



Perspectives on molecular signaling in cancer and update on therapeutic options for the treatment of metastatic cancer

As guest editor of this special series of the *Annals of Translational Medicine (ATM)*, I would like to share with the readers at large the contributions from invited authors who provided their unique and expert perspectives on the important and critical topics of this focus series titled “Cancer Metastasis: Molecular signaling and therapeutic options”. According to the latest data released by the world health organization, cancer is the second leading cause of death globally, accounting for almost 10 million deaths. In 2018, one in six deaths were caused by cancer. While Lung cancer is leading the death toll in both men and women, breast cancer is the most diagnosed cancer and second cause of death in women. The burden of cancer continues to grow globally and exerts significant physical, emotional and financial strain on individuals, families, communities and health systems. This burden is even higher in low- and middle-income countries, where health systems are least prepared to manage this stress-causing burden. Survival rates in cancer patients are however higher in countries where health systems are strong, compared to low and middle-income countries, thanks to easy access to early detection, treatment quality and survivorship care. In addition, and regardless of the economic status of the country of origin, a significant number of cancer patients still does not have access to good quality diagnosis and treatment, even in industrialized countries, therefore, highlighting the fact that cancer is a problem that is not unique to industrialized nations (1). Accordingly, this special series of the ATM provides perspectives on cancer not only in the United States (US), but also worldwide. We present a global, comprehensive view of innovative research, and reviews from basic science at the bench level to clinic interventions and public health considerations. This edition of ATM builds on much of the research and clinical care conducted by authors from the USA, along with articles from basic science researchers, clinicians and public health specialists from Europe and Asia.

The acquisition of the metastatic phenotype is responsible for the death of ~90% of breast cancer (BC) patients (2). In fact, metastatic BC is the 2nd leading cause of cancer-related deaths in women in the United States, annually accounting for more than 41,000 deaths and 260,000 new cases of invasive BC (2). Typically, metastases are incurable and result in a median survival of only 1.5 to 3 years for BC patients. Clinically, ~30% of BC patients diagnosed with early-stage, noninvasive disease will ultimately progress to late-stage, metastatic disease, an event that severely limits treatment options and associates with dismal clinical outcomes (3). This problem is exacerbated by the fact that BCs are heterogeneous and comprised of at least 5 genetically distinct subtypes (4-7). Amongst individual BC subtypes, those classified as triple negative BCs (TNBCs) are especially lethal due to their highly metastatic behavior and propensity to recur rapidly (6,7). As a group, TNBCs lack expression of hormone receptors (ER- α and PR) and ErbB2/HER2, which has prevented the development of FDA-approved targeted drug therapies effective against this BC subtype. Likewise, recurrent TNBCs frequently acquire resistance to standard-of-care chemotherapeutic agents (e.g., doxorubicin, cyclophosphamides, and taxanes) through mechanisms that remain incompletely understood. The reviews by Kansakar *et al.* describes how WAVE3 functions as a major driver of the invasion-metastasis cascade. WAVE3 is a member of the WASP/WAVE family of actin-cytoskeleton remodeling proteins (8,9), and plays an essential role in the regulation of cancer cell migration and invasion, through the regulation of the EMT program (10,11). The oncogenic activity of WAVE3 is also driven by its regulation and maintenance of the cancer stem cell (CSC) subpopulation in TNBC tumors (12). The activity of WAVE3 in cancer cells is also driven by phosphorylation downstream of PI3K, and that this post-translational modification enhances the WAVE3-mediated activation of cancer cell properties, including the activation of tumor growth, invasion, metastasis, and resistance to standard of care therapies (13). The review by Kansakar and colleagues discusses the recent literature highlighting the role of WAVE3 as major player in the oncogenesis of TNBC and other cancer types by regulating several hallmarks of cancer.

Kindlin-2 (K2) has been characterized as a novel regulator of metastatic progression and disease recurrence in cancer (14-16). Kindlins are a small gene family (3 members) of FERM domain-containing adaptor proteins that function as essential drivers of integrin activation (17-19). Moreover, aberrant Kindlin expression and activity is associated with several human pathologies, including cancer (17,18). K2 is the most widely expressed member of the Kindlin family; its homozygous deletion in mice is embryonic lethal, while mice heterozygous at the K2 locus exhibit overtly normal phenotypes that give way to defects in angiogenesis, hemostasis, and the cytoskeletal architecture upon stress induction (19). K2 expression is also

dysregulated in several human cancers, including those originating in the breast. The review by Wang *et al.*, provides an extensive update on the molecular mechanisms involving K2 in the activation of cancer cell behavior, and how therapeutic targeting of K2 may prove to be beneficial for the treatment of cancer.

One of the hallmarks of cancer is the activation of the invasive phenotype of cancer cells (20,21). Cancer cells acquire this invasive property in part through the formation of invadopodia structures or invadosomes, which are developed as cell membrane protrusions composed mainly of F-actin fibers and lipid rafts, but also contain a plethora of enzymatically active proteins (22). Invadopodia are very specialized structures with enhanced proteolytic activities that allow the degradation of the extracellular matrix (ECM) at the contact interface between the ventral surface of cancer cells, which is also enriched in adhesive structures like focal adhesions, and the ECM (23). The proteolytic and degradative activities of invadopodia are believed to play a major role in driving the invasion and metastasis of cancer cells (24,25). The review by Augoff *et al.* provides an in-depth analysis of how the formation and activity of invadopodia are regulated and how invadopodia regulate cancer cell invasion.

As noted above, our current understanding of how BCs become metastatic remains poor, as does our knowledge of how disseminated BCs escape clinical detection by remaining latent for years before reemerging as chemoresistant and incurable secondary tumors. Indeed, the mysteries of metastatic latency have been identified as 1 of the 10 most critical research gaps and translational priorities needed to be solved to alleviate BCs. Detecting disseminated malignancies has always been a big challenge and is highlighted by the fact that analysis of the majority of post-mortem pathology of trauma patients identified undiagnosed micrometastatic lesions (26,27), implying that dormant BC micrometastases play a pivotal role in the majority of BC-associated mortalities. The manuscript by Schiemann *et al.* discusses the role that cancer dormancy and the epigenetic-mediated regulation of dormancy play in cancer pathology, progression and metastasis.

Meanwhile, the manuscript by Yu *et al.* is a comprehensive review of emerging concepts on the role of mitochondrial metabolism in cancer metastasis. The mitochondrial system, in addition to its known status as the power generator for the body, recent data have also found the mitochondrion to be critical for several metabolic functions, such as oxidative phosphorylation, β -oxidation of fatty acids, tricarboxylic acid cycle, calcium handling, proline synthesis, and heme biosynthesis. Accordingly, dysregulation of the mitochondrial metabolic activities has been associated with several metabolic diseases, as well as cancer (28-30). Yu and colleagues discuss how mitochondrial metabolites derivatives and how mitochondrial metabolic plasticity, by adapting to anabolic functions can play a major role in driving oncogenesis and cancer metastasis. This review also highlights how mitochondrial metabolism can be a promising target for novel anticancer therapies.

The focus of the review by Horowitz *et al.*, on the other hand, is entirely dedicated to epithelial ovarian cancer (EOC) and potential treatment modalities for this type of cancer which is yet another devastating cancer to women at different stages of their lives (31). In 2020, it was estimated that more than 60,000 women will lose their lives to EOC (2). Standard of care treatments for EOC, like in TNBC, are still relying on the cytotoxic platinum-based chemotherapies, with very low response rate, mainly because of early disease recurrence and the development of chemoresistance, therefore, accounting for the overall poor outcome and survival rate (32,33). In this review, the authors discuss how the interplay between the different tumor components and the tumor microenvironment drives the chemoresistance phenotype of EOC. The authors also discuss the multitude of cellular and molecular pathways driving the chemoresistance phenotype as well summarize the ongoing clinical trials that are currently available to patients with EOC.

New advances in screening and treatment strategies in breast cancer (BC) has led to a significant decline in breast cancer-related mortality over that past 30 years. However, the improved outcome has not proven to be equitably distributed across the diverse populations world-wide (34). In the United States, specifically, the mortality rate within African American (AA) women is significantly higher (20% to 50%) compared to their European American (EA) counterparts (3,35), therefore, highlighting the deep problem of cancer health disparities between these two populations. This disparity in BC incidence and outcome is further higher in AA women with TNBC, where the probability of being diagnosed with TNBC is more than double in AA than in EA women (36). Furthermore, even though the TNBC represent less than 20% of all BC subtypes (4,5), the incidence rate of TNBC accounts for almost 50% of all BCs in AA (37). The review by Varadan *et al.* discusses how the interplay between the patients' socioeconomic status and the biology of their tumors contributes the health disparities in AA with TNBC.

Finally, Abraham *et al.* provide an in-depth perspective and an update of the therapeutic options that are currently available to patients with metastatic breast cancer and their impact on the patient survival. In addition to the traditional standard of care therapies, the authors also discuss the utility and efficacy of novel targeted and personalized therapies for patients that fail to respond to standard of care therapies.

The authors hope that these reviews will provide new and helpful information to readers with expertise in basic, translational and clinical cancer research, with special interest in metastatic cancer.

Acknowledgments

The author would like to thank all the authors that contributed to this focus series on Cancer Metastasis: Molecular signaling and therapeutic options. The author also tanks members of the Sossey-Alaoui lab for their critical reading and review of the manuscript.

Funding: This work was supported in part by NIH grant R01CA226921 to K. Sossey-Alaoui and by start-up funds from MetroHealth System to K Sossey-Alaoui.

Footnote

Provenance and Peer Review: This article was commissioned by the editorial office, *Annals of Translational Medicine* for the series “Cancer Metastasis: Molecular signaling and therapeutic options”. The article did not undergo external peer review.

Conflicts of Interest: The author has completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/atm-2019-cm-09>). The series “Cancer Metastasis: Molecular signaling and therapeutic options” was commissioned by the editorial office without any funding or sponsorship. KSA served as the unpaid Guest Editor of the series and serves as an unpaid editorial board member of *Annals of Translational Medicine* from Sep 2019 to Aug 2021. The author has no other conflicts of interest to declare.

Ethical Statement: The author is accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

Open Access Statement: This is an Open Access article distributed in accordance with the Creative Commons Attribution-NonCommercial-NoDerivs 4.0 International License (CC BY-NC-ND 4.0), which permits the non-commercial replication and distribution of the article with the strict proviso that no changes or edits are made and the original work is properly cited (including links to both the formal publication through the relevant DOI and the license). See: <https://creativecommons.org/licenses/by-nc-nd/4.0/>.

References

1. Ferlay J, Colombet M, Soerjomataram I, et al. Estimating the global cancer incidence and mortality in 2018: GLOBOCAN sources and methods. *Int J Cancer* 2019;144:1941-53.
2. Siegel RL, Miller KD, Jemal A. Cancer statistics, 2020. *CA Cancer J Clin* 2020;70:7-30.
3. Valastyan S, Weinberg RA. Tumor metastasis: molecular insights and evolving paradigms. *Cell* 2011;147:275-92.
4. Perou CM. Molecular stratification of triple-negative breast cancers. *Oncologist* 2011;16 Suppl 1:61-70.
5. Perou CM, Sorlie T, Eisen MB, et al. Molecular portraits of human breast tumours. *Nature* 2000;406:747-52.
6. Carey L, Winer E, Viale G, et al. Triple-negative breast cancer: disease entity or title of convenience? *Nat Rev Clin Oncol* 2010;7:683-92.
7. Carey LA. Directed therapy of subtypes of triple-negative breast cancer. *Oncologist* 2010;15 Suppl 5:49-56.
8. Sossey-Alaoui K, Head K, Nowak N, et al. Genomic organization and expression profile of the human and mouse WAVE gene family. *Mamm Genome* 2003;14:314-22.

9. Sossey-Alaoui K, Su G, Malaj E, et al. WAVE3, an actin-polymerization gene, is truncated and inactivated as a result of a constitutional t(1;13)(q21;q12) chromosome translocation in a patient with ganglioneuroblastoma. *Oncogene* 2002;21:5967-74.
10. Sossey-Alaoui K, Bialkowska K, Plow EF. The miR200 family of microRNAs regulates WAVE3-dependent cancer cell invasion. *J Biol Chem* 2009;284:33019-29.
11. Sossey-Alaoui K, Li X, Ranalli TA, et al. WAVE3-mediated cell migration and lamellipodia formation are regulated downstream of phosphatidylinositol 3-kinase. *J Biol Chem* 2005;280:21748-55.
12. Bledzka K, Schiemann B, Schiemann WP, et al. The WAVE3-YB1 interaction regulates cancer stem cells activity in breast cancer. *Oncotarget* 2017;8:104072-89.
13. Sossey-Alaoui K, Li X, Cowell JK. c-Abl-mediated phosphorylation of WAVE3 is required for lamellipodia formation and cell migration. *J Biol Chem* 2007;282:26257-65.
14. Sossey-Alaoui K, Pluskota E, Bialkowska K, et al. Kindlin-2 Regulates the Growth of Breast Cancer Tumors by Activating CSF-1-Mediated Macrophage Infiltration. *Cancer Res* 2017;77:5129-41.
15. Sossey-Alaoui K, Pluskota E, Szpak D, et al. The Kindlin2-p53-SerpineB2 signaling axis is required for cellular senescence in breast cancer. *Cell Death Dis* 2019;10:539.
16. Sossey-Alaoui K, Pluskota E, Szpak D, et al. The Kindlin-2 regulation of epithelial-to-mesenchymal transition in breast cancer metastasis is mediated through miR-200b. *Sci Rep* 2018;8:7360.
17. Plow EF, Das M, Bialkowska K, et al. Of Kindlins and Cancer. *Discoveries (Craiova)* 2016;4:e59.
18. Rognoni E, Ruppert R, Fassler R. The kindlin family: functions, signaling properties and implications for human disease. *J Cell Sci* 2016;129:17-27.
19. Plow EF, Qin J, Byzova T. Kindling the flame of integrin activation and function with kindlins. *Curr Opin Hematol* 2009;16:323-8.
20. Hanahan D, Weinberg RA. The hallmarks of cancer. *Cell* 2000;100:57-70.
21. Hanahan D, Weinberg RA. Hallmarks of cancer: the next generation. *Cell* 2011;144:646-74.
22. David-Pfeuty T, Singer SJ. Altered distributions of the cytoskeletal proteins vinculin and alpha-actinin in cultured fibroblasts transformed by Rous sarcoma virus. *Proc Natl Acad Sci U S A* 1980;77:6687-91.
23. Linder S. The matrix corroded: podosomes and invadopodia in extracellular matrix degradation. *Trends Cell Biol* 2007;17:107-17.
24. Revach OY, Geiger B. The interplay between the proteolytic, invasive, and adhesive domains of invadopodia and their roles in cancer invasion. *Cell Adh Migr* 2014;8:215-25.
25. Artym VV, Zhang Y, Seillier-Moisewitsch F, et al. Dynamic interactions of cortactin and membrane type 1 matrix metalloproteinase at invadopodia: defining the stages of invadopodia formation and function. *Cancer Res* 2006;66:3034-43.
26. Klein CA. Framework models of tumor dormancy from patient-derived observations. *Curr Opin Genet Dev* 2011;21:42-9.
27. Hensel JA, Flaig TW, Theodorescu D. Clinical opportunities and challenges in targeting tumour dormancy. *Nat Rev Clin Oncol* 2013;10:41-51.
28. Mishra P, Chan DC. Metabolic regulation of mitochondrial dynamics. *J Cell Biol* 2016;212:379-87.
29. Gorman GS, Chinnery PF, DiMauro S, et al. Mitochondrial diseases. *Nat Rev Dis Primers* 2016;2:16080.
30. Sotgia F, Whitaker-Menezes D, Martinez-Outschoorn UE, et al. Mitochondria "fuel" breast cancer metabolism: fifteen markers of mitochondrial biogenesis label epithelial cancer cells, but are excluded from adjacent stromal cells. *Cell Cycle* 2012;11:4390-401.
31. Lheureux S, Gourley C, Vergote I, et al. Epithelial ovarian cancer. *Lancet* 2019;393:1240-53.
32. Ledermann JA, Raja FA, Fotopoulou C, et al. Newly diagnosed and relapsed epithelial ovarian carcinoma: ESMO Clinical Practice Guidelines for diagnosis, treatment and follow-up. *Ann Oncol* 2018;29:iv259.
33. Morgan RJ Jr, Armstrong DK, Alvarez RD, et al. Ovarian Cancer, Version 1.2016, NCCN Clinical Practice Guidelines in Oncology. *J Natl Compr Canc Netw* 2016;14:1134-63.
34. Yedjou CG, Sims JN, Miele L, et al. Health and Racial Disparity in Breast Cancer. *Adv Exp Med Biol* 2019;1152:31-49.
35. DeSantis CE, Fedewa SA, Goding Sauer A, et al. Breast cancer statistics, 2015: Convergence of incidence rates between black and white women. *CA Cancer J Clin* 2016;66:31-42.
36. Troester MA, Sun X, Allott EH, et al. Racial Differences in PAM50 Subtypes in the Carolina Breast Cancer Study. *J Natl*

Cancer Inst 2018;110:176-82.

37. Scott LC, Mobley LR, Kuo TM, et al. Update on triple-negative breast cancer disparities for the United States: A population-based study from the United States Cancer Statistics database, 2010 through 2014. *Cancer* 2019;125:3412-7.



Khalid Sossey-Alaoui

Khalid Sossey-Alaoui, PhD

Department of Medicine, MetroHealth Medical Center and Case Western Reserve University School of Medicine, Rammelkamp Center for Research, Cleveland, OH 44109, USA. (Email: kxs586@case.edu; ksosseyalaoui@metrohealth.org)

Submitted May 27, 2020. Accepted for publication Jun 10, 2020.

doi: 10.21037/atm-2019-cm-09

View this article at: <http://dx.doi.org/10.21037/atm-2019-cm-09>

Cite this article as: Sossey-Alaoui K. Perspectives on molecular signaling in cancer and update on therapeutic options for the treatment of metastatic cancer. *Ann Transl Med* 2020;8(14):899. doi: 10.21037/atm-2019-cm-09