Immunosuppressive metabolites in tumoral immune evasion: redundancies, clinical efforts, and pathways forward

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ABSTRACT

Tumors accumulate metabolites that deactivate infiltrating immune cells and polarize them toward anti-inflammatory phenotypes. We provide a comprehensive review of the complex networks orchestrated by several of the most potent immunosuppressive metabolites, highlighting the impact of adenosine, kynurenines, prostaglandin E₂, and norepinephrine and epinephrine, while discussing completed and ongoing clinical efforts to curtail their impact. Retrospective analyses of clinical data have elucidated that their activity is negatively associated with prognosis in diverse cancer indications, though there is a current paucity of approved therapies that disrupt their synthesis or downstream signaling axes. We hypothesize that prior lukewarm results may be attributed to redundancies in each metabolites' synthesis or signaling pathway and highlight routes for how therapeutic development and patient stratification might proceed in the future.

INTRODUCTION

Tumors exhibit spectacular dexterity in evading immune response.¹ Cancer cells co-opt immune checkpoints, deactivating immune cells through receptor-ligand interactions and through the production of immunosuppressive metabolic byproducts. Within the last decade, seven checkpoint inhibitor antibodies have been approved by the Food and Drug Administration (FDA) to address suppressive receptor-ligand interactions between tumor and immune cells mediated by cytotoxic T-lymphocyte-associated antigen 4 (CTLA-4), programmed cell death 1 (PD-1), or programmed death-ligand 1 (PD-L1).² However, there are no therapies for cancer indications that address the activity of immunosuppressive metabolites, despite their contribution to tumorous immunosuppression and negative impact on patient prognosis. The synthesis and signaling pathways of the most potent immunosuppressive metabolites, including adenosine, kynurenine, prostaglandin E₂ (PGE₂), and norepinephrine and epinephrine are inherently redundant. Several metabolic enzymes or distinct

enzymatic pathways might catalyze their synthesis in the tumor microenvironment, and once created, they can agonize multiple receptors on immune cells. We hypothesize that such redundancies have been a key obstacle slowing development of effective pharmacological blockades against these metabolites.

To this end, we will review each immunosuppressive metabolite's synthesis and signaling networks, detailing how internal redundancies may contribute to difficulties developing effective therapeutic interventions. We will also highlight instances of interplay within and between these immunosuppressive networks that might be considered in future clinical and preclinical efforts. Finally, we will discuss several potential pathways forward, emphasizing broad-acting therapeutics, combination therapies, and biomarker-based patient stratification. We note that the immunosuppressive metabolites discussed here are not an exhaustive representation of those that can be found in the tumor. Other immunosuppressive metabolites like lactate and downstream catabolites of arginine, including nitric oxide and polyamines, have been reviewed elsewhere.^{3–6}

SYNTHESIS OF IMMUNOSUPPRESSIVE METABOLITES

In addition to mediating immune escape in cancer, adenosine, kynurenines, PGE₂, and norepinephrine and epinephrine have varied and widespread physiological roles, which may explain why redundant generation mechanisms exist for each. To wit, adenosine is an intermediate in nucleotide recycling that regulates sleep and influences cardiovascular plasticity.⁷ Regulation of the kynurenine pathway is essential to normal cerebral function, and its dysregulation has been linked to neurobiological and psychological diseases.⁸ PGE₂ influences hematopoiesis, neuronal signaling, and renal function,

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and norepinephrine and epinephrine (ie, epinephrine) are the classical ‘fight-or-flight’ hormones.^{9 10}

Adenosine: redundant enzymes and distinct biosynthetic pathways

Adenosine is a ubiquitous nucleoside that is primarily generated in the extracellular space. Normal levels are between 40 and 460 nM, but tumorous concentrations can reach 1–100 μM.^{11–13} In addition to cancer cells, several immune cells can synthesize adenosine, particularly tumor-resident Tregs, myeloid-derived suppressor cells (MDSCs), and tumor-associated macrophages (TAMs).¹⁴ Multiple metabolic pathways, each of which can employ multiple enzyme homologs, can produce adenosine.

The most studied pathway has two steps: hydrolysis of ATP to AMP via an ectonucleoside triphosphate diphosphohydrolase (ENTPDase), and then hydrolysis of AMP to adenosine by a 5′-nucleotidase (5′NTDase). The hypoxic tumor microenvironment promotes ATP release into the extracellular space by stressed, dead and dying cells, ensuring substrate availability. The membrane-anchored enzymes CD39 (ENTPDase) and CD73 (5′NTDase) have been broadly implicated in cancer and can be upregulated on cancer and immune cells, though numerous other ectoenzymes catalyze hydrolysis of ATP or AMP (figure 1).^{14 15} For example, alkaline phosphatases hydrolyze AMP into adenosine and are upregulated in certain cancers.^{16 17}

Adenosine can also be synthesized from nicotinamide adenine dinucleotide (NAD⁺) in two related extracellular metabolic pathways. In one, CD203a degrades NAD⁺ into nicotinamide mononucleotide and AMP, and in the other, CD38 converts NAD⁺ into nicotinamide and ADP ribose, which is then cleaved by CD203a into pyrophosphate and AMP.¹⁸ AMP can then be dephosphorylated into adenosine by a 5′NTDase, an AP, or the tartrate-resistant acid phosphatase enzyme.¹⁹ Connexin 43 allows for intracellular NAD⁺ to traverse the cell membrane.²⁰

Similarly, adenosine can be synthesized from cyclic AMP (cAMP) through two metabolic pathways (figure 1). In one, 10 adenylate cyclase isoforms catalyze the conversion of extracellular ATP to cAMP, and in the other, intracellular cAMP is secreted via the multidrug resistance proteins 4, 5, and 8 (MRP4,5,8).²¹ cAMP is converted to AMP through an ectophosphodiesterase (PDE) (likely to be an isoform from one of the PDE super families 4,7,8,10,11), and AMP is dephosphorylated into adenosine.²² Extracellular AMP may instead be deaminated by AMP deaminase to form inosine monophosphate, which can be dephosphorylated by CD73 to form inosine.²³ Adenosine deamination, catalyzed by either adenosine deaminase (ADA) 1 or 2, also yields inosine.²⁴

Adenosine synthesis can occur through multiple distinct pathways, in which each enzymatic step can be catalyzed by multiple enzymes. While the relative contribution of each pathway or enzyme is currently unknown, this ‘synthesis redundancy’ could make inhibiting tumorous production of adenosine a challenge.

Kynurenines: indoleamine 2,3-dioxygenase and tryptophan 2,3-dioxygenase redundancies

Ninety-five per cent of the essential amino acid tryptophan is catabolized through the kynurenine pathway (figure 1).²⁵ Tumorous metabolic reprogramming upregulates expression of the initial and rate limiting step of the kynurenine pathway, the oxidation of tryptophan to N-formyl L-kynurenine, skewing the pathways’ outputs toward production and secretion of kynurenine.²⁶ Tryptophan oxidation is catalyzed by any of three distinct enzymes, all of which promote cancer progression and correlate with poor patient outcomes: (1) interferon γ (IFN-γ)-induced indoleamine 2,3-dioxygenase (IDO1), which is normally expressed in the peripheral lymph organs, colon, and epididymis; (2) IDO2, a less catalytically active isoform of IDO1, that is partly induced by IFN-γ and is normally expressed by some antigen-expressing cells, and cells of the liver, kidney, and brain; or (3) tryptophan 2,3-dioxygenase (TDO), an enzyme normally only present in the liver and brain.^{27–30}

Formamidases then rapidly convert N-formyl L-kynurenine to kynurenine, which, in normal tissue, can proceed through intermediates before yielding the end products xanthurenic acid, and to a much lesser extent, NAD⁺ (figure 1).^{31 32} However, cancer and other cells in the inflammatory tumor environment secrete kynurenine into the extracellular space via the L-type/large neutral amino acid transporter 1 (LAT1) tryptophan/kynurenine antiporter and other transporters, where it can then be imported into T cells by LAT1, LAT2, or Proton-Coupled Amino Acid Transporter 4 (figure 2).^{33–35} Normal serum kynurenine levels are between ~1 and 3 μM.³⁴ While few measurements of kynurenine levels in the tumor microenvironment have been performed, concentrations surpassing ~37 μM have been measured in glioblastoma cell culture supernatant.^{34 36} MDSCs, TAMs, neutrophils, endothelial cells, mesenchymal stem cells, and dendritic cells (DCs) can also secrete kynurenine.²⁶

A metabolomic and transcriptomic analysis of 928 cell lines in the Cancer Cell Line Encyclopedia revealed that IDO1, IDO2, and TDO mRNA expressions levels are predictive of a cancer cell’s ability to produce and secrete kynurenine.³⁷ Of the cell lines that highly secreted kynurenine, roughly one-third expressed both IDO1 and TDO, one third expressed only IDO1, and one third only TDO, a heterogeneous expression pattern that was confirmed by transcriptomic data in The Cancer Genome Atlas (TCGA). Such heterogeneity could imply difficulty toward inhibiting only one enzyme as a method to prevent kynurenine synthesis.

Kynurenic acid, another potential immunosuppressive metabolite, is produced through the transamination of kynurenine by four kynurenine aminotransferases (KATs 1–4; figure 1).³⁸ All four KATs have been shown to be upregulated in solid and blood cancers, and urine kynurenic acid level is a biomarker for prostate cancer.^{38 39} Quinolinic acid, another immunosuppressive metabolite, is a downstream metabolite of kynurenine

Figure 1 The redundant synthesis networks of immunosuppressive metabolites. Kynurenine pathway (left—green). After being imported into a cell by amino acid transporters, tryptophan is oxidized by one of three enzymes—IDO1, IDO2, or TDO2—to n-formyl-L-kynurenine, which is then converted to kynurenine by formamidases. Kynurenine may exit the cell through the LAT1, which simultaneously imports tryptophan, or continue down the kynurenine pathway until converted into xanthurenic acid (XANA) (major route) or nicotinamide adenine dinucleotide (NAD⁺) (minor route). Four different kynurenine aminotransferase (KAT) enzymes can transaminate kynurenine into kynurenic acid. Alternatively, kynurenine can be converted into 3-hydroxy-kynurenine by kynurenine 3-hydroxylase (K-3-H), or into anthranilic acid by kynureninase. 3-hydroxy-kynurenine may be converted to XANA (by KATs) or 3-hydroxyanthranilic acid (3-OHA) by kynureninase. 3-OHA may also be produced from anthranilic acid by 3-hydroxyanthranilic acid 3,4-dioxygenase (3H-3,4-D), and quinolinic acid phosphoribosyl transferase mediates the conversion of quinolinic acid to NAD⁺. XANA and kynurenic acid can be transported out of the cell by the OAT1 and OAT3, while connexin 43 allows NAD⁺ transport. Adenosine synthesis pathways (top center—yellow): adenosine synthesis occurs extracellularly in the tumor microenvironment and can use either ATP or NAD⁺ as a pathway substrate. ATP (or ADP to a lesser extent) can be dephosphorylated by CD39 or 6 other NTPDases into AMP. NAD⁺ can be converted to AMP directly by CD203a or through an ADP ribose (ADPR) intermediate by CD38 (followed by conversion to AMP by CD203a). AMP can also be generated from cAMP. cAMP is synthesized from protein G_s-activated adenylate cyclase from ATP. Intracellular cAMP can be excreted through multidrug resistance proteins 4, 5, or 8 into the extracellular space where a family of ecto-phosphodiesterases convert cAMP into AMP. AMP is finally dephosphorylated to adenosine by six possible enzymes, most prominently CD73. Adenosine can be deaminated to produce inosine by adenosine deaminase 1 (ADA) (membrane-attached via CD26) or 2. AMP can also be deaminated into IMP by AMP deaminase, and IMP can be dephosphorylated by CD73 to generate inosine (not shown). (Nor)epinephrine synthesis pathways (bottom center—magenta): schematic adapted from *Molinoff and Axelrod*. Epinephrine synthesis begins with tyrosine (TYR) and typically proceeds through L-dopa, dopamine (DOP), and norepinephrine (NOR) intermediates, though a parallel biosynthetic pathway exists. (Nor)epinephrine are exported through an undetermined mechanism. Prostaglandin E2 synthesis pathways (right—Blue): phospholipase A2s cleave AA from the inner leaflet of the cell membrane. Free AA is converted to PGH2 by the cyclooxygenase enzymes (COX-1, COX-2, and in the brain, COX-3). PGH2 is then converted to PGE2 by mPGES-1, mPGES-2, or cPGES (or p23), though PGH2 is also a precursor for several other prostaglandin derivatives (indicated by multiple reaction arrows stemming from PGH2 into the cytosol). PGE2 can also be generated when 8-iso-PGE2 undergoes epimerization. PGE2 is exported by MRP 1, MRP 2, or MRP 4. All proteins are denoted by text within shapes while metabolites are free-floating text. Straight lines indicate an enzyme catalyzed reaction, curved lines indicate transport or are drawn for clarity, and curved dotted lines indicate an undetermined transport mechanism. Stacked proteins indicate the existence of more than one enzyme capable of performing the indicated metabolic reaction. Up to 19 phospholipase enzymes exist in mammals, but only 1 is shown for clarity. 3-OHK, 3-hydroxykynurenine; AA, arachidonic acid; AAD, aromatic acid decarboxylase; ADO, adenosine; ANTHA, anthranilic acid; cAMP, cyclic AMP; CD, cluster of differentiation; CFE, catecholamine-forming enzyme; cPGES, cytosolic prostaglandin E synthase; DBH, dopamine-beta-hydroxylase; EPI, epinephrine; GCAP, germ cell alkaline phosphatase; IDO, indoleamine 2,3-dioxygenase; IAP, intestinal alkaline phosphatase; IMP, inosine monophosphate; INO, inosine; KYNA, kynurenic acid; KYN, kynurenine; KYNase, kynureninase; LAT, L-type/large neutral amino acid transporter; mPGES, microsomal PGE synthase; MRP, multidrug resistance-associated protein; NFL KYN, N-formyl-L-KYN; NM EPI, N-methyl epinephrine; NMT, non-specific methyltransferase; NTPDase, ectonucleoside triphosphate diphosphohydrolase; OAT, organic ion transporter; OCT, octopamine; PDE, phosphodiesterase; PGH2, prostaglandin H2; PLAP, placental alkaline phosphatase; PNMT, phenylethanolamine-N-methyltransferase; QAPT, quinolinic acid phosphoribosyl transferase; QUINA, quinolinic acid; SYN, synephrine; TDO, tryptophan 2,3-dioxygenase; TNAP, tissue-nonspecific alkaline phosphatase; TRACP, tartrate-resistant acid phosphatase; TRP, tryptophan; TYRA, tyramine.

Figure 2 The redundant signaling networks of immunosuppressive metabolites. Kynurenine (KYN) pathway (left—green): KYN enters the cell via the LAT1, LAT2 or PAT4 transporters. In the cytoplasm, KYN binds the AhR-cPGES-Hsp90-AIP complex. AIP dissociates from the complex after KYN binds, and the KYN-AhR-cPGES-Hsp90 complex translocates into the nucleus. within the nucleus, cPGES and Hsp90 dissociate, and AhR forms a heterodimer with Arnt that modulates gene expression in a widespread manner. Kynurenic acid (KYNA) also agonizes the extracellular GPR35 receptor, which inhibits Akt, p38, and ERK1/2 phosphorylation, and quinolinic acid (QUINA) agonizes the NMDA receptors to increase intracellular Ca^{2+} and ROS and decrease IFN- γ and TNF- α production. Adenosine (ADO) (top center left—yellow): ADO can agonize four different G-protein coupled receptors (GPR35), of which $\text{A}_{2\text{A}}\text{AR}$ and $\text{A}_{2\text{B}}\text{AR}$ stimulate cAMP generation and downstream immunosuppression mediated by the cAMP-PKA-CREB pathway, shown in orange. (Nor)epinephrine (NOR) (top center right—magenta): NOR and epinephrine (EPI) can agonize either α -adrenergic or β -adrenergic receptors (β -AR). Agonism of all three β -AR elicits immunosuppressive signaling cascades through cAMP-PKA-CREB. Prostaglandin E2 (PGE2) (top right adrenergic—blue): similar to ADO, PGE2 can agonize four receptors, of which two (EP2 and EP4) elicit immunosuppressive signaling cascades through cAMP-PKA-CREB. cAMP-PKA-CREB signaling pathway: cAMP production is increased by ADO, NOR, and/or PGE2 signaling through G_s -mediated activation of adenylate cyclase. adenylate cyclase generates cAMP from ATP, and cAMP then activates PKA. A cAMP-PKA complex enters the nucleus where it phosphorylates the CREB transcription factor. phosphorylated CREB recruits HAT and CBP and then modulates target gene expression. All proteins are denoted by text within shapes while metabolites are free-floating text. Straight lines indicate an enzyme catalyzed reaction while curved lines indicate transport or are included for clarity. $\text{A}_\text{x}\text{AR}$, adenosine receptor; AhR, aryl hydrocarbon receptor; AIP, aryl hydrocarbon receptor-interacting protein; ARNT, aryl hydrocarbon receptor nuclear translocator; Ca^{2+} , calcium ion; cAMP, cyclic AMP; CBP, CREB binding protein; cPGES, cytosolic PGE synthase; CREB, cAMP-response element binding protein; EP $_x$, PGE2 receptor; ERK, extracellular signal-regulated kinase; G_x , protein G; HAT, histone acetyltransferase; Hsp, heat shock protein; IFN- γ , interferon gamma; LAT, L-type/large neutral amino acid transporter; NMDAR, N-Methyl-D-aspartate receptor; P, phosphate group; PAT, proton-assisted amino acid transporter; ROS, reactive oxidative species; TNF- α , tumor necrosis factor alpha.

hydroxylation and can be converted into NAD^+ .⁴⁰ Under inflammatory conditions, macrophages and DCs produce quinolinic acid in excess, although the consequences of such are not well understood.⁴⁰ Hydroxylated kynurenine is also converted into xanthurenic acid by a KAT, and xanthurenic and kynurenic acid can be excreted by organic ion transporters 1 and 3, then

filtered from the blood in the kidneys.^{41 42} In addition to feeding the kynurenine pathway, tryptophan is the metabolic precursor for serotonin, which may have roles in tumorigenesis and immune modulation.^{43 44} Circulating serotonin is largely generated in the intestine, then released into the bloodstream where it is stored by platelets.⁴⁵

PGE2: cyclooxygenase and PGE synthesis redundancies

PGE2 is the most common of the prostaglandins, which are ubiquitously synthesized fatty acid hormones.⁴⁶ In solid tumors, PGE synthesis (PGES) enzymes can be upregulated, leading to tumorous accumulation and a 2–3 fold increase in serum levels.^{47–49} Besides cancer cells, TAMs and tolerogenic DCs produce PGE2.⁵⁰ PGE2 synthesis occurs primarily through the three step cyclooxygenase (COX) pathway: first, arachidonic acid is cleaved from the cellular inner membrane by phospholipase A2 enzymes, then COX enzymes convert arachidonic acid to prostaglandin H2 (PGH2) via a PGG2 intermediate, and finally, prostaglandin synthases isomerize PGH2 to PGE2.⁵¹ PGE2 is exported into the extracellular space, where it can interact with immune cells, by the MRP1,2,4, which are upregulated in blood and solid cancers.⁵² Notably, PGH2 is also required to produce several other essential prostaglandins and their derivatives.⁵³

Multiple enzymes can catalyze each step of the COX pathway for PGE2 biosynthesis (figure 1). As many as 19 phospholipase A2 enzymes have been associated with mammalian PGE2 production.⁵¹ Two COX isoforms, COX-1 and COX-2, can be expressed by most cells, and COX-3, a COX-1 splice variant that retains the first intron, is expressed in the brain. COX-1 and COX-2 employ identical catalytic mechanisms yet differ in terms of transcriptional regulation and protein sequence, and both are upregulated in a variety of cancers.⁵⁴ COX-1 is expressed constitutively, while COX-2 expression is induced by NF- κ B or MAPK after exposure to a variety of cytokines or hypoxia.⁵¹ An amino acid difference between COX-1 and COX-2 (V509I) has allowed development of selective COX-2 inhibitors, with the goal of circumventing toxicity arising from complete inhibition of prostaglandin synthesis.⁵⁵ In the final step of PGE2 synthesis, three PGES can isomerize PGH2 to PGE2: microsomal PGES-1 (mPGES-1), mPGES-2, and cytosolic PGES (cPGES).⁵¹ mPGES-1 expression is inducible and often elevated in cancers, while mPGES-2 expression is constant but elevated in gliomas and colorectal cancer.^{51 56 57} cPGES, also referred to as p23, has a role of interest in addition to producing PGE2. It forms a complex with the aryl hydrocarbon receptor (AhR), the transcription factor in the kynurenine signaling pathway (discussed below).⁵⁸ Therefore, each step of the primary route of PGE2 synthesis can be catalyzed by multiple enzymes, such that inhibition of a specific enzyme may not fully prevent PGE2 synthesis.

Finally, evidence exists for an alternative PGE2 synthesis method that does not require enzymes. Specifically, free radical-catalyzed peroxidation of arachidonic acid can form the isoprostane, 8-iso-PGE2. 8-iso-PGE2 can undergo acid- or base-catalyzed epimerization to PGE2.⁵⁹

Norepinephrine and epinephrine: parallel metabolic pathways

Stress incites the canonical flight-or-fight response during which the tyrosine-derived catecholamines, epinephrine

and norepinephrine, are released from the adrenal medulla to interact with adrenergic receptors.¹⁰ Chronic stress and subsequent (nor)epinephrine production have been linked to poor patient prognosis due to the metabolites' action in promoting tumorigenesis and stifling immune response.⁶⁰ Human neuroblastoma, mammary adenocarcinoma, pancreatic ductal carcinoma, and colon adenocarcinoma cell lines, in addition to certain immune subsets, have been shown to secrete nor(epinephrine).^{61–64} For instance, pancreatic ductal carcinoma cells elevated epinephrine and norepinephrine secretion and increased cellular proliferation after exposure to nicotine, while macrophage, peripheral T cells, Tregs, and neutrophils are capable of norepinephrine secretion.^{63 65–68}

Canonical biosynthesis of norepinephrine and epinephrine begins with tyrosine, which is converted to 3,4-dihydroxyphenylalanine, then dopamine, norepinephrine, and epinephrine.⁶⁹ However, multiple parallel biosynthetic pathways ensure that the production of norepinephrine and epinephrine can occur despite any single enzyme deletion, as reviewed by Molinoff and Axelrod (figure 1).⁶⁹ For instance, norepinephrine synthesis can occur without a dopamine intermediate, and epinephrine synthesis can occur without norepinephrine.

SIGNALING REDUNDANCIES AND IMMUNOSUPPRESSION

As it is accepted that cancers engage several immune checkpoints simultaneously, the ability of metabolic intermediates to suppress anticancer immune response is of great interest. Kynurenines, adenosine, PGE2, and norepinephrine and epinephrine can all decrease the activity of cytotoxic T cells, key antitumor effector cells. Interestingly, just as multiple enzymes catalyze the biosynthetic pathways to produce these immunosuppressive metabolites (synthesis redundancy), these molecules often mediate their suppressive impact by agonizing more than one receptor (receptor redundancy). In addition to limiting the activity of immune cells, each of these metabolites promotes tumorigenesis through autocrine or paracrine signaling loops.

Adenosine: immunosuppressive signaling through A_{2A}AR and A_{2B}AR

Adenosine promotes tolerance in the tumor environment and inhibits anticancer immune cell activity. Adenosine can agonize four G protein-coupled extracellular receptors, A₁AR, A_{2A}AR, A_{2B}AR, and A₃AR.⁷⁰ In particular, agonism of the high affinity A_{2A} or low affinity A_{2B} adenosine receptors stimulates production of cAMP from ATP, which acts through Protein Kinase A (PKA) and cAMP-response element binding protein (CREB) to inhibit inflammatory signaling pathways and suppress immune responses (figure 2). Agonism of either A_{2A}AR or A_{2B}AR results in immunosuppression, and immune cells upregulate both receptors under hypoxic conditions. For

instance, in CD8⁺ T cells, A_{2A}AR agonism reduces cytotoxicity, inflammatory cytokine production, and TCR-mediated signaling.⁷¹ A_{2A}AR agonism further inhibits inflammatory cytokine production by neutrophils, macrophages, and DCs, and promotes immunosuppressive macrophage behavior.^{70,71} MDSCs also express A_{2A}AR in the tumor, and its agonism results in increased interleukin 10 (IL-10) production.⁷¹ IL-10 has been shown to behave in a dual role to lessen inflammatory response and drive tumor progression in a tumor-specific and context-specific manner.⁷² Adenosine agonizes the lower affinity A_{2B}AR receptor with similar effects, diminishing CD4⁺ T cell inflammatory function, reducing NK cytotoxicity, inhibiting neutrophil superoxide production and oxidative burst, and downregulating expression of nitric oxide synthase while increasing IL-10 production by macrophages.^{73–77}

Adenosine signaling through A_{2A}AR or A_{2B}AR further promotes differentiation of tolerogenic immune cells, while acting as a signaling molecule to help mediate their impact.⁷¹ Adenosine-A_{2A}AR signaling promotes Treg differentiation from CD4⁺ T cells, and Tregs experience a positive feedback loop for adenosine synthesis in which adenosine drives their expression of CD39 (figure 3).^{78,79} Tregs use adenosine to amplify their immunosuppressive effects on other immune cells even in death, as tumor oxidative stress causes apoptotic Tregs to release and convert excessive amounts to ATP to adenosine.⁸⁰ Adenosine-A_{2B}AR signaling promotes DC differentiation to a tolerogenic phenotype that expresses vascular endothelial growth factor (VEGF), IL-10, COX-2, transforming growth factor β (TGF- β), and IDO (figure 3).⁸¹ Finally, adenosine signaling within the tumor microenvironment promotes tumorigenicity, vascularization, and metastasis, for instance, via induction of VEGF and TGF- β production by cancer and tumor-associated immune cells.^{71,81} Because adenosine agonizes two receptors with similar impact, targeting only one therapeutically may not fully prevent its impact.

In addition to adenosine, cAMP and inosine can have immunomodulatory effects. cAMP-activation of PKA in immune cells can limit cytotoxic and effector functions.⁸² Excreted cAMP serves as an additional feedstock for adenosine generation, which could further agonize A_{2A}/A_{2B}AR receptors to promote cAMP production in a positive feedback loop.⁸³ Inosine has ~1000fold less affinity for A_{2A}AR than adenosine, but can still induce suppressive signaling cascades.^{84,85} However, recent studies have shown that glucose-deprived effector T cells can use inosine as a carbon source, and that inosine enhanced the efficacy of adoptive T cell transfer and checkpoint blockade in solid tumors.^{86,87} Therefore, more work is necessary to deconvolute the role of inosine in the tumor microenvironment.

Kynurenines signaling through AhR, GPR35, and N-methyl-D-aspartate receptor

Extensive studies have established the immunological effects caused by catabolism of tryptophan through the kynurenine pathway. IDO induction and the resultant tryptophan depletion was first demonstrated to cause T cell anergy through the GCN2-mediated starvation response.⁸⁸ Kynurenine, and to a lesser extent its downstream metabolic and transamination products (eg, 3'-hydroxykynurenine, quinolinic acid, and kynurenic acid) have now been shown to broadly suppress antitumor immune cell function by agonizing the AhR.^{36,89} In addition, kynurenine can undergo a spontaneous chemical conversion to create compounds, dubbed trace-extended aromatic condensation products, that bind AhR with picomolar affinity.⁹⁰

Ligand-free AhR resides in the cytoplasm in complex with AhR-interacting protein (AIP), heat shock protein 90, and cPGES/p23.⁵⁸ On ligand binding, the complex sheds AIP, allowing nuclear translocation. The complex fully dissociates in the nucleus, and AhR binds to the AhR Nuclear Translocator (ARNT) to form a heterodimeric transcription factor (figure 2). AhR/ARNT alters expression of thousands of genes that vary in a cell-type and ligand specific manner.⁹¹ In particular, the kynurenine-AhR axis inhibits the anticancer activity of CD8⁺ and CD4⁺ T cells. Signaling mediated by the kynurenine pathway further limits CD8⁺ T cell production of IL-2, decreases efficacy of CAR-T cell therapies, increases CD8⁺ T cell PD-1 expression, and diminishes the efficacy of α -PD-1, α -PD-L1, and α -CTLA4 treatments (figure 3).^{35,92–94} Culturing CD4⁺ T cells with kynurenine generates CD4⁺ FoxP3⁺ Tregs in an AhR and dose-dependent manner.⁹⁵

Similar to adenosine, kynurenines act to promote tumor survival in addition to inhibiting immune cells.³⁶ AhR is highly upregulated in cancers and can play a pivotal role in tumor development by dysregulating proliferation, differentiation, metabolism, apoptosis, and angiogenesis.⁹⁶

Certain immune cells may further drive autocrine/paracrine feedforward loops for AhR signaling. In DCs, IDO expression drives kynurenine production, and AhR activation maintains IDO expression (figure 3).⁹⁷ Similarly, MDSCs produce kynurenine by upregulating IDO expression, and kynurenine-AhR signaling promotes MDSC recruitment and activation.⁹⁸ Kynurenine-AhR signaling induces a suppressive feedforward loop in CD8⁺ T cells by upregulating transcription of kynurenine importers and increasing AhR expression.³⁵

Kynurenic acid, also signals through the AhR to reduce T cell activity, and separately is thought to agonize the G-coupled protein receptor, GPR35, which is expressed by T cells and some innate immune cells.^{89,90,99} GPR35 agonism inhibits phosphorylation of Akt, p38, and ERK 1/2, which may contribute to immune regulation.³⁸ For instance, kynurenic acid agonism of GPR35 reduces IL-4 release by invariant natural killer cells and inhibits tumor necrosis factor- α (TNF- α) secretion by monocytes.^{89,100}

Figure 3 Interactions within and between the immunosuppressive metabolite networks. The adenosine (ADO), kynurenine, (PGE2), (nor)epinephrine (NOR), and other immune checkpoint pathways interact and stimulate each other and may employ feedforward signaling/synthesis regimes. (1) Blockade of PD-1 or CTLA-4 appears to induce (IDO1) expression. (2) A_{2B} AR signaling induces (IDO1) expression. (3, 4) CD39 and CD73 can promote synthesis of (PGE2), and A_{2A} AR or A_{2B} AR signaling can induce expression of COX-2. (5) β -AR signaling upregulates (IDO1) expression. (6, 7) β 2-AR signaling has been linked to the upregulation of COX-2 and cPGES. (8, 9, 10) cPGES is a modulator of (AhR) activity, and COX-2 can influence expression of (TDO2) and (IDO1). (11, 12) (AhR) signaling can increase PD-1 expression and correlates with reduced efficacy of an α CTLA-4 antibody. (13) (AhR) signaling can promote expression of (CD39). (14) (AhR) signaling can promote expression of COX-2. (15) EP2/4 signaling can increase PD-1 expression. (16) (PGE2) can induce expression of CD73. (17, 18) β -adrenergic signaling correlates with diminished therapeutic efficacy of α PD-1 antibodies, and decreases the therapeutic efficacy of a 4-1BB blockade. (19, 20) (CD73) expression correlates with diminished therapeutic efficacy of α PD-1 antibodies and of an α CTLA-4 antibody. (21) CD38 is associated with tumorous resistance to anti-PD-1/PD-L1 antibody therapy. (22, 23) $A_{2A/B}$ AR signaling drives (CD73) expression, and A_{2A} AR signaling increases (CD39) expression. (24) PGE2-EP2/4 signaling can upregulate COX-2 expression. (25) (AhR) signaling drives upregulation of (IDO1) expression. In total, direct links between each pathway have been described with the exception of a relationship between ADO and NOR. Colors denoting ADO, kynurenine, (PGE2), and NOR pathways follow (figures 1 and 2). For brevity, the distinct cell types in which these interactions were described are not included. Curved lines between pathways indicate an interpathway interaction whereas curved semicircles indicate intrapathway regulation. A_x AR, adenosine receptor; AhR, aryl hydrocarbon receptor; β -AR, β -adrenergic receptor; CD, cluster of differentiation; COX, cyclooxygenase; cPGES, cytosolic prostaglandin E synthase; CTLA, cytotoxic T-lymphocyte-associated protein; EPI, epinephrine; PGE receptor; IDO, indoleamine 2,3-dioxygenase; PD, programmed cell death receptor; TDO, tryptophan 2,3-dioxygenase.

Another kynurenine pathway metabolite, quinolinic acid, is a potent agonist of the *N*-methyl-D-aspartate (NMDA) category of glutamate receptors expressed in the central nervous system and on immune subsets.^{41 101} NMDA

receptor (NMDAR) agonism leads to increased intracellular Ca^{2+} and reactive oxidative species concentrations in T and NK cells, and suppresses the ability of these cells to produce IFN- γ .¹⁰² Additionally, activated CD4⁺ T

cells upregulate NMDAR expression, and quinolinic acid suppresses their production of IFN- γ and TNF- α .¹⁰³

Finally, serotonin is produced from tryptophan in a distinct metabolic pathway, and appears to promote tumor growth or mediate inflammatory signaling in a tumor-specific and context-specific manner.^{43 44} For instance, serotonin promotes growth of colorectal tumors, but also stimulates the release of IFN- γ by lymphocytes and stimulates T cell proliferation.^{44 45 104} It is conceivable that competition for tryptophan precursor could limit either kynurenine or serotonin production.

PGE2 immunosuppressive signaling through EP2 and EP4

PGE2 can agonize four G protein-coupled receptors: EP1, EP2, EP3, and EP4. PGE2 signaling through EP2 and EP4 has been implicated in immunosuppression and cancer progression.¹⁰⁵ EP2 and EP4 agonism results in a downstream increase in intracellular cAMP levels followed by PKA/CREB activation, as seen with adenosine agonism of A_{2A}AR and A_{2B}AR (figure 2).¹⁰⁶ EP2 is ubiquitously expressed but has lower affinity for PGE2, while EP4 is a high affinity receptor whose expression is induced by hypoxia.⁹

PGE2 signaling through EP2 limits inflammatory T cell responses by interfering with T cell receptor signaling in CD4⁺ T cells and by promoting a tolerogenic phenotype in DCs that impairs CD8⁺ T cell function.^{107 108} Similarly, EP4 agonism promotes generation of immune cell tolerogenic phenotypes, stimulating differentiation of MDSCs and Tregs, and immunosuppressive functions by macrophages and neutrophils.^{109 110} In macrophages, EP2 agonism suppresses phagocytosis, enhances IL-10 production, and decreases TNF- α production, and PGE2 upregulates COX-2 expression, in an EP2/4 dependent manner, creating an autocrine feedforward loop (figure 3).^{111–113} Signaling through either EP2 or EP4 results in increased PD-1 expression in tumor-infiltrating CD8⁺ T cells of lung cancer patients (figure 3).¹¹⁴ Because PGE2-mediated immunosuppression occurs via two receptors, targeting one may not fully prevent its impact. In addition to inhibiting immune responses, PGE2-EP2/EP4 signaling contributes to tumorigenesis, metastasis, and angiogenesis.¹⁰⁵

Norepinephrine and epinephrine bind two GPCR classes

Norepinephrine and epinephrine, often abbreviated (nor)epinephrine hereafter, bind two distinct classes of G-coupled protein receptors: α -adrenergic receptors (α 1 and α 2) and β -adrenergic receptors (β 1, β 2, and β 3).¹¹⁵ Importantly, agonism of the three β -adrenergic receptors, all of which can be expressed by immune cells, promotes cAMP production and downstream signaling through the cAMP-PKA-CREB cascade (figure 2). Therefore, it is unsurprising that β -adrenergic signaling has inhibitory effects on antitumor immunity. While special attention has been paid to the immunoregulatory action of the β 2-adrenergic receptor, agonism of the β 1 and β 3 receptors also suppresses immune cell activity.¹¹⁶

For instance, in a murine B-cell lymphoma, non-selective β -adrenergic (β 1, β 2, and β 3) agonism suppressed CD8⁺ T cells, decreasing their proliferation, IFN- γ production, and cytotoxicity.¹¹⁷ β 2-adrenergic signaling has also been shown to promote differentiation of CD4⁺ Foxp3⁺ T cells into Foxp3⁺ Tregs, decrease IL-2 production by CD4⁺ T cells, suppress NK cell cytotoxicity, reduce inflammatory cytokine production by DCs, and increase expression of Arg1 and PD-L1 on tumorous MDSCs.^{118–122} Similarly, β 1 receptor agonism induces CD4⁺ differentiation toward a Treg phenotype, and selective β 3 blockade increased intratumoral CD8⁺ T cell and NK cell cytotoxicity and diminished Treg and MDSC counts in a murine melanoma model.^{123 124} The norepinephrine- β -adrenergic signaling axes can suppress the ability of T cells and macrophages to produce TNF- α , and more generally, psychological stress has been shown to correlate with MDSC levels in stage II and III breast cancer patients.^{125 126}

In addition to immune suppression, β -adrenergic signaling contributes to tumorigenesis by enhancing cancer proliferation, angiogenesis, metastasis, and drug resistance, and stress has long been considered to negatively impact prognoses.^{127 128} As seen with adenosine, kynurenine, and PGE2, these ‘fight-or-flight’ metabolites function through immunosuppressive paracrine signaling and tumorigenic autocrine signaling by agonizing multiple receptors on immune or tumorous cells.

INTERACTIONS BETWEEN IMMUNOSUPPRESSIVE METABOLITE AXES

As we have described, the metabolic and signaling networks of adenosine, kynurenine, PGE2, and nor(epinephrine) are often redundant, that is (1) there can exist multiple biosynthetic routes for each metabolite, or multiple enzymes can catalyze the same reaction within a specific metabolic pathway (figure 1), and (2) each metabolite might agonize multiple unique receptors to induce suppressive signaling cascades (figure 2). To wit, A_{2A}AR and A_{2B}AR for adenosine, AhR and GPR35 for kynurenines, EP2 and EP4 for PGE2, and β 1, β 2, and β 3 for (nor)epinephrine. In addition, agonism of different receptors by different metabolites might induce the same immunosuppressive signaling network, as seen with induction of cAMP-PKA-CREB signaling by A_{2A/B}AR, EP2/4, and β -adrenergic receptor agonism, as well as several other instances of crosstalk (figure 3). In this manner, the loss of immunosuppressive signaling mediated by one metabolite can be compensated for by another, and multiple metabolites might act in combination to amplify their impact. For instance, some Tregs can synthesize both adenosine and PGE2, and adenosine and PGE2 have been shown to cooperatively amplify immunosuppressive cAMP-PKA signaling to inhibit cytotoxic cells.^{129 130}

In a similar vein, the kynurenine and PGE2 pathways can amplify each other through a ‘cross-metabolite’ feedforward loop (figure 3). COX-2/PGE2 enhances TDO expression in glioma and drives IDO expression

in numerous cancers, while AhR associates with the COX-2 gene's promoter to drive its transcription.^{131–133} The PGE2 synthesis enzyme, cPGES (p23), also complexes with AhR in the cytoplasm to protect it from degradation.¹³⁴ As adenosine and PGE2 can stimulate feed forward synthesis loops by themselves in certain cells and cooperatively amplify cAMP-signaling, it is intuitive that they could stimulate each other's synthesis through another 'cross-metabolite' feedforward loop.^{129–131 135} Specifically, A_{2A}AR agonism and cAMP signaling upregulates COX-2 expression and PGE2 production, and PGE2 or adenosine can induce TAMs to express CD73 (figure 3).^{136 137} Norepinephrine and epinephrine also interact with the other cAMP-signaling immunosuppressive metabolites, though explicit cross-metabolite feed forward loops have yet to be demonstrated. It has been shown that norepinephrine agonism of the β 2-adrenergic receptor promotes expression of COX-2 and cPGES, resulting in elevated tumorous PGE2.¹³⁸ Macrophages incubated with epinephrine also upregulated expression of COX-2, IDO1, and IL-10.¹³⁹

Therefore, these metabolites are not only functioning in parallel within the tumor but may often be interacting to activate and enhance their immunosuppressive potential (figure 3). As such, attempting to alleviate immune suppression in the tumor microenvironment by targeting one component of a synthesis or signaling pathway, or even one entire pathway, may prove insufficient without patient stratification and proper therapeutic choices for the relevant checkpoints at hand.

CLINICAL IMPLICATIONS OF IMMUNOSUPPRESSIVE METABOLITES

Clinical data demonstrate that elevated immunosuppressive metabolite levels and increased expression of many of their synthesis enzymes correlate with worse prognosis. As such, the pharmacological blockade of these metabolites is of great interest and has driven the discovery of multiple small molecules that antagonize their cognate receptors or inhibitors that target their synthesis enzymes. Therefore, for each metabolite, we will highlight clinical correlations before briefly describing completed and ongoing clinical efforts that target its synthesis and signaling pathways.

For the adenosine axis, expression of CD39 and CD73 predict poor prognosis in numerous cancers, as does A_{2A}AR expression.^{140–142} Tumorous adenosine concentrations have rarely been directly determined, although a quantification method has been recently described.¹⁴³ In addition, an analysis of TCGA transcriptomic data from nearly 10,000 tumors showed that across all cancer types, an 'adenosine signaling gene expression signature' negatively correlates with survival and predicts reduced response to PD-1 and CTLA-4 blockade.¹² Preclinical results have also showed an association between CD38 and anti-PD-1/PD-L1 resistance, and in patients with ovarian

cancer, the frequency of PD-1⁺/CD38⁺ tumor-infiltrating lymphocytes (TILs) is negatively associated with disease stage, lymph node metastasis, and postoperative chemotherapy prognosis.^{144 145}

Several ongoing or completed trials have focused on various targets within the adenosine axis, and selected trials for the adenosine, kynurenine, PGE2, and nor(epinephrine) pathways are briefly highlighted in table 1. For instance, a phase I/II trial targeting CD38 (Isatuximab) and PD-1 (Cemiplimab) reported a manageable safety profile, reduction of CD38⁺ immune subsets in the tumor microenvironment, and enhanced activation of peripheral T cells, and that 20.8% and 65% of metastatic castration-resistant prostate cancer and non-small cell lung cancer (NSCLC) patients, respectively, achieved stable disease.¹⁴⁶ Phase I trials of CD73-targeting antibodies (Oleclumab) or A_{2A}AR antagonists (Ciforadenant, AZD4635) alone, or in combination with approved α PD-L1 checkpoint inhibitor antibodies (durvalumab, atezolizumab), showed manageable safety profiles,^{147–150} and combination therapy of CPI-006 (α CD-73 antibody) and Ciforadenant was also well-tolerated.¹⁵¹ Larger trials testing efficacy of A_{2A}AR inhibition and CD73 blockade are ongoing, and a dual A_{2A}AR/A_{2B}AR antagonist, etrumadenant, showed promising safety and PK/PD in a phase I clinical trial and is being evaluated further.^{152 153}

Kynurenine levels in clinical samples are measured more routinely than adenosine, and an elevated ratio of kynurenine to tryptophan in the serum correlates with worse prognosis across cancers, as does elevated IDO1, IDO2, and/or TDO expression.^{36 154–158} Kynurenine levels are expected to increase predominantly in the tumor microenvironment, but in some cases, there is an observable systemic increase.^{157 159} For instance, in diffuse large B-cell lymphoma patients, kynurenine serum levels were up to 10-fold higher than typical.¹⁵⁷ When treated with rituximab-cyclophosphamide-hydroxydaunorubicin-vincristine-prednisone regimen (R-CHOP) therapy, patients with high serum kynurenine levels (>1.5 μ M) experienced significantly reduced overall survival (58%) compared with the patients with normal levels (<1.5 μ M) (85%).¹⁵⁷ Elevated kynurenine:tryptophan ratios also correlate with reduced response to Nivolumab (α PD-1 antibody) in patients with renal cell carcinoma, as though kynurenine were acting as an adaptive mechanism of resistant to checkpoint inhibition.¹⁶⁰ Conversely, the IDO/AhR signaling axis has been proposed as 'upstream' of PD-1 expression, as heightened serum and tumor kynurenine levels correlate with increased AhR nuclear localization and PD-1 expression on tumor-infiltrating CD8⁺ T cells in breast and colon cancer (figure 3). Therefore, it remains unclear if kynurenine/AhR drives PD-1 expression or drives resistance to PD-1 blockade, raising doubt about the ability of PD-1 blockade and IDO1 inhibition to synergize.^{35 161}

Notably, the IDO1 inhibitor, epacadostat, provided no additional benefit to pembrolizumab treatment (α PD-L1 antibody) in a recent failed phase III clinical trial,

Table 1 Selected clinical trials for therapeutics targeting immunosuppressive metabolites

	Target	Therapeutic	NCT ID	Phase	Indications	Combination therapy	Results reported to date	Ref.
ADO	CD38	Isatuximab	03367819	I/II	mCRPC, NSCLC	Cemiplimab	Manageable safety profile ↓ CD38 + immune cells in TME ↑ Activated peripheral T cells	146
	A2AAR	Ciforadenant	02655822	I	RCC, mCRPC	Atezolizumab	mCRPC: 20% PR for monotherapy and 13% PR for combi therapy	147 148
		AZD4635	02740985	I	Advanced solid tumors, NSCLC, mCRPC, CRC	Abiraterone acetate docetaxel Durvalumab enzalutamide oleclumab	6% PR for monotherapy 5% CR & 11% PR for combination therapy	149
	A2AAR & A2BAR	Etrumadenant	03846310	I	NSCLC	Carboplatin pembrolizumab Pemetrexed zimerelimab*	43% PR Did not worsen safety profile	153
	CD73	Oleclumab	02503774	I	Solid tumors	MEDI4736	↓ Tumorous CD73 expression & ↑ CD8 + infiltration in 5/9 biopsies	150
KYN		CPI-006	03454451	I	NSCLC, RCC, CRC, TNBC, CC, OC, PC, EC, sarcoma, SCCHN, BC, mCRPC, NHL	Ciforadenant pembrolizumab	Treatment well-tolerated ↓ CD73 + B cells, ↑ CD73-CD4+ T cells, ↓ CD8 + T cells one monotherapy patient (9%) saw PR	151
	IDO1	Epacadostat	02752074	III	Melanoma	Pembrolizumab	PFS and OS not improved	162
	AhR	BAY2416964	04069026	I	Advanced solid tumors			

Continued

Table 1 Continued

Target	Therapeutic	NCT ID	Phase	Indications	Combination therapy	Results reported to date	Ref.
PGE2	COX-1 and COX-2 Aspirin	02394769	N/A	CRC		40%–50% of patients saw ↓ of urinary metabolite of PGE2 below threshold expected to ↓ recurrence	172
COX-2	Celecoxib	00062179	II	NSCLC	Carboplatin paclitaxel	Normalized PGE2 intratumoral levels Combination therapy: no PD, 17% CR, 48% PR Did not worsen safety profile	174
EP2 and EP4	TPST-1495	04344795	I	CRC, NSCLC, SCCCHN, UC, EC,			
EP4	Grapiprant	03658772	I	msCRC	Pembrolizumab		
NOR/EPI	β1, β2, and β3	02420223	II	MM		Accelerated engraftment after HCT reduced infection Inhibition of pathways associated with adverse outcomes	178
β1, β2, β3, and α1	Carvedilol	02944201	II	Prostate cancer			

Each therapeutic candidate is organized by the pathway it pertains to and its target within that pathway. For each therapeutic, an NCT was selected due to novelty and/or intriguing results. We note that several hundred other trials are ongoing that target these pathways, so this table is far from exhaustive.

*No results reported for the Zimberlimab combination treatment arm.

ADO, adenocarcinoma; BC, bladder cancer; CC, cervical cancer; CRC, colorectal cancer; EC, endometrial cancer; EPI, epinephrine; GA, gastric adenocarcinoma; GJA, gastroesophageal junction adenocarcinoma; mCRPC, metastatic castrate-resistant prostate carcinoma; NCT, National Clinical Trial; NHL, non-Hodgkin lymphoma; NOR, norepinephrine; NSCLC, non-small cell lung cancer; OC, ovarian cancer; PC, pancreatic cancer; RCC, renal cell cancer; SCCCHN, squamous cell carcinoma of the head and neck; TME, tumor microenvironment; TNBC, triple negative breast cancer; UC, urothelial carcinoma.

rendering the overall status of IDO1 blockade as a therapeutic strategy uncertain.¹⁶² Postmortem analyses have thoroughly detailed possible contributing factors toward Epacadostat's failure, including but not limited to: (1) lack of patient pre-screening for tumorous IDO1/IDO2/TDO expression, as IDO2/TDO activities might compensate for blockade of IDO1; (2) inappropriate dosing regimens and lack of confirmation that kynurenine-AhR signaling was sufficiently reduced; (3) the possibility that Epacadostat may activate AhR due to structural similarity to AhR agonists; (4) uncertainty if PD-1 and IDO1 inhibition act synergistically; and (5) restriction of enrollees to melanoma patients, as melanoma may be a non-responsive tumor type.^{161 163} Recent work has also shown that a large portion of tumors may express TDO.³⁷ Interestingly, Epacadostat's failure may have inadvertently cast light on the importance of broadly targeting an immunosuppressive metabolite's signaling or synthesis pathway. Despite its well-publicized disappointment, Epacadostat is still being tested in 18 ongoing trials. In addition, four other IDO1 inhibitors (Indoximod, BMS-986205, KHK2455, and MK-7162), a dual IDO1/TDO inhibitor (DN1406131), and two AhR inhibitors (KYN-175 and BAY2416964) are being evaluated in the clinic for various malignancies.

As with the kynurenine and adenosine synthesis pathways, the expression levels of several enzymes that catalyze production of PGE2 (such as phospholipases, COX-2, and PGES, and MRP transporters) correlate with poor clinical prognoses.^{52 164 165} While rarely measured directly, PGE2 levels are higher in gliomas and pituitary adenomas than in normal tissues, and higher in the serum of NSCLC patients than controls.^{49 166–168} Notably, non-steroidal anti-inflammatory drugs (NSAIDs) inhibit the COX enzymes. Retrospective analyses have shown that long term use of Aspirin, which inhibits both COX-1 and COX-2, and other COX-2-specific NSAIDs (eg, Celecoxib) reduce risk of developing numerous cancer types, and Aspirin reduces mortality risk after diagnosis.^{169–172} In addition, PGE2 signaling increases CD8⁺ T cell PD-1 expression in lung cancer (figure 3), and recent preclinical studies have demonstrated that Aspirin synergizes with PD-1 blockade to enhance tumorous control and rejection.^{114 173} In the clinic, Aspirin has shown promise as a cancer preventative modality by limiting PGE2 production.¹⁷² Currently, Celecoxib is being evaluated in 35 ongoing clinical trials (defined as trials listed as 'Not yet recruiting', 'Recruiting', 'Enrolling by invitation', and 'Active, not recruiting' in the NCT.gov database) across many cancer indications, some of which incorporate PD-1/PD-L1 blockade and/or chemotherapy as a combination therapy.¹⁷⁴ Other small molecule inhibitors of COX enzymes (etodolac, diclofenac, meloxicam) are also being assessed for cancer indications. Less effort has been aimed at addressing EP2-mediated and EP4-mediated signaling: a single EP4 antagonist, Grapiprant, is being tested in combination with pembrolizumab in a phase I trial, and a dual EP2/EP4 antagonist, TPST-1495,

is being tested in a phase I trial, also in combination with pembrolizumab.

Upregulated expression of norepinephrine or epinephrine-producing enzymes or their elevated levels also correlate with more advanced and/or aggressive disease.^{61 62} Analogous to NSAIDs, the use of β -adrenergic antagonists, commonly called β -blockers, has been retrospectively associated with a reduction in cancer diagnosis and mortality risk after diagnosis.^{175 176} 'Selective' β -blockers antagonize the β 1 receptor (eg, Metoprolol and Atenolol) while 'non-selective' β -blockers antagonize multiple adrenergic receptors, for instance, propranolol (β 1, β 2, and β 3 receptors) and carvedilol (β 1, β 2, β 3, and α 1 receptors). These retrospective analyses have routinely showed enhanced benefit from 'non-selective' β -blockers, suggesting that broad prevention of (nor) epinephrine-signaling is important for therapeutic benefit.^{177 178} The 'non-selective' antagonists propranolol (24 ongoing trials) and carvedilol (2 ongoing trials) are being broadly investigated as cancer therapies, while 'selective' agonists Metoprolol and Atenolol are being evaluated primarily for analgesia in cancer indications.

Increased expression of enzymes that catalyze the synthesis of the immunosuppressive metabolites adenosine, kynurenine, PGE2, and epinephrine correlates with worsened patient prognosis, immune suppression, and even poor response to checkpoint inhibitor antibody therapies. As such, numerous clinical trials and therapeutic efforts have been initiated to prevent their impact. However, the redundancies seen in and between immunosuppressive metabolic synthesis and signaling pathways makes preventing their activity a challenge.

CHALLENGES AND PATHWAYS FORWARD

The recent clinical failure of Epacadostat, an IDO1-specific inhibitor, put the status of kynurenine-targeting therapeutics in jeopardy. However, thorough postmortem analyses have granted insights toward how such a failure can be avoided in the future, potentially through patient stratification, dosing optimization, or by broadening therapeutic scope to account for IDO/TDO redundancies, or more generally, synthesis or signaling redundancies found in immunosuppressive metabolite pathways. Here, we will describe avenues to alleviate the impact of suppressive metabolites by accounting for each metabolites intranetwork- and internetwork redundancies. Avenues include patient stratification, combination therapies, development of broad-spectrum (pan) inhibitors or antagonists, and/or development of new therapeutics that target the 'weakest link' of a metabolite's pathways (eg, a receptor for which no redundancy exists). Pan-inhibitors/antagonists and therapeutics against targets that lack redundancies may prove particularly promising.

For kynurenine-mediated immune suppression, patient stratification to ensure IDO1⁺ & TDO⁺ expression has been proposed as a necessary measure to improve efficacy of IDO1-targeting monotherapies. However, such

baseline stratification prior to therapy may fail to account for the fact that the presence of activated T cells, such as could occur with PD-1 or CTLA-4 blockade, can itself induce IDO1 expression, while PGE2 has been shown to induce both IDO1 and TDO (figure 3).^{132 133 179} Induction of reactive IDO or TDO during therapy would not be detected by baseline screening but would still be immunosuppressive and a target for therapy. Hence, initial biomarker stratification might be helpful but may not be definitive and should be followed by metabolomics analyses to confirm in vivo reduction in kynurenine levels following intervention. Kynurenine metabolomic analyses are relatively routine, and they can be coupled with analysis of IDO1, IDO2, and TDO expression levels that might deduce the mechanism of resistance to IDO1 inhibition (eg, increased IDO1 expression, induced TDO expression) to adjust accordingly.¹⁶¹ Preclinical studies have shown that kynurenine synthesis can be targeted by combining TDO and IDO1 inhibitors, or by using dual IDO1/TDO and pan-IDO1/IDO2/TDO inhibitors. Such therapies have claimed durable benefits in murine tumor models, but pan-inhibition of the kynurenine pathway can (1) drastically raise levels of serotonin and other tryptophan metabolites that cause toxicities and neurological impacts and (2) prevent intracellular NAD⁺ synthesis if the NAD⁺ salvage pathway is inefficient or not functional. Therefore, the tolerability profile of pan-inhibitors must be fully investigated and should consider all of tryptophan and kynurenine metabolism.^{180 181}

AhR is the sole receptor for which kynurenine is known to be a ligand (figure 2). Thus, antagonizing AhR or depleting kynurenine directly would not be expected to be limited by IDO1/IDO2/TDO redundancies. In this vein, targeting kynurenine-mediated immune suppression with AhR antagonists has shown promise in preclinical models, and tumorous degradation of kynurenine using an engineered kynureninase enzyme (PEG-KYNase) has also shown efficacy.¹⁸²⁻¹⁸⁴ Still, expectations should be tempered for both therapeutic routes, as neither would prevent tryptophan depletion and any resulting immune suppression. In addition, AhR controls broad transcriptional programs impacting gut homeostasis, immunity, and numerous other processes, so antagonism could have unpredictable impacts.^{185 186} The PEG-KYNase utilized for preclinical studies is of bacterial origin, sparking immunogenicity and tolerability concerns, and its catalytic product, anthranilic acid, might re-enter the canonical kynurenine pathway if it were hydroxylated (figure 1).¹⁸⁷

Similar challenges and opportunities exist for targeting adenosine immunosuppression. In murine models, combination therapies against CD73 and CD39 or CD73 and A_{2A}AR improve response compared with CD73, CD39, or A_{2A}AR monotherapy. This improved response implies that single-target therapeutics do not fully prevent adenosine synthesis or signaling, as may be expected from the adenosine axis' redundancies. Complete inhibition of adenosine synthesis is very challenging due to the diversity of enzymes and metabolic pathways that contribute

toward its production, but dual A_{2A}AR/A_{2B}AR inhibitors might largely prevent adenosine signaling. To this end, ongoing clinical trials testing the antitumor activity and safety of Etrumadenant will begin reveal the potential of A_{2A/B}AR dual antagonism.¹⁸⁸ However, it is important to note that adenosine signaling through A_{2A}AR and A_{2B}AR is important for normal physiological functions such as maintenance of the sleep cycle and cardiovascular regulation, so non-tissue-specific blockade of adenosine signaling could have safety concerns.^{189 190} Alternatively, it may be possible to degrade the adenosine molecule itself using an enzymatic therapy, similar to the PEG-KYNase mechanism of action. Adagen, a PEG-conjugated enzyme from *Bos taurus*, can deaminate adenosine into inosine, and its repurposing for cancer has been proposed recently.⁸⁶ Because of its non-human origin, Adagen could suffer from the same tolerability and immunogenicity issues noted for PEG-KYNase, and efficacy-limiting anti-drug antibodies arise even when Adagen is used to treat immunocompromised Adenosine Deaminase Severe Combined Immunodeficiency (ADA-SCID) patients.¹⁹¹ Recently, a PEGylated, engineered ADA 2 showed anti-tumor efficacy in preclinical studies.¹⁹² Therefore, therapeutic efforts to target adenosine directly or to block both A_{2A}AR and A_{2B}AR signaling might be well positioned to counter the internal pathway redundancies seen in the adenosine axis.

Even if adenosine-mediated immunosuppression were eliminated, cancers have alternative routes to promote cAMP-mediated immunosuppressive signaling. However, established therapeutics that target PGE2 or the epinephrine axes exist and might be repurposed. As we've noted, the non-selective $\beta_1/\beta_2/\beta_3$ adrenergic receptor antagonist, propranolol, and the dual COX-1/2 inhibitor, aspirin, correlate with reduced cancer risk in retrospective analyses.^{193 194} By targeting multiple receptors/enzymes, these therapies may be less likely to be impacted by PGE2/(nor) epinephrine pathway redundancies. A dual EP2/EP4 antagonist is also being tested clinically, giving another broad-acting route toward preventing PGE2-signaling. Alternatively, inhibition of the adenylate cyclases which produce cAMP from ATP might be advantageous due to the position of cAMP as a focal point connecting various immunosuppressive signaling pathways. Considering that immune cells largely express adenylate cyclase isoform 7, delivery of a specific inhibitor to tumorous immune cells may thwart the immunosuppressive signaling contributions of the adenosine, PGE2, and (nor)epinephrine pathways simultaneously.¹⁹⁵

cAMP-mediated signaling also impacts numerous other aspects of disease progression, specifically promoting tumor growth and metastasis.¹⁹⁶ Thus, while retrospective studies provide valuable insights, each treatment or combination therapy must be evaluated for efficacy as immunotherapies in preclinical models and then in human studies. It will be important deconvolute the impacts of altering tumorous signaling versus improving antitumor immunity, and therapeutic benefit might be

improved for either impact by employing broad-spectrum blockade.

Promisingly, perioperative non-selective β -adrenergic antagonism (Propranolol) and COX-2 selective inhibition (via Etodolac) was well tolerated and improved immune function in a recent small clinical trial of colon cancer.¹⁹⁷ It will be particularly important to determine safety profiles associated these therapies. For instance, COX inhibition results in gastrointestinal and cardiovascular hazards due to reduced production of important prostaglandins (other than PGE₂), and β -adrenergic signaling contributes to the regulation of many bodily functions apart from immune response.^{10 53} Importantly, between 1999 and 2004, the COX-2 selective inhibitors Rofecoxib and Celecoxib caused more than 26 000 deaths compared with less than 10 000 deaths in patients taking 'traditional' non-selective NSAIDs (eg, aspirin).¹⁹⁸ Hence, treatment with traditional NSAIDs may be preferable to COX-2 selective inhibitors in terms of both patient safety and accounting for pathway redundancy. Pan-cAMP inhibition may also have other unintended consequences. For example, while exercise stimulated antitumor immunity and tumor regression in tumor-bearing mice, the benefits of exercise were negated on eliminating epinephrine-mediated signaling.¹⁹⁹

Patient stratification for adenosine, PGE₂, or nor(epinephrine) targeting therapies could be beneficial, however, will likely prove challenging because many enzymes catalyze their production, their signaling pathways intersect, and measuring PGE₂ and adenosine levels is uncommon. A promising alternative to direct adenosine measurement is the recently described 'adenosine signaling gene expression signature' which correlated well with in vivo tumorous adenosine levels in mice and may have utility as a proxy for adenosine levels in the clinic.¹² While developing robust methods to directly measure PGE₂ would be ideal, it may be also be possible to develop a similar 'PGE₂ signaling gene expression signature', or a pan-cAMP-signaling signature that encompasses adenosine, PGE₂, and nor(epinephrine). Interestingly, seven of the 14 genes that comprise the adenosine signaling signature are upregulated by PGE₂-mediated signaling (FOXP3, COL3A1, APP, FOS), part of the PGE₂ signaling pathway (CREB and MAPK1), or utilized for PGE₂ synthesis (PTGS2, that is, COX-2).^{200–205} Therefore, this signature likely encompasses aspects of PGE₂-signaling or general cAMP-mediated immunosuppression, and with fine-tuning, may be able to gage individual contributions. Certain patient stratification attempts may also prove more straightforward. Targeting PGE₂ in brain malignancies might preferentially employ acetaminophen due to its ability to: (1) traverse the blood-brain barrier (unlike many other COX inhibitors), and (2) inhibit COX-3, the primary COX isoform expressed in brain tissue.^{206 207}

Other therapeutic routes to prevent the impact of immunosuppressive metabolites are also being developed. CAR-T cell therapies might be engineered to

resist immunosuppressive metabolic signaling. For instance, A_{2A}AR signaling can limit the efficacy CAR-T cells, but silencing of A_{2A}AR expression afforded resistance in a murine model.²⁰⁸ CAR-T cells have also been engineered to express a peptide, RIAD, which inhibits cAMP-mediated signaling cascades by interacting with PKA, potentially blunting PGE₂, (nor)epinephrine and adenosine signaling.^{209 210} These 'metabolically armored' CAR-T cells demonstrated improved control of tumor growth and immune cell activity in murine models.²⁰⁹ Finally, ongoing clinical evaluations have been keen to demonstrate synergistic benefits from targeting synthesis enzyme or metabolite-receptors in combination with established checkpoint inhibitors but have almost universally been performed with PD-1/PD-L1 blockade. However, blockade of other protein-ligand interactions might also provide synergistic benefit. For instance, pharmacological blockade of CD73 enhanced the immunostimulatory efficacy of both PD-1 and CTLA-4 antibodies in mouse models, and β -adrenergic signaling has been shown to limit therapeutic efficacy of α PD-1 and α 4-1BB antibodies (figure 3).^{117 211}

CONCLUSION

Herein, we have detailed the pathway and signaling redundancies prevalent for immunosuppressive metabolites, as well as highlighting overlap between signaling pathways, each pathway's clinical relevance, and ongoing therapeutic efforts. Redundancy is not a novel concept in tumorous immune evasion; for instance, CD8⁺ TILs from mice treated with anti-PD-1 upregulate expression of CTLA-4.²¹² However, the level of redundancy inherent in the ability of each of these individual metabolites to suppress immunity is surprising. Because compensating metabolic pathways may limit efficacy of inhibitors of single enzymes (eg, IDO1), and several metabolites agonize multiple receptors (eg, EP2/EP4), we've emphasized the importance of therapies that broadly target each pathway. To this end, clinical trials are currently testing 'pan-antagonists' of each of the 'metabolic immunosuppressive receptor' subclasses that is, A_{2A/2B}AR, EP2/EP4, β -adrenergic receptors, and AhR, as well as broad-inhibitors of kynurenine (IDO1/TDO or IDO1/IDO2/TDO) and PGE₂ (COX-1 and COX-2) synthesis. In light of cross-metabolite pathway redundancies, most prominently cAMP-mediated signaling, it may also be important to consider combination therapies against multiple immunosuppressive metabolites. For instance, therapeutics that target the adenosine pathway might synergize with repurposed COX-1/COX-2 inhibiting NSAIDs and beta blockers, though more efforts are required to determine if these drugs truly function as 'metabolic immunotherapies'. Because of the similarity in each pathways synthesis and signaling redundancies, lessons learnt from targeting one immunosuppressive metabolic pathway might guide future successes (and help avoid potential pitfalls) for other metabolites. Finally, as safety concerns exist for

each single-target inhibitor or antagonist, determining the safety profile of more broadly acting therapies (and their combinations) will be of the utmost importance.

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