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The non-redundant role of N-WASP in podosome-mediated matrix degradation in macrophages

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Abstract

Wiskott-Aldrich Syndrome Protein (WASP) is a hematopoietic cell-specific regulator of Arp2/3dependent actin polymerization. Despite the presence of the highly homologous N-WASP (neural-WASP), macrophages from WAS patients are devoid of podosomes, adhesion structures in cells of the monocytic lineage capable of matrix degradation via matrix metalloproteases (MMPs), suggesting that WASP and N-WASP play unique roles in macrophages. To determine whether N-WASP also plays a unique role in macrophage function, N-WASP expression was reduced using silencing RNA in a sub-line of RAW 264.7 macrophages (RAW/LR5). Similar to reduction in WASP levels, cells with reduced N-WASP levels were rounder and less polarized. Interestingly, podosomes still formed when N-WASP was reduced but they were unable to perform matrix degradation. This defect was rescued by re-expression of N-WASP, but not by over-expression of WASP, indicating that these proteins play distinct roles in podosome function. Additionally, reducing N-WASP levels mistargets the metalloprotease MT1-MMP and it no longer localizes to podosomes. However, N-WASP was only found to co-localize with MT1-MMP positive vesicles at podosomes, suggesting that N-WASP may play a role on the targeting or fusion of MMP containing vesicles to podosomes in macrophage-like cells.

Keywords

WASP; N-WASP; podosomes; macrophages; matrix metalloproteases; matrix degradation; vesicle trafficking

Introduction

Macrophages play an important role in the immune response through phagocytosis of foreign pathogens and subsequent antigen presentation. However, macrophages may also destroy non-infected cells when they are recruited to sites of inflammation. In tumors, they may speed the progression of metastasis and their recruitment into breast tumors has been correlated with a poor prognosis (reviewed in (Pixley and Stanley 2004)).

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Wiskott-Aldrich Syndrome and X-linked thrombocytopenia are X-linked diseases caused by mutations in the gene encoding WASP and are characterized by immune system malfunctions including thrombocytopenia, eczema, recurring infections, and autoimmune disorders (reviewed in Thrasher and Burns (Thrasher and Burns)). However, while WASP is only found in hematopoetic cells, the ubiquitously expressed WASP homolog N-WASP is also present in these cells, albeit at low levels (Isaac et al. 2010; Snapper et al. 2001; Stamm et al. 2005; Suzuki et al. 2002), suggesting that N-WASP cannot compensate for the reduced WASP protein levels typically seen in WAS patients or in the WASP knockout mouse.

WASP is important in Arp2/3 mediated actin polymerization and, among other things, regulates the formation of podosomes, dynamic adhesion structures that are localized to the ventral surface of cells that mediate matrix degradation by localization of various enzymes including matrix metalloproteases (MMPs) (Linder 2007; Yamaguchi et al. 2006). It has also been shown that podosomes are absent in macrophages and DCs derived from WAS patients, with disorganized clusters of F-actin lacking the typical vinculin ring (Burns et al. 2001; Linder et al. 1999). Podosome-like structures are also present in non-hematopoietic cells such as endothelial cells, vascular smooth muscle cells, Src-transformed cells and in certain aggressive cancer cells where they are often called invadopodia. These structures have distinct organization and they contain N-WASP in the F-actin core instead of WASP (reviewed in (Linder 2007)).

N-WASP has 50% overall sequence homology with WASP, with conserved domain organization, including a WH1 domain that binds WASP interacting protein (WIP), a proline-rich region that binds SH3 proteins, a GBD domain that binds Cdc42-GTP, and a basic sequence that binds PIP_2 (Miki et al. 1996). At the C terminus, both proteins have a VCA domain, which binds actin via the V domain and the Arp2/3 complex via the CA region but N-WASP has an additional V domain. In vitro data using the purified VVCA domain of N-WASP showed that this domain alone was sufficient for actin polymerization and its activity was higher than the isolated VCA domain of WASP (Zalevsky et al. 2001). This suggests that N-WASP is a better inducer of actin polymerization due to its additional V domain. Additionally, WASP and N-WASP are differentially activated by SH3 proteins in vitro where Grb2 and Nck1 are more potent activators of N-WASP than WASP (Tomasevic et al. 2007), consistent with potentially different cellular functions. Also, studies using pathogenic microorganisms suggest WASP family proteins may not compensate for each other completely. For example, Shigella outer membrane protein VirG binds to N-WASP and not to WASP and the defect in Shigella motility in the absence of N-WASP cannot be restored by ectopic expression of WASP (Snapper et al. 2001; Suzuki et al. 2002). However, WASP does compensate for N-WASP in cytoplasmic vaccinia virus motility (Snapper et al. 2001) and Mycobacterium marinum can exploit either WASP or N-WASP interchangeably (Stamm et al. 2005). These studies suggest that WASP and N-WASP may serve redundant or non-redundant functions depending on the cellular context.

Due to the differences between WASP and N-WASP in their activation and cellular processes, it is speculated that they may play unique, nonredundant roles. While many studies have focused on WASP, the function of N-WASP in macrophages has not been clearly elucidated and is the subject of this short communication.

Material and Methods

Cells, Constructs, antibodies and other reagents

All cells were maintained at 37° C in a 5% CO₂ incubator. Lac Repressor expressing murine monocyte/macrophage RAW 264.7 cell line (RAW/LR5) (Cox et al. 1997) were grown in RPMI medium (Mediatech Inc, Manassas, VA) containing 10% new born calf serum

(Cambrex, Walkersville, MD), 100 U/ml penicillin and 100 ug/ml streptomycin (Sigma, Saint Louis, MO). RAW/LR5 cells were transfected using Fugene HD (Roche, Indianapolis, IN) with bovine Wild-N-WASP and N-WASP Δ V in pSR α (Miki et al. 1998), GFP-N-WASP (Yamaguchi et al. 2002), myc-tagged WT human WASP construct in pEFBos. GFP or mCherry tagged MT1-MMP was a generous gift from Jose Bravo-Cordero (Bravo-Cordero et al. 2007). RNA-mediated interference of WASP and N-WASP was performed according to Park and Cox (Park and Cox 2009). Chicken anti-N-WASP (Clone AE920) (Isaac et al. 2010) was generated based on a sequence indicated in (Miki et al. 1996). The following commercial antibodies were also used: monoclonal mouse anti- β -actin (Clone AC-15) and anti-vinculin (hVin1; Sigma), monoclonal mouse anti-MT1-MMP (Clone Ab-4; Calbiochem), mouse anti-GFP (Roche), rabbit anti-WASP (H250; Santa Cruz). Alexa Fluor 647-, 568-, and 488-conjugated secondary antibodies or phalloidin were from Molecular Probes, Eugene, OR).

Immunofluorescence microscopy

Cells were plated on either 12 mm glass coverslips or 35 mm uncoated glass bottom microwell dishes (MatTek Corporation, Ashland, MA) and were allowed to adhere overnight. Cells were fixed for 7 minutes in 3.7% formaldehyde in BWD buffer (20 mM HEPES, 125 mM NaCl, 5 mM KCl, 5mM glucose, 1mM KH₂PO₄, 10 mM NaHCO₃, 1mM CaCl₂, 1mM MgCl₂, pH 7.4), permeabilized for 5 minutes in 0.2% Triton X-100 (in BWD), and blocked in 10% milk in TBS (137 mM NaCl, 24.7 mM Tris, pH 7.4). For MT1-MMP staining, LR5 cells were simultaneously fixed and permeabilized with saponin to preserve membrane integrity, as described (Eddy et al., 2000). Cells were incubated with the primary antibodies of interest in TBS and then with fluorophore-conjugated secondary antibodies. Finally, coverslips were mounted in antifade mounting medium (50% glycerol, 1X PBS, 100mM n-Propyl gallate (Sigma)) for widefield or in PBS for TIRF microscope (Olympus, X150/1.45 NA, oil objective) coupled to a Sensicam cooled CCD camera.

Morphological parameters were analyzed as follows: a cell displaying migratory morphology was defined by an unambiguous leading and trailing edge (Wheeler et al. 2006). The elongation index was determined using Image J (http://rsb.info.nih.gov/ij/) by tracing the cells and measuring the ratio of the major axis over the minor axis of the fit ellipse. At least 60 cells per experiment were analyzed for each determination. Podosomes were identified as F-actin punctate structures surrounded by a vinculin ring. F-actin core intensity was obtained by identifying the area inside the vinculin ring from F-actin and vinculin merged image and circling the corresponding area on F-actin alone image for intensity measurement. At least 100 podosomes per cell type were analyzed for each experiment.

Matrix degradation assay

Degradation of fluorescently-labeled fibronectin (FN) by RAW/LR5 cells was determined by a protocol detailed in (Dovas et al. 2009). Briefly, cells were plated on either Alexa488 or Alexa568 fluorescently labeled fibronectin for 16 hours and areas of loss of fluorescence were measured as degradation area per cell. For rescue experiments shWASp or shN-WASP cells, were transfected with the indicated constructs followed by plating on Alex568 labeled fibronectin as described above. Transfected cells were identified by either GFP labeling or staining for Myc. At least 50 cells per experiment were analyzed.

Western blotting

Cells were lysed for 10 minutes in ice-cold lysis buffer (1% Triton X-100, 25 mM Tris, 137 mM NaCl, 2 mM EDTA, 1 mM sodium orthovanadate, 1 mM benzamidine, 10 μ g/ml aprotinin, 10 μ g/ml leupeptin, pH 7.4). Clarified whole-cell lysates were mixed with 5X

Laemmli buffer and boiled. Samples were run on an 8% SDS-PAGE gel, transferred onto PVDF membranes (Immobilon-P, Millipore), and blotted with appropriate antibodies. Blots were exposed using Kodak IS440 machine and bands were quantified using Kodak ID 3.6 (Kodak Imaging Systems).

Data analysis

Results were considered statistically different when analysis using a Student t-test resulted in differences between two means with a p value of less than 0.05. All graphs indicate mean values and error bars signify standard error of the mean.

Results and Discussion

N-WASP is required for macrophage polarization

It has been suggested in the field that WASP and N-WASP have non-redundant functions given the defect associated with the absence of WASP despite the presence of N-WASP (Calle et al. 2006). To determine if N-WASP has a distinct function from WASP, we initially determined N-WASP localization in a macrophage cell line (RAW/LR5) transiently transfected with GFP-N-WASP (Fig. 1). While WASP has been consistently localized to podosomes in human and mouse macrophages, dendritic cells and osteoclasts (Calle et al. 2004a; Calle et al. 2004b; Linder et al. 1999), N-WASP was observed in both podosomes (Fig. 1A, inset) and other cellular puncta (arrows; Fig. 1A). These puncta did not display distinct podosome characteristics, namely, an F-actin core surrounded by a vinculin ring, and are dispersed throughout the cytoplasm consistent with the vesicular localization of N-WASP observed in dendritic cells and in breast carcinoma cells (Calle et al. 2006; Parsons et al. 2005). It should be noted that while N-WASP was not present in dendritic cell podosomes, GFP-N-WASP significantly localized to macrophage podosomes (Fig. 1B). The localization of N-WASP to F-actin rich adherent structures was also confirmed using TIRF microscopy (Supplemental Fig. 1). This result may be due differences in cell types, dendritic versus macrophages, or to the over-expression of N-WASP. To minimize the latter possibility only cells that expressed low levels of GFP-N-WASP were analyzed. Attempts to localize endogenous N-WASP were unsuccessful since all of the antibodies available against N-WASP cross-reacted with WASP. However, the presence of N-WASP in cellular puncta suggests that it may perform unique functions in macrophages.

To determine the specific role of N-WASP in macrophages, N-WASP levels were reduced using silencing RNA alone or in combination with silencing RNA against WASP. Reduction of N-WASP using short hairpin RNA (shRNA) resulted in no stable, viable cells, suggesting that N-WASP is an essential protein. In mice, unlike WAS^{-/-}, N-WASP^{-/-} is embryonic lethal (Snapper et al. 2001). Therefore, a transient approach to reducing N-WASP was used by selecting in puromycin media for two days to remove non-infected cells. At this point N-WASP levels were reduced by approximately 80% and yet the cells were still viable as determined by trypan blue exclusion (data not shown). Under the same conditions WASP levels were reduced by more than 85% (Fig. 2A). Consistent with other reports using monocytes and macrophages from WAS patients (Badolato et al. 1998; Zicha et al. 1998), shWASP cells appeared to be more round and fewer cells showed a clear head to tail migratory phenotype than control cells. Reduced levels of N-WASP also decreased cell elongation and migratory morphology. Consistent with a role for N-WASP in the generation of a migratory morphology others have shown that N-WASP is required for carcinoma cells to extend polarized protrusions towards a chemoattractant (Desmarais et al. 2009) and epithelial cell chemotaxis (Yamaguchi et al. 2002). While reduction of both N-WASP and WASP did not enhance the effect on cell polarity and elongation greater than that caused by the reduction of WASP alone (Fig. 2B and C), this was most likely because the combined

shRNA treatment did not reduce the levels of each as effectively as use of single shRNA (Fig. 2A). The observed effects on cell morphology may be caused by the effects of WASP and N-WASP in lamellipodial and filopodial extension through their interaction with the Arp2/3 complex in macrophages (Hufner et al. 2002). In addition, this data also suggest that WASP and N-WASP may have partially overlapping functions in macrophage morphology.

Interestingly, it was observed that reduction of either WASP or N-WASP led to a 3 and 1.5fold increase in N-WASP or WASP expression, respectively (data not shown) suggesting the possibility that the cells were attempting to compensate for the loss of the expression of one WASP family member by enhancing the expression of remaining member. In a recent study we have found that WASP is expressed at fifteen times the level of N-WASP in primary macrophages, dendritic cells and RAW/LR5 macrophages and forced over-expression of N-WASP in RAW/LR5 cells with reduced levels of WASP restored the morphology defects observed in WASP deficient cells (Isaac et al. 2010).

N-WASP deficient macrophages produce podosomes, but they are non-functional

Even though cells with reduced N-WASP demonstrated clear morphological defects and GFP-N-WASP was localized to podosomes, cells with reduced N-WASP levels still contained podosomes (Fig. 3A). There was no significant difference in the number of cells containing podosomes but there was a slight decrease in the number of podosomes per cell (Fig. 3B and data not shown). To determine whether the podosomes present in cells with reduced N-WASP level are still functional, their ability to degrade a fibronectin substrate was analyzed. Consistent with previous results (Dovas et al. 2009), cells with reduced WASP levels displayed reduced podosomes and degradation. Unlike WASP, N-WASP deficient cells still contain podosomes, but they are unable to degrade matrix as efficiently as control cells (Fig. 3C). Subsequent rescue experiments found that expression of GFPtagged N-WASP, that was not recognized by the targeting siRNA, was able to restore the ability of shN-WASP cells to degrade matrix (Fig. 3D). Surprisingly however, expression of myc-WASP was unable to restore the degradation defect following N-WASP reduction (Fig. 3D). Myc tagged WASP was fully capable of restoring both podosome formation and function in shWASP cells (data not shown and (Dovas et al. 2009)). This data indicated that the role of N-WASP in matrix degradation was unique and was non-redundant with WASP.

To determine whether actin polymerization induced by N-WASP was required for degradation, shN-WASP cells were transfected with a mutant version of N-WASP containing a deletion in the V domain that results in decreased actin polymerization (Miki and Takenawa 1998). Unlike wild-type N-WASP, this mutant (NW- Δ V) was unable to fully restore matrix degradation but a partial rescue of degradation was observed (Fig. 3D), suggesting that other functions of N-WASP that are independent of actin polymerization may be involved. However, the Δ V mutant used is a deletion of only one of the V domains and still retains a partial ability to polymerize actin *in vitro* and this may be the reason for the partial recue observed (Co et al. 2007). In summary, the role of N-WASP in podosomes may only partially be associated with its role on the induction of actin polymerization and another function of N-WASP in podosomes may exist.

Several studies have implicated N-WASP in the trafficking of vesicles. For example, N-WASP has been shown to localize to endosomal and lysosomal vesicles in *Xenopus* oocyte extracts (Taunton et al. 2000). Additionally, actin-based vesicle movement induced by overexpression of phosphatidylinositol 5-kinase in fibroblasts was dependent on N-WASP (Rozelle et al. 2000). Expression of N-WASP truncation mutants in N-WASP defective cells indicated that the WH1 and polyproline, but not the GBD and basic, domains were important in recruitment to the vesicle surface potentially by recruiting SH3-SH2 adaptor proteins, Nck and Grb2, and WIP (Benesch et al. 2002). The physiological relevance of N-

WASP recruitment for endocytosis was demonstrated by a decrease in EGF internalization in N-WASP deficient fibroblasts (Benesch et al. 2005; Innocenti et al. 2005). N-WASP remains associated with actin filaments barbed ends after nucleation via the V domains and is involved in actin "comet tail" motility by linking the vesicle membrane to actin filaments *in vitro* (Co et al. 2007).

N-WASP is required to localize MMPs to podosomes

It has been shown in the literature that specific MMPs are localized at podosomes. In endothelial cells, MT1-MMP and MMP 9 have degradation activity beneath podosomes (Varon et al. 2006)(reviewed in (Poincloux et al. 2009). Additionally, the polarized exocytosis of MT1-MMP–containing vesicles has been demonstrated in invading breast cancer cells (Bravo-Cordero et al. 2007). Based on the above, it is possible that N-WASP may play a role in MMP trafficking. In addition to the delivery of MMPs to invadopodia and podosomes, it has been suggested that active MT1-MMP is localized to invadopodia by means of the recycling pathway (Itoh and Seiki 2004; Remacle et al. 2005; Wang et al. 2004). N-WASP activity was observed in vesicles and in particular recycling vesicles, using FRET to determine the interaction between Cdc42 and N-WASP in breast carcinoma cells (Parsons et al. 2005).

To explore the role of N-WASP on matrix degradation through MMP targeting to podosomes, colocalization experiments were performed with mCherry MT1-MMP. MT1-MMP demonstrated a punctate distribution with partial overlap with GFP-N-WASP both by mCherry MT1-MMP and by staining for endogenous MT1-MMP (Fig. 4A and data not shown). However, when examined further it was determined that MT1-MMP colocalized with GFP-N-WASP predominantly at F-actin rich puncta that appeared to be podosomes. MT1-MMP localization to podosomes was confirmed using vinculin as an additional marker for podosomes (Fig. 4B). Next, the effect of N-WASP on MT1-MMP localization to podosomes was analyzed. MT1-MMP localization to podosomes was inhibited by silencing of N-WASP (Fig. 5A and 5B). Additionally, MT1-MMP localization was altered in cells that overexpressed the ΔV N-WASP mutant that has been demonstrated to inhibit endogenous N-WASP activity (Miki and Takenawa 1998) (Fig. 5B). This was not simply due to over-expression since overexpression of wild-type GFP-N-WASP did not significantly alter MT1-MMP localization. In either case, there was reduced MT1-MMP at podosomes (Fig. 5B), explaining the reduced matrix proteolysis observed (Fig. 3D). These studies suggest that N-WASP may play a role in the recruitment of MMPs to podosomes.

In support of a role of N-WASP in mediating MMP recruitment, others have demonstrated a role for N-WASP in vesicle trafficking (Benesch et al. 2002; Co et al. 2007; Kovacs et al. 2006; Taunton et al. 2000). Therefore, studies were performed using immunofluorescence microscopy to determine the colocalisation of N-WASP with several vesicle markers, such as Rab 5, 8, 8b and 11 as well as the golgi marker TGN38. None of these markers appeared to colocalize with N-WASP (data not shown). However, it is possible that N-WASP is only transiently associated with vesicles and further studies using live cell imaging would be needed. This in conjunction with the association of N-WASP with MT1-MMP only at sites of podosomes supports the hypothesis that N-WASP is required for vesicle fusion during exocytosis.

Several recent studies have shown a role for the exocyst in MMP secretion in carcinoma cells. Cdc42 is required to recruit IQGAP, another actin binding protein to the exocyst [(Sakurai-Yageta et al. 2008) and the exocyst binds to Arp2/3 and both of these interactions are required for invadopodia function (Liu et al. 2009). Cdc42 activated N-WASP has been shown to stimulate secretion in neuroendocrine cells. N-WASP mutants with a VVCA deletion were not recruited to the cell periphery and exocytosis of growth hormone in PC12

cells was impaired. However, the VVCA or GBD domains alone were not sufficient for exocytosis suggesting that other domains of N-WASP are also required for exocytosis (Gasman et al. 2004). It is interesting to speculate that N-WASP dependent actin polymerization is required for MMP containing vesicles to target and/or fuse at the podosome similar to the role of actin polymerization in neuroendocrine cells (reviewed in (Malacombe et al. 2006)).

Alternatively, it is possible that N-WASP may play a role in podosome maturation. It has been shown by others that MMPs are only targeted to mature invadopodia (Artym et al. 2006) and only the static podosomes of macrophages degrade matrix (Kopp et al. 2006). While the fraction of cells containing podosomes is unaltered in cells with reduced N-WASP levels, it should be noted that these cells do have fewer podosomes per cell (17.8 \pm 1.1 versus 27.8 \pm 0.8 for control cells). The reduced number of podosomes in shN-WASP cells is not due to decreased podosome formation since podosomes reform with the same kinetics as control cells when recovering from cytochalasin D treatment (data not shown). Future studies are needed to determine whether N-WASP regulates podosome lifetime.

Conclusions

Thus, we find that in the RAW264.7 macrophage-like cell-line that WASP is necessary for podosome formation and N-WASP is required for podosome dependent degradation of ECM indicating that these proteins serve non-redundant roles in podosome function. Unlike invadopodia, matrix degrading structures in carcinoma cells, N-WASP is not necessary for podosome formation since podosomes are still present in shN-WASP cells. Collectively these data indicate that WASP is responsible for podosome formation and N-WASP is only required for matrix degradation, while N-WASP serves two functions in invadopodia; 1) actin polymerization required for invadodopodia formation and 2) maturation (Oser et al. 2009).

We speculate that N-WASP regulation of podosome degradation occurs via directed MT1-MMP transport. Consistent with other reports (Linder 2007), we find that MT1-MMP is localized to podosomes and importantly this localization is altered by N-WASP silencing. However, the precise role of N-WASP in the localization of MT1-MMP to podosomes is not clear. N-WASP may be involved in transport of nascent MT1-MMP to podosomes or it acts in the recycling of MT1-MMP vesicles to the cell surface via an intermediate recycling compartment. Further work will be needed to investigate the spatial involvement of N-WASP on MT1-MMP trafficking.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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ABBREVIATIONS LIST

WASP	Wiskott-Aldrich Syndrome Protein
N-WASP	neural-WASP
MMP	matrix metalloproteinase

siRNA

small-interfering RNA

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Figure 1. GFP-N-WASP localizes to podosomes

RAW/LR5 cells transiently transfected with GFP-N-WASP were fixed and stained with Factin (red) and vinculin (blue) (to indicate podosomes). Epifluorescence images are shown in (A) and the inset contains a magnified image of the boxed region (Bar, 20 μ m). (B) The colocalization of N-WASP with podosomes was analyzed using ImageJ and the fraction of podosomes where GFP-N-WASP was co-localized, adjacent to podosomes, or non-localized was determined. Error bars represent SEM. (n=17 cells). Nusblat et al.



Figure 2. Cells with reduced N-WASP are less polar and elongated

RAW/LR5 cells were transfected with control shRNA (Control), shN-WASP (shN-W), shWASP (shW), or both plasmids (shN-W/W) and selected for 2 days in puromycin. (A) Representative western blot is shown of whole cell lysates probed using an antibody that recognizes both WASP and N-WASP. The open arrow represents N-WASP and the closed arrow represents WASP. The fraction of cells with a polarized or migratory phenotype (B) or the elongation index (C) was determined as described in the Materials and Methods. Error bars represent SEM. * p<0.05, $n \ge 3$.

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Figure 3. N-WASP deficient cells produce podosomes, but are unable to degrade a fibronectin matrix

(A) Representative images of RAW/LR5 cells transfected with either control shRNA, shN-WASP, shWASP and stained for F-actin and vinculin (to identify podosomes) are shown. Image on right is a merged imaged of a 10 µm square shown in the inset. (Bar, 10 µm). (B-D) Cells transfected with control shRNA(Control), shN-WASP (shN-W), shWASP (shW), or both plasmids (shN-W/W) cells were plated on coverslips coated with fluorescently labeled fibronectin for 16 hrs and (B) the fraction of cells with podosomes or (C) the degradation area per cell were determined as described in Material and Methods. (D) Degradation area per cell was determined for control and shN-WASP cells or those rescued with GFP alone (GFP), GFP-N-WASP (NW), mutated N-WASP (NW Δ V) or Myc-WASP (Myc-W). Error bars represent SEM. * p<0.05, n ≥ 3.



Figure 4. N-WASP co-localizes with MT1-MMP at podosomes

RAW/LR5 cells transiently transfected with (A) mCherry MT1-MMP and GFP-N-WASP and (B) mCherry-MT1-MMP. Cells were fixed and stained with vinculin and F-actin as indicated and representative epifluorescence images are shown. (Bar, 10 µm)



Figure 5. N-WASP is required for MT1-MMP localization to podosomes

(A) mCherryMT1-MMP expressing shN-WASP cells were fixed and stained with vinculin and F-actin and representative epifluorescence images are shown. (Bar, 10 μ m) (B) The fraction of mCherry MT1-MMP containing podosomes was analyzed using ImageJ for control, shN-WASP cells (shNW), mutated N-WASP (NW Δ V) or GFP-N-WASP (GFP-NW). Error bars represent SEM. * p<0.05, n \geq 3.

shNW

NWAV GFP-NW

mock