

Invadopodia: proteolytic feet of cancer cells

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Abstract: The leading cause of death in cancer patients is metastasis. Invasion is an integral part of metastasis and is carried out by proteolytic structures called invadopodia at the cellular level. In this introductory review, we start by evaluating the definition of invadopodia. While presenting the upstream signaling events involved, we integrate current models on invadopodia. In addition, we discuss the significance of invadopodia in 2D and 3D and in vivo. We finally point out technical challenges and conclude with open questions in the field.

Key words: Metastasis, invadopodia

1. Metastasis

The leading cause of death in cancer patients is metastasis. Metastasis defines both the process of spreading of cancer cells from the primary tumor and the resulting secondary tumors. The primary tumor changes its place (meta + stasis) and new tumors form at distant sites. During metastasis of carcinoma (cancer of epithelial tissue), tumor cells proliferate in an uncontrolled fashion, induce angiogenesis (new blood vessel formation), degrade the underlying basement membrane and penetrate into the connective tissue, migrate towards blood vessels, intravasate (enter blood vessels), survive in the blood circulation, extravasate (exit blood vessels), and form secondary tumors in distant organs (Figure 1A). Therefore, cancer metastasis is a disease of altered cell adhesion, motility, and invasion.

2. Definition of invadopodia

Under physiological conditions such as bone resorption, cells invade into tissues in a tightly regulated manner. Normal bone osteoclasts form special cellular structures called podosomes to degrade and thus remodel the bone matrix. During cancer metastasis, tumor cells perform uncontrolled invasion using cellular structures called invadopodia (Figure 1B). The term invadopodia was first used by Chen (1989) to describe membrane protrusions involved in the local degradation of the extracellular matrix. After 25 years, the field has grown to be complex and rather complicated even in terms of definitions. There are 3 major structures in cells, each of which can be defined

in terms of their molecular components and the functions they carry out. These are focal adhesions, podosomes, and invadopodia. They do have similarities, but they are also distinct from one another. In attempts to clear up some of the confusion, podosomes and invadopodia have also been collectively called invadosomes. Although focal adhesions do share common protein markers with podosomes, they were thought to be more distinct from podosomes and invadopodia; however, recently proteolytic activity has also been observed for these structures, further blurring the borders between the definitions of these structures (McNiven, 2013). Available data raise the question of whether focal adhesions, podosomes, and invadopodia share a common precursor. A conservative comparison of focal adhesions, podosomes, and invadopodia is presented in the Table. Please note that this table is compiled conservatively to include the data most commonly agreed upon in the literature. There are also several reviews and milestone papers describing in detail various aspects summarized here (Ayala et al., 2006; Gimona and Buccione, 2006; Linder, 2007; Gimona et al., 2008; Caldieri et al., 2009; Linder, 2009; Yilmaz and Christofori, 2009; Linder et al., 2011a, 2011b; Murphy and Courtneidge, 2011; Oser et al., 2011; Yamaguchi, 2012). In particular, there are comprehensive reviews on the signaling mechanisms involved (Stylli et al., 2008; Destaing et al., 2011; Hoshino et al., 2013). In this review, we will focus on invadopodia, integrate current models, and point out technical challenges and open questions in the field.

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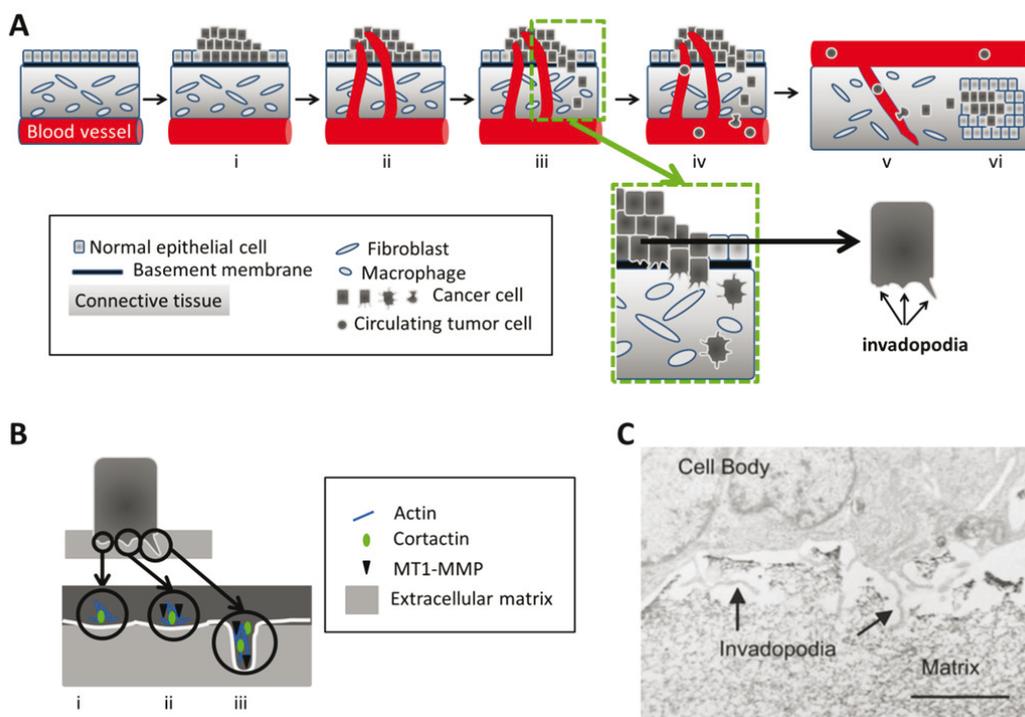


Figure 1. Metastasis and invadopodia. A. Metastasis comprises (i) uncontrolled proliferation, (ii) angiogenesis, (iii) invasion, (iv) intravasation, (v) extravasation, and (vi) secondary tumor formation. Invasion involves matrix degradation carried out by invadopodia. B. Invadopodia form and mature at multiple stages: (i) initiation, (ii) stabilization, and (iii) maturation (see also Figure 3). Initiation involves recruitment of actin and cortactin, MT1-MMP recruitment leads to stabilization. Maturation stage involves matrix degradation. When cells are on a thick matrix, invadopodia appear as membrane protrusions penetrating into the extracellular matrix. C. Electron micrograph of an MDA-MB-231 cell cultured on gelatin. The ultrastructure of invadopodia (arrows) is shown. Reprinted by permission from Macmillan Publishers Ltd: Oncogene (Bowden et al., 1999) copyright 1999.

Invadopodia are relatively dynamic, actin rich, proteolytic cellular structures formed by invasive cancer cells (Bowden et al., 2006; Buccione et al., 2009; Linder, 2009; Linder and Aepfelbacher, 2003) (Figure 2). Invadopodia can be from a few hundred nanometers to several microns wide and can be up to 8 micrometers if the underlying matrix is thick enough (Baldassarre et al., 2003). Invadopodia also form on stiff substrates such as glass and thus they can be studied with high resolution imaging (DesMarais et al., 2009). The molecular markers for invadopodia include actin and its associated proteins cortactin, Arp2/3, N-WASP, Nck1, and cofilin as well as matrix metalloproteinase MT1-MMP (Artym et al., 2006; Stylli et al., 2008; Oser et al., 2009, 2011). In addition, actin filaments, microtubules, and intermediate filaments cooperate during invadopodia elongation (Schoumacher et al., 2010). In melanoma cells, invadopodia contain $\alpha 3 \beta 1$ integrin at the core and $\alpha 5 \beta 1$ integrin at the periphery (Mueller et al., 1999). Integrins at invadopodia may function to signal and to focus degradation of the extracellular matrix (ECM) (Buccione et al., 2009; Mueller

et al., 1999). However, it is unclear if invadopodia have an adhesive function as they lack vinculin (Gimona et al., 2008; Linder, 2009). That is, whether invadopodia require local adhesion at the sites of formation is unknown. Preliminary results from our lab using nano-patterned surfaces suggest that invadopodia do not require local adhesion (unpublished data). Experiments using nano- and micro-patterned substrates can present valuable approaches to answer such questions and other aspects of invadopodia/podosome research such as dynamics of mechanical properties (Labernadie et al., 2010).

In images of cells forming invadopodia, the Golgi complex appears to be polarized and juxtaposed to the site of invadopodia, suggesting a link between proteolytic activity and membrane transport (Baldassarre et al., 2003; Buccione et al., 2009; Caldieri and Buccione, 2010). However, if and how the spatial positioning of invadopodia is controlled is not known. In addition, the position and orientation of the Golgi can be modulated by micrometer scale surface patterns (They et al., 2006). Therefore, if invadopodia are positionally linked to the

Table. Comparison of focal adhesions, podosomes, and invadopodia.

	Focal adhesions	Podosomes	Invadopodia
Cell type	virtually all cells	osteoclasts, macrophages, endothelial cells, smooth muscle cells	cancer cells
Function	adhesion, matrix degradation?	matrix degradation	matrix degradation
Cellular localization	cell periphery	distributed	leading edge and proximal to Golgi
Composition	Actin Vinculin Talin Paxillin Focal adhesion kinase Integrin	Actin Vinculin WASP Grb2 MT1-MMP	Actin Arp2/3 Cortactin N-WASP Nck1 Cofilin Tks5 MT1-MMP
Shape	ellipse	ring	dot
Size	<20 μm	<1 μm \times 4 μm	<8 μm \times 5 μm
Number per cell	<400	20–100	1–40
Stability/ Persistence	stable/several hours	highly dynamic/2–12 min	dynamic/up to 3 h

Golgi, changing the position of the Golgi by culturing cells on different micrometer scale surface patterns should also change the localization of invadopodia. Thus micro-patterned substrates present themselves as valuable tools for the question at hand.

3. Upstream of invadopodia

Growth factors act as intercellular signaling molecules that promote various processes such as cell growth, proliferation, differentiation, and motility. In addition, growth factor receptors and integrins are known to cross-talk (Moro et al., 2002; Yamada and Even-Ram, 2002; Alam et al., 2007; Gilcrease, 2007). Growth factors can be soluble, transmembrane, or ECM bound proteins (Ruoslahti et al., 1992; Massague and Pandiella, 1993; Taipale and Keski-Oja, 1997). Epidermal growth factor (EGF) is 1 of the 7 ligands of EGF receptor (EGFR also known as ErbB1), which is in turn the most studied member of the ErbB receptor family (Cohen, 1962; Carpenter and Cohen, 1990; Harris et al., 2003; Singh and Harris, 2005). Furthermore, EGFR expression correlates with poor prognosis in breast cancer (Sainsbury et al., 1985; Lewis et al., 1990; Memon et al., 2006). EGF is known to induce motility and invadopodia formation in breast cancer cells (Yamaguchi et al., 2005). However, whether EGFR is present at invadopodia and acts directly and locally or not is not known.

In terms of signal transduction, growth factor receptor tyrosine kinase and integrin initiated upstream events have been shown to promote invadopodia formation through phosphorylation of cortactin via a Src and Arg dependent

pathway (Stylli et al., 2008; Oser et al., 2010; Destaing et al., 2011; Mader et al., 2011; MacGrath and Koleske, 2012). $\beta 1$ integrin has been shown to promote metastasis, invadopodia maturation, and matrix degradation through Arg (Beatty et al., 2013). Local changes in pH induced by NHE1 are also shown to regulate cortactin phosphorylation (Magalhaes et al., 2011). Furthermore, small GTPases are shown to be spatiotemporally regulated at invadopodia where RhoC is inactivated at the center of invadopodia and is activated at its periphery so that cofilin is active at the center and is inactive at the periphery (Bravo-Cordero et al., 2011, 2012). Here, RhoC is shown to act through ROCK, which phosphorylates LIMK, which in turn phosphorylates and inactivates cofilin.

4. An integrated model of invadopodia

Over the years, valuable research has produced models that describe invadopodia. An integrated model is presented in Figure 3. One of the early studies classified invadopodia formation into 4 stages: I. Invadopodia initiation, II. Preinvadopodia, III. Mature invadopodia, and IV. Late invadopodia. Cortactin levels are at their maximum at stages II and III and subside afterwards, while actin levels reach a peak at stage III. MT1-MMP reaches a maximum at stage II and is stable thereafter, while matrix degradation saturates at stage III (Artym et al., 2006).

Later on, a more detailed model was presented by Oser et al. (2009), pointing out the central role of cortactin in invadopodia. Here, cortactin was shown to regulate cofilin and N-WASP activities and thus control the stages

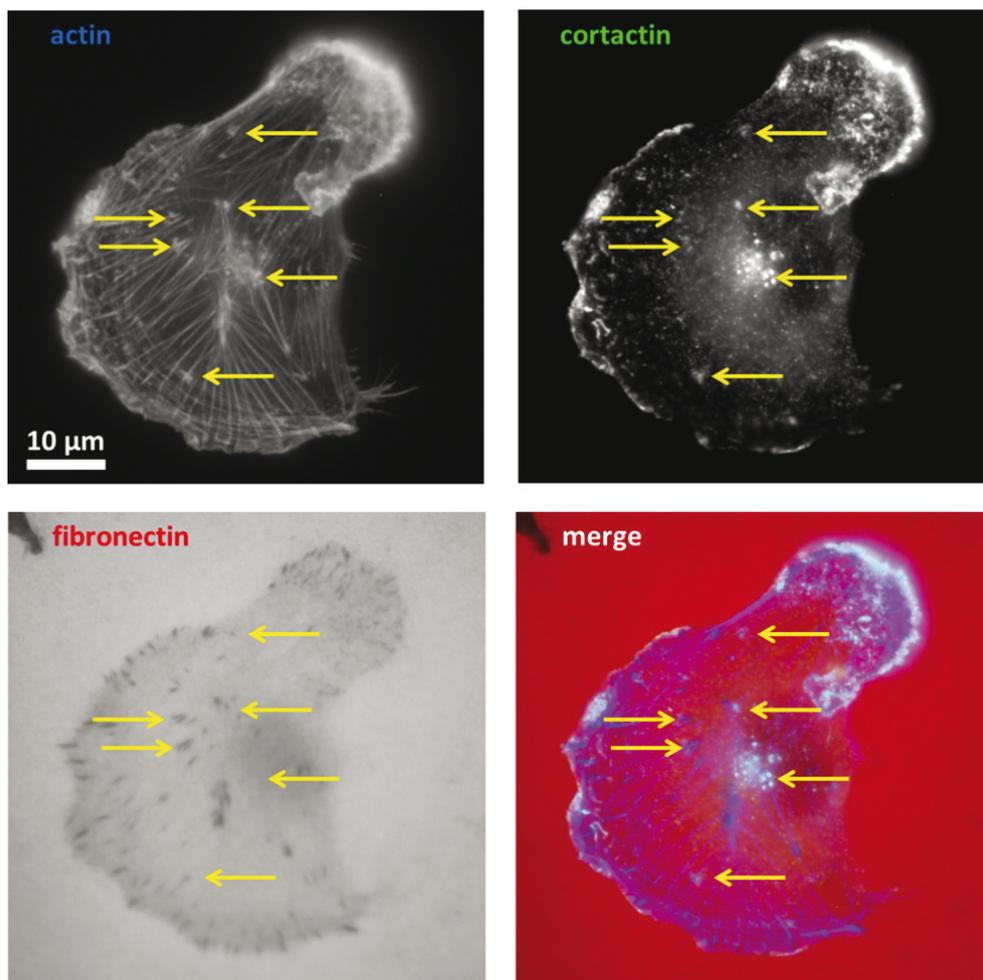


Figure 2. Immunofluorescence images of invadopodia in MDA-MB-231 breast cancer cells cultured on fibronectin, an extracellular matrix protein. Actin, cortactin, and fibronectin were labeled using blue-fluorescent phalloidin, cortactin specific antibodies followed by green-fluorescent secondary antibodies, and fibronectin specific antibodies followed by red-fluorescent secondary antibodies, respectively. Yellow arrows point to invadopodia. Cortactin and actin colocalize at invadopodia. At mature invadopodia, the underlying matrix of fibronectin is also degraded. In the merged image, cortactin, actin, and fibronectin are shown in green, blue, and red, respectively.

of invadopodia formation. Four stages were redefined here: Stage I – Precursor formation: Cortactin, N-WASP, cofilin, and Arp2/3 form a complex. Stage II – Activation of actin polymerization: Nck1 joins the precursor complex while phosphorylation of cortactin activates cofilin's severing activity, which in turn provides free barbed ends for Arp2/3 for new actin polymerization. Stage III – Stabilization: Cortactin is dephosphorylated, cofilin re-joins the complex, and invadopodia precursors are stabilized. Stage IV – ECM degradation: MT1-MMP is recruited to invadopodia and ECM is degraded.

The model by Oser et al. was then refined in terms of involvement of Tks5 and SHIP2, which are shown to be required for invadopodia maturation but not initiation

(Sharma et al., 2013). The integrated model we present here comprises 3 stages: initiation, stabilization, and maturation. Initiation here describes a combination of the previously described stages I and II and involves structural complex formation and actin polymerization. Stabilization includes MMP recruitment. At the maturation stage, MMPs are activated and matrix degradation takes place.

To recapitulate, the first events in the signal transduction pathways that result in invadopodia formation are integrin and/or growth factor activation. Although the intermediates are not entirely known, activation of Src is the key event for invadopodia formation. Src in turn activates Arg, which phosphorylates cortactin. While unphosphorylated cortactin, unphosphorylated

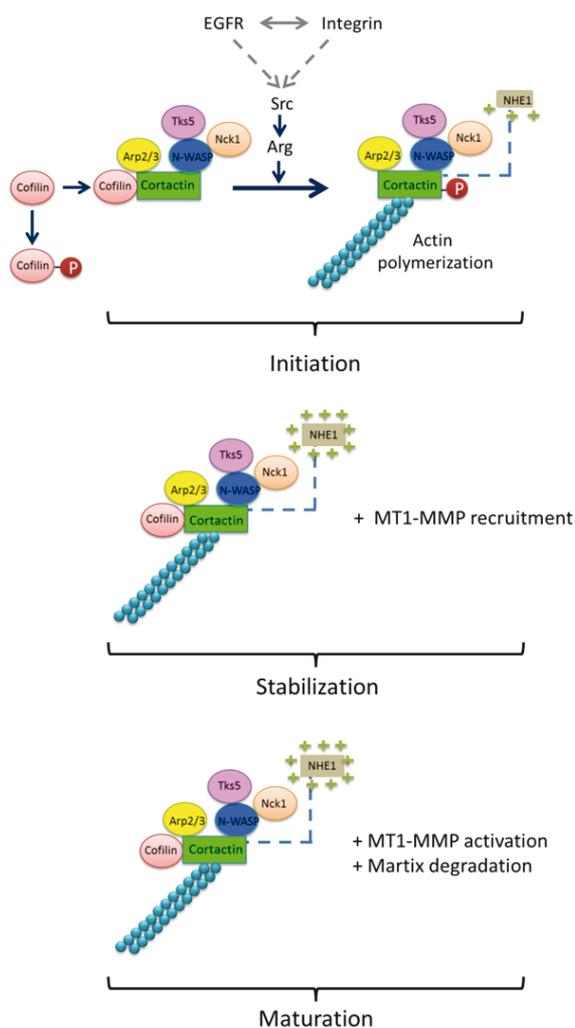


Figure 3. Integrated model for the initiation, stabilization, and maturation of invadopodia. Presented is a combination of various models published in the literature through the years. Initiation: Growth factor and/or integrin initiated signaling cascades result in phosphorylation of cortactin by Arg, which in turn is activated by Src. Phosphorylation of cortactin releases cofilin from the invadopodial complex comprising Arp2/3, N-WASP, Nck1, and Tks5. Release of cofilin promotes actin polymerization. Stabilization: Cortactin is dephosphorylated and cofilin is re-recruited to the complex. NHE1 induces local decrease in pH and MT1-MMP is recruited. Maturation: MT1-MMP is activated and matrix degradation takes place.

cofilin, N-WASP, Tks5, Nck1, and Arp2/3 coexist in the pre-initiation complex, after phosphorylation of cortactin, cofilin leaves the complex and enables actin polymerization. At the same time, NHE1, which causes a decrease in the local pH, is recruited. During stabilization, cofilin is dephosphorylated and binds back to cortactin, and MT1-MMP is recruited to invadopodia. Finally, in

the maturation stage MT1-MMP is activated and matrix degradation takes place.

5. Invadopodia in 2D and 3D and in vivo

Invadopodia are observed in both 2D and 3D cell cultures in vitro. Most research has been carried out in 2D cell culture, while studies in 3D conditions are increasing. In 2D cell culture, invadopodia are found at the ventral surface of cells. They even form on substrates such as glass without a matrix coating. If the substrate underneath cancer cells is degradable and thick enough, invadopodia extend as proteolytic protrusions into the matrix. When cancer cells are embedded in a 3D matrix, the definition of a ventral surface dissolves and cancer cells can form proteolytic protrusions at various points. As in 2D, how the cellular localization of invadopodia is determined in 3D is unknown. However, invadopodia are composed of actin, cortactin, cofilin, Tks5, and MT1-MMP in both 2D and 3D cultures and in vivo settings (Blouw et al., 2008; Lizarraga et al., 2009; Magalhaes et al., 2011; Gligorijevic et al., 2012; Yu Machesky, 2012).

Invadopodia have been observed in 2D and 3D in vitro settings. However, the in vivo and physiological relevance has only been recently clarified. N-WASP expression is shown to increase in invasive breast cancer (Yu et al., 2012). Expressions levels of cortactin have been positively correlated with aggressiveness of head and neck squamous cell carcinoma and breast carcinoma (Buday and Downward, 2007; Clark et al., 2009). Expression level of Tks5 has been shown to increase in breast cancer and glioma (Seals et al., 2005; Stylli et al., 2012). MMP expression levels are known to be differentially regulated in various cancers (Kessenbrock et al., 2010). Early MMP inhibitors failed in clinical trials, and their nonproteolytic functions are speculated to be one of the reasons. However, there are still promising candidates such as MT1-MMP (Chen et al., 1994; Chen, 1996; Sabeih et al., 2009). In addition, upstream players that induce invadopodia formation such as Mena^{INV}, Arg, IL-6, EGFR, and faciogenital dysplasia protein Fgd1, are known to have increased expression levels in various cancers (Ayala et al., 2009; Clark et al., 2009; Li et al., 2010b; Wang et al., 2004, 2007; Gil-Henn et al., 2013). Thus both upstream regulators and structural components of invadopodia present vital opportunities for diagnosis and therapy.

6. Technical bottlenecks for research on invadopodia

A technical limitation for research on invadopodia has been the limited number of assays for proteolytic activity. Fluorescently labeled gelatin or fibronectin is commonly used in addition to DQ-collagen, which becomes fluorescent upon degradation. In addition, Packard et al. (2009) have used a substrate that shows sites of degradation

by MT1-MMP. Fluorogenic peptide substrates have also been utilized to assay matrix degradation by MMPs (Leight et al., 2013). The field would greatly benefit from novel assays that allow the determination of matrix degradation, particularly for 3D culture and in vivo settings.

Another bottleneck has been the difficulty in analyzing invadopodia in a quantitative manner. Counting invadopodia in a cell or in a field of view based on co-labeling of actin and cortactin, for instance, requires either brute force of manual counting or elegant image processing approaches. An alternative approach has been to quantify the area of matrix degraded by invadopodia rather than counting individual structures (Li et al., 2010a). Well-designed image processing approaches would be greatly beneficial for research on invadopodia.

7. Conclusions and open questions

In conclusion, while our understanding of invadopodia continues to improve despite confusion even at the definitions level, the field requires the incorporation of new technologies and there are many open questions waiting to be answered, such as: Do focal adhesions, podosomes, and invadopodia share a common precursor? Do invadopodia have an adhesive function? How is the cellular localization of invadopodia controlled? Is EGFR present at invadopodia and does it act directly or indirectly? How can we better assay the proteolytic activity of invadopodia? How can we improve the quantitative analysis of invadopodia? How

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can we better exploit upstream regulators and structural components of invadopodia for the diagnosis and therapy of cancer?

Glossary

Cortactin: Cortical actin binding protein. Cortactin promotes actin polymerization.

Cofilin: An actin binding protein. Cofilin severs actin filaments.

N-WASP: Neural Wiskott–Aldrich syndrome protein. N-WASP promotes actin nucleation.

Tks5: Tyrosine kinase substrate 5, a scaffold protein.

Nck1: Noncatalytic region of tyrosine kinase adaptor protein 1. Nck1 is an adaptor protein involved in signal transduction.

MT1-MMP: MT1-MMP is a matrix metalloproteinase also known as MMP-14.

Src: The first described proto-oncogene. A nonreceptor tyrosine kinase.

Arg: A member of the Abelson family of nonreceptor tyrosine kinases.

NHE1: A Na⁺/H⁺ exchanger.

RhoC: A member of the Rho family of small GTPases.

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