

RESEARCH ARTICLE

Involvement of SARA in Axon and Dendrite Growth

Cristina Isabel Arias^{1‡}, Sebastián Omar Siri^{1,2,3}, Cecilia Conde^{1,2,3*}

1 Laboratorio Neurobiología, INIMEC-CONICET, Córdoba, Argentina, **2** Universidad Nacional de Córdoba, Córdoba, Argentina, **3** Instituto Universitario de Ciencias Biomédicas de Córdoba, Córdoba, Argentina

‡ Current Address: Departamento de Biotecnología, Laboratorio de Hemoderivados, Universidad Nacional de Córdoba, Córdoba, Argentina

* cconde@immf.uncor.edu



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Abstract

SARA (Smad Anchor for Receptor Activation) plays a crucial role in Rab5-mediated endocytosis in cell lines localizing to early endosomes where it regulates morphology and function. Here, we analyzed the role of SARA during neuronal development and tested whether it functions as a regulator of endocytic trafficking of selected axonal and membrane proteins. Suppression of SARA perturbs the appearance of juxtannuclear endocytic recycling compartments and the neurons show long axons with large growth cones. Furthermore, surface distribution of the cell adhesion molecule L1 in axons and the fusion of vesicles containing transferring receptor (TfR) in dendrites were increased in neurons where SARA was silenced. Conversely, SARA overexpression generated large early endosomes and reduced neurite outgrowth. Taken together, our findings suggest a significant contribution of SARA to key aspects of neuronal development, including neurite formation.

Introduction

In mammal cells, endocytic membrane traffic plays an essential role in delivering membrane components, receptor-associated ligands and solute molecules to intracellular destinations. This requires significant coordination between spatially segregated sorting compartments that function to determine the cellular fate of cargos. After internalization, a cargo is transported to early endosomes (EE) where sorting decisions are made [1]: proteins targeted for degradation shift to late endosomes and lysosomes, whereas proteins recycled to the cell surface through recycling endosomes (RE) are subject to slow recycling, or fast recycling if travelling directly from early endosomes for later reinsertion into the plasma membrane [2–3]. The fate of the endocytic cargo is determined by the activity and molecular nature of the endosomal sorting machinery.

The endosomal pathway is known to play a decisive role in many neurodevelopment processes, including migration, polarization and synaptic function [4–7]. Neurons are among the best examples of polarized cells, having two functionally different structural domains: a single long axon, and multiple short highly branched dendrites. In neurons, the regulation of endosomal trafficking is particularly complex, since the generation of asymmetric domains

requires specialized membrane trafficking not only to promote neurite outgrowth but also to ensure differential distribution of components to the axonal or somatodendritic domains [8–11]. Dysfunction of proteins involved in endocytic trafficking has been linked to the development of neurodegenerative diseases, implicating at the membrane trafficking control machinery as a critical factor in neuron function [12–16].

SARA is a FYVE protein (Fab1, YOTB, Vac1 and EEA1, [17]) that binds to PI3P (phosphatidylinositol 3-phosphate), is highly enriched in endocytic membranes and is involved in membrane trafficking. [18]. SARA also contains a Smad-binding domain (SBD) required for the interaction with the transcription factors Smad2 and Smad3 [19] and a C-terminal region that interacts with the type I TGF β receptor (TGF β -RI) [17]. It has been suggested that SARA has a crucial function in the recruitment of Smad to the TGF β receptor, ensuring appropriate subcellular localization of the activated receptor-bound complex. The FYVE domain directs the ligand TGF β to EE, where it interacts with both TGF β receptors and Smads [17]. Recent data suggests that SARA is dispensable for functional TGF β -mediated signaling, because in various B-cell lymphomas no correlation was found between SARA expression and the levels of TGF β -induced phosphorylation of Smads. Moreover, knock-down of SARA in HeLa cells does not interfere with TGF β -induced Smad activation, Smad nuclear translocation, or induction of TGF β target genes [20]. These data suggest that SARA may regulate other events. For example, it has been shown that SARA overexpression causes enlargement of EE, and significantly delays transferrin recycling. These alterations resemble the defects caused by overexpression of the Rab5 mutant (Rab5Q79L) and suggest that SARA plays an important functional role downstream of Rab5-regulated endosomal trafficking [21, 22]. Interestingly, SARA also interacts with ubiquitin ligase RNF11, participates structurally and functionally in the ESCRT (endosomal sorting complexes required for transport) and regulates degradative EGFR trafficking [23].

Recently, Chang et al. have suggested a protective role of SARA in skin carcinogenesis, showing that SARA is not involved either in the activation process of TGF- β signal transduction or mouse development [24]. Moreover, SARA has been proposed as a novel vesicle-tethering molecule capable of interacting with membrane proteins such as rhodopsin and syntaxin 3 in axonemal vesicles [25], suggesting that SARA may play a role in neuronal morphogenesis. In the present study, we analyzed the consequences of the expression or knockdown regulation of SARA on neuronal development. We provide novel evidence suggesting a key role for SARA in several neuronal morphogenetic events, including neurite formation, development and polarized growth. We propose that some of these functions may be linked to the role of SARA in regulating the trafficking of membrane components between EE and RE, both in axons and in dendrites.

Materials and Methods

DNA constructs

Expression vectors encoding SARash-HcRed, SARash-GFP and scrambled control-sh plasmid were described previously [25]. The SARA-GFP or SARA-Flag plasmids were described in Hu et al. [21]. Rab8-GFP, Rab11a-GFP and Rab5-GFP were generously gifted by Dr J.L. Daniotti (National University of Córdoba, CIQUIBIC); these were generated by Dr J.L. Daniotti, Dr M. Colombo (National University of Cuyo) and Dr. J. Bonifacino (NICHD, National Institutes of Health, Bethesda, MD, USA), respectively. Rab4-GFP was a kind gift from Dr. G. Yudowski (Departments of Psychiatry, University of California at San Francisco) and TfR-GFP from Dr. Thierry Galli (INSERM, Institut Jacob Monod, France).

Antibodies

The following primary antibodies were used in this study: mAb against Tau protein, an axonal marker (clone Tau-1) and mAb against MAP2, a dendritic marker (clone AP-20) (both were a generous gift of Dr. L. I. Binder, Northwestern University, Chicago, USA) diluted 1:1000 and 1:400, respectively; mAb against tyrosinated α -tubulin (Tyr-Tubulin; Sigma Ch. Co. USA) diluted 1:8000 for IF or 1:10000 for WB; polyclonal antibody (pAb) against SARA (H-300) (sc-9135, Santa Cruz Biotechnology, Inc. CA, USA) diluted 1:20 for IF or 1:200 for WB; an affinity purified rabbit polyclonal antibody raised against L1 (provided by Dr. Hiroyuki Kamiguchi); mAb against GFP (ROCHE) diluted 1:400 for IF and 1:1000 for WB; rabbit polyclonal antibody raised against Flag (Sigma Aldrich) diluted 1:500 for IF or 1:2000 for WB and a rabbit polyclonal antibody against LIMK1 (Santa Cruz Biotechnology) diluted 1:1000 for WB. For some experiments rhodamine-phalloidin (1:1000; Molecular Probes, Eugene, OR, USA) was used to stain filamentous actin (F-actin).

Culture, transfection, immunofluorescence and imaging

Pregnant rats (Wistar, Charles River 251 Ballardvale Street, Wilmington, MA01887) were obtained from the institutional (INIMEC-CONICET) specific pathogen free (SPF) vivarium. All animal procedures and care were approved by the institutional animal care committee (INIMEC-National Research Council and Universidad Nacional de Córdoba, Argentina) and the National Department of Animal Care and Health (SENASA-Argentina). Efforts were made to minimize animal suffering and to reduce the number of animals used.

Embryonic day-18 rat embryos (euthanized by CO₂ gas in chamber with gas inflow precisely regulated) were used to prepare primary hippocampal cultures as described [26–28]. SARA-KO mice were generated as described by Chang et al, 2014 [24] and were generously provided by Dr C.H. Sung (Cornell University, USA). Transient transfection of cultured neurons was performed with Lipofectamine 2000 [26] and the constructs used at concentrations ranging from 2–4 μ g/ml. Neurons were fixed 16 h later (for overexpression experiments) or 20–24 h later (for silencing experiments) with 4% PFA in 4% sucrose-containing PBS, and permeabilized in 0.2% Triton X-100 in PBS for 5 min prior to antibody incubation. CHO-K1 cells were cultured in high glucose DMEM containing L-glutamine, supplemented with 10% fetal bovine serum, penicillin and streptomycin (Invitrogen, Carlsbad, CA).

Cells were visualized using a conventional (Zeiss Pascal) inverted confocal microscope. Images were processed using Adobe Photoshop. Neuronal shape parameters were evaluated as described by Bisbal et al. [29]. Briefly, montages showing the complete neuronal arbor of transfected neurons were created from confocal images (maximal projections) acquired through a 60 X 1.4 NA oil objective lens and the number and length of neuritic processes measured using ImageJ software. Manders coefficients, m1 and m2, are proportional to the amount of fluorescence of the colocalizing objects in each component of the image, relative to the total fluorescence in that component. From the distribution of green (EEA1) / red (SARA) or red (SARA) / green (EEA1) pixels, m1 and m2, respectively, were calculated. Statistical analysis was performed with Statistica software (StatSoft, v8.0). Data are presented as the mean \pm s.e.m. from at least three representative independent experiments for each DNA construct. Differences among experimental groups were analyzed by one-way ANOVA and post-hoc Tukey test. Statistical significance was defined as $p < 0.05$, 0.01 or 0.001.

Western blot assays

Levels of endogenous SARA, ectopically expressed SARA and tubulin after shRNA treatment in Chinese hamster ovary (CHO-K1) cells or in cultured hippocampal pyramidal neurons were

analyzed by Western blotting as described [27, 30]. Immunoblotting assays using extracts of transfected CHO-K1 cells or brain lysates were carried out essentially as described by Bisbal et al [30].

Analysis of cell-surface proteins

To label cell-surface proteins, CHO-K1 cells were washed three times with ice-cold PBS, pH 8, and then incubated with the membrane impermeable sulfo-NHS-biotin (Pierce, Rockford, IL, USA) at a final concentration of 0.3 mg/ml in PBS, pH 8, at 4°C for 20 min. After biotinylation, the cultures were washed with ice-cold PBS and harvested with RIPA buffer. Biotinylated proteins were collected with avidin-agarose beads (Pierce) and analyzed by Western blot with indicated antibodies.

Time-lapse video microscopy, total internal reflection fluorescence microscopy (TIRFM) and Fluorescence recovery after photobleaching (FRAP)

For time-lapse fluorescence microscopy and total internal reflection fluorescence microscopy (TIRFM), transfected cells were cultured in special Petri dishes [31]. After transfection, the dishes containing the attached cells were placed on a Harvard microincubator located on top of the stage of a fully motorized Nikon TE-2000 E inverted microscope equipped for differential interference contrast, epifluorescence, and TIRFM. For conventional time-lapse fluorescence microscopy, cells were imaged using a 60X or a 100X oil-immersion objective, an ORCA II ER camera (Hamamatsu) and MetaMorph software (Molecular Devices) [32]. We used TIRFM to image single vesicular insertion events under low intensity and superficial light exposure conditions that minimize phototoxicity [33]. Cells were visualized with a 60X 1.45 NA objective, equipped for through-the-objective TIRF illumination using a 488 nm argon laser. Neurons were imaged in Neurobasal medium supplemented with 30 mM HEPES buffer, pH 7.2, and maintained at 37°C. Time-lapse sequences were acquired at a continuous rate of 2 or 5 frames/s during 2–4 min. Pyramidal neurons were selected by morphological criteria (in wide-field images) before imaging in TIRFM mode. Live images shown represent raw data with simple background subtraction of the averaged blank field intensity. Fusion events were detected semiautomatically using maximum pixel intensity and centroid analysis to define sites of insertion [33]. FRAP experiments were performed in CHO-K1 cells previously transfected with Rab11-GFP and HcRed or sh-SARA and analyzed 20 h later. These experiments were performed in an Olympus FluoView FV1000 confocal microscope (Olympus, Japan) using a 20X UPlanApo oil immersion/0.8 NA objective. In these experiments, the pinhole was fully opened in order to minimize changes in fluorescence detection due to Z-drift occurring during time lapse imaging. Before performing FRAP experiments, we standardized bleaching protocols to avoid significant reversible photobleaching using standard protocols [34]. FRAP methodology and quantitative image analyses were those previously described [35–36].

Results

SARA distributes in somatodendritic and axonal compartments in hippocampal neurons

Most of the experiments designed to study the function of SARA have been performed in cultured cells (HEK and MDCK cell lines) [21] or human mesangial cells [37]. However, little is known about the subcellular distribution of SARA in neurons. We sought to address this issue

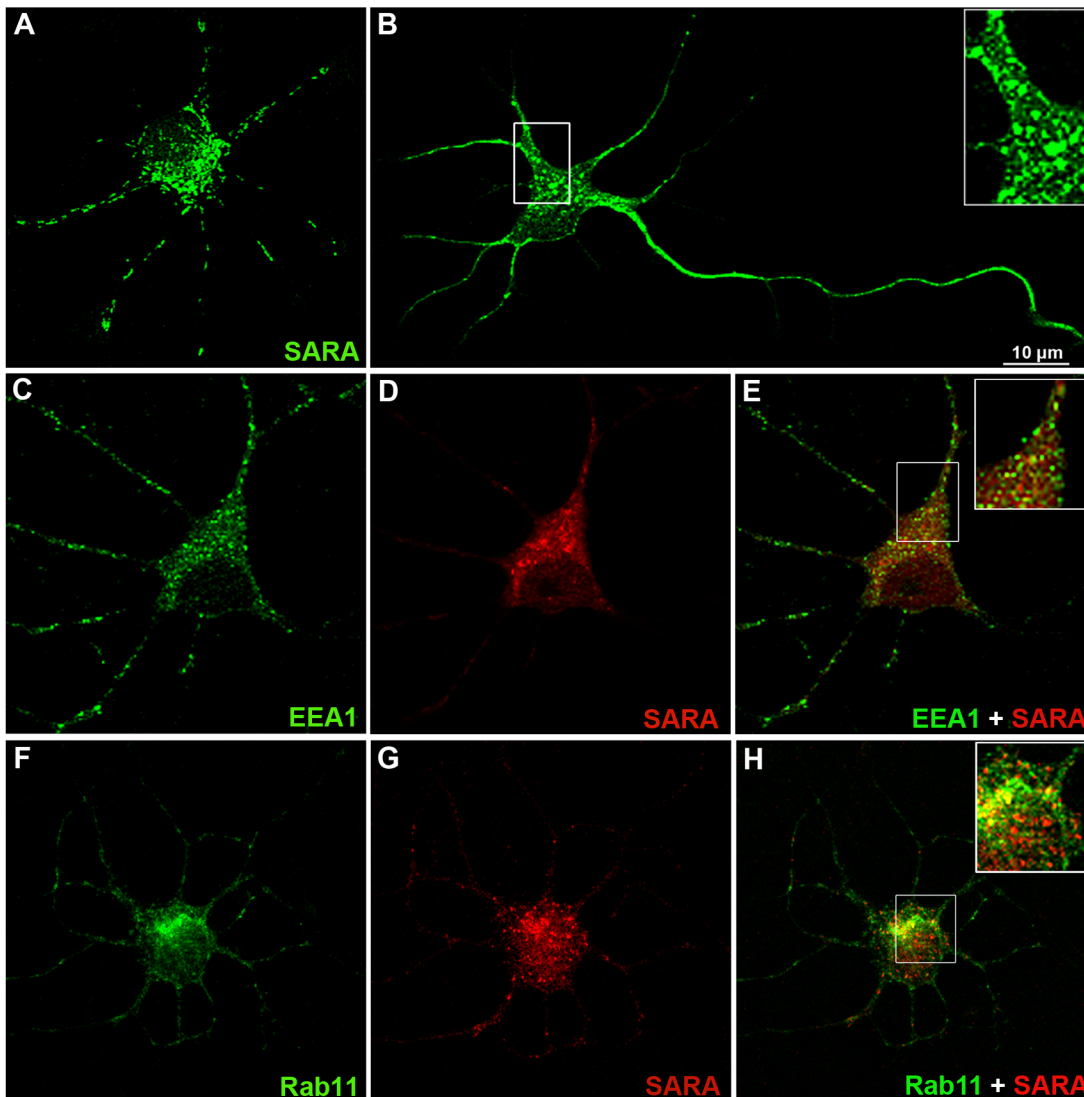


Fig 1. SARA subcellular distribution in young cultured hippocampal pyramidal neurons. (A) Confocal image showing an example of stage 2 neurons (1DIV) or (B) stage 3 neurons (2-3DIV) immunostained for SARA (green). Note that SARA immunolabeling is present in the soma, in all types of neurites (minor process and axon) and that it extends into neuritic tips. SARA is associated with early and recycling endosomal compartments. (C-H) Neurons double-labeled, using antibodies against EEA1 (green, C) and SARA (red, D) or Rab11 (green, F) and SARA (red, G). The SARA-labeled punctae displayed partial colocalization with the early endosomal marker (E, inset) or recycling endosomal marker (H, inset). Different heterogeneous populations of endosomes were observed: one containing only EEA1 or Rab11, a second with EEA1 or Rab11 and SARA, and a third containing only SARA endosomes. All cells were imaged by confocal microscopy. Data is typical of over 10–20 neurons imaged for each condition.

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using neuronal cultures prepared from rat embryonic hippocampi, a widely used model for studying neuronal morphogenesis [38].

First, we examined the cellular localization of SARA in young cultured hippocampal neurons (during the first 48 hours in vitro), when they either display a symmetric array of short neurites (minor undifferentiated processes, stage 2, Fig 1A) or have already developed an axon-like neurite (stage 3, Fig 1B). Staining with a SARA antibody revealed punctated (tubulo-vesicular), endosome-like labeling throughout the neuron. Confocal fluorescence microscopy revealed that, in neurons in stages 2 and 3, SARA is evenly distributed throughout the cell body, minor neurites, the axon and growth cones (Fig 1A and 1B). The SARA-labelled punctae

displayed a partial but significant colocalization with the early endosomal markers EEA1 (Fig 1C–1E) or Rab5 (not shown) and with the recycling endosome marker, Rab11 (Fig 1F–1H). Quantitative estimation of colocalization was further performed using the JACoP plugin of ImageJ. The values for Pearson correlation (r) revealed a positive correlation between SARA and EEA1 or Rab11 (0.70 and 0.67, respectively). The Manders coefficients, $m1$ and $m2$ [39], had values of 0.80 and 0.65, respectively, indicating that at least 65% of SARA-positive endosomes colocalize with EEA1-positive endosomes. However, $m1$ and $m2$ values for Rab11 were 0.77 and 0.8, indicating that nearly 80% of SARA endosomes colocalize with Rab11-positive endosomes.

To avoid the use of the high concentrations of antibodies required to detect endogenous Rabs, as a second approach, neurons were transiently transfected with GFP tagged Rabs. These GFP fused for Rab4, Rab5, and Rab11 correctly localized to their target membranes [40]. Confocal images of neurons expressing Rab5-GFP and stained for endogenous SARA revealed that both proteins partially colocalized (S1 Fig). The quantitative analysis showed results for SARA and EEA1 similar to those we previously found using antibodies to detect endogenous Rabs ($r = 0.76$; $m1$ and $m2 = 0.77$).

Rab4 is a protein associated with sorting endosomes, a tubular mature compartment of EE [41–42]. Our results show that Rab4-GFP partially colocalizes with endogenous SARA in vesicle-like structures (S1 Fig). The colocalization indexes obtained were $r = 0.8$, $m1 = 0.86$ and $m2 = 0.87$ for Rab4-GFP and endogenous SARA.

Our results above described revealed that, whereas some neuronal domains have endosomes with partial or good colocalization indexes between EEA1 or Rab5 and SARA, other domains show endosomes differentially enriched with either EE markers or SARA. This suggests that the remaining endosomes non-colocalizing would be either in the same endosome but in a different domain [23, 40] or in different endosomes, and they would probably have different functions [43].

In summary, these experiments show that SARA is distributed in neurons in all the types of neurites where it partially localizes to different functional domains of EE and RE. These observations also raise the possibility of SARA having a role in neuronal development.

SARA suppression changes recycling endosome distribution

To study the effect of SARA overexpression, neurons in culture were transfected with a plasmid driving the expression of a SARA-Flag fusion. Consistently with previous reports [23, 44–46], we found that ectopically expressed SARA distributes in a larger vesicular compartment also positive for Rab5 (Fig 2A–2C). Similar results were obtained with Rab4-GFP (Fig 2F–2H), suggesting that SARA overexpression in neurons results in early endosome expansion, probably due to an excess membrane and homotypic fusion of endosomes.

Endocytic membrane traffic in many cell types has an essential role in directing membrane components to their correct location [6, 47]. Growth, remodeling and maintenance of the axonal and dendritic compartments in neurons rely heavily on membrane trafficking. In this scenario, Rab11 is a key protein involved in the transport of material from peripheral EE to the RE [48–49]. We therefore tested the effect of SARA on recycling endosome distribution and morphology. In cells overexpressing SARA, Rab11-GFP endosomes lost the main juxtannuclear distribution usually observed in control neurons (Fig 2K), and Rab11A-positive recycling endosomes appeared as a scattered fluorescent signal throughout the neuronal soma (Fig 2L and 2M).

In the next series of experiments, hippocampal neurons in culture were then transfected with a specific shRNA designed to knockdown SARA, (HcRed-tagged sh-SARA; S2 Fig) and

the morphology/distribution of EE or RE was analyzed in cells co-expressing Rab5-GFP, or Rab4-GFP or Rab11-GFP constructs. No apparent alterations were detected in EE (Rab5-GFP distribution) in SARA suppressed neurons compared to control neurons (Fig 2A vs. 2D and 2E), but Rab4-GFP endosomes changed their distribution (Fig 2F vs. 2I and 2J; S3 Fig) completely occupying the neuronal perikaryon. In cells where SARA was knocked down, Rab11-positive recycling endosomes also occupied the neuronal perikaryon area, spreading beyond the usual region where they are predominantly founded in control neurons (Fig 2K vs. 2N and 2O). Similar results were found when we analyzed Rab4 and Rab11-endosomes distribution in SARA-depleted neurons from SARA-KO mice (Fig 3).

It is known that Rab5 stimulates the fusion between EEs [50] and that Rab4 activity facilitates the fission of tubules emanating from EEs and preferentially regulates direct recycling from the sorting endosomes [41; 51]. The multiple and dynamic organization of Rab5, Rab4 and Rab11 constitutes a mosaic of membrane domains involved in functional communication between contiguous Rab domains ensuring that the flow of cargo molecules through distinct domains on endosomes is properly regulated [52–53].

Our data suggest a new role for SARA as a regulator of the pathways connecting early, sorting and recycling endosomes into hippocampal neurons. We propose that changes in SARA expression would lead to shifts in those Rab regulated pathways.

SARA regulates neuronal development

Recycling of the plasma membrane is known to regulate axon extension [54–55]; however, the specific molecules and molecular mechanisms underlying membrane dynamics during

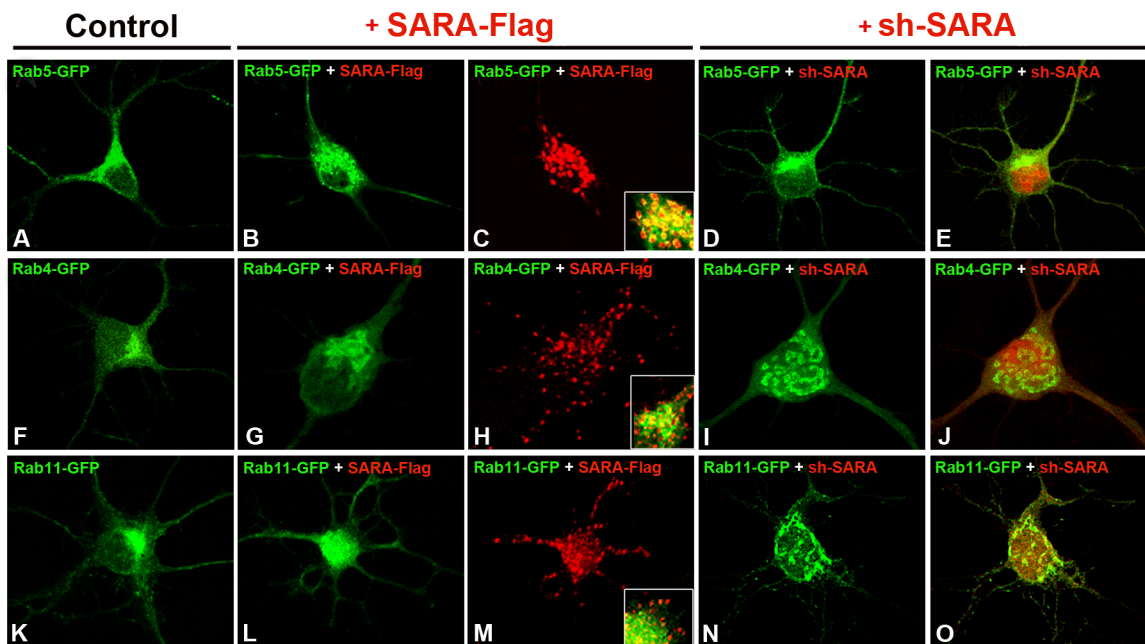


Fig 2. Characterization of Rabs distribution in neurons with modified SARA expression. Hippocampal pyramidal neurons at 7DIV were transfected with (A) Rab5-GFP, (F) Rab4-GFP or (K) Rab11-GFP where usual Rabs localization was observed. In neurons cotransfected with Rab5-GFP + SARA-Flag (B-C, inset), Rab5 labeling was extensively colocalized with SARA-Flag on enlarged early endosomal membranes; (G-H) no obvious alterations were observed for Rab4 endosomes after SARA overexpression. (L-M) Images showing intracellular distribution of Rab11-positive endosomes when SARA was overexpressed. Images of neurons transfected with (I-J) Rab4-GFP or (N-O) Rab11-GFP and cotransfected with sh-SARA (red); Rab4- and Rab11-positive recycling endosomes were distributed occupying the neuronal perikaryon area compartment compared to their distribution in controls (I vs F; N vs K). (D-E) No changes were observed in Rab5-GFP neurons cotransfected with sh-SARA. This suggests that there was a change in the Rab4-positive and Rab11-positive vesicular compartments in EE and RE after SARA suppression.

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neuronal migration and morphogenesis have not been fully characterized. Interestingly, expression of dominant-negative Rab5 mutants enhanced NGF-mediated neurite outgrowth in PC12 cells, whereas a constitutively active Rab5 mutant or Rabex-5 inhibited this process [56].

Gain-of-function studies were subsequently performed to examine the phenotypes of neurons with ectopic expression of SARA. In these experiments, neurons were transfected with SARA-GFP or SARA-Flag and the morphology of the cells was analyzed. The results obtained showed that neurons overexpressing SARA-GFP (Fig 4C and 4D) displayed abnormally short

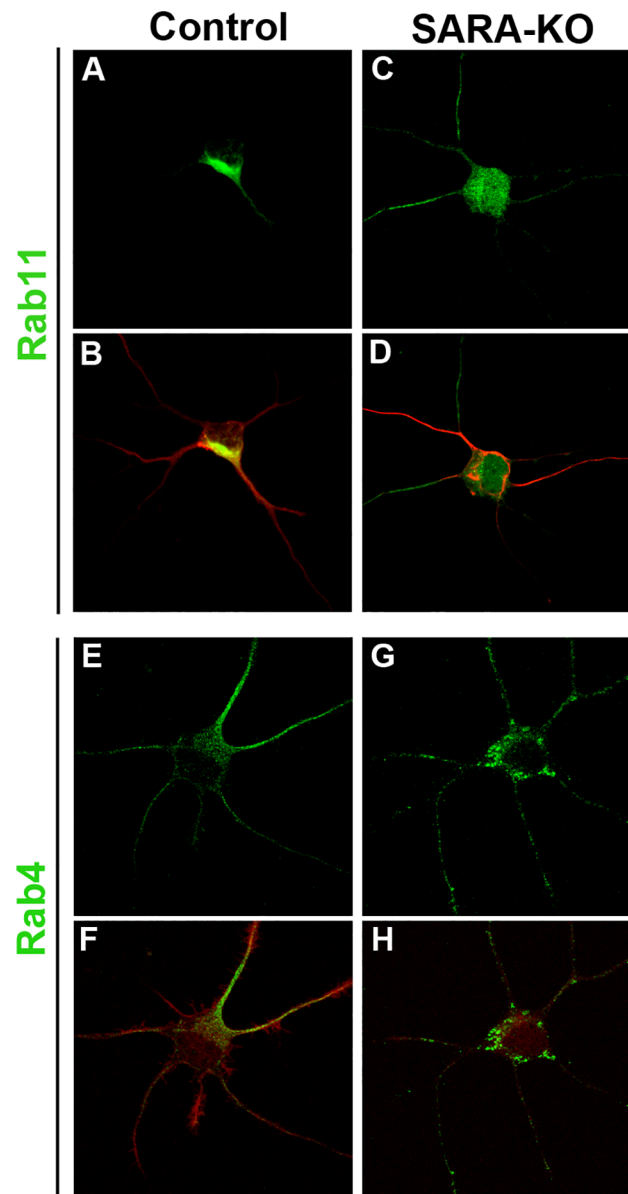


Fig 3. Rab11 and Rab4 positive endosome distribution in SARA-KO neurons. (A-D) Double-stained neurons for Rab11 and Tyr-Tubulin. Tyr-Tubulin labeling is used to show the general morphology of the neuronal cells. (E-H) Double-stained neurons for Rab4 and Rhodamine-Phalloidin. Phalloidin stains F-actin and shows integrity of actin cytoskeleton. Note that Rab11- and Rab4-positive recycling endosomes appear as a scattered fluorescent signal throughout the neuronal soma in SARA-KO neurons (C, G) vs juxtannuclear localization in control neurons (A, E). All cells were imaged by confocal microscopy. Data is typical of over 10 neurons imaged for each condition.

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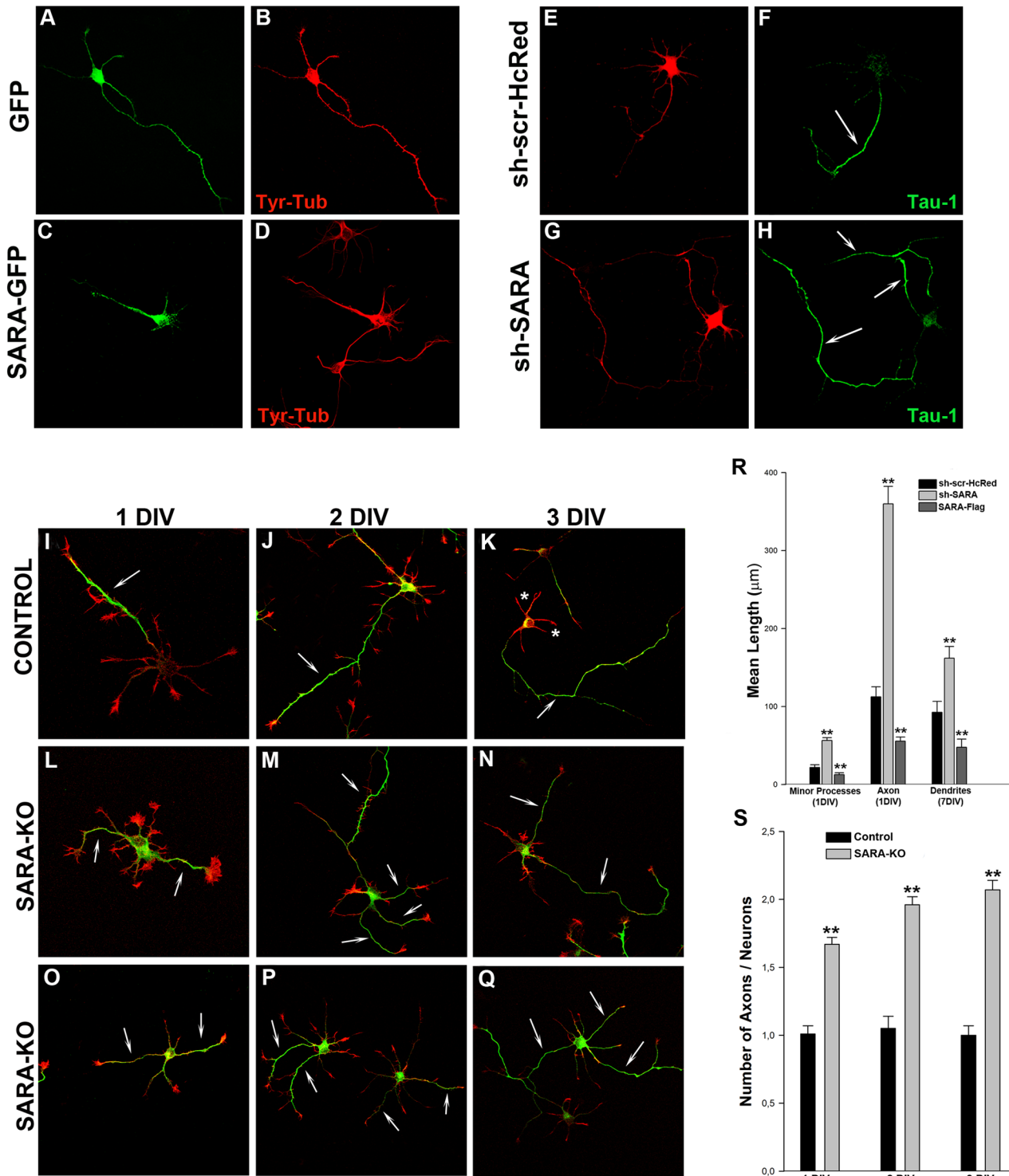


Fig 4. SARA suppression enhances axon elongation and induces formation of supernumerary axons. (Upper panel, left) Morphology of 1 DIV neurons transfected with GFP (A) or SARA-GFP (C), double stained with a mAb against Tyr-Tubulin (B and D). Note that neurons with overexpressed SARA have their neuritic development arrested or delayed. (Upper panel, right) Images showing examples of neurons treated with control sh-scr-HcRed (control; E-F) or sh-SARA (G-H). Cultures were transfected with the corresponding plasmids (1–2 μg DNA each) 24 h after plating and counterstained with a mAb against the axonal marker Tau-1 (green, F-H). Note that SARA suppression increases axonal length and produces the formation of supernumerary axons (arrows in H), contrasting with the single axon in control neurons (arrow in F). Quantification of neurite length in knockdown SARA neurons reveals a significant increase in minor processes, axons and dendrites (** $p < 0.001$; R), respect to the control or SARA overexpressed neurons. For dendrite

measurement, neurons were transfected with the same constructs but at 7 DIV and counterstained with a mAb against MAP2, a marker of neuronal dendrites. (Bottom panel) Hippocampal pyramidal neurons from control mice (I-K) or SARA-KO mice (L-Q) fixed at 1, 2 or 3 DIV and stained with a mAb against Tau-1 (green) and Rhodamine-Phalloidin (red). Note that the most of SARA-KO neurons show two or more axons (arrows). Confocal images were from independent experiments, litter mice, and different pregnant females. Bar graph shows average number of Tau-1-positive axons / 50 neurons analyzed (**p<0.001; S).

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neurites when compared to mock-transfected control cells, expressing either GFP (Fig 4A and 4B) or Flag (not shown). Morphometric analysis revealed a 50% reduction in the length of minor processes, axons and dendrites in neurons over expressing SARA (Fig 4R; Table 1). Thus, SARA overexpression inhibits neurite extension and increases the homotypic fusion of EE.

To analyze SARA participation in neuronal morphogenesis, we evaluate the consequences of its suppression on neurite outgrowth and axon extension in cultured neurons. To this end, neurons of 1 DIV were transfected with sh-SARA and examined 20–24 h later by costaining with Tau-1 antibody, to identify the axon. The majority (90%) of SARA-silenced neurons (Fig 4G and 4H) reached stage 3 of neuritic development, displaying much longer and branched axon-like neurites (Tau-1 +) than control cells, transfected either with HcRed alone or scrambled sh-RNA (sh-scr-HcRed) (Fig 4E and 4F). Identical results were observed of hippocampal pyramidal neurons cultured from SARA-KO mice, stained with Tau-1 antibody at 1, 2 or 3 DIV. These neurons exhibited two or more Tau-1+ axons (Fig 4I–4Q and 4S). Quantitative measurements confirmed these observations and revealed that SARA suppression disrupts the polarity of these neurons, resulting in a 2-3-fold increase in the length of minor neurites and number of axons, (Fig 4R; Table 1). Our results suggest that SARA silencing promotes axon and neurite outgrowth and abolishes neuronal polarity.

All together, these results constitute the first indication of SARA being a regulator of neuronal morphology and development.

SARA suppression perturbs somatodendritic and axonal protein delivery

We next assessed whether the enhanced neurite outgrowth observed in SARA-depleted cells could be explained through differential membrane delivery in axonal and somatodendritic compartments. FRAP (fluorescence recovery after photobleaching) experiments were performed to measure the mobility of fluorescent Rab11-positive recycling endosomes and their behavior when SARA was depleted. CHO-K1 cells were transfected using Rab11-GFP + sh-scr-HcRed for control cells (Fig 5A) or Rab11-GFP + sh-SARA (Fig 5B) and analyzed 20h later. The stronger area of Rab11-GFP signal was photobleached by confocal laser microscopy and monitored for fluorescence recovery. These experiments show that, in SARA knockdown cells fluorescence recovery in Rab11-positive recycling endosomes was significantly faster than in control cells (Fig 5C). These data suggest that the imbalance caused by a decrease in the levels

Table 1. Effect of SARA suppression or overexpression on neuronal shape parameters.

	Minor Processes Length 1DIV (µm)	Axonal Length 1DIV (µm)	Dendrites 7DIV (µm)
GFP	21.77 ± 3.51	112.47 ± 12.79	92.60 ± 13.93
sh-SARA	56.45 ± 3.61	359.92 ± 22.39	161.82 ± 15.23
SARA-GFP	12.49 ± 2.51	55.59 ± 5.32	47.66 ± 14.62

Neurons were transfected with 1–2 µg of plasmid for the experiments. Each value represents the mean ± s.e.m. of at least 30 cells for each experimental condition. Statistical analysis was determined as described in Materials and Methods.

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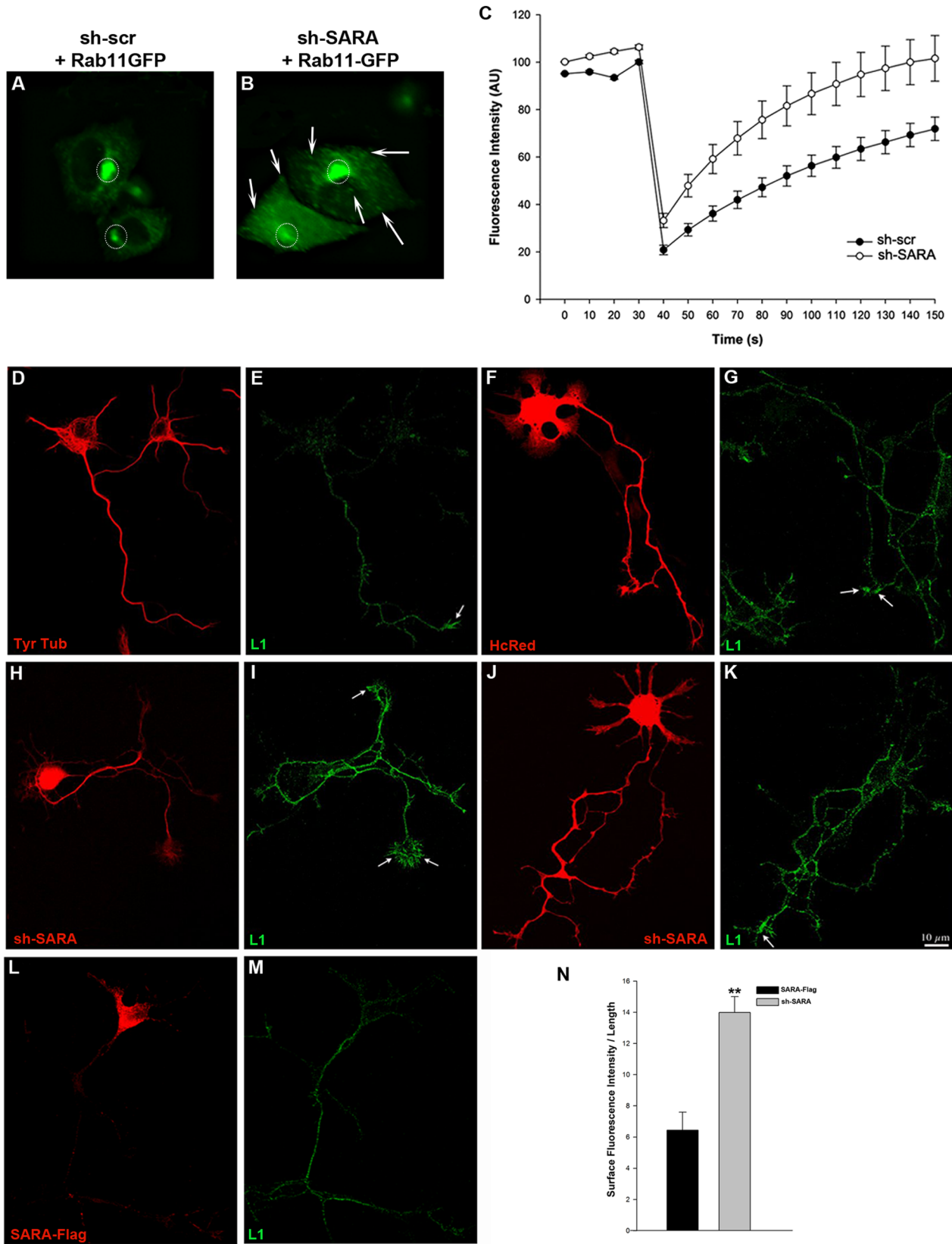


Fig 5. SARA as regulator of membrane delivery. (Upper panel) Images from FRAP experiments in CHO-K1 cells. RE was photobleached in the regions of interest (ROI, circles) in cells expressing Rab11-GFP with sh-scr-HcRed (A) or co-transfected with sh-SARA (B). Arrows indicate Rab11-positive recycling endosomes dispersed over all cells. Recovery of fluorescent molecules from the surrounding area into the photobleached region was monitored over the 150 sec during which the frames were taken. The graph quantifies the recovery into the photobleached area as the fluorescence intensity obtained in three different FRAP experiments (5 control or sh-SARA cells were analyzed in each experiment). SARA suppression significantly increases the movement of Rab11-GFP endosomes into photobleached areas (C). (Bottom panel) Suppression of SARA alters the surface distribution of the axonal membrane protein L1. Confocal images showing the surface distribution of L1: (D-E) control neurons stained for Tyr-Tubulin or transfected with: (F-G) HcRed, (H-K) sh-SARA, (L-M) SARA-Flag. Neurons were exposed to primary antibody against L1 without permeabilization. (H-K) Images show apparent increase of L1 in the axonal membrane compared to the L1 level in control or SARA over-expressed neurons. Quantification of fluorescence intensity (IF) of 3 different regions of the axons ($L = 10 \mu\text{m}$ each region) and the IF/L ratios were averaged to obtain a single value by neuron ($n = 12$ control and sh-SARA neurons). The graph confirms the observation finding more L1 on the axonal surface of SARA silenced neurons (N) (** $p < 0.001$).

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of SARA at the EEs may result in a higher number of vesicles directed to the RE or vesicles that move faster to the RE, probably avoiding the regulatory processes mediated by Rabs.

L1 is a cell adhesion molecule highly enriched in the axonal compartment [5], and it has been demonstrated that its endocytosis and trafficking are required for axonal adhesion, growth cone advance and migration [54, 57]. Alterations in RE cause an increase of membrane delivery and an accumulation of L1 at the soma membrane of PC12 cells [58–60]. We therefore analyzed whether the alterations in RE observed after SARA suppression produced changes in L1 endocytosis at the axonal membrane. Neurons were transfected with sh-SARA or HcRed and surface L1 was visualized by incubation of fixed cells with an anti-L1 antibody before detergent extraction. We quantified the amount of L1 protein added at the growth cone periphery, as an indirect measure of the membrane that is aggregated when SARA expression is modified. SARA suppressed neurons showed higher levels of L1 at the axonal and growth cone membranes (Fig 5H–5K) than HcRed controls or SARA-overexpressing neurons (Fig 5D–5G, 5L and 5M, respectively). Quantitative fluorescent measurements confirmed these observations (Fig 5N), suggesting a function of SARA in the regulation of membrane trafficking to the axonal compartment, the main source for its growth and elongation.

To evaluate a role of SARA as a regulator of membrane traffic to the somatodendritic compartment, we assessed the dynamics of transferrin receptor (TfR), a dendritic membrane protein. Selective sorting and delivery are responsible for the dendritic targeting of TfR [61]. We used TIRFM to evaluate how the expression of SARA alters the fusion of TfR-GFP-containing vesicles with the somatodendritic plasma membrane. TfR-GFP fusion events were defined as those vesicles approaching to the plasma membrane (rapid spread), followed by fluorescence scattering and subsequent disappearance of the vesicle. Examples of fusion events are shown in Fig 6A and 6B. Quantification of the total number of fusions every 2 min revealed that SARA overexpression significantly reduced the number of TfR-GFP positive vesicles fusing to the somatodendritic membrane, while in neurons transfected with sh-SARA the number of fusion events were 3–4 fold higher than in control neurons (Fig 6C).

These results were confirmed using a biochemical approach. SARA expression was altered in CHO-K1 cells and cell-surface proteins were labeled with a sulfo-biotin reagent. Cells were cotransfected with SARA-Flag + TfR-GFP or HcRed + TfR-GFP or sh-SARA + TfR-GFP and cell-impermeant sulfo-biotin was added to the culture medium at 4°C, at which temperature the endocytosis was markedly retarded. After biotinylation, cultures were lysed and incubated with avidin beads to obtain two fractions by centrifugation: the pellet with cell-surface biotinylated proteins and the supernatant with intracellular un-biotinylated proteins. Western blot analysis revealed that the total amount of TfR-GFP in membrane was significantly higher in SARA depleted cells than in control or SARA-overexpressing cells (Fig 6D–6F). These experiments show a regulatory effect of SARA on membrane expansion in the somatodendritic and axonal compartments in hippocampal neurons.

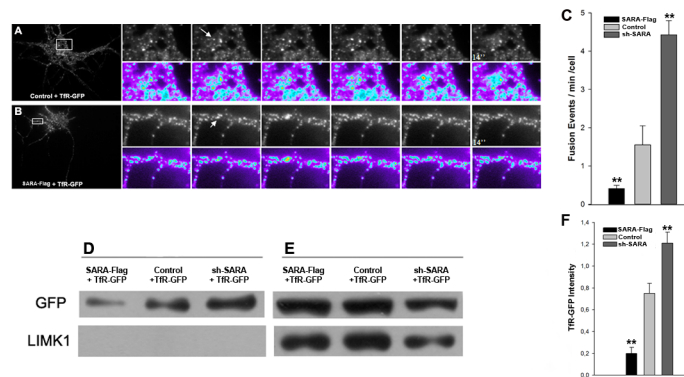


Fig 6. SARA suppression perturbs somatodendritic protein delivery. Time lapse images by TIRFM showing fusion events of vesicles to the plasma membrane in neurons of 7 DIV, transfected with Tfr-GFP plus HcRed (A) or SARA (B). Images were taken every 2 seconds for at least 2 minutes. The graphic shows a significant reduction of fusion events at the somatodendritic plasma membrane in cells transfected with SARA plus Tfr-GFP and significantly more fusion events in sh-SARA transfected neurons (C; n = 17 neurons; **p<0.001). The Tfr-GFP cell-surface protein increases after SARA silencing. CHO-K1 were transfected with Tfr-GFP alone or plus SARA-Flag or sh-SARA. The cultures were incubated with the membrane-impermeable sulfo-NHS-biotin at 4°C to prevent endocytosis. Then the cells were harvested and biotinylated proteins were collected with avidin-agarose beads (Pierce). Membrane (D) and intracellular (E) proteins were isolated by centrifugation and analyzed by Western Blot with the indicated antibodies. Absence of cytosolic contamination was corroborated with anti-LIMK1 antibody. Graph showing the increase of Tfr-GFP fluorescence intensity relative to total protein load (labeled with Tubulin; F) in SARA suppressed neurons respect to control or SARA overexpressed neurons. Data are means of three independent experiments.

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Discussion

In this study, we show a novel and fundamental role for SARA in axon/dendrite development in hippocampal neurons, a phenomenon that probably involves SARA participation in the regulation of endosomal trafficking and membrane expansion. Ectopic expression of SARA leads to inhibition of axonal growth and the formation of enlarged early endosomes. On the other hand, neurons in which SARA was depleted developed multiple and long Tau+ axon-like processes, longer neurites and large growth cones. Furthermore, SARA acute silencing or knockout in culture neurons also disturbed the Rab11-positive recycling endosomes. It is possible that, in neurons where SARA was silenced, there may be higher levels of stable microtubules, thereby producing neurons with multiple axons (polarity alteration) and RE that are more dispersed than in normal conditions. Lin et al. [62] showed that, in cells with a higher level of stable, detyrosinated (Glu) microtubules, the RE were more dispersed, suggesting that stable microtubules play a role in the organization of RE and in facilitating the movement of vesicles from the RE to the cell surface. Consistently, changing the microtubule dynamics is sufficient to alter neuronal polarization. Low doses of the microtubule-stabilizing drug taxol shift polymerizing microtubules from neurite shafts to process tips and lead to the formation of multiple axons [63].

Here we demonstrate that SARA partially colocalizes with EE markers (EEA1, Rab5, and Rab4), which suggests that SARA may play a role linking these domains on EE. Through its interaction with Rab5, SARA could recruit cargo into vesicular clusters of EE and, when interacting with Rab4, it would regulate the entry of cargo to tubular regions of EE to be later directed to the sorting endosomes. Rab proteins associate with different subcompartments generating membrane-specific domains [40]. Rab5 is associated with EEs in which it regulates endocytic vesicle fusion with preexisting EEs and homotypic fusion between early endosomes

[64]. During maturation, EEs extend tubules (sorting endosomes Rab4 + [65]) that become the RE. Rab4 distributes into EEs and REs and it has been postulated to control a fast recycling pathway [64; 66–67]. Recently Falk et al. suggested that Rab5 and Rab4 participate in a recycling loop within the growth cone that controls the speed of axon elongation [68]. The present observations are consistent with this proposal and place SARA as a novel regulator of early endosomal trafficking and axon formation in developing neurons.

Our results also suggest a role for SARA in the regulation of Rab11-positive endosomes. Rab11 regulates membrane trafficking between TGN and RE, recycling of TfR [69] and AMPA receptors on the cellular surface in neurons [70]. The existence of a regulated route through various endosomes endows them with the capacity to finely coordinate the distribution of receptors and the extent of signaling [53]. Lazo et al. described how after the addition of BDNF, Rab11-positive structures were less mobile and significantly larger, suggesting that there is an increase of Rab11-positive vesicular compartments in dendrites after dendritic branching stimulation with BDNF [71]. Constitutively active and WT forms of Rab11A significantly increased the axonal outgrowth of cortical neurons [72]. Considering the principal role of active Rab11 in promoting trafficking from recycling endosomes to the plasma membrane, it is conceivable that Rab11-positive vesicular regions or domains observed in SARA-suppressed neurons may function as “Rab11-stations” which stimulate axonal outgrowth by providing axons and dendrites with extra membranes.

Recently, Chang et al. reported that adult SARA-CKO mice (conditional KO) developed normally and described no significant abnormalities compared with WT mice. The histopathological analysis of these animals was not shown in the manuscript [24]. However, reinforcing our findings of neurons transfected with sh-SARA, our results in cultured hippocampal neurons from SARA-KO mice showed supernumerary axons, which imply significant alterations in neuronal polarity. Whether this has an impact on behavior or neurophysiology in the SARA-KO animals remains an open question.

Alterations of SARA expression in neurons not only caused morphological changes but also resulted in functional alterations: SARA silencing resulted in an increased number of TfR containing vesicles that fuse with cellular surface and ectopic SARA expression decreased them. The importance of correct delivery of membrane and its associated proteins for maintenance of the identity of axonal and dendritic compartments in neurons has been recognized [73–74]. Using TfR and L1 proteins, we demonstrate that SARA regulates membrane delivery to somatodendritic and axonal compartments. These effects were also observed in cell lines in which it was proposed that SARA oligomerization could contribute to early endosomal membrane fusion and/or expansion [21].

Together, these findings point up the importance of regulated endosomal trafficking in a membrane expansion process in which SARA would play a role as a new regulator protein. Our data indicated that SARA could work both as a positive regulator of homotypic and heterotypic EE fusion and as a negative regulator of membrane release toward the RE in neurons. We propose that unusually high homotypic endosomes fusion occurs after SARA overexpression, keeping proteins (e.g. TfR) at the EE and thus preventing them being delivered to the RE and plasmatic membrane by the slow recycling pathway or directly to the membrane by the fast recycling pathway. These neurons overexpressing SARA show short neurites and low amounts of TfR and L1 on the surface. On the other hand, in SARA KD /KO neurons, EEs are unable to mature because normal homotypic endosome fusion is necessary for EE maturation, and so there is no regulation/sorting trafficking between EE-RE compartments. It may be that Rab11-Rab4 positive endosomes disperse all over the neuron, which in turn increases TfR (by fast or slow recycling) and L1 (by slow recycling) to the cell surface, generating more axons and larger growth cones.

Finally, the endosomal machinery controls the withdrawal of adhesion molecules in proximal regions and their subsequent insertion at the front of the cell, all of which ensures the traction forces needed to maintain cell morphology and migratory patterns [75]. It has been recently shown that Rab5 and Rab7, two GTPases known to be involved in membrane trafficking and protein recycling control, are important to regulate normal migration [76]. Since SARA activity is downstream of Rab5 direction, it would be interesting to evaluate neuronal migration through in vivo experiments when SARA expression is altered.

Multiple classes of proteins are responsible for ensuring the specificity of the sorting, budding, transport and fusion events during endosomal trafficking in neuronal or non-neuronal cells [77–81]. The endosomal pathway in neurons is not yet completely characterized, but there are many labs actively involved in research and new insights are emerging constantly. The precise roles that various proteins play in neuronal membrane trafficking are being investigated; from now on SARA could be one of these. Our results demonstrate that changes in SARA activity alter neuron development.

Supporting Information

S1 Fig. Colocalization between Rab5-GFP and SARA. Rab5-GFP transfected neurons were stained with SARA antibody. The images (A-C) show partial colocalization. *m1* and *m2* maps generated with Costes colocalization mask (red color in the scale 100% colocalization, blue no colocalization condition; D-E). Similar results for Rab4-GFP transfected neurons (F-G). (TIF)

S2 Fig. SARA expression after sh-SARA treatment. (A) Neurons transfected with sh-SARA (red) for 20hs and followed by immunostaining for SARA (green; B). Arrows show lower expression level of SARA protein in sh-SARA-transfected neuron compared with normal level of SARA (asterisk). Immunoblotting of an equal amount of proteins extracted from neurons treated with SARA-Flag plus control or sh-SARA for 20 hr (C) was revealed using the indicated antibodies. (TIF)

S3 Fig. Rab4-positive endosomes change their distribution after SARA suppression. (A-B) Rab4-GFP endosome localization in control and (C-D) sh-SARA neurons. (B, D) Histograms showing Rab4-positive endosomes flooding the soma in sh-SARA neuron. (TIF)

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Author Contributions

Conceived and designed the experiments: CIA CC. Performed the experiments: CIA SOS CC. Analyzed the data: CIA CC. Contributed reagents/materials/analysis tools: CIA SOS CC. Wrote the paper: CC. Supervised all steps of the research, including formulation of the working hypothesis, data analysis, and writing of the manuscript: CC.

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