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▶ To cite this version:

Colleen Sweeney, Gwendal Lazennec, Christoph Vogel. Environmental exposure and the role of AhR in the tumor microenvironment of breast cancer. Frontiers in Pharmacology, 2022, 13, pp.1095289. 10.3389/fphar.2022.1095289 . hal-03918758

HAL Id: hal-03918758 https://hal.science/hal-03918758v1

Submitted on 18 Oct 2023

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1 Environmental exposure and the role of AhR in the tumor microenvironment of breast cancer

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Keywords: AhR₁, Breast Cancer₂, Air Pollution₃, Particulate Matter₄, Health Disparities₅, Environmental Injustice₆

15 Abstract

Activation of the aryl hydrocarbon receptor (AhR) through environmental exposure to chemicals 16 17 including polycyclic aromatic hydrocarbons (PAHs) and polychlorinated dibenzo-p-dioxins (PCDDs) 18 can lead to severe adverse health effects and increase the risk of breast cancer. This review considers 19 several mechanisms which link the tumor promoting effects of environmental pollutants with the AhR signaling pathway, contributing to the development and progression of breast cancer. We explore 20 21 AhR's function in shaping the tumor microenvironment, modifying immune tolerance, and regulating 22 cancer stemness, driving breast cancer chemoresistance and metastasis. The complexity of AhR, with 23 evidence for both oncogenic and tumor suppressor roles is discussed. We propose that AhR functions as a "molecular bridge", linking disproportionate toxin exposure and policies which underlie 24 environmental injustice with tumor cell behaviors which drive poor patient outcomes. 25

26 1 Introduction – Environmental Exposure and Breast Cancer

27 Air pollution and occupational exposure studies have reported positive associations with the risk of

- developing breast cancer (Amadou et al., 2021). Air pollution and ambient particulate matter (PM)
- 29 contain a complex mixture of compounds, including polycyclic aromatic hydrocarbons (PAHs) and
- 30 various metals (e.g. iron, nickel, copper), which may induce reactive oxygen species (ROS) and
- 31 inflammation (Rückerl et al., 2007, Grunig et al., 2014) and stimulate the progression of breast
- cancer (Romaniuk et al. 2017). PAHs are generated during combustion processes and derive from
 various sources such as indoor fireplaces, wildfires, industrial activities, and vehicular traffic and the
- exposure to PAHs has been identified as a risk factor for breast cancer (Lichtiger et al., 2021,
- 35 Gamboa-Loira et al., 2022). Importantly, a stronger association of breast cancer risk was found with
- 36 traffic related air pollution (TRAP) and higher PAH exposure intensity and duration of exposure (Lee
- 37 et al. 2019, Large and Wei 2017, Mordukhovich et al. 2016, Nie et al. 2007, Shen et al. 2017, Brody
- et al. 2007). Vehicular traffic is a major ambient source of PAH exposure and the PAH
- 39 benzo[a]pyrene (BaP) is classified as a human carcinogen by the International Agency for Research

40 on Cancer (IARC, 2010). Furthermore, BaP and other PAHs have been identified as ligands of the

- 41 aryl hydrocarbon receptor (AhR), a ligand-activated transcription factor, belonging to the bHLH-PAS
- family, which regulates multiple target genes and is best known for its role as a xenobiotic receptor 42
- 43 (Boonen et al. 2020, Vogel et al. 2020). The activation of the AhR signaling pathway via
- 44 environmental pollutants including dioxins and PAHs has been associated with the development of
- breast cancer (Narasimhan et al. 2018, Kolluri et al. 2017, Murray et al 2014, Warner et al. 2011, 45
- 46 Birnbaum and Fenton 2003, La Merrill et al. 2010, Donovan et al. 2018, Gearhart-Serna et al. 2020).
- 47 Reports from our team and other groups suggest an important role of AhR as an immune-modulator 48
- and mediator of toxic responses triggered by particulate matter (PM) derived from TRAP (O'Driscoll
- et al. 2013, O'Driscoll et al. 2018, Castaneda et al. 2018, Yuan et al. 2020). 49
- 50 Recent studies confirmed an increased risk of breast cancer with vehicular-specific PM exposure
- 51 among African American and Japanese American women living near major roads, highlighting the
- 52 link between environmental injustice and health disparities (Niehoof et al. 2020, Cheng et al. 2019).
- 53 Indeed, residential proximity to major roadways is a recognized risk factor beyond breast cancer, in
- 54 cardiovascular disease (Kulick et al. 2018, Kirwa et. al 2014, Kingsley et. al 2015, Hart et. al 2014)
- and renal disease (Lue et. al 2013). Further, it disproportionately impacts racial and ethnic 55
- 56 minoritized groups and those of lower socioeconomic status, the legacy of the widespread practice of
- 57 redlining in the United States (Hwa Jung et. al 2022, Swope et. al 2022). While the Fair Housing Act
- 58 of 1968 prohibited racial discrimination in housing and lending, exclusionary zoning and other
- 59 practices such as gentrification has perpetuated residential segregation
- (https://www.brookings.edu/research/neighborhood-segregation-persists-for-black-latino-or-hispanic-60
- 61 and-asian-americans/). In a study of Hillsborough County in Florida, Stuart et al found that blacks,
- 62 Hispanics, and people living below the poverty line are much more likely to reside close to sources of air pollution but further from air quality monitoring sites while whites were found to live closer to
- 63 64 monitoring sites but significantly further from pollution sources (Stuart et. al 2009). Wu et al found
- 65 that particulate matter (PM) collected near a major Los Angeles freeway (compared to an urban
- 66 background location) induced significantly higher production of the cytokines IL-6, IL-8, and TNF-α,
- (Wu et. al 2014), suggesting a link between AhR activation, AhR-driven inflammation (Vogel et. al 67
- 68 2011, Wu et. al 2021, Vacher et. al 2018) and proximity to pollution. The interaction between
- environmental exposure, socio-economic related stress and psychosocial stress in under-resourced 69
- neighborhoods has been termed the environmental "riskscape" by Morello-Frosch et al (Morello-70
- 71 Frosch & Shenassa, 2006, Morello-Frosch & Lopez, 2006). As noted by Morello-Frosch, the
- 72 Institute of Medicine recognizes this as a type of "double jeopardy" in which elevated stress impairs
- 73 the ability of individuals living in under-resourced neighborhoods to endure the myriad health
- 74 consequences of chronic environmental exposures (https://www.scientificamerican.com/article/end-
- 75 double-jeopardy/#).
- 76

77 2 **Role of AhR in Breast Cancer**

78 Approximately two decades ago, AhR was found to be overexpressed in mammary cancer in rats 79 (Trombino et al. 2000) sparking curiosity as to its role in breast cancer progression. Several studies

80 have since shown that chemical exposure and AhR activation affect processes of mammary gland

- differentiation, disrupting pregnancy-related differentiation and milk production, and increasing the 81
- 82 risk of breast cancer (Warner et. al 2002, Vorderstrasse et al. 2004, Lew et al. 2011, Belton et al.
- 83 2018, Kay et al. 2022). Further studies have elucidated AhR's molecular contribution to carcinogenic
- 84 progression and ratified the oncogenic role of AhR in breast cancer cells (Wang et al. 2017, Wang et

al. 2020). In support of its role as a breast cancer oncogene, AhR activation is sufficient to transform

- 86 human mammary epithelial cells and promote their migration, invasion and epithelial-to-
- 87 mesenchymal transition (EMT) (Brooks and Eltom 2011). Work from our group showed that chronic
- exposure of MCF10AT1 and MCF-7 cells to estradiol (E2) resulted in AhR overexpression and
- 90 et al. 2009) accompanied by increased proliferation, invasion, and apoptosis resistance. The
- 91 resistance to apoptosis was also demonstrated in human breast cancer cell lines treated with the
- prototypical AhR ligand TCDD when apoptosis was induced by chemotherapeutics (doxorubicin,
 lapatinib and paclitaxel) (Bekki et al. 2015). Treatment with PAH mixtures which bind to and
- lapatinib and paclitaxel) (Bekki et al. 2015). Treatment with PAH mixtures which bind to and
 activate AhR also increased cell proliferation and expression of antiapoptotic proteins in MCF-7 cells
- 94 activate Anic also increased cell promeration and expression of95 via AhR signaling (Gearhart-Serna et al. 2020).
- 96 Several studies have reported AhR overexpression in human breast cancer (Li et al. 2014, D'Amato 97 et al. 2015, Vacher et al. 2018). Using samples from breast cancer patients, we found that AhR is
- 98 frequently over-expressed in ER-negative human breast tumors, and this is closely correlated with
- 99 elevated expression of the NF-κB subunit RelB and inflammatory markers such as IL-8 (CXCL1 in
- 100 mouse) and COX-2 (Vogel et al. 2011). This was also observed by Vacher et al. with significant
- 101 overexpression of cytokines, including IL-8, in AhR high expressing tumors (Vacher et al. 2018). We
- 102 demonstrated that C/EBPβ serves as a key transactivator for AhR-mediated COX-2 gene induction
- 103 (Vogel et al. 2000, Vogel et al. 2004). Interestingly, COX-2, CXCL1, and IL-8 have been identified
- as critical genes that mediate breast cancer invasion and metastasis to lung and lymph nodes (Freund
- et. al 2003, Minn et al. 2005, Ahmed et al. 2021). A recent report suggests that inhibition of COX-2
 expression reduces mammary tumor multiplicity and size in the polyoma middle T antigen (PyMT)
- 107 mouse model (Esbona et al. 2016). In our recent study we demonstrated that overexpression of AhRR
- 108 (Aryl Hydrocarbon Receptor Repressor) suppresses AhR-driven (TCDD-stimulated) growth of
- 109 syngeneic mammary tumors as well the onset, growth and metastasis of spontaneous mammary
- 110 tumors in PyMT mice (Vogel et al. 2021). In human breast cancer, high expression of AhRR, the
- 111 dedicated AhR repressor, independently predicts prolonged metastasis-free survival (Vacher et al.
- 112 2018), in agreement with our findings in PyMT mice (Vogel et al. 2021). Interestingly, knockdown
- 113 of AhRR in normal human mammary epithelial cells resulted in anchorage-independent cell growth
- suggesting that the AhRR may function as a tumor suppressor gene (Zudaire et al. 2008).
- 115 In a mouse model of BRCA1-associated breast cancer, AhR was found to transcriptionally induce the
- 116 EGF receptor ligand, Amphiregulin, driving tumor growth and macrophage infiltration. Of note, this
- 117 was inhibited by the combination of an AhR inhibitor and an EGF receptor inhibitor, suggesting new
- 118 therapeutic possibilities for this type of breast cancer (Kubli et. al 2019). The relationship between
- 119 AhR activation and breast cancer-related death was recently assessed using an artificial intelligence
- 120 tool to analyze the scientific literature, with strong evidence that AhR activation is an adverse
- 121 outcome pathway in breast cancer (Benoit et al. 2022).
- 122 Interestingly, many studies have also provided evidence for a tumor suppressor role for AhR, with
- evidence that AhR can inhibit tumor growth (Fritz et al. 2007, Feng et al. 2020, Jin et al. 2014) while
- inhibition of AhR or AhR deficiency promotes tumor development (Abdelrahim et al. 2003, Safe et al. 2017). For example, in the ApcS580/+; KrasG12D/+ mouse model of colon tumorigenesis,
- al. 2017). For example, in the ApcS580/+; KrasG12D/+ mouse model of colon tumorigenesis,
 intestinal epithelial specific AhR knockout promoted tumorigenesis through enhanced Wnt signaling
- 120 Intestinal epitienal specific Ank knockout promoted unforgenesis through enhanced witt signaling 127 (Han et. al 2021). In p53 deficient mice, AhR knockout significantly increased incidence of thymic
- 128 lymphomas and sarcomas and decreased survival (Phillips et. al 2022). In a mouse model of sonic
- hedgehog type-medulloblastoma, AhR deletion in cerebellar granule cell progenitors accelerated
- 130 tumorigenesis through increased TGF β -SMAD3 signaling (Sarić et. al 2020) with high AhRR

- 131 expression linked to decreased patient survival. Further, in an unbiased functional genomics screen,
- 132 AhR was identified as metastasis suppressor in a lung cancer model (Nothdurft et. al 2020). In in
- 133 vitro studies, AhR was demonstrated to cooperate with the Rb tumor suppressor to prevent S-phase
- 134 cell cycle entry (Puja et al. 2000) while activation of AhR by the prototypical ligand TCDD inhibited
- the growth of MCF7 breast cancer cells (Vogel and Abel 1995). David Sherr's team investigated
- 136 AhR agonists and antagonists in a direct comparison and concluded that the sustained activation of
- 137 AhR drives the later, more lethal stages of some cancers, but that AhR agonists under some
- 138 circumstances can counteract tumor development and may also serve as cancer therapeutics
- 139 (Narasimhan et al. 2018). In this vein, O'Donnell (O'Donnell et. al 2021) and others (Rowland et. al
- 140 2019) have pursued SMAhRTs, Select Modulators of AhR-regulated Transcription, to specifically
- exploit the anti-cancer functions of AhR. Notably, they identified a modulator which induced AhR dependent Fas ligand expression and breast and liver tumor cell apoptosis without increasing
- expression of the prototypical AhR target gene, CYP1A1, suggesting that AhR transcriptional
- 144 activity can be fine-tuned, to specifically unlock its function as a tumor suppressor.

145 **3** Cytokines and Chemokines in Breast Cancer and the Tumor Microenvironment

- 146 The tumor microenvironment (TME) corresponds to the fact that tumor cells are surrounded in close
- 147 proximity by a number of non cancerous cells including cancer associated fibroblasts (CAFs),
- 148 mesenchymal stem cells (MSCs), adipocytes, myeloid derived suppressor cells (MDSCs), tumor
- 149 associated macrophages (TAMs), tumor associated neutrophils (TANs), tumor infiltrating
- 150 lymphocytes (TILs), and endothelial cells (Binnewies et al. 2018, Joyce and Pollard 2009, Lazennec
- and Lam 2016). In addition to direct contact with tumor cells, TME cells will interact with tumor
- 152 cells though a number of different soluble factors including cytokines and chemokines, which will
- reshape TME to support cancer initiation, progression, and metastasis (Ali and Lazennec 2007,
- 154 Mancini et al. 2021, Lazennec and Richmond 2010) (Fig. 1).
- 155 In breast cancer, many chemokines and cytokines have been analyzed and identified as important
- 156 factors contributing to the development of breast tumors (Masih et al. 2022, Narita et al. 2016). In
- 157 particular the CXCR4/CXCL12 axes has been reported to control breast cancer metastasis and the
- 158 involvement of CAFs (Muller et al. 2001, Orimo et al. 2005). The CAF-driven CXCR4/CXCL12 axis

159 may also stimulate the accumulation of protumorigenic lipid associated macrophages which supports

- 160 an immunosuppressive microenvironment in breast cancer (Timperi et al. 2022).
- 161 CCL2 and CCL5 have also retained attention in breast cancer, as they are expressed by cancer cells
- and promote the recruitment of TAMs and metastasis by inducing Th2 polarization of CD4+ T cells
- 163 (Brummer et al. 2018, Zhang et al. 2015, Soria and Ben-Baruch 2008, Chavey et al. 2007). In
- addition, the ligands of CXCR2 (CXCL1, 2, 3, 5, 6, 7, 8) have been shown in a number of studies to
- 165 be involved in the aggressiveness of triple negative breast cancers (TNBC) (Chavey et al. 2007,
- 166 Bieche et al. 2007, Acharyya et al. 2012). The genes of these chemokines are encoded by a small
- region of chromosome 4q21 and have been found to be coregulated in TNBC (Bieche et al. 2007).
- 168 Moreover, cancer cells expressing high levels of CXCL1 and CXCL2 acquire an advantage in terms
- 169 of survival in metastatic sites and favor the recruitment of TANs (Acharyya et al. 2012).
- 170 Interestingly, CXCR2 itself is also playing a major role in the aggressiveness of TNBC, in particular 171 through its expression on TANs (Boissiere-Michot et al. 2020, Boissiere-Michot et al. 2021).
- 1/1 Inrough its expression on TAINS (Boissiere-Michot et al. 2020, Boissiere-Michot et al. 2021).
 172 Although the levels of CXCR2-expressing neutrophils is correlated to high grade breast cancers, its
- role is rather to counteract tumor progression (Boissiere-Michot et al. 2021), as it is correlated with a
- better survival of the patients and its deletion favors tumor growth and metastasis (Timaxian et al.
- 175 2021). There are many links between AhR and CXCR ligands in particular. For instance, we have

176 shown that the complex of AhR and NFkB RelB was able to bind to a specific binding elements of chemokines including the CXCL8 promoter and to promote its activation though protein kinase A 177 (Vogel et al. 2007). RelB/AhR complex is also involved in the overexpression of CXCL8 in breast 178 179 cancer (Vogel et al. 2011, Bekki et al. 2015). A significant elevated level of CXCL8 mRNAs expression (56-fold) has also been found in tissue samples of high stage compared to low stage 180 patients and adipose-derived stem cells (Razmkhah et al. 2010). AhR may also interact with NFkB 181 RelA causing the upregulation of c-myc and stimulation of tumorigenesis in MCF-7 cells (Kim et al. 182 2000). Further HER2 overexpression in MCF-7 cells resulted in pro-inflammatory signaling and 183 184 induction of IL-6 enhancing mammosphere formation in an AhR-dependent manner (Zhao et al. 185 2013). The role of AhR as a mediator of chronic inflammation in breast cancer has been recently 186 reviewed elsewhere (Guarnieri 2020). Moreover, a recent study by Kubli et al. has shown that AhR 187 was induced by reactive oxygen species (ROS) in mammary epithelial cells, which in turn enhance AREG (amphiregulin) production (Kubli et al. 2019). In basal-like and BRCA1-related breast 188 189 cancers, ROS expression was correlated with AhR levels and the expression of the chemokines 190 CXCL1, CXCL2 and CCL5. Targeting AhR or AREG reduced the recruitment of macrophages in 191 tumors in mouse models and AREG expression was associated with the density of macrophages in 192 human tumors. Another cytokine upregulated by AhR activation is IL-22 which is an important 193 factor controlling host defense and gut immunity. However, dysregulation of IL-22 may contribute to 194 the development of TNBC and the pathology in breast cancer (Katara et al. 2020, Wang et al. 2018, 195 Kim et al. 2014, Voigt et al. 2017). IL-22 has also been described to mediate macrophage infiltration 196 in the TME and the migration of breast cancer cells (Kim et al. 2020). Results from MCF-7 cells co-197 cultured with preadipocytes and an in vivo zebrafish model showed that prototypical AhR ligand 198 TCDD enhanced the invasive and metastatic potential of MCF-7 cells implicating the importance of

199 AhR in the TME (Koual et al. 2021).

200 4 AhR as a Critical Player in the Tumor Microenvironment of Breast Cancer

201 The development of metastatic disease, which accounts for greater than 90% of cancer mortality, requires collaboration between tumor cells and their environment. Recent studies reveal that the TME 202 203 possesses remarkable cellular heterogeneity with an important role of immune cells in the 204 development and progression of breast cancer (Place et al. 2011, Ben-Baruch 2002). The TME also consists of an acellular component (e.g., soluble cytokines, chemokines, and growth factors), that 205 206 forms part of the stromal structure as described above. TAMs and MDSCs are tumor-associated 207 myeloid cells (TAMCs) and have been identified as key players in breast cancer progression and metastasis (Cha and Koo 2020). MDSCs are myeloid cells at earlier stages of differentiation and 208 209 serve as precursor of TAMs (Coffelt et al. 2009). Their presence and frequency have been directly correlated with tumor aggressiveness and is associated with poor survival rates in breast cancer 210 211 (Zhang et al. 2012, Zhao et al. 2017, Mukhtar et al. 2011a, Mahmoud et al. 2012, Leek et al. 1996, 212 Qiu et al. 2018). They have been found to drive cancer progression via immune regulatory functions creating a tolerogenic environment allowing the tumor to progress (Fig. 1). TAMCs inhibit tumor 213 immune responses by blocking T cell functions and proliferation, but they also directly trigger tumor 214 215 growth by promoting cancer stemness, angiogenesis, EMT and metastasis formation. In breast cancer patients, levels of MDSCs in peripheral blood were found to be about 10-fold higher compared to 216 217 healthy control individuals (Safarzadeh et al. 2019). Moreover, they found a direct relationship 218 between MDSC levels and tumor stage of breast cancer patients. The study underlines the importance 219 of MDSCs in tumor progression and invasion which was supported by Diaz-Montero et al. (Diaz-220 Montero et al. 2009) showing that MDSC levels are associated with the clinical stage and metastatic disease burden in patients with breast cancer. MDSCs possess strong immunosuppressive activities 221 222 and interact with other immune cells to regulate their functions. The number and abundance of TAMs

and MDSCs is considered to be an important factor in the clinical success of cancer immunotherapy,

underlining their critical role in suppression of immunity in breast cancer patients (Gnant et al. 2011,Gomez-Roca et al. 2015).

226 AhR plays a critical role in carcinogenesis and tumor immunity (Murray et al. 2014, Xue et al. 2018). 227 Activation of AhR via Kynurenine (Kyn) produced by the immunosuppressive enzyme indoleamine 228 2, 3-dioxygenase (IDO) in glioblastoma cells has been found to induce the accumulation of TAMs 229 (Takenaka et al. 2019, McKay et al. 2021). They reported that the AhR ligand Kyn is able to activate 230 AhR in TAMs, leading to an increased expression of the chemokine receptor CCR2 by TAMs, which 231 enhances the recruitment of TAMs in response to CCL2. Moreover, AhR stimulates the production of 232 the exonucleotidase CD39 by TAMs, interfering with the function of cytotoxic CD8+ T cells 233 (Takenaka et al. 2019). In melanoma patients, high levels of IDO1 are associated with high levels of 234 Kyn and immunosuppression (Campesato et al. 2020). Using a melanoma model, it was shown that 235 tumors expressing high levels of IDO1 present an enrichment of TAMs and selective inhibition of 236 AhR decreases tumor progression, by inhibiting the immunosuppression mediated by IDO1. Another 237 link of AhR with immune response in cancer is highlighted by the fact that AhR mediates the induction of the poliovirus receptor CD155 by IL-4 and LPS in macrophages, as CD155 is 238 239 suppressing T cell function (McKay et al. 2021). In the same line, the inhibition of AhR activity in a 240 model of pancreatic cancer promotes the infiltration of CD8+ T cells and improves the response to 241 immune therapy (Hezaveh et al. 2022). This study also showed that AhR is highly expressed in 242 TAMs, involved in their polarization, and associated with a reduction of iNOS, CCL4, and TNFa 243 levels. Further, Neamah et al. (Neamah et al. 2019) found that treatment with the AhR ligand TCDD induces the accumulation of MDSCs in the peritoneal cavity. Interestingly, we found an 244 245 accumulation of CD11b+ F4/80+ and CD11b+ F4/80- Ly6G+ cell subsets in adipose tissue 246 associated with a significant increase of the chemokine CXCL5 in TCDD-treated mice (Vogel et al. 247 2016) which indicates accumulation of TAMs and MDSCs (Ugel et al. 2015). Although TAMs and 248 MDSCs are regarded as separate populations, some markers including CD11b are shared among 249 TAMs and MDSCs (Ugel et al. 2015). There are specific markers (e.g. Ly6G and Ly6C) that can be 250 used to distinguish them. Further, MDSCs and TANs express high levels of S100A9 and the 251 immunosuppressive enzymes IDO and arginase 1 (Arg1) which are specific for their immune-252 suppressive activity in TME of breast cancer (Fridlender et. al 2009, Ostrand-Rosenberg 2016).

253 The polarization of TAMs and MDSCs within the TME is highly dependent on the local milieu of 254 immune regulatory factors (e.g. C/EBPß and S100A9) and cytokines and chemokines which can 255 originate from stromal cells (Fig 1). Recently, we identified C/EBPB as a critical transcription factor 256 in AhR-dependent induction of S100A9 after treatment with PM rich in PAHs (Dahlem et al. 2020). 257 The S100 calcium binding protein S100A9 has been shown to play a critical role in mediating the 258 expansion of MDSCs in breast cancer models (Zhao et al. 2012). Moreover, S100A9 can act as a 259 transcriptional coactivator during breast cancer development (Song and Struhl 2021) and promotes 260 the immune-suppressive activity of MDSCs (Ostrand-Rosenberg 2016). Regardless of any direct 261 lineage link and distinction between MDSCs, TANs and TAMs, the most important criteria for their 262 role in carcinogenesis are their immune-suppressive and pro-tumoral activities. Importantly, the AhR 263 has been demonstrated to regulate the expression of immune-regulatory markers including Arg1, 264 IDO, IL-10, COX-2, C/EBPB, and S100A9 (Neamah et al. 2019, Vogel et al. 2008, Vogel et al. 2013, 265 Dahlem et al. 2020, Benson and Shepherd 2011, Simones and Shepherd 2011, Bankoti et al. 2010), 266 which are critical factors in the pathogenesis of breast cancer (Dey et al. 2021, Yu et al. 2014, Yu et 267 al. 2013). Moreover, TCDD increased the activity of the immunosuppressive enzyme IDO which mediates tumor immunity in breast cancer cells (Bekki et al. 2015). Interestingly, AhR as well as 268 269 NFkB RelB have been shown to induce IDO expression (Vogel et al. 2008, Yu et al. 2014), which is

also critically involved in the immunosuppressive mechanisms of myeloid-derived suppressor cells

- 271 (MDSCs) in breast cancer (Yu et al. 2013). The number and frequency of TAMs and MDSCs have
- been directly correlated with tumor aggressiveness, and indirectly correlated with clinical outcome in
- breast cancer (Mukhtar et al. 2011b). The literature also shows that accumulation of TAMCs is a
- significant prognostic factor in breast cancer (Zhao et al. 2017). A significant heterogeneity of
- TAMCs in mammary tumors has been described (Movahedi et al. 2010) and the activation of AhR
- has been shown to activate TAMs (Takenaka et al. 2019) and induce the accumulation of MDSCs
- 277 (Neamah et al. 2019). The mechanisms that are driving the polarization of immune-suppressive
- 278 TAMCs in the TME by AhR signaling activated through the exposure to PM, PAHs, and dioxin like
- 279 chemicals are not clear yet. In summary, data from the literature strongly suggest AhR's critical role
- 280 in the microenvironment of mammary tumorigenesis promoting tumor progression and metastasis.

281 5 The Intersection Between Environmental Exposure and Cancer Stemness

282 Breast cancer stem cells (BCSCs), a small but highly plastic subpopulation of tumor cells, have taken

- 283 center stage in the interplay between chemoresistance, recurrence, and metastasis (Shan et. al 2021).
- BCSCs, capable of both self-renewal and recapitulation of tumor heterogeneity, are multidrug-
- resistant (MDR) and highly immune-evasive. MDR is due in part to robust expression of the ABCG2
- efflux protein, also known as Breast Cancer Resistance Protein (BCRP) (Zhou et. al 2001, Zattoni et.
- al 2022), a direct transcriptional target of AhR (Tan et. al 2010). Substantial efforts have focused on
- strategies which will lead to the effective elimination of BCSCs, however it is recognized that
- standard endocrine and chemotherapy regimens paradoxically enrich for BCSCs with mesenchymal
- 290 features, driving tumor recurrence (Li et. al 2008, Creighton et. al 2009, Famta et. al 2022).

291 AhR has been implicated in cancer stemness and immune evasion in various tumor types serving as a 292 "molecular bridge" between environmental exposure and poor patient prognosis. In lung and 293 nasopharyngeal carcinoma cells, AhR was shown to drive the expression of a panel of stemness 294 genes, including ABCG2 (Yan et. al 2018). Interestingly, ABCG2 has been directly implicated in 295 expanding the stem population in osteosarcoma cells (Zhou et. al 2001). In non-small cell lung 296 carcinoma, the deubiquitinase UCHL3 promoted cancer stemness through stabilization of AhR 297 (Ouyang et. al 2020). Recently, activation of AhR by the endogenous ligand kynurenine was linked 298 to colon cancer stemness, immune evasion through PD-L1 induction and metastasis (Miyazaki et. al 299 2022). In an oral squamous cell carcinoma model, tumor cell- and immune cell-expressed AhR 300 collaborated to promote tumor immune evasion with AhR knockout in tumor cells restoring anti-301 tumor immunity (Kenison et. al 2021).

302 In breast cancer, tranilast, a tryptophan metabolite and AhR agonist, was shown to inhibit the BCSC 303 population in MDA-MB-231 (triple negative) breast cancer cells and abrogate metastasis in a tail 304 vein injection model (Prud'homme et. al 2010), in an AhR dependent manner. In agreement with 305 these findings, several studies reported that AhR activation inhibits the BCSC population (Yamashita 306 et. al 2021, Saito et. al 2021). In MCF7 (ER+) cells expressing a constitutively active AhR or treated 307 with the AhR agonists 3-Methylcholanthrene (3MC) or β -naphthoflavone (β -NF), the BCSC 308 population was decreased (Zhao, S. et. al 2012). Most recently, camalexin, an indole phytoalexin 309 and AhR agonist was shown to decrease the BCSC population of MCF7 and T47D (ER+) breast cancer cells (Yamashita et. al 2022). Conversely, AhR activation by the potent agonists TCDD and 310 311 DMBA was found to increase the breast cancer stem cell population and was implicated in doxorubicin resistance of MCF-7 breast cancer cells (Al-Dhfyan et. al 2017). In Tamoxifen-resistant 312 313 MCF7 cells, AhR antagonism inhibited the BCSC population and also inhibited tumor growth 314 (Dubrovska et. al 2012). In Hs578T (triple negative) and SUM149 (inflammatory) breast cancer

- cells, AhR was shown to augment the BCSC population, and its inhibition decreased tumor growth
- and sensitized cells to both adriamycin and paclitaxel (Stanford et. al 2016). This study also found a
- 317 significant correlation between AhR activity and "cancer stem cell- and migration/invasion-
- associated gene sets" in an analysis of 79 human breast cancer cells lines and more than 1,850 human
- 319 breast cancers. In inflammatory breast cancer, AhR was linked to BCSC maintenance through the
- 320 Wnt5a/β-catenin signaling pathway (Mohamed et. al 2018). AhR crosstalk with Wnt/β-catenin
- 321 signaling in the regulation of CSCs has been reported in several studies (Al-Dhfyan et. al 2017,
- 322 Akhtar et. al 2022).
- 323 The role of AhR in cancer stemness and breast cancer stemness more specifically is complex,
- influenced by mode of AhR activation, engagement with various signaling pathways and cell context.
- 325 Nevertheless, the collective evidence strongly suggests that AhR activation by environmental toxins
- 326 and endogenous ligands (Ala 2021) aligns with chemoresistance, recurrence and metastasis, the
- 327 hallmarks of cancer stemness. This places AhR at the intersection between racial/ethnic and
- 328 socioeconomic disparities in toxin exposure in under-resourced neighborhoods, as discussed
- 329 previously, and cancer stemness, undermining response to cancer therapy, worsening the riskscape
- that an individual must navigate. In a recent review by Lagunas-Rangel, the authors pose the question
- 331 "Can Exposure to Environmental Pollutants Be Associated with Less Effective Chemotherapy in
- 332 Cancer Patients?" The authors summarize evidence which strongly supports this hypothesis, which
- includes toxins which activate AhR (Lagunas-Rangel et. al 2022). Therachiyil examines this from the
- 334 perspective of gynecological cancers. (Therachiyil et. al 2022).

335 Conclusion

- 336 Collectively, the body of literature indicates that the role of AhR in cancer is complex, with
- ample evidence for both an oncogenic and tumor suppressor function, depending on cell and
- tissue context and mode of AhR activation. However, exposure studies indicate that
- 339 environmental pollutant-mediated activation of AhR is consistently oncogenic, highlighting the
- 340 potential for cautious therapeutic intervention. The data from human and in vivo studies, as well
- 341 as in vitro experiments suggest that exposure to environmental pollutants especially PAHs and
- dioxin-like chemicals, potent ligands for AhR, increases breast cancer risk and worsens
- 343 outcome through chemoresistance, immune evasion, EMT, tumor cell proliferation, and
- metastasis, linked functional outcomes of cancer stemness (Fig. 2). Some critical questions
- 345 remain, including how AhR activation modulates the tumor microenvironment. This review 346 also highlights the role of AhR at the interface between historical and existing systemic
- 346 also highlights the role of AhR at the interface between historical and existing systemic 347 practices - which reinforce residential segregation and environmental injustice - and the
- molecular drivers of aggressive tumor biology. While policies and molecules are not frequently
- in the same conversation, greater dialogue is needed and opportunities for "upstream" disease
- in the same conversation, greater dialogue is needed and opportunities for "upstream" dise
- 350 prevention through systemic change should be prioritized.

351 Author Contributions

Review concept: CS, GL and CV. Review design: CS and CV. Manuscript preparation: CS, GL and CV. Manuscript editing: CS and CV. Manuscript review: CS, GL and CV.

354 Conflict of Interest

- 355 The authors declare that the research was conducted in the absence of any commercial or financial
- 356 relationships that could be construed as a potential conflict of interest.

357 Funding

- 358 This work was supported by pilot funding from the Molecular Oncology Program at the UC Davis
- 359 Comprehensive Cancer Center and R21ES030419 (CF).

360 Acknowledgements

361 C.S. gratefully acknowledges the dedicated mentorship of Dr. Jann Murray-Garcia.

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1044 **Figure Legends**

1045 Figure 1: Schematic of proposed mechanisms by which traffic-associated air pollution (TRAP)

activates AhR to induce the accumulation of TAMCs (1) and expression of immune suppressive 1046

1047 factors (2) leading to a tumor promoting microenvironment (3) and growth and metastasis of breast 1048

cancer.

1049 Figure 2: Overview of the link between systemic exclusion, environmental injustice, and AhR-driven

1050 tumor biology. Discriminatory housing and lending policies (ex: redlining) drove neighborhood

1051 racial/ethnic and socioeconomic segregation which persists today due to ongoing systemic 1052 discrimination and gentrification. Resources were and are disproportionately allocated to wealthier

1053 neighborhoods, contributing to neighborhood disinvestment. The proximal neighborhood has more

1054 traffic, pollution-generating factories and dump sites. It has less green space for stress relief and

1055 exercise, worsening the riskscape that individuals in the proximal neighborhood must navigate.

1056 Individuals living in the proximal neighborhood are chronically exposed to environmental toxins,

- tipping the scales of environmental justice against them. Polycyclic aromatic hydrocarbons (PAHs), 1057
- Polychlorinated dibenzo-p-dioxins (PCDDs), and Polychlorinated biphenyls (PCBs) are generated by 1058

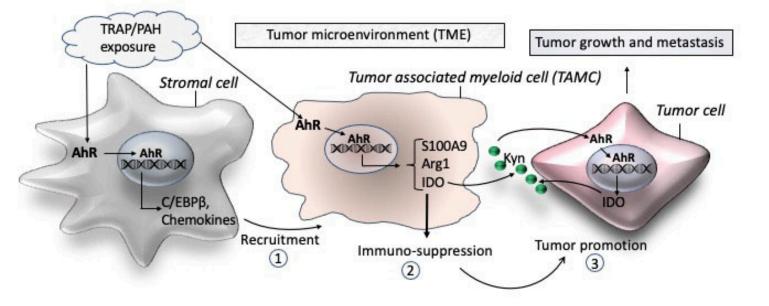
1059 combustion processes as components of ambient particulate matter (PM) derived from urban areas

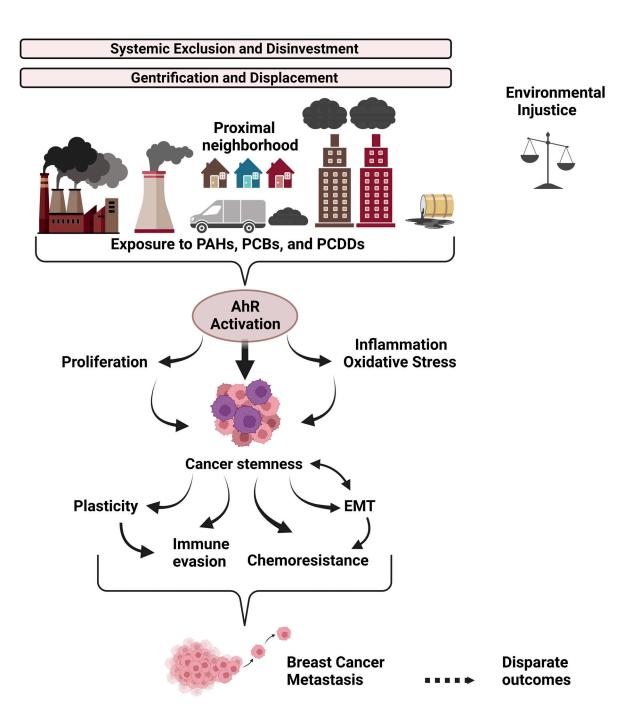
1060 and industrial activities. PAHs, PCDDs and PCBs robustly activate the AhR signaling pathway,

promoting cancer stemness and interrelated functional outcomes, including plasticity. 1061

1062 chemoresistance, EMT and immune evasion, which synergize to drive breast cancer metastasis and

1063 disparate outcomes for individuals in proximal neighborhoods.





Distal neighborhood