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Nuclear transport of galectin-3 and its therapeutic implications

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Abstract

Galectin-3, a member of β -galactoside-binding gene family is a multi-functional protein, which regulates pleiotropic biological functions such as cell growth, cell adhesion, cell-cell interactions, apoptosis, angiogenesis and mRNA processing. Its unique structure enables it to interact with a plethora of ligands in a carbohydrate dependent or independent manner. Galectin-3 is mainly a cytosolic protein, but can easily traverse the intracellular and plasma membranes to translocate into the nucleus, mitochondria or get externalized. Depending on the cell type, specific experimental conditions in vitro, cancer type and stage, galectin-3 has been reported to be exclusively cytoplasmic, predominantly nuclear or distributed between the two compartments. In this review we have summarized the dynamics of galectin-3 shuttling between the nucleus and the cytoplasm, the nuclear transport mechanisms of galectin-3, how its specific interactions with the members of β -catenin signaling pathways affect tumor progression, and its implications as a therapeutic target.

Galectin-3: structure and function

Galectin-3 is a unique ~31kD lectin belonging to the evolutionarily conserved family of galectins that share a unique carbohydrate recognition domain (CRD). Galectin-3 is a chimera protein, which contains a collagen- α -like domain and an N-terminal domain in addition to the CRD [1]. Each of the three structural domains of galectin-3 is associated with at least one specific function: (a) the NH2 terminal domain contains a serine phosphorylation site, which is important in regulating its nuclear localization; (b) the

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proline-rich collagen- α -like sequence is cleavable by matrix metalloproteinases and this cleavage acts as a surrogate diagnostic marker for MMP-2 and MMP-9 activities in the tissue; and (c) a COOH terminal contains a single carbohydrate-recognition domain and the NWGR anti-death motif. Although the presence of galactose is essential for all galectins' binding, its affinity for the monosaccharide ligand is rather weak, with K_d values in mM range. The binding affinity of galectins increases if galactose is attached to other saccharides e.g. N-acetylglucosamine forming N-acetyllactosamine [2]. Galectin-3 is mainly a cytosolic protein, but there is ample evidence to confirm its presence on the cell surface, in the conditioned media of some cell lines, in the extracellular matrix and in the biological fluids and sera. This suggests that galectin-3 is a shuttling protein and may have multiple functions accordingly [3-5]. Galectin-3 lacks the classical secretion signal sequence and does not pass through the standard ER/Golgi pathway [6]. Still it can be transported into the extracellular milieu via a non-classical pathway [7].

Cells differ widely in their capacity to secrete galectin-3. While J774.2 macrophage cells secrete 30–45% of their galectin pool [8], BHK and MDCK cells export 10–15% [9, 10] and WEHI-3 mouse macrophages secrete a very small percentage of galectin-3 in the conditioned medium [8]. The exact mechanism of galectin-3 secretion is not yet known. Various groups have proposed different possible mechanism for its secretion such as via vesicular release [8-11], release as a component of the exosome [12], or by mechano-transduction mechanism [13]. Once it is released into the extracellular matrix, because of its ability to bind to glycosylated proteins, it can interact with a myriad of partners such as EGFR, integrins, N-CAM, fibronectin, and laminin [14-18]. Thus, galectin-3 plays an integral role in multiple biological processes including, but not limited to, cell–cell or cell–matrix adhesion, signal transduction, inhibition of cell receptor internalization, induction of T-cell apoptosis, and induction of angiogenesis [14, 19-23].

Galectin-3 also plays significant roles when expressed intra-cellularly. For example, it binds to members of the serine (S)- and arginine (R)-rich splicing factor family (SR proteins) and form spliceosome complexes within the nucleus [24, 25]. Galectin-3 translocate to the perinuclear membrane in breast cancer cells following a variety of apoptotic stimuli such as cisplatin, staurosporine, or serum withdrawal [26-28]. Over-expression of cytoplasmic galectin-3 in the tongue cancer patients was associated with a decreased disease-free survival [29]. Expression of galectin-3 in the nucleus of human prostate cancer correlated with decreased cell proliferation, while its over-expression in the cytoplasm was reported to promote its anti- apoptotic activity as well as increased cell proliferation, tumor growth, invasion, and angiogenesis [30, 31]. Nuclear staining of galectin-3 was correlated to the lobular type of invasive carcinoma, while tumor stromal expression represented high-grade malignancy in human breast carcinoma [32, 33]. In colorectal cancer an increased cytoplasmic expression of galectin-3 was observed in more advanced stages [34].

Numerous cytosolic molecules have been identified as galectin-3 ligands, which include several molecules involved in the apoptotic pathway: Bcl-2 [35, 36], CD95 (APO-1/Fas) [37], Nucling [38], Alix/A1P1 [39]. Galectin-3 has been shown to translocate either from the cytosol or from the nucleus to the mitochondria following exposure to apoptotic stimuli [27] and to block changes in the mitochondrial membrane potential thereby preventing apoptosis

[40]. The involvement of cytosolic galectin-3 in regulation of cell proliferation, differentiation, survival and cell death was additionally confirmed by the findings that it affects K-ras [41, 42] protein and Akt protein [43, 44]. Synexin (annexin VII) a Ca^{2+} and phospholipid binding protein mediates the translocation of galectin-3 to perinuclear mitochondrial membranes indicating its involvement in cellular trafficking [27].

In this review we will focus on the nuclear/cytoplasmic transport of galectin-3 and its implications as a therapeutic target.

The nuclear pore complex and nucleo-cytoplasmic protein transport

The nuclear envelope is mostly recognized as a diffusion barrier between the cytoplasm and the nucleoplasm. The nuclear envelope consists of two phospholipid bilayers, called the inner and the outer nuclear membrane. The inner nuclear membrane faces the nucleoplasm and contains proteins that interact with the chromatin [45, 46]. The outer nuclear membrane is continuous with the endoplasmic reticulum of the cell and has partially overlapping functions in the transport, synthesis and folding of proteins, and the synthesis of lipids [47]. The inner and the outer nuclear membrane of the nuclear envelope are penetrated by a large macromolecular structure nuclear pore complex (NPC) [48, 49].

NPCs are the only gateway between nuclear contents and cytoplasm. The NPCs mediate selective bidirectional nucleocytoplasmic transport [50, 51]. Small molecules, ions and metabolites below a molecular weight of roughly 20 kDa can diffuse through NPCs; however, macromolecules greater than 40–60 kDa need to be actively transported [52]. The overall structure of the NPC has been determined by distinct electron microscopy approaches and is evolutionarily conserved from yeasts to mammals [53, 54]. Vertebrate NPCs display an eightfold rotational symmetry, have an outer diameter of ~ 120 nm, and are large, ~ 125 MDa structures embedded in the nuclear membrane. The complex can be minimally characterized as having three substructures: the cytoplasmic fibrils, a central core, and the nuclear basket [51, 52, 54]. In higher eukaryotes, NPCs are found at a similar density, ~ 4000 NPCs/nucleus in cultured human cells, although it is strongly influenced by the cell size and the level of biosynthetic activity [55].

The NPC is composed of approximately 30 different proteins termed nucleoporins (Nups; Fig. 1) [56]. Nucleoporins have a very limited set of domains, restricted to β -propellers, α -solenoids, phenylalanine-glycine (FG) repeats, coiled-coil and transmembrane domains, and can be subdivided into three groups [57-59]. The first group comprises the transmembrane proteins that anchor the NPC to the nuclear envelope and reside at the boundary between the central framework and the pore membrane, including gp210, Ndc1 and POM121. The second group represents the structural scaffold of the NPC embedded into the nuclear envelope, including the Nup107-160 complex and the Nup93-205 complex. The third group includes the nucleoporins that form peripheral components of the NPC and contain repetitive FG motifs and/or coiled-coil motifs, which facilitate nucleocytoplasmic transport through the nuclear pore. This third group can be further subdivided into the cytoplasmic filaments including Nup358 and Nup214, the central framework including Nup98 and the Nup62 complex, and the nuclear basket including Nup153, Nup50 and Tpr [57-59]. The

nucleocytoplasmic transport is mediated by factors that belong to nuclear transport receptors termed karyopherins (importins/exportins; Fig. 2) [60, 61]. Karyopherins bind their protein cargoes in the cytoplasm by binding specific nuclear localization signal (NLS). The karyopherin-cargo complex translocates into nucleus through the NPC via interactions with nucleoporins. Once in the nucleus, RanGTP binds to karyopherin which triggers the dissociation of the import complex, whereby the karyopherin is recycled back to the cytoplasm. Conversely, a protein containing a nuclear export signal (NES) forms a trimeric complex with an exporting karyopherin and RanGTP. The export complex passes through the NPC and is dissociated in the cytoplasm by hydrolysis of Ran-bound GTP [50, 60, 61]. All karyopherins bind directly to nucleoporins containing FG repeats as they pass through the NPC [54]. Several studies have demonstrated that various karyopherins have different affinities for specific nucleoporins [54, 62].

As described above, the main function of NPCs is to help transport of molecules between the cytoplasm and the nucleus. A tight regulation of nucleocytoplasmic transport is essential for cell homeostasis. In addition to its role in normal physiology, loss of NPC or nucleoporins function has been implicated in several diseases including cancer and autoimmune disease [59, 63, 64]. Recently gp210 has been demonstrated to be unregulated and to play a crucial role during myogenic and neuronal differentiation [65]. Nup133, a component of the Nup107-160 complex, is required for neuronal differentiation during mouse development [66]. Nup155 is shown to have a role in the physiology of the heart; a Nup155 homozygous missense mutation causes atrial fibrillation which causes sudden death, and loss of Nup155 function appears to disrupt nuclear pore function and develop cardiac arrhythmia [67]. Nup62 mutation causes autosomal recessive infantile bilateral striatal necrosis [68]. The expression of Nup98 is induced by interferons and Nup98 plays important roles in mRNA export from the nucleus, which is targeted by viruses and regulated by interferon [69]. Deletion of Nup98 in mice leads to lethality at embryonic day 6.5 to 7.5, and suggested a role for Nup98 in gastrulation [70]. Nup358 is implicated in myogenesis [71] and knockout of Nup358 leads to increased aneuploidy in mice [72]. Interestingly, Nup358 is an active E3 ligase in the SUMOylation reaction [73], thus, NPCs are indirectly involved in the regulation of numerous cellular processes through the SUMO pathway. Nup153 is also shown to bind to SUMO protease SENP2 [74]. In cancer patients, elevated levels of several nucleoporins were observed; Nup88 was found to be overexpressed in ovarian, breast, colorectal cancer, etc. [75-77]. Rae1 was elevated in breast and lung cancer [78, 79]. Chromosomal translocations between Nup98 and HOX (homeobox transcription factor) gene family have been reported in acute myelogenous leukemia [80, 81]. Nup98-Hox fusions are identified as oncogenes and display strong transcriptional activity and leukemogenic potential. Nup214 and DEK (DNA-binding protein) fusion gene also contributes to leukemia development [81, 82]. It has recently been demonstrated that Tpr interacts with tumor suppressor p53 and regulates autophagy in cancer cells [83]. Moreover, nucleoporins have been described as having important roles during mitosis, which is involved in cellular functions and cancer development [84-86]. Although recent studies have demonstrated nucleoporins are implicated in many biological and physiological functions, the detailed mechanism of nucleoporins in diseases such as cancer still remains unclear.

Galectin-3 shuttling between the nucleus and the cytoplasm

Galectin-3 exhibits pleiotropic biological functions; extracellular galectin-3 mediates cell migration and cell adhesion by interacting with cell surface and extracellular matrix glycoproteins and glycolipids, and intracellular galectin-3 regulates signaling pathways by interacting with cytoplasmic and nuclear proteins [87-89]. Interestingly, galectin-3 shuttles between the cytoplasm and the nucleus [3]. Nuclear and cytoplasmic galectin-3 is likely to be linked with proliferation and differentiation, respectively [87-89]. Nuclear galectin-3 has been associated with pre-mRNA splicing [90] and gene expression of cyclin D1 and c-myc [91, 92]. Thus, understanding the mechanism of galectin-3 transportation between the nucleus and the cytoplasm might provide a therapeutic modality to modulate gene expression related to cancer.

There have been many studies reported on the correlation between galectin-3 subcellular distribution and prognosis in various cancers. In normal cells, nuclear galectin-3 is presumably associated with cell proliferation [90]. Galectin-3 localization in the nucleus promotes cell proliferation by the induction of cyclin D1 expression in human breast epithelial cells [91]. In lung carcinoma, the expression of nuclear galectin-3 is a predictive factor of recurrence and/or a worse clinical outcome [93, 94]. In papillary thyroid cancer cells, nuclear galectin-3 interacts with the thyroid-specific TTF-1 transcription factor and enhances the transcriptional activity to promote the proliferation of the cells [95]. The elevated expression of galectin-3 in the nucleus is a significant pathological parameter related to histological differentiation and vascular invasion in esophageal squamous cell carcinoma patients [96]. In contrast, galectin-3 expression in the nucleus is greatly decreased in colon and prostate carcinomas [30, 97, 98]. In an experimental prostate cancer cell, nuclear galectin-3 suppresses malignancy whereas cytoplasmic galectin-3 promotes tumorigenicity [30, 98]. The levels of nuclear galectin-3 are markedly decreased during the progression from normal to cancerous states in tongue carcinomas [29]. Cytoplasmic galectin-3 expression translocated from the nucleus also exhibits anti-apoptotic activity by interacting with Bcl-2 [99]. It remains still unclear why galectin-3 expression is regulated differently among organs during cancer progression.

It is important to understand the transport mechanism of galectin-3 between the nucleus and the cytoplasm for the development of regulation of galectin-3 function. Although galectin-3 lacks a typical NLS, some reports have mentioned the essential sequences of galectin-3 for nuclear localization. There are several reports which have described the specific sequence responsible for galectin-3 nuclear transport; the first 11 amino acids of human galectin-3, which contains a Ser⁶ phosphorylation site, is required for nuclear distribution [100]; deletion of the 103 N-terminal amino acid residues of hamster galectin-3 resulted in the nuclear localization of galectin-3 [101]; deletion of C-terminal domain of mouse galectin-3 lost the nuclear localization pattern, suggesting that nuclear translocation of galectin-3 is dependent on the IXLT type NLS in the end of C-terminal domain, the critical residues include I²⁵³, L²⁵⁵ and T²⁵⁶ fitting the IXLT motif identified to be important for the nuclear localization of the *Drosophila* protein Dsh [3]; incomplete forms of carbohydrate recognition domain (CRD) region of galectin-3 abrogate the nuclear accumulation, whereas N-terminal domain of galectin-3 promotes the nuclear transportation [102, 103]. There seems to be an

inconsistency in these reports, probably because of the use of different species of galectin-3 or different cells, but also suggesting that mechanism of galectin-3 nuclear import is complicated and very specific. The presence of both phosphorylated and non-phosphorylated endogenous galectin-3 has been reported. The phosphorylated form of galectin-3 was found in both cytosolic and nuclear fractions, whereas the non-phosphorylated form was presented exclusively in the nucleus [104]. The phosphorylation of galectin-3 seems to be necessary and essential for its functions, and galectin-3 has been shown to be phosphorylated at the residue of N-terminal Ser⁶ by casein kinase 1 [105]. The galectin-3 phosphorylation is reportedly important for its nuclear export into the cytoplasm [99, 106].

There are at least two pathways for the translocation of galectin-3 into the nucleus; a passive diffusion and an active transport through the NPC (Fig. 3). Galectin-3 is synthesized as a monomer, and partially forms a dimer and even a pentamer in some situation [107, 108]. Therefore, galectin-3 might be imported into the nucleus by passive diffusion as a monomer, or by active transport as a dimer or a pentamer form due to its size. In active transport into the nucleus, galectin-3 translocates by the importins-dependent transport system [102]. Although there is no classical NLS in galectin-3 sequence, an NLS-like sequence ²²³HRVKKL²²⁸ in the C-terminal region of galectin-3, which resembles sequences of p53 and c-Myc NLSs has been identified [109-111]. Importin- α is the receptor subunit that recognizes the NLS, and associates with importin- β via the importin- β binding domain [60, 61]. Interactions between importin- β and the nucleoporins containing FG repeats are essential in translocation through the NPC [54]. Galectin-3 directly binds to importin- α but not to importin- β , although the functional interaction of importin- β is required for galectin-3 nuclear translocation, and imported into the nucleus via the importin- α/β complex [102]. Furthermore, galectin-3 export mechanism has been also demonstrated. The galectin-3 NES exists in C-terminal region, and regulates the export of galectin-3 into the cytoplasm [103, 106]. During the export, nucleoporin Nup98 specifically regulates the galectin-3 transportation through NPC [112]. The N-terminal domain of Nup98 (1-505aa) interacts with CRD region of galectin-3. Galectin-3 directly binds to Nup98 but not to a nuclear export receptor CRM1 (exportin 1), although Nup98 is a cofactor for CRM1-mediated nuclear export. As described above, a few mechanisms of the interaction between NPC and galectin-3 have been established to date. Further studies are required to examine how galectin-3 shuttles between the nucleus and cytoplasm. Understanding the mechanism and control of galectin-3 transportation would provide new therapeutic target/modality for treating disease such as cancer.

Galectin-3 and β -catenin signaling through nuclear translocation

Interactions of galectin-3 with a plethora of ligands both in the intracellular and extracellular compartments have been reviewed in details previously [88, 89, 113]. In the nucleus, galectin-3 acts as a pre-mRNA splicing factor and is involved in the spliceosome assembly (reviewed in [114]). In the current review, we will focus on its interactions with β -catenin pathway resulting in the regulation of β -catenin regulated transcriptional activity. β -catenin is a downstream component of the Wnt signaling pathway. In the absence of Wnt stimulation, the levels of cytoplasmic β -catenin are low since it is ubiquitinated and

constantly degraded in the proteasome [115]. It is phosphorylated by glycogen synthase kinase-3 β (GSK-3 β) and casein kinase 1 α (CKI α) in a multi-protein complex that also contains adenomatous polyposis coli (APC) and scaffold protein axin [116-120]. Phosphorylated β -catenin is recognized by E3 ubiquitin ligase complex, ubiquitinated and degraded [121, 122]. In activation of the Wnt signaling pathway, binding of Wnt ligand to frizzled receptor and co-receptor LRP5/6 triggers the association of destruction complex with phosphorylated LRP inhibiting the degradation pathway consequently leading to stabilization of cytoplasmic β -catenin and its translocation into the nucleus, where it binds to the transcriptional factor Tcf/Lef and serves as a co-activator of Tcf/Lef to stimulate transcription of the Wnt target genes including c-myc, cyclin D1, cyclooxygenase-2, matrix metalloproteinase-7, gastrin, and ITF-2 [123-128]. Activating mutations in Wnt pathway components, including loss-of-function mutations of *APC* or less frequently in *CTNNB1* (which encodes β -catenin) and *AXIN*, increase β -catenin protein levels and have been found in numerous human cancers including colorectal, gastric, and ovarian cancer [129]. However, mutations of Wnt pathway proteins are not the only factors that contribute to β -catenin activation [130]. Shimura et al [92, 131] reported that in breast cancer cell line BT-459, galectin-3 forms a complex with β -catenin independent of either APC or β -catenin mutations and it forms a ternary complex with Tcf-4. NH₂ terminus of β -catenin was reported to interact with COOH terminus of galectin-3 and could be inhibited by lactose. These authors also showed that galectin-3 is phosphorylated, like β -catenin, by CKI [131] and GSK-3 β and like β -catenin, its nuclear import-export is phosphorylation dependent [131]. While phosphorylation of galectin-3 at S6 by CKI serves as a molecular switch for sugar binding [132], and regulation of nuclear export [99], the phosphorylation of β -catenin by CKI α and GSK-3 β promotes its proteasomal degradation. Shi and colleagues [133] showed that inhibition of both Wnt-2 and galectin-3 in colorectal cancer cells had synergistic effects on suppressing Wnt signaling and inducing apoptosis. Song et al [134] showed that in colon cancer cells, galectin-3 mediates β -catenin expression and Tcf-4 activity by regulation of GSK-3 β phosphorylation and activity via the PI3K/Akt pathway. It was shown that down-regulation of galectin-3 resulted in Akt and GSK-3 β dephosphorylation and increased GSK-3 β activity, which resulted in increased β -catenin phosphorylation and degradation [134]. Using galectin-3 knockout mice Mendonca et al [135] recently showed that galectin-3 was an important partner for GSK-3 β inactive form (phosphorylated at Ser9) to drive oncogenic transformation. We recently showed that Nup98 promotes nuclear export of galectin-3 [112], while its nuclear import is regulated by karyopherins [102, 103]. Depletion of Nup98 resulted in nuclear translocation of galectin-3, where it interacts with β -catenin and reduces its transcriptional activity [112]. Ferrazzo et al [136] however showed that nuclear localization of galectin-3 in adenocarcinoma of salivary gland may be related to a more aggressive behavior and did not seem to affect cyclin D1 expression by β -catenin. Thus, there seems to be a discrepancy on the relationship of nuclear galectin-3 to β -catenin activity, which could be explained by the observation of Kim et al [137]. A germline variation in galectin-3 gene at position 191 resulting in H⁶⁴ or P⁶⁴ was reported by us earlier [138]. Kim et al analyzed this variation in gastric cancer patients. Presence of H⁶⁴ and not P⁶⁴ galectin-3 enhanced nuclear accumulation of β -catenin as well as increased expression of Tcf-4 target genes such as fascin-1 and c-myc through augmented

promoter binding activity of Tcf-4 [137]. It is therefore important to first check the H/P⁶⁴ status of the cells before analyzing its collaboration with β -catenin.

A more detailed understanding of the mechanisms by which galectin-3 augments Wnt signaling may facilitate the development of chemo-preventive and therapeutic strategies for various cancers such as colorectal, prostate, breast, multiple myeloma and acute myeloid leukemia, where its regulatory roles are well established.

Galectin-3: a therapeutic target?

Numerous studies have focused on the molecular mechanisms of galectin-3 involved in cancer cell chemo-resistance, which have been reviewed in details [139-142]. A number of investigators have used forced expression of galectin-3 or its down regulation or by anti-sense treatment to study its effects on drug resistance. While some other groups have used a more direct approach to show the role of galectin-3 in chemo-resistance. Cheng et al [143] reported an up-regulation of galectin-3 in cisplatin and LY294002 (a phosphatidylinositol 3-kinase inhibitor) surviving chronic myeloid leukemia cells. These authors also reported a direct correlation between GSK-3 β and galectin-3. Mazurek et al [144] showed an up-regulation of galectin-3 in TRAIL resistant sub-population of metastatic colon cancer cell line LS-LiM6. Silencing of galectin-3 restored TRAIL sensitivity. It was reported that galectin-3 impedes trafficking of death receptor by anchoring them in glycan nano-clusters, blocking the execution of the apoptosis signal caused by TRAIL [144]. Galectin-3 silencing was also reported to augment gemcitabine and cisplatin-induced apoptosis in pancreatic cancer cell lines and inhibited migration and invasion through degradation of beta-catenin [145, 146]. Lin et al [147, 148] used a highly specific small molecule inhibitor of galectin-3 (Td131_1) and showed a synergistic activity with doxorubicin in papillary thyroid cancer cell lines 8505-C and TPC-1. In chronic myelogenous leukemia cells enforced expression of galectin-3 activated Akt and Erk, induced accumulation of Mcl-1 and promoted multi-drug resistance to tyrosine kinase inhibitors, doxorubicin, cytarabine, etoposide, and vincristine as a result of impaired apoptosis induction [149].

The vast majority of anti-cancer drugs currently used act by inducing apoptosis via the intrinsic pathway. Numerous mechanisms underlie cancer chemo-resistance [150]: galectin-3 appears to suppress cell apoptosis and hence, decreases sensitivity of cancer cells to chemotherapeutic drugs [151]. Several investigators have attempted to re-sensitize the cells to chemotherapeutic drugs by targeting galectin-3. Modified citrus pectin (MCP) was reported to inhibit galectin-3 functions by inhibiting its interactions with its glyco-conjugate ligands [152-154]. Johnson et al [155] showed that galectin-3 targeting via MCP or via lactosyl-L-leucine (LL) decreased malignant endothelial cell proliferation by themselves and sensitized these cells to the cytotoxic effect of doxorubicin. Treatment of metastatic cells MDA-MB-435 with MCP and LL also increased their sensitivity to taxol [151]. MCP/GCS100 also induced calpain activation in prostate cancer cells that led to their sensitization to cisplatin treatment [156]. In multiple myeloma cells an increased efficacy of bortezomide and dexamethasone on apoptosis in the presence of MCP/GCS100 was observed [157]. In a recent study, Lee et al demonstrated that inhibition of galectin-3 enhanced the efficacy of anticancer drug epirubicin in colon cancer caco-2 cells. Galectin-3 knockdown increased the

intracellular accumulation of epirubicin, suppressed the mRNA and protein expression of b catenin, cyclin d1, c-myc, p-glycoprotein, MDR-associated proteins 1 and 2 and increased the mRNA levels of GSK-3 β , Bax, caspase3 and caspase9 indicating that silencing of galectin-3 sensitizes the MDR cells to epirubicin by inhibiting ABC transporters and activating the mitochondrial pathway of apoptosis through modulation of b catenin/GSK-3 β pathway in human colon cancer cells [158].

Concluding remarks

There is ample evidence now to indicate the efficacy of galectin-3 as a therapeutic target. Due to its unique chimeric structure, two main types of interactions of galectin-3 have been reported. Some of the interactions occur via its carbohydrate binding domain, which can be interrupted by specific sugar competitive molecules such as MCP/GSC100 or small sugar analogs. A majority of its intracellular interactions occur via protein-protein interactions and are not inhibited by lactose.

Nuclear transport of galectin-3 is phosphorylation dependent, although, the domain responsible for the nuclear transport is debatable. However, it has been confirmed by various investigators that the phosphorylated galectin-3 is instantly exported to the cytoplasm, where it protects the cells from drug induced apoptosis [99]. The galectin-3 nuclear export proceeds via leptomycin-inhibitable pathway (155). Although no studies as yet have targeted the nuclear transport of galectin-3 as a cancer preventive mechanism, it would be an interesting approach. Prevention of galectin-3 phosphorylation will prevent its nuclear export, but it will not be an easy task because of lack of a specific inhibitor. Furthermore, disruption of β -catenin and galectin-3 interactions may also prove useful. Another important aspect to investigate while using galectin-3 as a therapeutic target is the germline mutations in galectin-3. Presence of H⁶⁴ in galectin-3 is associated with a more malignant phenotype. This variant is sensitive to cleavage by MMP-2 and -9, is associated with increased breast cancer incidence and also shows nuclear localization and increased β -catenin activity compared to the P⁶⁴ variant. In summary the precise regulatory mechanisms of galectin-3 expression in different cell types under different physiological and pathophysiological conditions need to be analyzed, which may help in development of new strategies to fight cancer.

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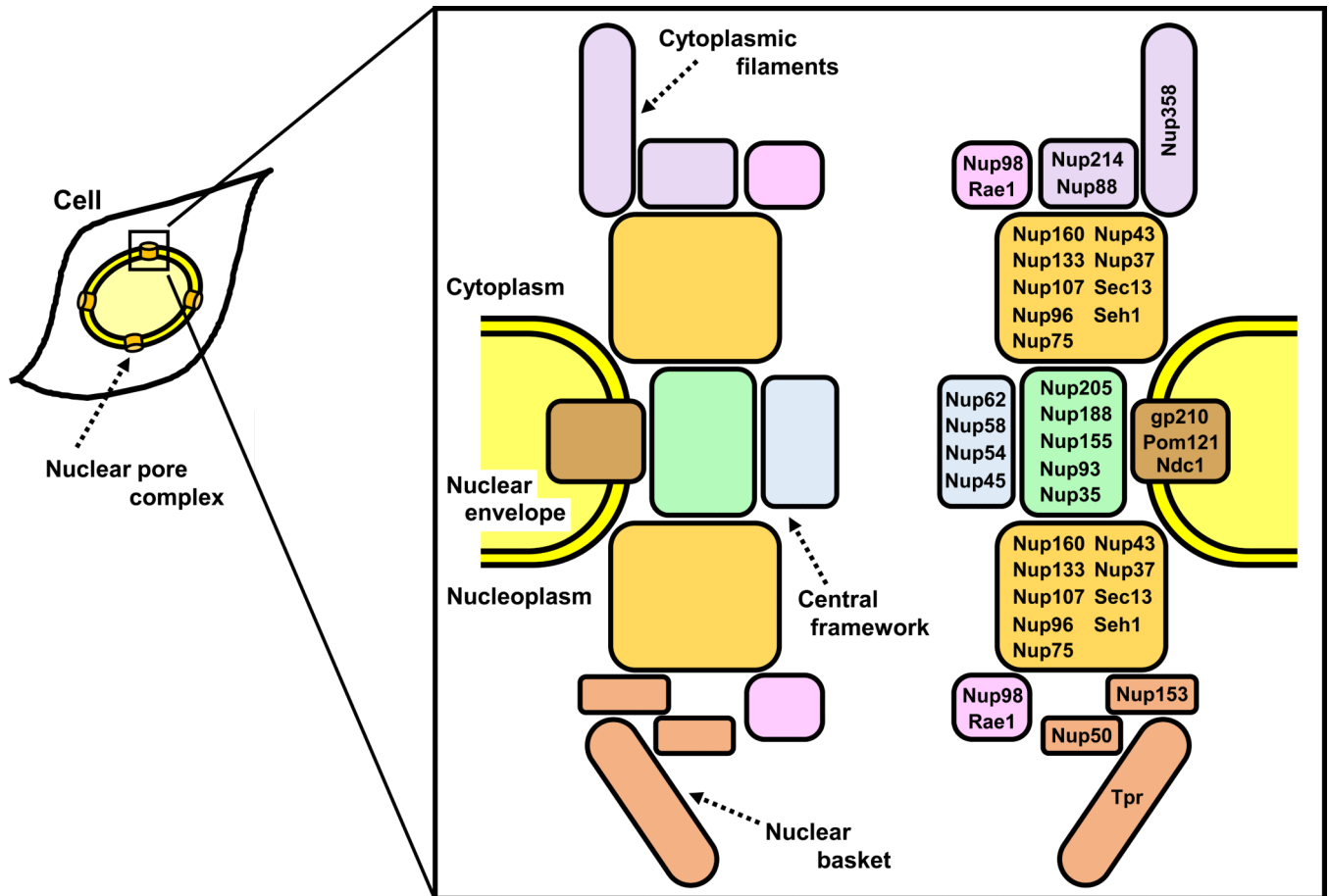


Figure 1.

Nuclear pore complex (NPC) structure and nucleoporins. In the box, schematic representation of the main structural components of NPC (left) and subcomplexes of nucleoporins within NPC (right) are shown. NPC essentially consists of cytoplasmic filaments, a central framework and a nuclear basket. Nucleoporins can be subdivided into different subcomplexes; Nup214 complex (Nup214, Nup88), Nup107-160 complex (Nup160, Nup133, Nup107, Nup96, Nup75, Nup43, Nup37, Sec13, Seh1), Nup62 complex (Nup62, Nup58, Nup54, Nup45), Nup93-205 complex (Nup205, Nup188, Nup155, Nup93, Nup35) and Nup98 complex (Nup98, Rae1).

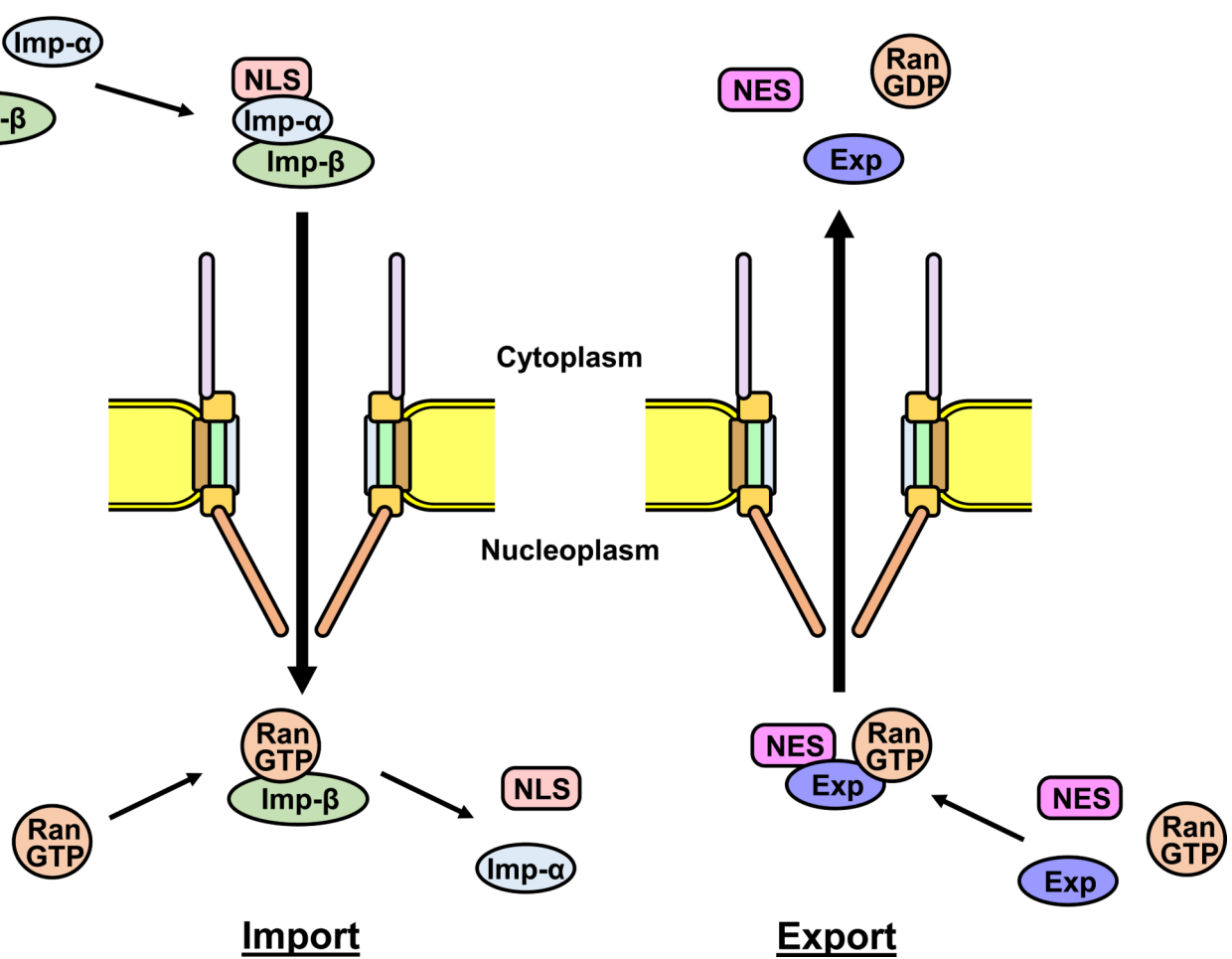


Figure 2.

Nucleocytoplasmic transport pathways through NPC. Karyopherin functions in nuclear import (importins) or nuclear export (exportins) are shown. Left: Importin- α recognizes and binds nuclear localization signal (NLS)-containing cargos. The importin- α forms a heterodimer with importin- β . The import complex docks at the NPC, and mediates import of cargos through NPC. In the nucleus the import complex encounters RanGTP, is disassembled and releases the cargos. Right: Exportin also recognizes and binds nuclear export signal (NES)-containing cargos in the nucleus in the presence of RanGTP. The export complex docks at the NPC, translocates to the cytoplasm and releases the cargos. Imp- α : importin- α ; Imp- β : importin- β ; Exp: exportin.

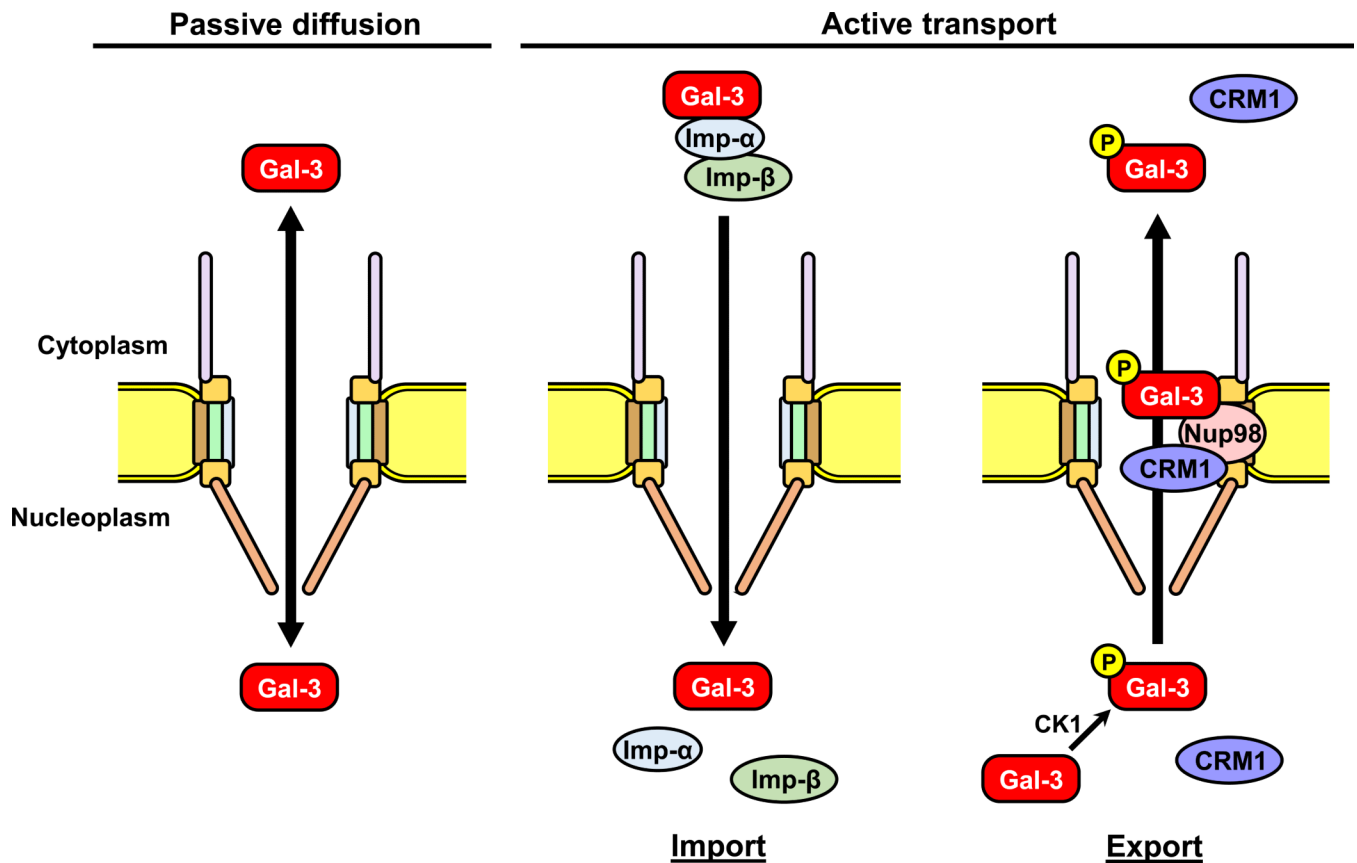


Figure 3. Schematic diagram of galectin-3 nucleocytoplasmic transport pathways. Galectin-3 is possibly translocated between nucleus and cytoplasm by both passive diffusion and active transport. During active nuclear transport, galectin-3 NLS binds to importin- α , followed by binding to importin- β in the cytoplasm. The galectin-3–importin- α/β complex docks at nucleoporins and enters the nucleus. The complex dissociates in the nucleus, releasing the galectin-3. In active transport into the cytoplasm, Nup98 plays an essential role in the galectin-3 export, with nuclear export protein CRM1 (exportin 1). Phosphorylation of galectin-3 by casein kinase 1 is required for the export. Gal-3: galectin-3; CK1: casein kinase 1.