

Complement membrane attack complexes activate noncanonical NF- κ B by forming an Akt⁺NIK⁺ signalosome on Rab5⁺ endosomes

Dan Jane-wit^a, Yulia V. Surovtseva^b, Lingfeng Qin^c, Guangxin Li^c, Rebecca Liu^d, Pamela Clark^d, Thomas D. Manes^d, Chen Wang^d, Michael Kashgarian^e, Nancy C. Kirkiles-Smith^d, George Tellides^c, and Jordan S. Pober^{d,e,1}

^aDivision of Cardiovascular Medicine, Yale University School of Medicine, New Haven, CT 06520; ^bYale Center for Molecular Discovery, Yale University, New Haven, CT 06516; ^cDepartment of Surgery, Yale University School of Medicine, New Haven, CT 06520; ^dDepartment of Immunobiology, Yale University School of Medicine, New Haven, CT 06520; and ^eDepartment of Pathology, Yale University School of Medicine, New Haven, CT 06520

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Complement membrane attack complexes (MACs) promote inflammatory functions in endothelial cells (ECs) by stabilizing NF- κ B-inducing kinase (NIK) and activating noncanonical NF- κ B signaling. Here we report a novel endosome-based signaling complex induced by MACs to stabilize NIK. We found that, in contrast to cytokine-mediated activation, NIK stabilization by MACs did not involve cIAP2 or TRAF3. Informed by a genome-wide siRNA screen, instead this response required internalization of MACs in a clathrin-, AP2-, and dynamin-dependent manner into Rab5⁺ endosomes, which recruited activated Akt, stabilized NIK, and led to phosphorylation of I κ B kinase (IKK)- α . Active Rab5 was required for recruitment of activated Akt to MAC⁺ endosomes, but not for MAC internalization or for Akt activation. Consistent with these *in vitro* observations, MAC internalization occurred in human coronary ECs *in vivo* and was similarly required for NIK stabilization and EC activation. We conclude that MACs activate noncanonical NF- κ B by forming a novel Akt⁺NIK⁺ signalosome on Rab5⁺ endosomes.

inflammation | signaling | endothelial cell | complement | membrane attack complex

The complement system contributes to both host defense and immunopathology (1). Using serum with high-titer panel reactive antibody (PRA) from sensitized transplant candidates as a source of complement-fixing human anti-HLA alloantibodies, we identified noncanonical NF- κ B signaling as a novel effector pathway used by complement membrane attack complexes (MACs) to induce proinflammatory changes in endothelial cells (ECs) that resulted in increased recruitment and activation of alloreactive effector memory T cells (2).

Noncanonical NF- κ B signaling is generally initiated by ligand engagement of TNF receptor (TNFR) superfamily members (3, 4). In the absence of ligand, NF- κ B-inducing kinase (NIK) protein is persistently translated and sequestered into a complex containing cIAP1, cIAP2, TRAF2, and TRAF3, where it is polyubiquitinated by cIAP2 and targeted for proteasomal degradation (5). Following ligand activation, TRAF3 along with its associated proteins are recruited to the cytoplasmic tail of the activated receptor (5–7), where the ubiquitin ligase activity of cIAP2 is diverted away from NIK to TRAF3 (8), causing degradation of TRAF3 accompanied by release and stabilization of NIK, which initiates signaling. In this sequence, TRAF3 degradation is necessary and sufficient for NIK accumulation (9, 10). In contrast to canonical NF- κ B activation by TNF- α or IL-1, which occurs rapidly within 15 min, noncanonical signaling typically requires ~8 h for initial detection (11). Here we demonstrate that MAC stabilization of NIK occurs within 30 min independent of the cIAP2-TRAF3-containing signalosome, and that furthermore, an endosome-based Akt- and Rab5-dependent signaling mechanism mediates NIK stabilization by MACs.

Results

Stabilization of NIK by MACs Occurs Rapidly and Independently of cIAP2-TRAF3 Signalosome Formation. We recently reported that cultured human umbilical vein ECs treated with high-titer PRA from sensitized transplant candidates selectively activated noncanonical but not canonical NF- κ B signaling in a mechanism requiring formation of MACs (2). We compared noncanonical NF- κ B activation by PRA with LIGHT, a cytokine activator of this pathway (12, 13). As expected, LIGHT caused a gradual increase in NIK (Fig. 1A), concurrent with cIAP2 induction and TRAF3 degradation (12). In contrast, PRA up-regulated NIK within 30 min without TRAF3 degradation (Fig. 1A). NIK induction occurred posttranscriptionally as NIK mRNA levels were unchanged (Fig. S1A), and NIK protein was decreased by inhibition of protein synthesis (Fig. S1B). Consistent with our previous results (2), NIK-dependent inflammatory genes were unaffected in human umbilical vein ECs (HUVECs) following siRNA-mediated knockdown of p65, but were reduced with NIK or RelB siRNA (Fig. S1C).

We then compared the effects of NIK stabilization by LIGHT and PRA following siRNA-mediated knockdown of TNFR superfamily-associated signalosome components. Consistent with previous reports (9, 10), knockdown of TRAF3 or cIAP2 (BIRC3) was sufficient to stabilize NIK in the absence of ligand (Fig. 1B,

Significance

Complement activation contributes to host defense and immunopathology. We recently discovered that membrane attack complexes (MAC), the terminal effector mechanisms of complement, activate proinflammatory functions in human endothelial cells (ECs) via noncanonical NF- κ B signaling. Here we elucidate the initial steps of how MACs activate this pathway. MACs formed on the surface of human ECs are rapidly internalized via clathrin-mediated endocytosis into Rab5⁺ endosomes, which subsequently recruit activated Akt in a Rab5-dependent manner. Akt recruitment results in NIK protein stabilization on the surface of the endosome within 30 min, initiating noncanonical NF- κ B signaling. MAC internalization in ECs lining human coronary arteries *in vivo* similarly activates noncanonical NF- κ B signaling. Our findings suggest new therapeutic targets for controlling complement-mediated inflammation.

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¹To whom correspondence should be addressed. Email: jordan.pober@yale.edu.

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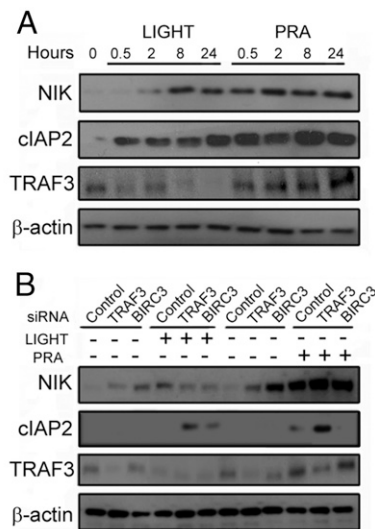


Fig. 1. Stabilization of NIK by MACs occurs rapidly and independently of cIAP2-TRAF3 signalosome formation. (A) ECs were stimulated with LIGHT (100 ng/mL) or PRA (1:5 dilution) for the indicated times. (B) HUVECs were treated with control, TRAF3, or cIAP2 (BIRC3) siRNA and treated with either LIGHT for 24 h or PRA for 4 h. Each experiment was repeated two to five times, with similar results.

top row, lanes 2, 3, 8, and 9). NIK protein expression induced by knockdown of TRAF3 or cIAP2 was not enhanced by the addition of LIGHT (Fig. 1*B*, top row, lane 2 vs. lane 5 and lane 3 vs. lane 6, respectively), suggesting that knockdown of these components and LIGHT affect the same pathway. In contrast, PRA robustly augmented NIK levels in ECs lacking TRAF3 or cIAP2 (Fig. 1*B*, top row, lane 8 vs. lane 11 and lane 9 vs. lane 12, respectively). Of note, at the time point examined, cIAP2, but not TRAF3, protein expression was actually increased by LIGHT as well as PRA treatments, consistent with previous reports indicating that cIAP2 is transcriptionally induced in response to noncanonical NF- κ B activation (11). Taken together, these data suggest that the cIAP2-TRAF2-TRAF3 signalosome is not involved in the stabilization of NIK induced by MAC.

A Genome-Wide siRNA Screen Identifies Endocytosis and Vesicular Trafficking Genes Involved in MAC-Induced NIK Stabilization. To elucidate alternative mechanism(s) of NIK stabilization by MACs, we performed a genome-wide siRNA screen. ECs were stably transduced with an NF- κ B-responsive luciferase reporter responsive to both canonical (TNF- α) and noncanonical (LIGHT and PRA serum) NF- κ B activators (Fig. S2*A*). Normal serum lacking significant anti-HLA IgG, the IgG⁺ fraction of PRA serum alone, or the IgG⁺ fraction of PRA combined with serum deficient in C6 lacking the capability to form MACs were all significantly weaker agonists than PRA serum (Fig. S2*B*). These data recapitulate our previous results identifying MACs as the primary activators of noncanonical NF- κ B (2).

Transduced ECs were screened as diagrammed in Fig. S2*C*. Duplicate screening showed a highly reproducible Pearson's r coefficient of 0.86 (Fig. S2*D*, Left) and a nearly normal distribution of robust z-scores (Fig. S2*D*, Right). Genes with an average robust z-score of ≥ 2 ($n = 831$; Fig. S2*D*, Right) with cell viability $\geq 90\%$ were defined as hits in the primary screen. From these hits, we identified 141 genes for counterscreening with TNF- α to select genes specific for noncanonical NF- κ B (Fig. S2*E*). Counterscreening identified genes concurrently showing $\geq 50\%$ siRNA-mediated inhibition of luciferase activity by PRA but $\leq 50\%$ inhibition of the activity by TNF- α (gray dots in Fig. S2*E*). Many of these genes were functionally annotated to endocytosis and vesicular trafficking ($n = 11$). Individual siRNAs in

the original pool of four were tested, and 10 of 11 genes showed $\geq 50\%$ inhibition by two or more individual siRNAs (Fig. S2*F*).

MACs Are Internalized into Clathrin⁺ Vesicles and Transferred to Rab5⁺ Early Endosomes. Based on siRNA screening results, we assessed the role of endocytosis of MACs in NIK stabilization. An average of 68 ± 27 total MACs were deposited per EC and $42 \pm 16\%$ of MACs were found intracellularly at 30 min after the addition of PRA (Fig. 2*A*). At this time point, MACs extensively colocalized with clathrin⁺ (Fig. 2*B*, Upper) and transferrin⁺ (Fig. 2*B*, Lower) vesicles ($65 \pm 3\%$ and $69 \pm 5\%$, respectively). Immune-electron microscopy (EM) of C9 at 5 min showed MACs within electron-dense surface invaginations anatomically consistent with clathrin-coated pits (arrowheads, Fig. 2*C*, first column) and showed colocalization of C9 within electron-dense coated vesicles at 30 min (Fig. 2*C*, second and third columns). Dual immune-EM on PRA-treated HUVECs confirmed colocalization of MAC and clathrin (Fig. 2*D*).

We then analyzed the endosomal distribution of C9 using Rab5 (early endosomes), Rab7 (late endosomes), Rab11 (recycling endosomes), and LAMP-1 (lysosomes) (Fig. S3*A* and Table S1). At 30 min, when NIK was stabilized, a much greater percentage of Rab5⁺ endosomes colocalized with C9 ($40 \pm 2\%$; Fig. 2*E*) compared with Rab7⁺, Rab11⁺, or LAMP-1⁺ endosomes. We observed concomitant high percent colocalization of two Rab5 effectors associated with early endosomes, APPL1 and EEA1 ($41 \pm 7\%$ and $44 \pm 9\%$, respectively; Fig. S3*B*) with MAC⁺ vesicles. Dual detection by immune-EM at this time point confirmed colocalization of Rab5 (Fig. 2*F*, arrows) and C9 (Fig. 2*F*, arrowheads).

Clathrin-Mediated Endocytosis of MACs Is Required for NIK Stabilization.

To examine the connection between clathrin-mediated endocytosis (CME) of MACs and noncanonical NF- κ B signaling, we treated ECs transfected with control siRNA or clathrin siRNA with PRA and analyzed them for the percent colocalization of C9 with Rab5⁺ vesicles (Fig. S4*A*). An $86 \pm 12\%$ reduction of clathrin (as assessed by Western blot analysis) reduced the number of MAC⁺Rab5⁺ vesicles per EC to 19 ± 3 , compared

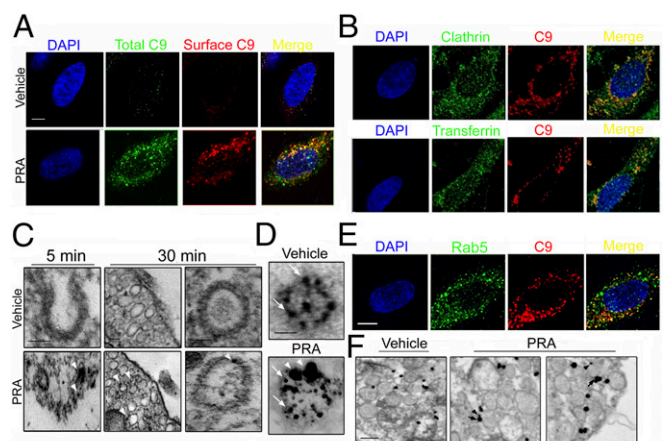


Fig. 2. MACs are internalized into clathrin⁺ vesicles and transferred to Rab5⁺ early endosomes. (A) Total and surface C9 were quantified after treatment with PRA for 30 min. (B) C9, clathrin, and transferrin levels were assessed by IF after a 30-min treatment with PRA. (C and D) ECs treated with PRA for 5 min (C, arrowheads, first column) or 30 min (C, arrowheads, second and third columns) were subjected to immune-EM for C9. Dual immune-EM staining of vesicles was done following 30 min of PRA for C9 (D, arrowheads) and clathrin (D, arrows). (Scale bars: 10 μ m in A and B; 100 nm in C and D). (E) ECs were treated with PRA for 30 min and stained as indicated. (F) Immune-EM of C9 (arrowheads) and Rab5 (arrows) in ECs treated with PRA for 30 min. Each experiment was repeated three to six times with microscopic analysis of ≥ 10 cells per experiment, with similar results. (Scale bar: 100 nm).

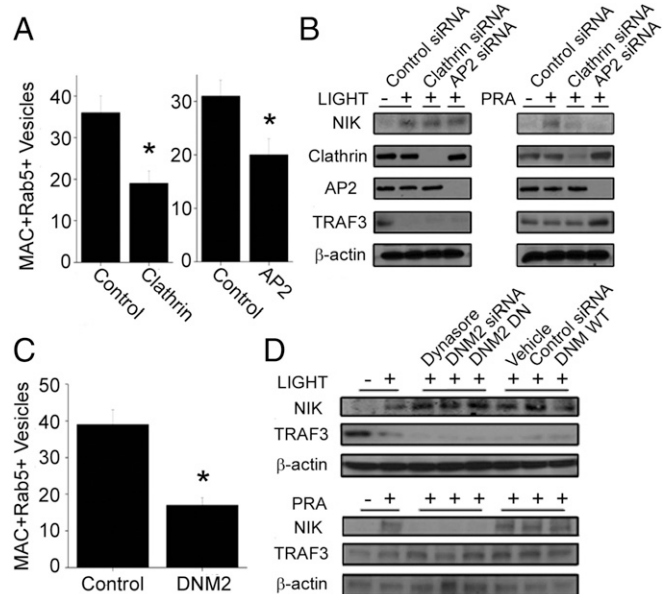


Fig. 3. CME of MACs is required for NIK stabilization. (A) HUVECs treated with control, clathrin, or AP2 siRNA were analyzed for C9 colocalization with Rab5. (B) Western blot analysis of ECs treated with LIGHT (100 ng/mL) or PRA (1:5 dilution) for 24 h or 4 h, respectively, in the presence of control, clathrin, or AP2 siRNA as indicated. (C) Following treatment with control or dynamin siRNA, colocalization between C9 and Rab5 was quantified. (D) ECs were treated with dynasore, DNM2 siRNA, or DNM2 DN and the relevant control and analyzed after 4 h of PRA treatment. Each experiment was repeated two to four times, with similar results. Quantitation of the data in A and C are represented as mean \pm SEM. * $P < 0.05$. (Scale bars: 10 μ m.)

with 36 ± 4 per EC transfected with control siRNA ($P = 0.03$; Fig. S4A and Fig. 3A) and dramatically reduced the levels of NIK protein (Fig. 3B) without affecting TRAF3 levels.

Because clathrin also may participate in vesicle trafficking from internal membranes (14–16), we performed siRNA knockdown of clathrin adaptor protein, AP2 (17, 18), which exclusively regulates

endocytosis of clathrin-coated vesicles from the cell surface (19–21). Western blot analysis revealed a $90 \pm 14\%$ reduction in AP2 protein levels. AP2-deficient ECs, like clathrin-deficient ECs, showed significantly reduced numbers of MAC⁺Rab5⁺ endosomes compared with controls (20 ± 3 vs. 31 ± 3 ; $P = 0.02$; Fig. S4A and Fig. 3A). AP2 knockdown similarly resulted in a marked reduction in NIK without affecting TRAF3 levels (Fig. 3B). In contrast, knockdown of AP2 or clathrin in LIGHT-treated HUVECs showed little effect on PRA-induced NIK stabilization (Fig. 3B).

Similarly, a $93 \pm 8\%$ siRNA-mediated reduction of dynamin, which mediates scission of clathrin-coated vesicles from the plasma membrane (22), also significantly reduced the number of MAC⁺Rab5⁺ vesicles compared with controls (17 ± 2 vs. 39 ± 4 ; $P = 0.002$; Fig. S4B and Fig. 3C) and ablated NIK induction (Fig. 3D). We verified these results in ECs treated with dynasore, a pharmacologic dynamin inhibitor, and in ECs transfected with a dynamin-2–dominant negative (DN) construct. Compared with respective controls, these treatments ablated NIK stabilization without affecting TRAF3 levels (Fig. 3D). In contrast, similar treatments in LIGHT-treated ECs did not significantly alter NIK or prevent TRAF3 degradation (Fig. 3D).

A MAC⁺NIK⁺ Signalosome Forms on Rab5⁺ Endosomes. Given our finding that CME of MACs is required for NIK stabilization, and because MAC heavily colocalized with Rab5⁺ endosomes, we hypothesized that NIK might be stabilized on the surface of these vesicles. We first investigated whether the cIAP1–cIAP2–TRAF2–TRAF3 signalosome is formed on MAC-containing vesicles. cIAP1 and cIAP2 showed a predominantly cytoplasmic distribution with some nuclear colocalization, whereas TRAF2 and TRAF3 staining appeared punctate, suggestive of vesicular colocalization. However, we did not observe any significant colocalization of C9⁺ endosomes with cIAP1, cIAP2, TRAF2, or TRAF3 ($8 \pm 1\%$, $8 \pm 1\%$, $14 \pm 2\%$, and $13 \pm 2\%$, respectively; Fig. S5A). In contrast, we found extensive colocalization of both NIK and phosphorylated IKK- α (pIKK- α), a downstream NIK effector, with C9⁺ vesicles ($64 \pm 9\%$ and $74 \pm 6\%$, respectively; Fig. S5B, Middle and Bottom). Subsequent three-color confocal microscopy showed colocalization of C9, NIK, and Rab5 in $11 \pm 2\%$ of Rab5⁺ vesicles (Fig. 4A). A similar percentage was observed for C9⁺Rab5⁺pIKK- α ⁺ vesicles

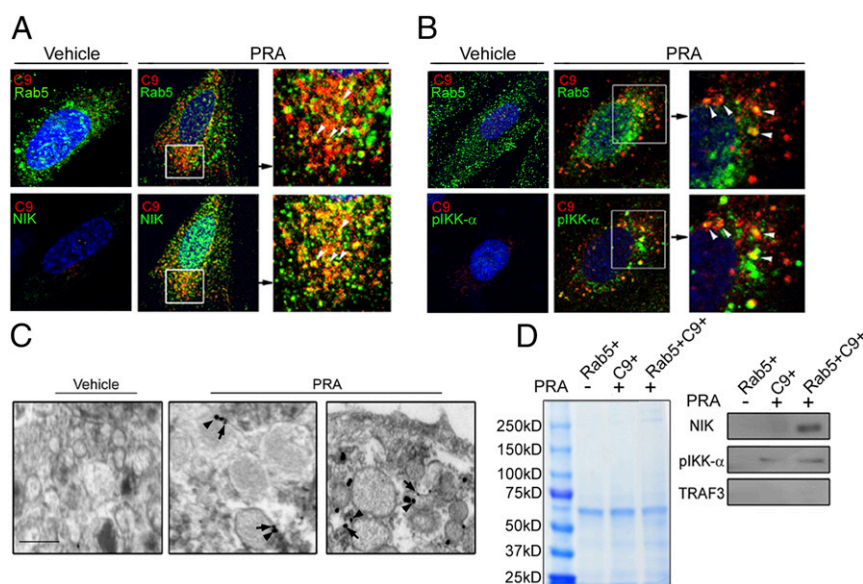


Fig. 4. MAC⁺NIK⁺ signalosome forms on Rab5⁺ endosomes. (A and B) Three-color staining of NIK, C9, and Rab5 (A) and pIKK- α , C9, and Rab5 (B). (C) Immunofluorescence of C9 (arrowheads) and NIK (arrows) in ECs treated with PRA for 30 min. (Scale bar: 100 nm.) (D) ECs were transfected with Rab5-GFP and treated with MACs containing labeled C9-Alexa Fluor 647. Subcellular fractions containing endosomes were analyzed by Coomassie blue staining (Left), followed by Western blot analysis (Right). Each experiment was repeated two to three times, with similar results.

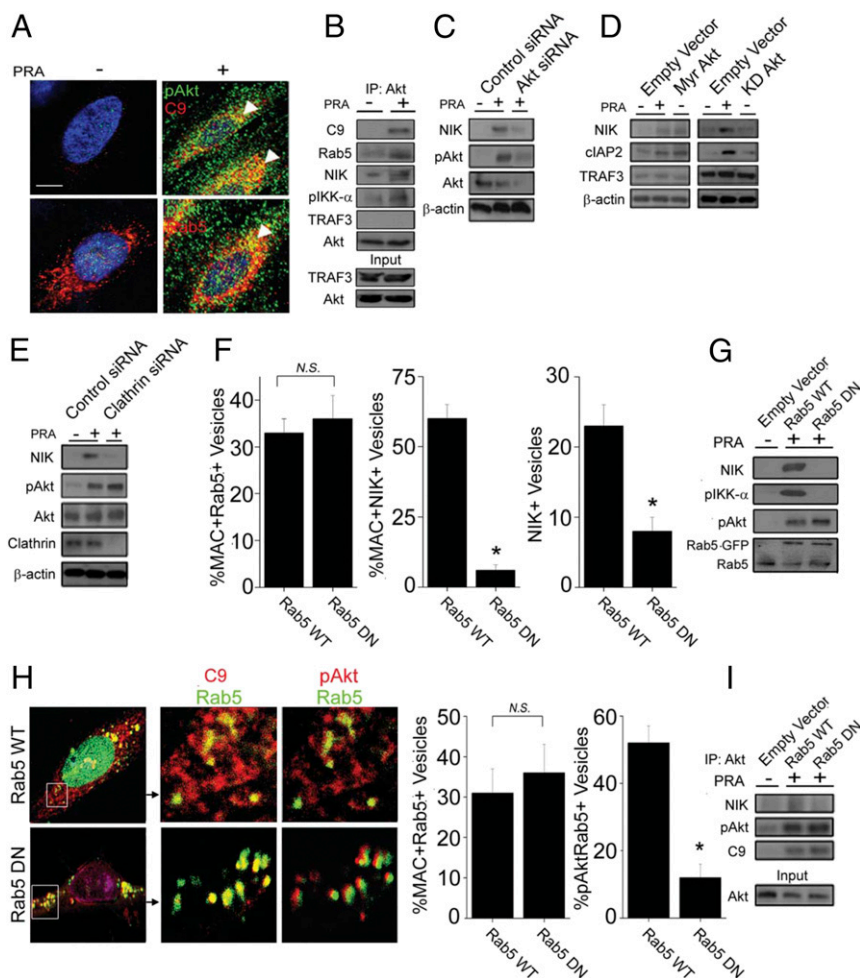


Fig. 5. Rab5-dependent recruitment of activated Akt to MAC⁺Rab5⁺ endosomes is required for NIK stabilization by MACs. (A) ECs treated with PRA for 30 min were evaluated by IF for phosphorylated AktSer473 (pAkt), C9, and Rab5. (B) ECs were treated with vehicle or PRA for 45 min, Akt was immunoprecipitated, and lysates were probed as indicated. (C) ECs transfected with control or Akt siRNA were treated with PRA for 4 h and probed as indicated. (D) ECs transfected with empty vector, myr-Akt, or KD-Akt were probed by Western blot analysis as indicated. (E) ECs transfected with either control or clathrin siRNA were treated with PRA for 4 h and probed as indicated. (F) ECs transfected with either Rab5 WT or Rab5 DN constructs were analyzed by confocal microscopy for C9⁺Rab5⁺ and C9⁺NIK⁺ vesicles following treatment with PRA for 30 min. (G) ECs transfected with empty vector, Rab5 WT, or Rab5 DN were probed by Western blot analysis as indicated. (H) Rab5 colocalization with C9 and pAkt in Rab5 WT and Rab5 DN mutants was assessed in ECs treated with PRA for 30 min (Left), and results were quantified (Right). (I) ECs transfected with empty vector, Rab5 WT, or Rab5 DN constructs were treated with vehicle or PRA for 45 min. Akt was immunoprecipitated, and lysates were probed as indicated. Each experiment was repeated two to five times, with similar results. Data are presented as mean \pm SEM. * $P < 0.05$. (Scale bar: 10 μ m.)

($13 \pm 4\%$; Fig. 4B). These findings were corroborated by dual immune-EM, which revealed vesicles that costained for C9 and NIK ($3 \pm 5\%$; Fig. 4C).

We further confirmed these morphological observations through biochemical analyses of isolated vesicles (23). ECs were transfected with Rab5-GFP and then treated with vehicle or PRA serum containing Alexa Fluor 647-labeled C9. The intact cell and subcellular fractions were identified (Fig. S64), and the GFP⁺ and Alexa Fluor 647⁺ subcellular fractions were isolated by FACS, analyzed by Coomassie blue staining (Fig. 4D, Left), and subjected to Western blot analysis. C9⁺ fractions (lanes 2 and 3) from PRA-treated ECs, but not from vehicle-treated ECs (lane 1), contained both NIK and pIKK- α , but not TRAF3 (Fig. 4D, Right). Equivalent levels of IKK- α and TRAF3 were found in control and PRA-treated fractions of intact cells (Fig. S6B). These data suggest that a MAC⁺Rab5⁺ compartment that excludes conventional noncanonical NF- κ B signaling components functions as a platform for NIK stabilization and recruitment of downstream noncanonical NF- κ B effectors like pIKK- α .

Rab5-Dependent Recruitment of Activated Akt to MAC⁺Rab5⁺ Endosomes Is Required for NIK Stabilization by MACs. Akt is implicated in IKK- α phosphorylation (24, 25). Furthermore, Rab5 endosomes, the sites at which MAC stabilizes NIK (Fig. 4A), are enriched in phosphatidylinositol 3,4,5-trisphosphate, to which Akt binds. Immunofluorescence (IF) confocal microscopy revealed colocalization of phosphorylated Akt (pAkt) with both C9 and Rab5 ($58 \pm 6\%$ and $31 \pm 7\%$, respectively; Fig. 5A). Moreover, Akt coimmunoprecipitated with C9, Rab5, NIK, and pIKK- α , but not with TRAF3, in PRA-treated ECs (Fig. 5B). Subsequent knockdown of Akt unexpectedly blocked NIK (Fig. 5C). Transfection of ECs with a constitutively active form of Akt, myristolated Akt (myr-Akt), but not a kinase-dead form of Akt (KD-Akt), induced NIK stabilization and cIAP2 induction without TRAF3 degradation (Fig. 5D), indicating that Akt induces NIK through its kinase activity. Blocking CME in ECs with clathrin-specific siRNA had no effect on Akt phosphorylation (Fig. 5E), indicating that MAC internalization of MAC and Akt activation are not coupled during MAC-induced NIK stabilization.

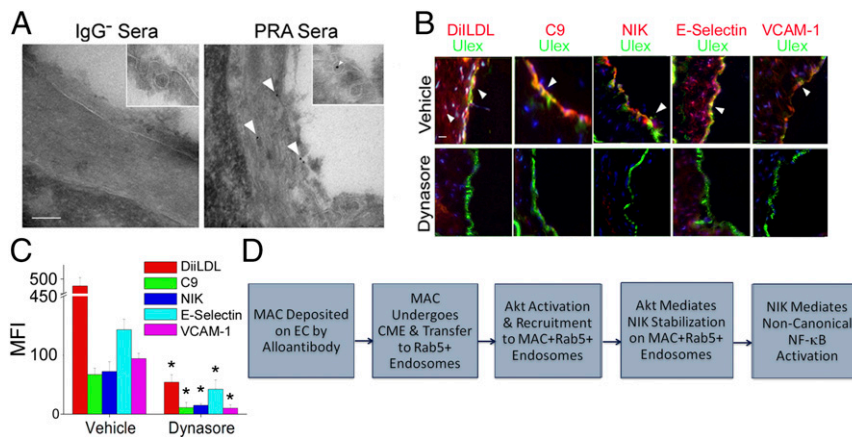


Fig. 6. MAC internalization occurs in human coronary arteries and is required for NIK stabilization in vivo. (A) SCID/beige mice bearing human coronary arteries were treated with PRA serum depleted of IgG (IgG⁻ sera) or PRA serum. Grafts were harvested at 18 h after injection and analyzed by immune-EM ($n = 2$ pairs of mice). (Scale bar: 10 nm.) (B and C) Grafts were harvested at 18 h after PRA injection (B), and MFI of the indicated stains was analyzed by IF (C) ($n = 5$ pairs of mice). (D) Schematic of events triggered by MACs to activate noncanonical NF- κ B. Quantitation of the data is represented as mean \pm SEM. * $P < 0.05$. (Scale bars: 200 μ m.)

To determine whether Rab5 GTPase activity is required for forming the MAC⁺Rab5⁺Akt⁺NIK⁺ signalosome, we transfected ECs with either Rab5-GFP wild-type (WT) or DN and assessed for MAC and/or NIK colocalization with C9⁺ vesicles (Fig. S4C and Fig. 5F). Compared with Rab5 WT EC, Rab5 DN ECs showed similar levels of C9⁺Rab5⁺ vesicles but dramatically decreased percentages of C9⁺ vesicles colocalizing with NIK ($6 \pm 2\%$ vs. $60 \pm 5\%$; $P < 0.001$; Fig. S4C and Fig. 5F). In fact, the overall numbers of NIK⁺ vesicles in Rab5 DN vs. Rab5 WT was reduced (4 ± 2 vs. 23 ± 3 ; $P = 0.005$). Furthermore, on Western blot analysis, ECs transfected with Rab5 DN showed reduced levels of both NIK and pIKK- α , but not of pAkt (Fig. 5G), compared with ECs transfected with Rab5 WT. These data suggest that active Rab5 is not essential for MAC delivery to Rab5⁺ endosomes but is required to stabilize NIK once MACs reach Rab5⁺ compartments.

We next examined by confocal microscopy whether Rab5 activity was required for Akt recruitment to C9⁺ vesicles. The number of pAkt⁺ vesicles was not significantly different between Rab5 DN and Rab WT (38 ± 7 vs. 50 ± 9 ; $P = 0.09$), consistent with our observation that Rab5 DN did not affect pAkt protein levels (Fig. 5G). However, at high magnification, we observed that pAkt⁺ vesicles in Rab5 DN EC samples were closely apposed but were not colocalized with Rab5⁺ vesicles (Fig. 5H, Bottom), thereby significantly reducing the percentage of pAkt colocalizing with Rab5 DN endosomes compared with Rab5 WT controls ($12 \pm 4\%$ vs. $52 \pm 5\%$; $P = 0.004$) (Fig. 5H, Right).

To verify these results, we pulled down Akt in PRA-treated HUVECs carrying Rab5 WT or Rab5 DN constructs. Compared with Rab5 WT ECs, NIK did not coimmunoprecipitate with activated Akt and C9 in Rab5 DN ECs (Fig. 5I). Collectively, these data show that Rab5-independent activation and Rab5-dependent recruitment of Akt to MAC⁺Rab5⁺ vesicles are separate but necessary processes for assembly of the endosome-associated signalosome mediating NIK.

MAC Internalization Occurs in Human Coronary Arteries and Is Required for NIK Stabilization in Vivo. Finally, to determine whether MAC internalization is required for NIK stabilization in ECs in vivo, we administered i.v. injections of PRA to immunodeficient SCID/beige mice in which a human coronary artery xenograft had been interpositioned into the infrarenal aorta, a treatment that deposits murine MACs on human ECs, induces NIK, and activates inflammatory genes (2). Immune-EM revealed

internalized MACs within PRA-treated graft ECs (5 ± 4), but not in grafts treated with IgG-depleted PRA serum (1 ± 0 ; Fig. 6A).

To test whether MAC internalization is required for NIK, we pretreated grafts with vehicle or dynasore before i.v. injection of PRA serum. We validated the ability of dynasore to block PRA-induced inflammatory changes in ECs in vitro, along with pitstop 2, a cell-permeable clathrin inhibitor (26). Pretreatment of ECs with dynasore or pitstop 2 significantly reduced NF- κ B-dependent luciferase activity (Fig. S7A) and attenuated NIK-dependent inflammatory gene induction (Fig. S7B). Dynasore was subsequently chosen based on its efficacy in vitro and its tolerability in vivo (27). We then tested the ability of dynasore to inhibit CME in vivo. Mice bearing human artery grafts were treated with dynasore or vehicle and then injected with PRA and with DiI-LDL, after which grafts were harvested for analysis. In grafts from animals pretreated with vehicle, we observed uptake of DiI-LDL in the intima and media (Fig. S7C, Upper Row). In contrast, grafts from dynasore-treated animals showed significantly reduced mean fluorescence intensity (MFI) of DiI-LDL without concurrent intimal loss, as measured by MFI of human CD31 staining (Fig. S7C, Lower Row). These data indicate functionally successful, noncytotoxic drug dosing.

Finally, we tested whether inhibiting endocytosis could block MAC-induced NIK and EC activation. In hosts treated with PRA, MACs were detected in the intima of arterial grafts along with NIK and markers of EC activation, including E-selectin and VCAM-1 adhesion molecules (Fig. 6B, Upper). Compared with controls, dynasore-treated animals had significantly attenuated levels of NIK and adhesion molecules (Fig. 6B, Lower), as indicated by MFI values (Fig. 6C). Interestingly, MAC surface levels were significantly reduced in dynasore-treated hosts compared with controls. We conclude that internalization of MACs by ECs is required for NIK stabilization and EC activation in vivo.

Discussion

MAC-induced noncanonical NF- κ B is known to potentiate EC activation and exacerbate allograft rejection in human coronary arteries (2). Here we investigated the unknown signaling pathway connecting MAC formation to NIK stabilization. Alloantibody-induced MACs on EC surfaces were rapidly internalized via clathrin-mediated endocytosis and transferred to Rab5⁺ endosomes. Akt was then activated in a Rab5-independent fashion and recruited to MAC⁺Rab5⁺ endosomes in a Rab5-dependent manner. Activated Akt isoforms localized on endosomes then mediated NIK stabilization, IKK- α phosphorylation, and proinflammatory

gene signaling (Fig. 6D). This novel mechanism contrasts with the widely held view that MACs initiate signaling from the cell surface via electrolyte (e.g., calcium) flux (28).

Our studies reveal a requirement for activated Akt in stabilizing NIK, a step upstream of IKK- α activation. Rab5-dependent Akt localization on endosomes suggests that Rab5 allows active Akt to phosphorylate substrates on MAC⁺Rab5⁺ endosomes, which may include NIK and other undefined substrates. Thus, Akt may represent a critical point for regulating MAC-induced non-canonical NF- κ B.

Materials and Methods

Cell Culture, Reagents, and Culture Treatments. All protocols were approved by the Yale University Institutional Review Board. HUVEC culture, treatment with PRA, Western blot analysis, real-time quantitative RT-PCR (qRT-PCR), and IF staining were performed as described previously (2). The protocol is described in detail in *SI Materials and Methods*.

Animal Studies. All protocols were approved by Yale University's Institutional Animal Care and Use Committee. Human coronary arteries were interposed into the descending aortae of adult female C.B-17 SCID/beige mice (Taconic) as described previously (2). Details of the IF staining procedure are provided in *SI Materials and Methods*.

Statistical Methods. Statistical analyses were performed using Origin software (OriginLab). Absolute numbers and percentages of vesicles were analyzed using the Student *t* test and χ^2 test, respectively. A *P* value < 0.05 was considered statistically significant. SEs are reported throughout the text. Endosome numbers were quantified using Volocity software (PerkinElmer). Colocalization was quantified using ImageJ image analysis software with the Just Another Colocalization Plugin (JACoP; National Institutes of Health).

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