



Immune Evasion Mechanism and AXL

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Extensive interest in cancer immunotherapy is reported according to the clinical importance of CTLA-4 and (PD-1/PD-L1) [programmed death (PD) and programmed death-ligand (PD-L1)] in immune checkpoint therapies. AXL is a receptor tyrosine kinase expressed in different types of cancer and in relation to resistance against various anticancer therapeutics due to poor clinical prognosis. AXL and its ligand, i.e., growth arrest-specific 6 (GAS6) proteins, are expressed on many cancer cells, and the GAS6/AXL pathway is reported to promote cancer cell proliferation, survival, migration, invasion, angiogenesis, and immune evasion. AXL is an attractive and novel therapeutic target for impairing tumor progression from immune cell contracts in the tumor microenvironment. The GAS6/AXL pathway is also of interest immunologically because it targets fewer antitumor immune responses. In effect, several targeted therapies are selective and nonselective for AXL, which are in preclinical and clinical development in multiple cancer types. Therefore, this review focuses on the role of the GAS6/AXL signaling pathway in triggering the immunosuppressive tumor microenvironment as immune evasion. This includes regulating its composition and activating T-cell exclusion with the immunesuppressive activity of regulatory T cells, which is related to one of the hallmarks of cancer survival. Finally, this article discusses the GAS6/AXL signaling pathway in the context of several immune responses such as NK cell activation, apoptosis, and tumor-specific immunity, especially PD-1/PDL-1 signaling.

OPEN ACCESS

Edited by:

Diana Boraschi, Shenzhen Institutes of Advanced Technology (CAS), China

Reviewed by:

Douglas Graham, Emory University, United States Carlo Cenciarelli, National Research Council (CNR), Italy

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Specialty section:

This article was submitted to Pharmacology of Anti-Cancer Drugs, a section of the journal Frontiers in Oncology

> Received: 10 August 2021 Accepted: 08 October 2021 Published: 28 October 2021

Citation:

Son H-Y and Jeong H-K (2021) Immune Evasion Mechanism and AXL. Front. Oncol. 11:756225. doi: 10.3389/fonc.2021.756225 Keywords: PD-1/PD-L1, Gas6/Axl signaling, immune checkpoint, immune evasion, tumor microenvironment (TME), AXL

THE GAS6/AXL SIGNALING PATHWAY

Like all TYRO3, AXL, and MER (TAM) receptors, AXL is composed of two immunoglobulin-like domains, two fibronectin III (FN III) domains, a transmembrane domain, and an intracellular kinase domain (1). GAS6 is a ligand for TAM receptors, with the highest affinity for AXL (2). The gamma-carboxy glutamic (GLA) regions of GAS6 have four epithelial growth factor (EGF)-like domains and modules similar to C-terminal sex hormone-binding globulin (SHBG) that are required to activate TAM receptors. These GLA regions bind to phospholipid phosphatidylserine (3) tethered to the extracellular surface of apoptotic cells or displayed on the outer parts of photoreceptors. Phosphatidylserine stabilizes the interaction between TAM and its ligands such as GAS6 by increasing the binding affinity and slowing the rate of GAS6 dissociation from receptors

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(4). GAS6 activates all three receptors with different affinities (AXL > TYRO3 >>> MER) based on the ability of each ligand to activate TAM receptors (5). Upon GAS6–AXL interaction, the complex dimerizes with another GAS6–AXL complex to form a 2:2 homodimerized complex with no direct AXL/AXL or GAS6/GAS6 contacts (6). Studies have demonstrated that genetic and pharmacological inhibition of AXL affects downstream signaling pathways, which include JAK-STAT (Janus kinase/signal transducers and activators of transcription), PI3K-AKT (phosphatidylinositol 3-kinase), AKT (protein kinase B), and RAS-RAF-MEK-ERK (Rat sarcoma virus-Rapidly Accelerated Fibrosarcoma-Mitogen-activated protein kinase-Extracellular signal-regulated kinase) (7, 8).

GAS6 AND AXL EXPRESSION IN THE TUMOR MICROENVIRONMENT

The tumor microenvironment changes continuously during cancer progression by regulating oncogenic signals such as secreted factors and tumor-promoting cells to induce construction of tumor cells' own niche (9). While AXL expression in tumors is readily recognized, it is less well known that AXL is expressed by various cells found in the tumor microenvironment, which include several immune cell types (10), fibroblasts (11), osteoclasts (12), and endothelial cells (13– 15). Furthermore, the unique tumor microenvironmental conditions may modulate AXL and GAS6 expression in both tumor and immune cells to promote aggressive and protumorigenic features. The tumor microenvironment can regulate AXL expression in various cells, and AXL seems to have a potential role in tumor development, progression, and metastasis.

AXL in Host Cells

In endothelial cells, AXL expression is involved in vascularization; i.e., when it is inhibited in tumor-bearing mice, it leads to the inhibition of tumor-induced angiogenesis (16–19). The interactions between the tumor and host immune cells in the tumor microenvironment can induce the expression of AXL and GAS6 to promote a cancerous microenvironment. Tumor cells may induce the expression of AXL and GAS6 in monocytic myeloidderived suppressor cells (M-MDSCs) and polymorphonuclear myeloid-derived suppressor cells (PMN-MDSCs) (20). Moreover, generally, AXL is expressed on bone marrow-derived cells (21–24), dendritic cells (DCs) (25, 26), macrophages (27, 28), monocytes (23), natural killer (NK) cells (29), and platelets (30).

GAS6 in Host Cells

GAS6 is expressed by luminal progenitor and basal cells around the ductal lining of mammary tissue (31). In the bone microenvironment, GAS6 is secreted by osteoblasts, which are involved in bone formation (32, 33). It was demonstrated that osteoblast-derived GAS6 induces AXL expression in tumor cells (34), which suggests that paracrine GAS6/AXL signaling promotes survival, inhibits apoptosis, and mediates homing of tumor cells to the bone. In the tumor microenvironment, cancer-associated fibroblasts (CAFs) and CD45-expressing tumor-infiltrating leukocytes (TILs) express GAS6 (35–38), and CD45+ cells from the bone marrow or peripheral blood express significantly less GAS6 than TILs (38). Besides, macrophages and dendritic cells express high levels of GAS6 (37, 38), which can be further promoted by various cytokines (36). Especially in macrophages, *in vitro* studies suggested that tumor cells or tumor cellconditioned media induce GAS6 expression and secretion (37, 39). Stromal cell-derived GAS6 was also shown to promote tumor cell migration, invasion, survival, and proliferation (36, 37). Potential downstream effectors of GAS6/AXL signaling through macrophage-derived GAS6 include pAKT and pSTAT3 (37).

AXL-MEDIATED TUMOR-SPECIFIC IMMUNE RESPONSE

AXL Changes Tumor Immune Microenvironment Components

During the past decades, modulating immune responses has been considered a tremendous potential therapeutics to treat cancer. Each patient's tumor immune microenvironment (TIME) seems to be related to this treatment responsiveness. It is becoming clear that both intrinsic and extrinsic factors modulate the composition of the TIME. Specifically, several immune cells in TIME have been reported to support tumor cells' survival through immune-suppressive functions (40, 41). Furthermore, cancer cells alter the expression of cell surface molecules to avoid detection by residential immune cells.

Several studies have revealed that GAS6/AXL signaling plays a vital role in promoting the immune-suppressive tumor microenvironment. Firstly, this signaling alters the expression level of several factors, including major histocompatibility complex I (MHC-I) and programmed death ligand-1 (PD-L1) in neoplastic cells (42). However, detailed changes are different depending on cell types and research conditions. Lung adenocarcinoma cell lines (PC9 and H1975) subjected to AXL inhibition by either bemcentinib or BGB324 significantly decreased PD-L1 (42). Pharmacologic AXL inhibition using a selective AXL inhibitor (R428 or SGI-7079) in tumor cells of C57BL/6 mice significantly increased the expression of PD-1 and MHC-I molecules (43). TAM knockout mice had increased MHC-I expression of myeloid cells. In the MCF10A cells, overexpression of TAMs did not increase PD-L1 expression, but in the PD-L1-expressing MDA-MB-231 cells, treating GAS6 liposomes increased PD-L1 expression and induced AXL phosphorylation (42). Hence, further studies are warranted to understand the detailed mechanism.

Next, the GAS6/AXL signaling pathway is involved in the recruitment of both myeloid and lymphoid lineage cells, which are involved in innate and adaptive immune responses, respectively (40). It promotes the secretion of immunosuppressive cytokines, including CCL3-5, G-CSF, and TGF β (44, 45), that are involved in the infiltration of several immune cells [macrophages and myeloid-derived suppressor cells (MDSCs)], which make it possible to escape immune surveillance (42) (**Figure 1**). Myeloid-derived suppressor

cells have a certain role regarding angiogenesis, cell invasion, metastasis, and suppression of CD8+ T cells (46, 47), and the number of these cells is reduced along with AXL knockdown (48). Lymphoid lineage cells including T cells, B-cells, and NK cells are increased when pharmacologic and genetic inhibitions of AXL are treated to cancer cells (42) (**Figure 2**). However, detailed parts are still to be further demonstrated because the number of tumor-infiltrating CD8+ T-cells is increased after AXL inhibition (49), while the other research showed that AXL inhibition does not affect the number of them (21, 46, 50).

AXL Contributes to T-Cell Exclusion

In addition to regulating TIME components, AXL also regulates T cell activity at different points. Usually, this helps control the excessive inflammatory response to protect normal cells; however, tumor cells take advantage of this protective mechanism by eliminating the T cell immune response toward cancer cells (51). In the study, not only the number of CD4+ and CD8+ T cells significantly increased, but also the gene expression associated with type 1 T-cell recruitment and functionality enhanced when AXL is inhibited in R428-treated tumor-bearing mice (43). Mainly, AXL receptor tyrosine kinase plays a particular role in T cell exclusion. AXL increases tumor cell

invasion and metastasis by promoting T cell exclusion, acting as an inducer of tumor cell plasticity (52, 53). Genetic deletion of AXL resulted in up to 20-fold enhanced T-cell infiltration and sensitization of tumor cells to radiotherapy and checkpoint immunotherapy of a transgenic mouse model (50).

Relationship Between AXL and Programmed Death 1

One of the immune checkpoints that are related to T cells is programmed death 1 (PD-1) and its ligand programmed cell death ligand 1 (PD-L1) (54). PD-L1 is expressed in several tumor cell types, and the interaction between PD-L1 and its receptor activates signaling pathways to prevent T-cell activation (55). Specifically, the expression of PD-L1 can serve as a dynamic mechanism for escaping host immune responses (56). For instance, neoplastic cells expressing PD-L1 have been reported to avoid cell death and continue to proliferate in the tumor microenvironment (42).

This PD-L1 immune checkpoint strongly interacts with AXL as AXL inhibition affects the PD-L1 pathway and activates the antitumor effect (43). When AXL is suppressed, the level of PD-L1 is decreased in lung adenocarcinoma and human triple-negative breast cancer cell lines (57, 58). However, tumor-



FIGURE 1 | The mechanisms of TAM (in particular AXL) regulation of immune evasion. (A) Modulation of the tumor-immune microenvironment: promotes secretion of various immunosuppressive chemokine. After AXL inhibition, cytokines are decreased (CXCL9, CXCL10, CXCL11), increased (CCL-2, CCL-3, CCL-4, CCL5, CXCL1, CXCL2, CXCL5), or there could be no difference (CXCL12). These changes lead to regulation in the recruitment of specific immune cells (monocytes, macrophages, CD8+ T cells, NK cells). (B) Promoting T cell exclusion. AXL expression is significantly correlated with the expression of genes encoding CXC chemokine receptor 6 (CXCR6) and PD-L1 which prevent T cell activation. Tumors treated with the combination of pharmacological inhibition of AXL and anti-PD-1 presented an increased number of CD8 T cells. (C) Enhancing the immune suppressive activity of regulatory T cells (Tregs). Tumor-specific Tregs can suppress antitumor immune responses against a broad range of tumor antigens, even after being activated by just one tumor-associated antigen. GAS6 induces CD4+ CD25+ Tregs to express CTLA-4 and Foxp3 especially with AXL. These activated Tregs increase the consumption of IL-2 or suppression of IL-2 production to block the activity of T lymphocytes. (D) TAM signaling is involved in the overall stage of NK cell differentiation. Especially, the GAS6/AXL pathway promotes FLT3 ligand-induced human NK cell development and cooperative interaction between the GAS6/AXL pathway and IL-15 signaling promotes NK cell differentiation. In the absence of AXL, IL-15 failed to activate several downstream signaling pathways, including PI3K, AKT, and ERK1/2. (E) Suppressing the genes encoding the cytokine suppressors SOCS1 and SOCS3. In case of DCs, they serve as "presentation platforms" for GAS6 which triggers STAT1-dependent cascade with type I IFN. TAM also activates phagocytosis of DCs and macrophages to clear apoptotic cell corpses which could induce immune responses. (F) Inhibit apoptosis. The GAS6/AXL pathway is important i



effectors including pAkt and pStat3 that lead to several changes in the tumor microenvironment and immune system. 1) Modulate the expression of MHC-1 and PD-L1. 2) Regulate the secretion of chemokines that are involved in recruiting several immune cells including monocytes, macrophages, and CD8+ T-cells. 3) Promote NK cell development through IL-15 signaling. As interruption of the GAS6/AXL pathway resulted in a reduction of FLT3 phosphorylation, this pathway may induce differentiation of NK cells by positively regulating FLT3 activation. 4) Inhibit apoptosis and activate phagocytosis involving several other pathways including activating the Akt and PI3K pathways. 5) Increase the immune-suppressive effect of Tregs. GAS6-induced CTLA-4 and Foxp3 expression are abrogated by blocking AXL.

infiltrating CD8+ T and CD4+ T cells subjected to AXL inhibition showed a noticeable induction of PD-1 on their surface. This relationship seems to be a systemic long-term memory immune response to tumor antigens (43). Furthermore, combining pharmacological inhibition of AXL with anti-PD-1 in a preclinical model of breast cancer reduces the primary tumor and metastatic burden, which is not shown when only one of them is treated. Tumors treated with the combination of these two therapeutic agents presented an increased number of CD8 T cells, with more activation of the NK cells (59). Moreover, other studies have linked AXL/PI3k signaling with increased expression of PD-L1 by tumor cells, and AXL inhibition potentiates PD-1 blockade in ID8 graft models (43, 60). This interaction is also demonstrated in lung adenocarcinoma, which showed that AXL expression significantly correlated with the expression of genes encoding PD-L1 and CXC chemokine receptor 6 (CXCR6) (57). Therefore, AXL receptor kinase is highly related to the PD-L1 immune checkpoint, contributing to immune evasion.

This relationship between AXL and PD-L1 may involve several immune cells for reactions. The analysis of TAM expression within the activated lymphoid compartment revealed that MERTK, but not AXL or TYRO3, is expressed on activated B lymphocytes and CD4+ and CD8+ T cells (61). Therefore, after apoptotic cells display PD-L1, other immune cells such as dendritic cells or macrophages sense PD-L1 involving AXL (4), and this signaling could be transferred to T-cells to take action toward immune evasion.

REGULATORY T CELLS' IMMUNE EVASION WITH AXL

Modulation of Immune Evasion With the Immune-Suppressive Activity of Regulatory T Cells

Regulatory T cells (Tregs) mediate immune evasion, which is considered a major mechanism of escaping immune surveillance (62). Especially, tumor-derived Tregs have a relatively more effective suppressive activity than naturally occurring Tregs (63, 64). These Tregs are guided to the tumor microenvironment by tumor cell-mediated chemokine production (65, 66). After that, Tregs suppress many physical and pathological immune responses, which are crucial in sustaining self-tolerance and immune homeostasis (67).

Contribution of Treg to Immune Suppression Through Antigen

Naturally occurring Tregs are produced in the thymus and occupy 5%–10% of the total CD4 + T cells in the peripheral

blood, and induced Tregs derive from naive T cells under certain conditions (68). The functional importance of Tregs in a tumorbearing host is shown in murine models of melanoma in which depleting Tregs temporarily induce an immune response against tumors and improve tumor clearance (69).

The generation and maintenance of Tregs to regulate autoimmunity require target antigens and T-cell receptor activation (70). Natural and induced Tregs independently contribute to tumor-specific tolerance. In the case of naturally occurring Tregs, an extensive unrestricted $\alpha\beta$ repertoire specific for a broad range of self-antigens, including tumor-associated antigens, is required which implies that the cells exercise their effect in an antigen-non-specific manner (71). One study on mice showed that induction of antigen-specific Tregs from naive cells in the tumor microenvironment does not seem to be intrinsically related to naturally occurring Tregs (72). These induced Tregs seemed to profoundly inhibit T-cell responses against tumors in an antigen-non-specific manner after being activated by a specific antigen (67). This implies that tumor-specific Tregs can suppress both medically induced and naturally occurring antitumor immune responses against a broad range of tumor antigens, even after being activated by just one tumor-associated antigen (62).

Controlling the Effect of Tregs by GAS6, Especially With the AXL Receptor

The suppressive effect of Tregs is increased by GAS6 mainly through the AXL receptor (5). Comparing how much the GAS6-induced CTLA-4 and Foxp3 expression in CD4+ CD25+ Tregs is abrogated, blocking the AXL receptor was more effective than blocking TYRO3 and MER. In addition, GAS6 has a stronger binding affinity to AXL than TYRO3 and MER (2, 73). Therefore, GAS6 has a direct role in the functions of Tregs, which enhances the suppressive activity mostly with the AXL receptor. These interactions gain credence through IL-2, a potent T cell growth factor. When CD4+CD25-T cells pretreated with GAS6 protein are cocultured with CD4+CD25 +Tregs, the proliferative activity of T cells was significantly decreased with consistent suppression of IL-2. Also, the elevated expression of CTLA-4 and Foxp3 in Tregs after Gas6 stimulation was abrogated after Axl knockdown by siRNA, and this group also showed an IL-2 level increase. These results indicate that Gas6 enhances suppression of CD4+T cells by increasing Tregs' ability to consume IL-2 or suppress IL-2 production (74).

THE ROLE OF TAM IN NK CELL ACTIVATION FOCUSING ON AXL AND IL-15

The innate immune system is usually known to recognize pathogens directly, but it also senses and destroys cells infected with pathogens. This arm of the innate response is mainly conducted by NK cells (75). Hematopoietic stem cells (HSCs) from bone marrow differentiate into common lymphoid progenitors and develop into NK cells, followed by maturation into NK cells (76, 77). Several studies have shown that NK cells are related to tumor progression in several ways, including

immune evasion. Pre-metastatic niches are promoted by suppressing NK cell functions under hypoxia (78).

TAM Controls Natural-Killer-Cell Activation, Especially at the Differentiation Stage

TAM signaling plays a pivotal role in regulating the activity of NK cells (79). When NK cells are activated, they kill their targets by secreting the CD95 ligand and TNF-related apoptosis-inducing ligand (80-84), but NK cells from TAM-deficient mice have inferior cytotoxic activity (4). Mice lacking TAM possessed NK cells that have a defective function in both IFN- γ production and cytotoxicity. This activity impairment is proportional to how effectively TAM genes are inactivated as all three TAM receptors are expressed by immature NK cells in the bone marrow (79, 85). The number of NK cells generated from human CD34+ HPCs reduced after blocking GAS6 binding to AXL by AXL-Fc or warfarin (86).

After proving that TAM regulates NK cell activity, the specific stage that TAM is mainly involved in is considered. When immature cells are grown with NIH3T3 fibroblasts expressing GAS6, stromal cells can restore their ability to drive NK-cell maturation *in vitro*. Furthermore, the mice showed normal perforin and granzyme B levels even when NK cells lack all three TAM receptors (4). However, these cells do not secrete IFN γ , which is produced predominantly by NK cells and exhibit a 10-fold lower killing ability against target cells than wild-type NK cells after stimulation. This means that these cells do not fully demonstrate the expression of activation and inhibition receptors expressed by cytotoxic NK cells (4). Thus, TAM signaling is involved in the terminal stage of NK cell differentiation.

Importance of Interleukin-15 and AXL in NK Cell Activation

Interleukin-15 (IL-15) is another critical factor for NK cell development, which contributes to the differentiation, survival, and function of NK cells (87). When the mice are deficient in Interleukin-2 (IL-2) and IL-15 receptor, which is required only for the actions of IL-2 and IL-15 rather than other growth factors, NK cells are deficient. However, a normal level of NK cells is observed in mice deficient in IL-2 or IL-2R α , the private receptor used by IL-2, suggesting that IL-15 is an essential factor in the differentiation of NK cells from uncommitted progenitors (88). Furthermore, the upregulation of NK cell activity was markedly reduced by the addition of monoclonal antibodies to IL-15, but not by antibodies to other cytokines such as IFN- α , IFN- γ , TNF- α , TGF- β , and IL-2 (89). This evidence supports the suggestion that IL-15 is essential in the activation of NK cells.

Several studies have shown that the IL-15 and GAS6/AXL pathways partially overlap intracellular signaling molecules (90). In the absence of AXL, IL-15 failed to activate several downstream signaling pathways, including PI3K, AKT, and ERK1/2 (91).

Enhancing the Role of AXL and IL-15 Through FMS-Like Tyrosine Kinase 3

The link between IL-15 and AXL could be extended to FMS-like tyrosine kinase 3 (FLT3), one of the receptor tyrosine kinases (RTK).

Mice with genetic disruption at the FLT3 locus presented a reduction of the numbers of B-lymphoid progenitors, dendritic cells, and NK cells (92). The FLT3 ligand can enhance NK cell differentiation in the presence of IL-15 (93). To activate the FLT3 and its ligand (FL) pathway, the phosphorylation of FLT3 is essential, which follows the binding of its ligand FL (94, 95). Interruption of the GAS6/AXL pathway resulted in a marked reduction of FLT3 phosphorylation even in the presence of FL. This implies that the GAS6/AXL pathway promotes FL-induced human NK cell development by positively regulating FLT3 activation (96) (**Figure 1**). As FLT3 can induce differentiation of NK cells with IL-15 and the interaction between AXL and FLT3 is demonstrated, AXL and IL-15 could be highly related to each other in various immune cell activation steps.

REGULATION OF INNATE IMMUNE CELLS BY AXL

In addition to its role in NK cell activation, TAM has another role in regulating the innate immune response. TAM signaling is generally activated by Toll-like receptor (TLR) and type I interferon signaling, which is part of the innate inflammatory response in dendritic cells (DCs) and macrophages (5). The AXL receptor was found to be upregulated when DCs are cultured with type I IFNs (10, 26, 97). Additionally, the co-expression of TAM and type I IFN receptors in macrophages and dendritic cells (DCs) inhibits the inflammatory response of the innate immune system through induction of the genes encoding the cytokine suppressors SOCS1 and SOCS3. More specifically, the binding of the apoptotic cells to DCs is immunosuppressive and serves as "presentation platforms" for GAS6. This triggers a STAT1-dependent cascade with type I IFN, and this signaling also induces SOCS1 and SOCS3 expressions, which inhibit downstream signaling pathways of TLRs and cytokine receptors (5). Thus, the innate immune mechanism is proceeded and dependent on TAM receptors.

REGULATION OF IMMUNE EVASION BY AXL THROUGH APOPTOSIS AND PHAGOCYTOSIS

Several pieces of research have shown the relationship between AXL and apoptosis. Apoptosis produces materials that can induce an immune response, which should be prevented for tumor immune evasion.

Inhibition of Apoptosis by GAS6 With AXL Kinase

Apoptosis of vascular smooth muscle cells (VSMCs) has been identified in the physiological remodeling of the vasculature. Cell death with cell proliferation, migration, and matrix turnover may induce changes in vascular architecture during development and diseases such as atherosclerosis (98). This apoptosis of VSMCs is coupled with GAS6 binding to the AXL receptor, and GAS6 inhibits apoptosis in cultured VSMCs through AXL phosphorylation (99). GAS6 and AXL are increased after vascular injury, and these molecules play an essential role in neointima formation by suppressing apoptosis (100). Besides, it is speculated that the GAS6/AXL pathway is vital to limiting VSMC apoptosis by activating AKT and PI3K along with several other pathways, including phosphorylation of BAD (BCL2-associated agonist of cell death) and activation of ERK1/2 (101–103).

Non-small cell lung cancer (NSCLC), a prevalent and devastating disease, shows overexpression and activation of MER and AXL. MER or AXL knockdown also improved *in vitro* NSCLC sensitivity to chemotherapeutic agents by promoting apoptosis. Also, AXL inhibition induces apoptosis by abating survival signaling mediated by AKT, CREB, BCL-XL, and survivin (104).

Removal of Apoptosis Remains by Activating Phagocytosis

Furthermore, some cancers develop specific mechanisms to clear apoptotic cells to regulate immune responses. Defects in the clearance of apoptotic cells can induce an immune response. Macrophages and DCs must remove many apoptotic cells corpses, but this form of homeostatic phagocytosis is impaired in TAM-deficient mice (24, 105, 106). This immunosuppressive effect of DCs is exerted through TAM signaling, and this immune response may be reinforced, especially in cancer cells as they overexpress AXL. Hence, TAM contributes to immune evasion by activating phagocytosis to remove these remains.

However, there is a possibility that TAM signaling induced by apoptotic cells is autocrine which means macrophages and DCs themselves produce GAS6, not from the apoptotic cells. In this case, phosphatidylserine is the primary stimulant that stabilized the interaction between TAM and its ligands (4). Thus, further research is needed to understand the exact mechanism.

CLINICAL TRIAL SUPPRESSING AXL

Targeting AXL for cancer treatment is under the spotlight, and several clinical studies involving the use of anti-AXL have been conducted. The first clinical trial treating an anti-AXL-specific small molecule inhibitor called BGB324 was performed in 2013. BGB324 blocks auto-phosphorylation of AXL on the COOH-terminal multiple docking sites Tyr821 with the subsequent activation of AKT and SFK phosphorylation (107). After entering phase I clinical trials in 2013, it is currently under phase II study to assess the safety of BGB324 when given in up to 77 patients advanced adenocarcinoma of the lung previously treated with pembrolizumab (108). Since then, several newly synthesized inhibitors specific for AXL receptors are being tested at a clinical stage.

Many small-molecule inhibitors in the clinical stage do not solely target AXL. Some inhibitors work as AXL pathway modulators that target factors such as MET, TYRO3, and FLT3 along with AXL and are used with several immune checkpoints inhibitors (ICIs), including nivolumab, pembrolizumab, durvalumab, and avelumab (109).

Small molecules are being developed mainly, and other therapeutic agents such as a monoclonal antibody or nucleotide

TABLE 1 | AXL inhibitor under development.

Compound	Target	Phase of development	Sponsor
		< Small-molecule inhibitors > (113)	
SP2215 Gilteritinib)	AXL, FLT3, ALK	Phase 3 (acute myeloid leukemia)	Astellas Pharma.
GB324 R428)	AXL (selective)	Phase 2 (triple-negative breast cancer, lung cancer metastatic)	Rigel PharmaceuticalsBerGenBIO
MS-777607 ASLAN002)	AXL, RON, MET, TYRO3, MER, FLT3	Phase 2 (advanced solid tumors)	Aslan Pharma. and Inventive Health Clinical
P3975	AXL	Preclinical	Deciphera Biotech
SK1363089/ L880 Foretinib)	AXL, MET, VEGFR2, RON	Phase 2 (breast cancer/carcinoma, renal cell etc.)	GlaxoSmithKline
DC1267	MET, AXL, TYRO3	Preclinical	Lead Discovery Centre
. Y2801653 Merestinib)	MET, MSTIR, DDRI, TIEI, MER, TYRO3, AXL	Phase 2 (carcinoma, non-small-cell lung/active, not recruiting)	Eli Lilly and Co.
IGCD 265	MET, AXL, VEGFR2	Phase 2 (non-small cell lung cancer)Phase 2 (carcinoma, non-small-cell lung/active, not recruiting)	Mirati Inc.
/IGCD516 Sitravatinib)	MET, AXL, RET, TRK, DDR2	Phase 3 (non-small cell lung cancer, advanced or metastatic solid malignancies)	Mirati Inc.
IP-470 Amuvatinib)	KIT, PDGFR1, FLT3, RET, AXL	Phase 1 (solid tumors)Phase 2 (small cell lung carcinoma)	Astex Pharma.
IPS-1034	AXL, DDR1, FLT3, KIT, MEK, MET, ROS1, TIE1	Preclinical	NeoPharma
PF-02341066 Crizotinib)	ALK, MET, RON, AXL	Approved for non-small-cell lung cancerPhase 2 (solid tumors)	Pfizer, NCI, EORTC etc.
GI-7079	MET, MER, YES, RET, FLT3, AXL	Preclinical	Astex Pharma
KI-606 Bosutinibm)	BCR-ABL, ABL, SRC, YES, MEK, AXL, BMX	Phase 3 (chronic myeloid leukemia/recruiting)Phase 2 (breast neoplasms)	Pfizer
U11248	KIT, FLT3, PDGFR,	Approved for renal cell carcinoma, imatinib-resistant gastrointestinal stromal	Pfizer/Further clinical sponsors include NCI,
Sunitinib)	VEGFR2, AXL	tumor, and metastatic pancreatic neuroendocrine tumorsPhase 3 (different solid tumors)	Baylor Breast Care Center, M.D. Anderson Cancer Center, etc.
P-0903	JAK2, ALK, ABL, AXL, MER	Phase 2 (acute myeloid leukemia)	Huntsman Cancer Institute/Tolero Pharmaceuticals
INC2025	MER, FLT3, AXL, TYRO3	Preclinical	University of North Carolina
(L184 Cabozantinib)	VEGFR2, MET, MEK, KIT, RET, AXL	Approved (medullary thyroid cancer)Phase 2, 3 for different solid tumors	Exelixis
		< Receptor monoclonal antibody >	
W327.6S2	AXL	Preclinical (non-small cell lung cancer)	(111)
9	AXL	Preclinical (pancreatic cancer)	(114)
8	AXL	Preclinical (pancreatic cancer)	(114)
		< Nucleotide aptamer >	. ,
L21.T (RNA Datamer)	AXL	Preclinical (non-small cell lung cancer)	(111)
NA AXL- patamar	AXL	Preclinical (ovarian cancer)	(17)

aptamer are also in preclinical progress. The YW327.6S2 phagederived monoclonal antibody binds to human AXL with high affinity, which blocks the GAS6 binding to the receptor and downregulates receptor expression (110). Aptamers are short structured single-stranded RNAs or DNAs that bind to a specific target molecule. They are promising alternatives with great potentials because of their low cost, lower toxicity, and higher affinity (111, 112). A selective RNA-based aptamer, GL21.T, binds the extracellular domain of AXL at high affinity and inhibits its catalytic activity. This includes ERK and AKT phosphorylation and inhibited *in vivo* lung tumor formation in mouse xenografts (111) (**Table 1**).

AUTHOR CONTRIBUTIONS

H-YS contributed to the conception and design of the study. H-YS and H-KJ wrote sections of the manuscript. All authors contributed to the manuscript revision and read and approved the submitted version.

FUNDING

This work was supported by the grant no. 04-2021-0570 from SNUH Research Fund.

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