them contribute to cancer development and dissemination.

The aim of the present review is to provide an overview of the current knowledge about the Wnt signalling pathway in tumour development and progression. Tumours from different origin are discussed. Although the canonical and the non-canonical Wnt/β-catenin pathway work together to control physiological and pathological processes[2], data related to each one are independently
deemed. Finally, the contribution of extracellular vesicles (EVs) in triggering the Wnt/β-catenin cascade is also analyzed.

**Wnt canonical pathway: β-catenin dependent**

The canonical pathway turns around the β-catenin intracellular level (Figure 1). In the absence of Wnt proteins the β-catenin “destruction complex” keeps low β-catenin in the cell. The “destruction complex” mainly consists of two kinases: casein kinase 1α (CK1α), glycogen synthase kinase 3 β (GSK-3β) and two scaffolds: axis inhibition (Axin), and adenomatous polyposis coli (APC). Firstly, β-catenin undergoes phosphorylation by CK1α at serine 45 (Ser45), Ser33, Ser37 and threonine 41 (Thr41) by GSK-3β. Then, the E3 ubiquitin ligase, denoted as β-transducin repeat–containing protein (βTrCP), marks β-catenin ubiquitination and degradation [1]. This prevents β-catenin nuclear translocation while allows histone deacetylation and chromatin compaction by the Groucho repressor, translating into the inhibition of gene transcription[6] (Figure 1a).

![Figure 1. The Canonical Wnt signalling pathway. (a) OFF STATE. In the absence of Wnt ligands β-catenin moves to the “destruction complex” consisting of casein kinase 1α (CK1α), glycogen synthase kinase 3 β (GSK-3β) and two scaffolds: axis inhibition (Axin), and adenomatous polyposis coli (APC). β-catenin undergoes phosphorylation at Ser45 residue by CK1α and at Ser33, Ser37 and Thr41 residues by GSK-3β. Then, the E3 ubiquitin ligase β-transducin repeat–containing protein (βTrCP), marks β-catenin ubiquitination and proteasomal degradation. This prevents β-catenin nuclear accumulation while allows chromatin compaction and Groucho-mediated promoter repression. (b) ON STATE. The Wnt ligands bind to frizzled (FZD) receptor and the low-density-lipoprotein-related protein 5/6 (LRP5/LRP6), this results in dishevelled (DVL) phosphorylation and β-catenin release from the “destruction complex”, allowing β-catenin accumulation and nuclear translocation. In the nucleus, the Groucho repressor undergoes displacement allowing β-catenin to interact with T-cell factor/lymphoid enhancer factor (TCF/LEF), chromatin remodeling and transcription of genes such as c-myc and cyclin D1.](image-url)
Once in the nucleus, the engagement of β-catenin transiently converts the TCF/LEF into transcriptional activators which displace Groucho and induces chromatin remodelling and transcriptional activity (Figure 1b).

A number of genes are targeted by Wnt-β-catenin. Among them, genes involved in positive- and negative-feedback regulation, cell-cycle progression, and stem cell homeostasis are the most commonly included genes.

**Wnt non-canonical pathways: Wnt/planar cell polarity (PCP) and Wnt/Ca²⁺**

To date, the canonical Wnt/β-catenin pathway is much better characterized than the non-canonical one (Figure 2).

![Figure 2. The Wnt non-canonical signalling pathways.](image)

In the non-canonical PCP pathway, Wnt ligands bind to FZD receptors and co-receptor RAR-related orphan receptor (ROR) or the receptor like tyrosine kinase 7 (PTK7), RAR-related orphan receptor (ROR) or the receptor like tyrosine kinase (Ryk) and convey the signal to DVL. On the one side, DVL forms the disheveled associated activator of morphogenesis 1 (DVL-Daam-1) complex, which triggers RhoA, Rho and ROCK to control cytoskeletal rearrangement. On the other hand, DVL triggers RAC, JNK and AP-1 involved in cell motility and polarity.

In the calcium-dependent pathway Wnt ligands bind to FZD and activate the phospholipase C (PLC) which hydrolyses the phosphatidylinositol (4,5)-biphosphates (PI(2)) to inositol (1,4,5)-triphosphates (IP3) and diacylglycerol (DAG). This translates into intracellular calcium release and the activation of CaN and CamKII. The calmodulin activation stimulates TAK-1 and NLK activity. CaN activates the NFAT, which moves to the nucleus and modulates the expression of genes involved in the control of gastrulation, ventral cell fate and tissue homeostasis.

In the PCP pathway, Wnt ligands bind to FZD receptors and co-receptor protein tyrosine kinase 7 (PTK7), RAR-related orphan receptor (ROR) or the receptor like tyrosine kinase (Ryk) and convey the signal to DVL. On the one side, DVL forms the disheveled associated activator of morphogenesis 1 (DVL-Daam-1) complex, which triggers a small guanosine-5'-triphosphate (GTP) GTase, such as ras homolog gene family member A (RhoA), RHO and RHO-associated kinase (ROCK). DVL also triggers ras-related C3 botulinum toxin substrate (RAC), JUN-N-terminal kinase (JNK) and the activator protein-1 (AP-1).[7] The PCP pathway is involved in the cytoskeletal rearrangement, cell motility and co-ordinates cell polarity. In vertebrates, the PCP pathway is also required for morphology and migration of dorsal mesodermal cells undergoing gastrulation, hair follicle organization, and orientation of stereocilia in the sensory epithelium of the inner ear [8] (Figure 2a).

In the calcium-dependent pathway Wnt ligands bind to FZD and activate the phospholipase C (PLC) which hydrolyses the phosphatidylinositol (4,5)-biphosphates (PI(2)) to inositol (1,4,5)-triphosphates (IP3) and diacylglycerol (DAG). This translates into the release of the intracellular calcium and the activation of both calcineurin (CaN) and calcium/calmodulin-dependent kinase II (CamKII). Moreover, the activation of calmodulin promotes the activation of the TGF-β-Activated kinase 1 (TAK-1) and nemo-like kinase (NLK), thereby antagonizing and neutralizing the canonical Wnt/β-catenin cascade. CaN activates the nuclear factor of activated T-cells (NFAT), which moves to
the nucleus and regulates the expression of target genes [7] (Figure 2b). The calcium-dependent pathway plays a crucial role in several processes, including early pattern formation during gastrulation [2], ventral cell fate [9], dorsal axis formation [10], and tissue homeostasis [11].

120 COLORECTAL CANCER

121 Colorectal cancer (CRC) is one of most common cancer worldwide and represents a deep cause of cancer mortality [12] with a rapid increase in incidence and death rate [13]. Dienemann et al. [14] established a new classification of CRCs into four consensus molecular subtypes (CMSs). Among them CMS2, CMS3, and CMS4 have a higher rate of APC mutations (over 50%) compared to CMS1. Each CMS has unique features: CMS1 (MSI Immune, 14%): hyper- mutated, microsatellite instability, strong immune activation; CMS2 (Canonical, 37%): epithelial, chromosomally unstable, marked Wnt and myc signalling activation; CMS3 (Metabolic, 13%): epithelial, metabolic dysregulation; and CMS4 (Mesenchymal, 23%): a prominent transforming growth factor β (TGFβ) activation, stromal invasion, and angiogenesis. Samples with combined features (13%) represent transition phenotypes or are supposed to reflect the intra-tumour heterogeneity [14].

121 The heterogeneous genetic ground underlying CRC initiation and progression mainly involves gene fusion, deletion or amplification, somatic gene mutations and epigenetic alterations. Wnt/β-catenin signalling has emerged as one of the most significant biological pathways in both physiological setting and CRC development. Almost all CRC are characterized by a hyper-active Wnt/β-catenin pathway, which, in many cases, is considered the most critical cancer initiating and driving event. Proteins and miRNAs guiding the Wnt/β-catenin pathway and proposed as potential CRC therapeutic targets are discussed.

128 Canonical Wnt/β-catenin pathway and CRC

129 Ring finger protein 6 (RNF6) is an oncogene frequently upregulated by gene amplification in primary CRC. Moreover, APC mutation and RNF6 copy number amplification were commonly found in CRC patients. RNF6 is a RING-domain E3 ubiquitin ligase and exerts its pro-metastatic effects by promoting CRC cell growth, cell-cycle progression, and epithelial to mesenchymal transition (EMT).

130 Furthermore, RNF6 expression and its gene amplification have been considered independent patients’ prognostic factors. RNF6 mediates the polyubiquitination of the transducin-like enhancer of split 3 (TLE3), a transcriptional repressor of the β-catenin/TCF4 complex, and its proteasome degradation. The lack of TLE3/TCF4/LEF interaction enhances the Wnt/β-catenin transcriptional activity, and the expression of its downstream target genes [15] (Table 1).

148 Table 1. Proteins/EVs involved in several tumours, their alteration, targets, and impact on tumours.

<table>
<thead>
<tr>
<th>PROTEIN</th>
<th>RELATED CANCERS</th>
<th>EXPRESSION LEVEL</th>
<th>PATHWAY INTERACTION</th>
<th>IMPACT ON TUMOUR</th>
<th>REF.</th>
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<td>progression</td>
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<td>EMT</td>
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<td>Non-canonical</td>
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<td>β-catenin</td>
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<td>BC</td>
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<td>β-catenin</td>
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<td>ALCAM</td>
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<td>p62/SQSTM1 (PCP)</td>
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<tr>
<td>HePTP</td>
<td>TNBC</td>
<td>Upregulated</td>
<td>-GSK-3β</td>
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**tumour progression**

- invasion
- proliferation
- chemoresistance
- metastasis
- stemness
- migration
- invasion
- progression
- proliferation
- metastasis
- CSC maintenance
- invasion
- proliferation
- cell cycle progression
- colony formation
- migration
- migration
- lactate production
- cell proliferation
- aggressiveness
- vessel invasion
- tumour size
- chemoresistance
- migration
- anchorage-dependent and independent cell proliferation
- metastasis
<table>
<thead>
<tr>
<th>Tumour Type</th>
<th>Cell Line</th>
<th>Regulation</th>
<th>β-catenin</th>
<th>Functional Implications</th>
<th>Reference(s)</th>
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<td>DLCA-3</td>
<td>TNBC</td>
<td>Downregulated</td>
<td>β-catenin</td>
<td>proliferation, colony formation, migration, invasion</td>
<td>[92],[94]</td>
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<td>DKK1</td>
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<td>[100],[102]</td>
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<td>SFRP1</td>
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<td>β-catenin</td>
<td>cell growth, stem-like phenotype</td>
<td>[104],[103]</td>
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<tr>
<td>AXIN2</td>
<td>OC</td>
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<td>[144],[103]</td>
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<td>LGR6</td>
<td>OC</td>
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<td>stemness, chemoresistance</td>
<td>[105,106],[107,108]</td>
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<td>[97,107],[109,110]</td>
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<td>FZD7</td>
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<td>[132,133]</td>
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The leucine-rich repeat-containing G-protein coupled receptor 5 (LGR5) is a Wnt/β-catenin target gene implicated in cancer cell proliferation and migration. It has been reported that LGR5 is highly expressed in CRC tissues compared to the healthy ones. A decline in β-catenin and c-myc mRNA expression were detected by knocking-down LGR5 expression, suggesting that it may regulate the Wnt/β-catenin activity by modulating the expression of β-catenin. Furthermore, since targeting LGR5 improves the response to chemotherapy, LGR5 has been proposed as a novel therapeutic target in CRC [16] (Table 1).

Table 1. miRNAs involved in the tumours, their alteration and tumour impact.

<table>
<thead>
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<td>Regulation</td>
<td>Function(s)</td>
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</table>

Sun and co-workers [25] identified miR-144-3p as a new biomarker for CRC diagnosis and response to treatment. miR-144-3p was found downregulated and associated with CRC pathological stages in CRC patients. Interestingly, miR-144-3p overexpression reduced CRC cell proliferation by delaying G1/S phase transition in tumour cells. On the contrary, the B-cell lymphoma 6 protein (BCL6), a nuclear protein belonging to the BTB/POZ/zinc finger (ZF) family of transcription factors, was found upregulated and surprisingly post-transcriptionally regulated by miR-144-3p. Previous studies revealed that BCL6 is involved in the control of cell cycle progression and differentiation [26,27]. Indeed, miR-144-3p/BCL6 co-operate to inhibit cellular proliferation, development, and progression of CRC by interfering with c-myc and cyclin D1 expression [25] (Table 1).

miR-377-3p displays an ambiguous role in CRC. Liu and colleagues [28] uncovered that upregulation of miR-377-3p promotes G1-S phase transition, cell expansion and EMT, while represses apoptosis in CRC patients. Moreover, GSK-3β, a direct miR-377-3p target, was found upregulated upon miR-377-3p overexpression. These data suggest that a complex regulatory network boosting tumour progression is associated with the expression of miR-377-3p in CRC.

Conversely, in a recent study, Huang et al. [29] have shown that miR-377-3p, significantly reduced in CRC patients, is involved in the control of proliferation, migration and chemo resistance, particularly at advanced tumour stage. The authors investigated miR-377 functions and mechanism of action in CRC cells. The zinc finger E-box binding homeobox 2 (ZEB2) and the X-linked inhibitor of apoptosis protein (XIAP) are two positive regulators of the Wnt/β-catenin cascade [30,31]. In CRC, ZEB2 enables tumour progression and invasion, whereas XIAP promotes cell proliferation and chemoresistance. De facto, miR-377-3p overexpression was found to suppress the malignant CRC phenotype, as well as cell proliferation, invasion and drug resistance by directly targeting the 3' UTR sequence of both ZEB2 and XIAP mRNAs. Since miR-377-3p/ZEB2-XIAP inhibited CRC progression by reducing Wnt/β-catenin-associated gene expression (e.g. cyclin D1, Axin2, TCF1, SOX2, c-myc, matrix metalloproteinase-2 (MMP-2), MMP-9, CD44, vascular endothelial growth factor (VEGF), and Twist) approaches increasing its expression have been proposed for novel therapeutic options (Table 1).

Functional experiments showed that miR-520e plays a pivotal role in regulating CRC cell proliferation, colony formation and invasion [32]. Moreover, it has been reported that low miR-520e expression is associated with the increased CRC growth and migration. The astrocyte elevated gene-1 (AEG-1), which acts as an oncogene [33], is a direct miR-520e target in CRC. Cells overexpressing miR-520e displayed lower GSK-3β phosphorylation and β-catenin expression. Mechanistically, it was found that miR-520e regulates cancer cell behaviour by targeting AEG-1 which on turn inactivate the Wnt/β-catenin signalling and the transcription of its downstream genes. Hence, miR-520e overexpression could represent a promising therapeutic target in CRC by AEG-1 suppression.

Approximately 40–50% of CRC patients develop metastasis, mostly to the liver and lung. In cancer patients, metastases are associated with 90% of all cancer-related death, thereby the mechanisms accounting for the metastatic spread have been deeply investigated. Zhang et al. [34] demonstrated that the rhomboid domain containing 1 (RHBDD1) plays a crucial role in driving metastasis formation in CRC patients, via the Wnt/β-catenin pathway. It has been shown that RHBDD1 is able to influence the Wnt/β-catenin cascade by increasing the phosphorylation of β-catenin at the Ser552 and Ser675 residue without affecting its nuclear translocation. Moreover, it promotes EMT, stemness, migration and invasiveness. RHBDD1 also improves the expression of the β-catenin target gene, ZEB1. Furthermore, the protein level of RHBDD1 positively correlated with ZEB1. Thereby, RHBDD1 has been proposed as a novel therapeutic target and/or a clinically useful biomarker for metastatic CRC (Table 1).
SLC35C1, or GDP-fucose transporter 1, is a member of the solute carrier (SLC) superfamily of solute carriers. The Deng’s group [35] explored the mechanism throughout SLC35C1 regulates the canonical Wnt/β-catenin pathway in CRC. They demonstrated a reduction of SLC35C1 and an increase of β-catenin at all tumour stages. Indeed, silencing SLC35C1 resulted in the increased release of Wnt3a and c-myc, Axin2 and cyclin-D1 expression. This suggests that SLC35C1 is involved in the control of the canonical Wnt/β-catenin pathway, and thereby in tumour cell proliferation and tumour progression (Table 1).

Neuronal pentraxin 2 (NPTX2) is a member of the neuronal pentraxin family and is essential for the formation of synapsis. NPTX2 was found overexpressed at both mRNA and protein level in CRC, particularly in metastatic lesions [36]. NPTX2, which was found to positively correlate with tumour stages, lymphatic invasion, distant metastasis, and poor patients’ outcome, promotes β-catenin nuclear translocation and the expression of c-myc, cyclin D1, Snail, and N-cadherin. No NPTX2 receptors have been identified in CRC, however, its cellular internalization was found mediated by the Wnt/β-catenin receptor, FZD6. Additionally, it has been reported that NPTX2/FZD6 interaction translates in cancer cell proliferation and metastasis formation by triggering the Wnt/β-catenin pathway [36] (Table 1).

Aberrant gene expression and DNA methylation profiles are considered hallmarks of CRC initiation and progression [37]. Due to the APC inactivating mutations, the Wnt/β-catenin pathway plays a key role in CRC metastatic spread [38][39]. Bruschi el al. [39] investigated the early transcriptional and epigenetic changes resulting from APC inactivation in intestinal crypts in crypt base columnar (CBC) cells. The authors have found that APC disruption rapidly induces changes in DNA methylation, indicating that focal remodelling of the DNA methylation profile occurs early and concomitantly with the first oncogenic event. Moreover, it has been demonstrated that the hyper-activation of the Wnt/β-catenin pathway associated with the APC loss-of-function turns out in a rapid increase of intestinal stem cell commitment towards differentiation. Again, it was correlated with the remodelling of the DNA methylation profile. This study unveils that early changes in DNA methylation are crucial for the impaired fate decision program associated with APC loss-of-function.

The kelch-like family member 22 (KLHL22) is a tumour suppressor protein involved in the development/progression of several cancers [40]. Low expression of KLHL22 was found in CRC tissues. KLHL22 overexpression was associated with decreased migration, invasion and reduced expression of the EMT markers, vimentin, N-cadherin, Twist1 and Snail1. Intriguingly, KLHL22 knockdown led to β-catenin and LEF increased expression, while KLHL22 overexpression translates into GSk-3β upregulation and β-catenin downregulation [40] (Table 1).

Non-canonical Wnt pathway and CRC

The canonical and non-canonical Wnt family members play discrete roles in CRC. The activation of the Wnt/calcium pathway turns into stimulation of sensitive proteins such as CamKII and PKC [38][41]. A Ror family of receptor tyrosine kinases, the ROR2 has been shown to act as a Wnt5a receptor or co-receptor [42]. Wnt5a has different roles in CRC. It can act as antagonist or agonist of the canonical Wnt/β-catenin pathway, depending on the cellular context. Lee et al. [43] noticed that the antagonism between the canonical and the non-canonical Wnt/β-catenin signalling pathways is linked to Wnt5a. Mechanistically, Wnt5a suppressed the canonical Wnt/β-catenin cascade by acting as ligand on the RORα [42]. After PKCa-mediated phosphorylation, RORα modifies its affinity and interacts with the armadillo repeat domains of β-catenin, thus suppressing its transcriptional activity.

Three relevant goals have been recently achieved by Voloshanenko et al. [44] supporting the role of Wnt5a/b in cell growth, via the non-canonical β-catenin pathway. First, they identified the procollagen-lysine,2-oxoglutamate 5-dioxygenase 2 (PLOD2), the hydroxacyl-CoA dehydrogenase (HADH), ligand-dependent corepressor (LCOR) and the receptor expression-enhancing protein 1 (REEP1) as candidate genes regulated by the non-canonical Wnt/β-catenin pathway. Second, these genes were found regulated by Wnt5a/b, as well as by ROR2, the DVL2, the activating transcription factor 2 (ATF2) and ATF4 in a non-canonical Wnt/β-catenin independent manner. Lastly, Wnt5a/b silencing was found to impair cancer cell proliferation.
Among several soluble Wnt proteins, Wnt11 was found upregulated in CRC patients [45]. Recently, Gorroño-Etxebarria and colleagues [46] have shown that increased Wnt11 and its FZD6, RYK, PTK7 receptors, positively correlate with poor prognosis. Additionally, Wnt11 downregulated β-catenin transcriptional activity and increased ATF2 via the non-canonical Wnt signalling pathway. Thereby, Wnt11 has been proposed as a prognostic biomarker and therapeutic target in CRC patients.

Tumour micro environment (TME) has a pivotal role in cancer development [47]. Liu et al.[48] reported that, unlike CRC cells, tumour associate macrophages (TAMs), and in particular M2-like cells express Wnt5a. Furthermore, it has been shown that Wnt5a positive TAMs regulate macrophages infiltration, tumour cell proliferation and migration. Wnt5a pro-tumour activity was found associated with the overexpression of the C-C motif chemokine ligand 2 (CCL2) in Wnt5a-treated macrophages. Consistently, Wnt5a knockdown reduced CCL2 expression in TAMs and their cancer-promoting activity. In Wnt5a-treated macrophages both CaMKII and ERK1/2 undergo phosphorylation and lead to CCL2 secretion. This study provided evidence for a new role of Wnt5a in CRC and describes a potential novel therapeutic target (Table 1).

** BREAST CANCER**

Breast cancer (BC) is the most diagnosed cancer in women [46-49], the first cause of cancer death in women worldwide [47][50], and one of the most expensive health care cost [46][49]. Both the canonical and non-canonical Wnt/β-catenin pathways are essential for mammary gland development [51] and for BC growth and dissemination [52]. Hyper-active Wnt/β-catenin was reported in breast tumours [53][53]. In human BC, elevated intracellular β-catenin level has been associated with high tumour grade [54] and poor prognosis. Moreover, up to 90% of metaplastic carcinomas and non-metastasizing fibromatosis have been associated with the highest β-catenin expression level [55]. Moreover, proteins such as Wnt3a [56] and xenopus frizzled 7 (Xfz7) [57] have been involved in the activation of both the canonical and the non-canonical Wnt signalling pathways.

**Canonical Wnt pathway and BC**

Dysregulation of the Wnt/β-catenin cascade has been associated with cancer initiation and metastasis formation [56][58]. Moreover, high β-catenin expression has been reported in basal-like BC subtype [50][53]. Additionally, it has been demonstrated that loss of secreted frizzled-related protein 1 (sFRP1) is an early event in BC patients and is associated with poor prognosis [59]. Furthermore, the activation of the Wnt/β-catenin cascade has been associated with radio resistance of progenitor cells. Thereby, the Wnt/β-catenin pathway has been proposed as a target to harm the self-renewal potential of stem/progenitors [60].

A recent study demonstrated that high β-catenin level is associated with miR106a overexpression and involved in BC cell growth. Additionally, high level of miR106a was reported to reduce cisplatin sensitivity. Major results were obtained exploiting the Wnt inhibitor, FH535. In fact, FH535 treatment reduced the expression of β-catenin, cyclin D1, c-myc and Ki67, impaired tumour growth and induced apoptosis [61].

In a different study [62], the impact of the Wnt/β-catenin canonical pathway in cisplatin resistance was investigated by silencing β-catenin via small interfering RNA (siRNA). The authors demonstrated that upon β-catenin silencing, the cells become more sensitive to cisplatin treatment. These effects were associated with the increased expression of the apoptotic proteins caspase 3/9.

A recent study demonstrated that miR-5188, aberrantly expressed in breast cancer patients, positively correlates with poor prognosis. The molecular analyses revealed that miR-5188 directly targets the forkhead box protein O1 (FOXO1). In physiological setting, FOXO1 binds β-catenin and induces its degradation. This implies that miR-5188 overexpression leads to β-catenin nuclear accumulation and transcription of its downstream target genes, mainly involved in EMT, tumour cell proliferation, metastasis formation and chemo resistance. Moreover, the authors elegantly showed that miR-5188 expression is under the control of c-Jun, which directly binds to its promoter region. This on turn generates a positive loop accelerating tumour progression. Clinically, miR-5188 has been proposed as a diagnostic or prognostic factor and/or a direct target for anti-cancer therapy [63].
The upregulation of the lncRNA hoxa transcript at the distal tip (HOTTIP) has been also linked to poor prognosis in BC patients. Overexpression of HOTTIP correlates with the expansion of breast CSCs (BSCSs) and the expression of the stem cell markers, OCT4 and SOX2. Han et al. [64] demonstrated a reduced expression of differentiation markers, such as CK18 and CK14 and that miR-148a inhibits BC cell migration and invasion by directly targeting Wnt1. Moreover, it has been reported that HOTTIP controls miR-148a-3p by acting as a competing endogenous RNA (ceRNA).

Thereby, HOTTIP promotes expansion of CSCs in vitro and tumorigenesis in vivo by regulating the miR-148a-3p/Wnt1/β-catenin axis [64]. These data are summarized in Table 2.

The LGR4 was identified as a prognostic marker in breast tumours displaying poor prognosis [65]. A tight molecular interplay between LGR4 and Wnt/β-catenin signalling has been reported to control stemness. Indeed, LGR4 binding to the soluble R-spondin proteins eases the Wnt/β-catenin cascade [64] [66]. Previous studies have proven that upregulation of ZEB1 by SLUG (the protein product of SNAI2), increased EMT [67]. As a matter of fact, LGR4 knockdown leads to SLUG and ZEB1 downregulation, thereby impairs invasion and metastasis [68]. A correlation with poor outcome and the expression of the LGR4 homolog LGR5 was also reported. LGR5 maintains the pool of BCSCs and promotes tumour progression and invasiveness by activating the Wnt/β-catenin canonical pathway [68] (Table 1).

Wang et al. [69] first demonstrated that the expression of the suppression of tumorigenicity 7 like (ST7L) is downregulated in BC cells, and more importantly, that ST7L acts as an antitumor supervisor by reducing GSK-3β phosphorylation and inducing β-catenin degradation. However, the mechanisms through which ST7L controls GSK-3β phosphorylation is still missing (Table 1).

A recent study [70] reported the overexpression of the transmembrane emp24 domain (TMED) in BC and its correlation to poor prognosis. Aberrant level of TMED boosts cell cycle progression, colony formation, migration and invasion and the expression of CDK2, CDK4, CDK6, cyclin E, β-catenin, cyclin D1, c-myc, MMP-7 and TCF4. Conversely, silencing TMED3 drastically reduced migration and invasion. Moreover, the observation that β-catenin knockdown translates in the reduction of its regulated genes supports the notion that the oncogenic effect of TMED goes through the Wnt/β-catenin pathway (Table 1).

Cryptotanshinone (CTS) is an herbal medicine derived from roots of salvia miltiorrhiza which displays anti-tumour properties. It has been shown that in vitro CTS reduces tumour cell growth, migration and invasion by downregulating the pyruvate kinase muscle isozyme M2 (PKM2), a protein involved in glycolysis, and more importantly in β-catenin activation [71].

Wnt non-canonical pathway and BC

Among the Wnt ligands, the most extensively studied ligand, activating the β-catenin independent pathway, is Wnt5a. However, its different biological actions are enlightened by the observation that it can also initiate the canonical β-catenin signalling cascade [20] [72].

Wnt5a is an evolutionarily conserved Wnt ligand, which plays an important role in developmental processes. Wnt5a+/- knockout mice showed perinatal lethality, due to developmental defects [73].

In tumorigenesis, Wnt5a signalling is central and displays multiple intriguing and opposite roles mainly acting as a β-catenin antagonist. These data are discussed.

The Wnt5a suppressive properties detected in tumours connoted by β-catenin hyper-activation has been linked to the shift towards the stimulation of the β-catenin independent signalling pathway.

Foxy5 is a Wnt5a mimicking hexapeptide able to decrease BC cell migration and invasion [74]. More recently Prasad et al. [75] confirmed these data and added new information on the role of Wnt5a in the regulation of the expression of the phosphofructokinase platelet-type (PFKP). They have shown that low PFKP level correlates to cancer cell migration and poor patients’ survival. The growth and expansion of tumour cells also rely on glucose consumption resulting in the accumulation of lactate. Cancer cell metabolism was also associated with β-catenin activation [76]. At this regard, it has been shown that Wnt5a affects the aerobic glycolysis by inhibiting the activation of β-catenin. Therefore, an onco-suppressive role was proposed for PFKP.
According to the study of Borcherdining et al. [77], Serra Roarty et al. [26][78] demonstrated that the paracrine activity of Wnt5a suppresses the expression of both β-catenin and cyclin D1. The authors have shown that Wnt5a supports TGF-β-mediated tumour suppressive functions by antagonising Wnt/β-catenin signalling and limiting tumour cell proliferation.

Moreover, Leris and colleagues [79] proved that Wnt5a mRNA level was significantly lower in tumour than in normal tissues, particularly in those displaying a more aggressive behaviour. Again, this observation has suggested a suppressive role of Wnt5a in cancers. It has been also reported that loss of Wnt5a associates with a higher histological tumour grade, increased risk of recurrence, and a shorter recurrence-free survival in invasive BC [80] (Table 1).

On the contrary, Kobayashi et al. [81] reported that Wnt5a is expressed in ER-positive BC cells and positively associates to vessel invasion, tumour size and migration. Mechanistically, Wnt5a induces the expression of the activated leukocyte cell adhesion molecule (ALCAM), a protein involved in migration and invasion. Knockdown of either Wnt5a or ALCAM inhibited tumour cell migration, confirming the role of the Wnt5a/ALCAM axis in the migratory phenotype of ER-positive BC (Table 1).

A relevant role of Wnt5a in reprogramming the TME was also described [82]. It has been shown that under pro-inflammatory conditions the non-canonical Wnt protein induces the expansion of the CD163(+) immunosuppressive macrophages translating in the release of IL-10 and the inhibition of the classical TLR4-NF-kB signalling pathway [82].

Moreover, higher level of Wnt5a was found in human monocye-derived myeloid dendritic cells (Mo-mDCs) than in normal monocytes and macrophages. Wnt5a was found to inhibit the generation of Mo-mDCs by stimulating BC cells to produce IL-6. In addition, the presence of IL-6 in the conditioned media of Wnt5a stimulated BC cells was found involved in the inhibition of Mo-mDC differentiation [83]. Consistently, overexpression of Wnt5a mRNA was detected in metastases derived from primary BC cells and in BC cell lines [84].

Wnt5a signalling is also able to modify the CD44-AKT signalling pathway, leading to a reduced BC cell migration and invasion. In epithelial BC cells, silencing of Wnt5a drives EMT-like changes without altering the expression of common EMT markers. On the contrary, it interferes with CD44 expression and induces pAKT downregulation, thereby acting via a EMT-independent mechanism [85].

The dual activity of Wnt5a has been also ascribed to the Wnt5a isoforms. Bauer et al. [86] have shown that the Wnt5a gene encodes for two distinct isoforms: the Wnt5a-long (Wnt5a-L) and Wnt5a-short (Wnt5a-S) isoform. When analysed in several cell lines Wnt5a-L reduced tumour progression, while Wnt5a-S promoted tumour growth.

Overall, Wnt5a may play multiple roles. Whether it acts as a tumour suppressor or a tumour promoter remains elusive and depends on the availability of essential receptors, the TME, and the activation of discrete signalling pathways.

TRIPLE-NEGATIVE BREAST CANCER

Triple-Negative Breast Cancer (TNBC) is an invasive type of breast carcinoma that lacks the expression of estrogen and progesteron receptor as well of the human epidermal growth factor receptor 2 (HER2) [87] and accounts from 10 to 15% of all BC [88].

TNBC patients have poor outcome due to the high grade of proliferation, early tumour dissemination, and the lack of targeting approaches [89,90]. The malignancy is associated with earlier age of onset, aggressive clinical course, and dismal prognosis [88]. TNBC gained attention due to the aggressiveness and the lack of effective treatment options. Therefore, the most relevant data on this breast cancer subtype are independently discussed.

Gene expression omnibus (GEO) databases were applied by Shen et al. [91] to gather gene expression data in TNBC patients who underwent chemotherapy. They reported that co-expression of NIMA-related kinase 2 (Nek2) and β-catenin correlated with patients’ poor prognosis. β-catenin binds to and is phosphorylates by the Nek2B isomer. Thereby, in TNBC, Nek2B functions as a β-catenin regulator by activating the Wnt signalling pathway and its downstream target genes. In
addition, it has been suggested that Nek2B and β-catenin may synergize to promote resistance to chemotherapy. However, further studies are required to better elucidate the relationship between β-catenin and Nek2 and its possible implications in cancer development (Table 1).

TNBC aggressiveness also relies on the activation of the non-canonical Wnt/PCP pathway. Indeed, the aberrant activation of downstream genes activated by the non-canonical Wnt/PCP pathway has been implicated in tumour growth and poor prognosis. Results from Puviarajasinghe and colleagues [92] revealed that van gogh-like 2 (VANGL2), a core Wnt/PCP component, plays a crucial role in cancer cell migration, anchorage-dependent and independent cell proliferation, as well as in tumour growth. Since, the scaffold p62/SQSTM1, a VANGL2-binding partner, has a key role in VANGL2–p62/SQSTM1–JNK pathway, the possibility to exploit p62/SQSTM1 as a potential therapeutic target has been proposed. This would be of particular relevance since the JNK targeting approaches are associated with major side effects in clinical setting (Table 1).

Yu and colleagues [93] demonstrated that the hematopoietic protein tyrosine phosphatase (HePTP) stabilizes β-catenin in the cytoplasm and allows its nuclear translocation by regulating the phosphorylation of GSK-3β. This results in the transcriptional activation of target genes leading to cell migration and invasion. Since knockdown of HePTP significantly suppresses metastases formed by TNBC cells, HePTP has been also proposed for therapeutic approaches in TNBC (Table 1).

Recently, Kong et al. [94] have shown that a Rho-GTPase-activating protein, the deleted in liver cancer gene 3 (DLC-3), is downregulated in TNBC and its expression is linked to lymphatic metastases. DLC-3 overexpression leads to β-catenin and c-myc downregulation as well as in reduced in vitro cell proliferation, colony formation, migration, and invasion. Hence, a tumour-suppressor role related to the inhibition of the Wnt/β-catenin signalling pathway has been postulated (Table 1).

Liu and colleagues [95] have reported a low expression of miR-6838-5p in TNBC compared to normal cells. miR-6838-5p overexpression reduced cell invasion, migration, EMT, β-catenin, c-myc and cyclin D1 expression by post-transcriptionally controlling Wnt3a expression.

Recently, miR-27a-3p was found overexpressed in tumour cells and linked to poor prognosis in TNBC patients. miR-27a-3p leads to the activation of Wnt/β-catenin cascade and enhances cell proliferation and migration by directly targeting the 3’-UTR region of GSK-3β [96] (Table 2).

OVARIAN CANCER

Ovarian Cancer (OC) is a global issue representing the fourth most common cancer in the female population, particularly in developed countries [97]. The poor survival rate is mainly due to the lack of screening methods at the early stages along with the absence of effective treatment options for advanced stages [96][98]. Among different OC subtypes, the epithelial subtype (EOC) holds about 90% of the overall ovarian malignancies [97][99].

Canonical Wnt pathway and OC

Wnt/β-catenin signalling pathway play a crucial role in carcinogenesis of all OC subtypes [98][100]. In particular, several transcription factors, proteins and miRNAs acting on this pathway have been explored [98][101].

Chen and co-workers [102] investigated the role of the Wnt/β-catenin pathway antagonist dickkopf-related protein 1 (DKK1). They showed that DKK1 is involved in the control of OC stemness. Mechanistically, it has been shown that STAT3 directly activates the transcription of miRNA-92a, translating in DKK1 downregulation [102]. Moreover, overexpression of miR-1207 was found to correlate with high nuclear β-catenin level [103]. Wu et al. [103] investigated the effects of miR-1207 on the expression of the SFRP1-AXIN2 and the inhibitor of β-catenin and T cell factor 4 (ICAT). They found that miR-1207 overexpression was associated with a reduced SFRP1-AXIN2 and ICAT expression and the appearance of a stem-like phenotype (Table 1).

Salem et al. [104] proved that miR-590-3p promotes OC growth and metastasis, by targeting FOXA2. Moreover, it has been shown that miR-590-3p upregulation significantly increase cell growth, migration, and invasion in EOC cells, both in vitro and in vivo [105]. Similarly, FOXA2, which exhibits suppressive activity on EOC cells, has been identified as a miR-590-3p target [105]. The cyclin
G2 gene (CCNG2) has been also reported to display several repressive actions on EOC-derived tumor cell lines. It inhibits cell proliferation, migration, invasion and EMT. Thereby, since miR-590-3p post-transcriptionally regulates FOXA2, FOXO3, CCNG2 and DDK1 expression, miR-590-3p has been proposed as a potential target in EOC patients [105]. A crucial role of SFRP1 in OC growth has been also proposed. Since miR-1180 is highly expressed in neoplastic tissues, Hu et al. [106] explored the relationship between miR-1180 and the SFRP1/Wnt/β-catenin signalling pathway in this context, demonstrating that miR-1180 triggers the activation of the Wnt/β-catenin cascade by targeting SFRP1.

The members of the R-spondin ligand family have been reported as positive effectors of the Wnt/β-catenin signalling [107]. LGR4-6 plays crucial roles in the activation of the Wnt/β-catenin cascade [107,108]. Moreover, Ruan et al. [107] have reported that LGR6 induces stemness and chemoresistance via the Wnt/β-catenin pathway in OC cells. Restraining the stem phenotype and increased sensitivity to chemotherapy have been proved by LGR6 silencing (Table 1).

A recent study established that the overexpression of the Rab GTPase family member, Rab14, regulates GSK-3β phosphorylation and β-catenin nuclear accumulation [109,110]. Moreover, high level of Rab14 was found associated with higher expression of Wnt/β-catenin target genes including MMP-7 and c-myc [110](Table 1).

Jiang et al. [97] have demonstrated that tetrandrine (TET) enhances the anti-tumour effect of paclitaxel (PTX) by decreasing c-myc and cyclin D1 and increasing p21 expression, resulting in cell cycle arrest. The pro-apoptotic effects of PTX+TET have been also investigated. TET was found to inhibit β-catenin downstream target genes by enhancing PTX activity and conferring sensitivity to PTX in resistant cells [97].

Barghouth and co-workers [111] demonstrated a more active Wnt/β-catenin signalling in carboplatin-resistant cells than in sensitive ones. Unlike the Wnt ligands, the negative Wnt regulators DKK1, SFRP1, and the FRZB have been found downregulated in cisplatin-resistant cells. These findings suggest that Wnt/β-catenin blockade may be effective on resistant EOC.

Non-canonical Wnt pathway and OC

The FZD7 is highly expressed in OC [112] and its overexpression in mesenchymal (Mes) and Stem-A OC subtypes, has been associated with the induction of EMT. The PCP pathway, which activates the Rho-ROCK axis, was found involved in the activation of actomyosin contractility, cadherin-based cell-cell adhesion and migration, while the Wnt/calcium pathway in the metastatic spread and cytoskeleton changes in this clinical setting [112]. Therefore, it has been proposed that the FZD7 controls both cell cycle progression and cell migration via the non-canonical Wnt/PCP pathway (Table 1).

The integrin beta 1 subunit (ITGBL1) was found highly overexpressed in OC [113]. It has been shown that ITGBL1 promotes cell migration and adhesion via Wnt/PCP, RhoA, the focal adhesion kinase, and the steroid receptor coactivator (FKα/src) pathway (Table 1).

The PTK7 which interacts with Wnt5A, LRP6 and FZD7 [114,115] may act as tumour suppressor or oncogene [116,117]. In EOC, PTK7 downregulation is indeed associated with a poor prognosis [116].

Luo and colleagues [118] have investigated the role of the alkaline phosphatase (ALPL) in OC. They demonstrated that ALPL overexpression inhibits EMT, migration and invasion of high grade serous OCs (HGSOC) and FZD2 correlates with a poor survival rate [118]. Mechanistically they have shown that ALPL overexpression represses Wnt5a/FZD2–mediated EMT activation possibly by interfering with STAT3 activation [118] (Table 1).

WNT PATHWAY AND OTHER CANCERS

Glioma is an aggressive tumour of the nervous system displaying rapid progression and poor prognosis. Zhao et al. [119] have found that overexpression of β-catenin and cyclin D1 is associated with high level of the long noncoding RNA, FGD5 antisense RNA 1 (IncRNA FGD5-AS1). A close relationship between them was straitened by the observations that inhibition of FGD5-AS1 reduced...
β-catenin and cyclin D1 expression while β-catenin downregulation decreases lncRNA FGD5-AS1 expression. This results in the impaired tumour cell migration and invasion.

Prostate cancer (PCa) is among the most common tumour in male. A recent study by Situ et al. [120] provided evidence for the involvement of the microRNA-939 (miR-939) in PCa. Downregulation of miR-939 was found in tumour tissues at advanced tumour stage, in distant lesions as well as associated with poor prognosis. Molecularly, it was demonstrated that miR-939 upregulation interferes with the Wnt/β-catenin cascade by directly targeting the hepatoma-derived growth factor (HDGF).

Osteosarcoma (OS) is a common bone paediatric tumour displaying high rate of lung metastasis. The inhibition of β-catenin activation, metastasis formation and chemoresistance were found modulated by tegavivint (a Wnt/β-catenin inhibitor) which has been proposed as an alternative therapeutic option in OS [121].

Melanoma is among the most immunogenic tumours displaying increased lymphocytic infiltration. Low 1α,25-dihydroxyvitamin D3 and vitamin D receptor (VDR) level correlates to increased cancer incidence and melanoma progression, respectively. Recently, it has been shown that high VDR expression correlated with the inhibition of tumour growth, low Wnt/β-catenin activation and the induction of the immune response [122] (Table 1).

The long non-coding RNA00261 (Linc00261) has been shown to display onco-suppressor properties in Pancreatic Cancer (PC). Linc00261 overexpression inhibits PC cell proliferation, invasion, EMT and metastasis. Bioinformatics analysis revealed that Linc0026 inhibits the activation of the β-catenin/TCF4 cascade and the metastatic spread by regulating the miR-552 5p/FOXO3 axis [123].

EXTRACELLULAR VESICLES AND THE WNT PATHWAY

EVs are heterogeneous small membrane-bound carriers with complex cargoes released under both physiological and pathological conditions. Almost any cell can release EVs, which act as intercellular mediators modifying target cell fate at closed or distant sites [124]. Based on the biogenesis, size, content, mechanisms of release and function, three discrete EV subtypes are recognized: microvesicles (MVs), exosomes, and apoptotic bodies [124].

EVs-mediated transfer of specific molecules are known to dictate the phenotype of the recipient cell. They can act on proliferation, motility, EMT, migration, invasion, immune evasion, chemoresistance, and TME reprogramming (Figure 3).
Figure 3. Schematic representation of cell-to-cell communication in the TME by EVs. EVs are released by almost all cell types in the TME. EVs serve as intercellular mediators transferring specific molecules (proteins including Wnt ligands and β-catenin, and miRNAs) to recipient cell thus promoting tumour expansion, cancer cell dedifferentiation in CSCs, chemo-resistance, and neovessel formation. CCs: cancer cells; CSCs: cancer stem cells; TEC: tumour-derived endothelial cell; CAF: cancer associated fibroblasts.

Moreover, EVs derived from serum or other biological fluids have been proposed as tumour biomarkers. More importantly, EVs have gained attention as anti-cancer tools. Indeed, EVs can be used as drug delivery systems or potential cancer vaccines. Moreover, the transfer of Wnt ligands or β-catenin via EVs has been proposed as a Wnt signalling activation mechanism.

Kalra et al. [125] have shown that EVs released by CRC cells and containing the mutant β-catenin and high Wnt/β-catenin activity boost the expression of target genes as c-myc and cyclin D1 when transferred to recipient cell (Table 43).

Table 3. EVs involved in several tumours, their alteration, targets, and impact on tumours

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<tr>
<td>14-3-3ζ in EVs</td>
<td>HEK293T</td>
<td>COS-7, SW480</td>
<td>CRC</td>
<td>Upregulated</td>
<td>β-catenin, GSK-3β, DVL2</td>
<td>survival, migration</td>
<td>[126]</td>
</tr>
<tr>
<td>Wnt ligands in EVs</td>
<td>CAFs</td>
<td>CRC</td>
<td>CRC</td>
<td>Upregulated</td>
<td>β-catenin</td>
<td>dedifferentiation, drug resistance, colony formation</td>
<td>[127, 128]</td>
</tr>
<tr>
<td>β-catenin in EVs</td>
<td>milk</td>
<td>HCC</td>
<td>HCC</td>
<td>Silenced</td>
<td>β-catenin</td>
<td>proliferation, tumour growth</td>
<td>[131, 132]</td>
</tr>
</tbody>
</table>
The 14-3-3 are conserved molecules displaying regulatory functions and promoting cancer progression [126]. The 14-3-3ζ isoform which binds both β-catenin and GSK-3β, leads to the nuclear translocation and accumulation of β-catenin and enhance cell motility. Moreover, EVs enriched in 14-3-3ζ and β-catenin, after internalization, promote cell survival and migration by activating the Wnt/β-catenin cascade [126] (Table 13).

Hu et al. [127] have investigated the mechanism of drug resistance in CRC and have proven that EVs released by fibroblasts drive dedifferentiation of CRC cells towards CSCs (Figure 3a). Additionally, they found that EVs derived from fibroblasts contain the Wnt ligands which activate the Wnt/β-catenin pathway in target cells, induce transdifferentiation of CRC cells into CSCs and increase drug resistance. Furthermore, it has been reported that collagen accumulation and the APC mutation in CRC cells stimulate the release of EVs and, under hypoxia conditions, fibroblast derived EVs boost CRC colony formation [128] (Table 13).

Accumulating evidence shows that EVs enriched in miRNAs are key determinants of human cancer cell growth, invasion and metastasis [129][129]. CAF-derived EVs enclose miR-92a-3p which contribute to cancer progression, stemness, EMT, and drug resistance. Moreover, miR-92a-3p enriched EVs correlated with the activation of the Wnt/β-catenin pathway [129] (Figure 3a).

Long non-coding RNA-APC1 (IncRNA-APC1) is a negative regulator of CRC. Low level of IncRNA-APC1 correlates with metastasis, advanced clinical stage and poor prognosis in CRC patients. APC, via IncRNA-APC1 promotes cell-cycle arrest and suppresses angiogenesis by lowering the release of CRC cell-derived EVs. Finally, it has been shown that EV-derived from CRC are enriched in Wnt1 and enhance CRC cell proliferation and migration via the non-canonical Wnt/PCP signalling [130].

Hepatocellular carcinoma (HCC) is one of the most common cause of cancer-related deaths worldwide. Constitutive activation of the Wnt/β-catenin pathway turns into the expression of the epithelial cell adhesion molecule (EpCAM) [131]. Ishiguro et al. [132] provided evidence that loss in β-catenin and reduced proliferation and invasion can be obtained by EpCAM positive liver cancer stem cells (LCSC) targeted by EVs engineered with a β-catenin specific siRNA (Table 13).

Multiple myeloma (MM) is a hematopoietic malignancy associated with an altered homeostasis of bone formation/resorption. MM-derived EVs enriched in DKK-1 were found to boost the Wnt/β-catenin signalling and contribute to the abnormal osteogenesis. The inhibition of EV shedding
combined with chemotherapy were found to impair tumour load, angiogenesis and osteolysis [133] (Table 1).

Furthermore, a recent study noticed that the release of EVs from HCC cells is increased in hypoxic conditions and linked to cancer cell proliferation, migration, invasiveness and EMT. Mechanistically they have shown that miR-1273f enriched in EVs activates the Wnt/β-catenin signalling cascade by targeting the Wnt/β-catenin inhibitor LHX6 [134].

Chen et al. [135] proved that EVs released from oral squamous cell carcinoma (OSCC) cells correlate with the increased level of β-catenin, the expression of several oncogenic markers, the reprogramming of normal gingival fibroblasts into CAFs, the increased metastasis, stemness reprogramming, chemoresistance, and patients’ poor survival (Table 1).

Xia et al. [136] have demonstrated the uptake of EVs and the delivery of functional miRNAs in different cell lines. The exosomal-miR-1260b was found crucial for the activation of the Wnt/β-catenin signalling and the invasiveness of lung adenocarcinoma cells.

Harada et al. [137] purified and characterized Wnt5b-associated EVs. In pancreatic PANC-1 and colorectal Caco-2 cell lines Wnt5a carried by EVs displays the ability to enhance cancer progression (Table 1).

Luga et al. [138] demonstrated that EV shedding by fibroblasts boosts BC cell growth and motility via the Wnt/PCP signalling. CAF-derived EVs were found crucial drivers of cell migration during metastasis formation. Moreover, they found that EVs secreted from fibroblast L cells promote the autocrine Wnt11-PCP cascade in tumour cells increasing their motility and metastatic properties (Table 1).

Lombardo et al. [139] provided evidence that EVs released by tumour-derived endothelial cells (TECs-EVs) boost in vivo TEC-derived neovessels. Mechanistically they showed that EV released by naïve TECs-EVs regulate the expression of APC, GSK-3β and drive β-catenin nuclear accumulation via miR-214-3p and miR-24-3p (Figure 3b). Overall, this study revealed a key role of the Wnt/β-catenin cascade in TEC-derived neovessel formation. Moreover, they recently showed that naïve TEC-EVs were also able to boost TNBC metastatic spread and lung metastasis formation when injected intravenously [140] (Table 2).

Overall these data indicate a crucial contribute of EVs released by different cell sources in driving tumor development and dissemination. Several data suggest that these effects mainly rely on the transfer of their specific cargo into target cells. Therefore, approaches able to modify their cargo, particularly miRs and proteins involved in their tumor promoting action, have been proposed as useful therapeutic options. EV engineering by using siRNA for mutated protein has been tested and their effectiveness demonstrated in pancreatic cancer [141]. This suggests that using siRNA for mutant β-catenin should be considered as an alternative option for CRC. Likewise, siRNA for different Wnt proteins or rearrangement of dysregulated EV miRs can be used to targeting the Wnt/β-catenin cascade. Alternatively, EVs loaded with Wnt/β-catenin inhibitors can be used as naturally delivery tools.

CONCLUSIONS

Cell-to-cell communication is part of the evolutionary processes. Wnt ligands are essential for the homeostasis and in the last 30 years genetic, biochemical, and molecular investigations have uncovered several Wnt signalling components [2,3]. Driving interest on this topic mainly relies on dysregulation of the Wnt/β-catenin signalling and cancer development/progression [3]. Moreover, Wnt/β-catenin cascade seems to contribute to the TME shape, which play a crucial role in the control of tumour progression and immune regulation. Many different Wnt proteins have been described, and among them Wnt5a, plays a critical role taking part in both the canonical and the non-canonical Wnt/β-catenin pathway [77,78].

The identification of specific tools able to interfere with the Wnt/β-catenin cascade has been a hotspot for many years. This is particularly true for CRC, in which almost 70% of CRC patients display APC mutations [15]. Apart from CRC, the Wnt/β-catenin pathway is gaining attention in several malignancies, such as breast, ovarian, melanoma, prostate and paediatric osteosarcoma [117–
119]. At this regard, BC and in particular TNBC are featured by the abnormal activation of both the
canonical and non-canonical Wnt/β-catenin pathway [89,90]. Likewise, a hyper-active Wnt/β-catenin
cascade has been shown to play a crucial role in the progression, stemness, and drug resistance in OC
cancer [101,105]. Several miRNAs have been identified to modulate this cascade and thereby widely studied
as screening markers or targets in different tumour settings [142].

In the TME, intercellular communication has been recently reported as mediated by the transfer
of EV molecular cargo and revised in [143]. Their cargo also includes a number of Wnt components.
Of note, wild-type and mutant β-catenin able to promote survival and proliferation of recipient cell
and, in several instance dedifferentiation towards a CSC phenotype, have been detected in EVs
(Figure 3a). Moreover, their role in mediating drug resistance has been reported. Furthermore, since
EVs are released within the TME their contribute in cancer growth and progression has been
extensively investigated [144]. EV shedding, blockade, or engineering have been proposed as
innovative anti-tumour instrument to fine-tuning the Wnt/β catenin pathway [142,145].

In the last decades several efforts have been directed to the development of Wnt/β catenin
targeting approaches in order to interfere with tumour progression. However, these efforts have been
limited by the crucial role of the Wnt/β catenin pathway in preserving tissue homeostasis. Therefore,
future energies should be directed to clearly dissect the mechanisms driving the unbalanced Wnt/β
catenin pathway in cancer, and EV mechanism of action should be considered amid them. Should
identified, targeting approaches would become a suitable anti-cancer option.

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