

(Patho-)physiological relevance of PINK1-dependent ubiquitin phosphorylation

Fabienne C Fiesel¹, Maya Ando¹, Roman Hudec¹, Anneliese R Hill¹, Monica Castanedes-Casey¹, Thomas R Caulfield¹, Elisabeth L Moussaud-Lamodière¹, Jeannette N Stankowski¹, Peter O Bauer¹, Oswaldo Lorenzo-Betancor¹, Isidre Ferrer^{2,3}, José M Arbelo⁴, Joanna Siuda⁵, Li Chen^{6,7}, Valina L Dawson^{6,7,8,9,10}, Ted M Dawson^{6,7,9,10,11,12}, Zbigniew K Wszolek¹³, Owen A Ross^{1,14}, Dennis W Dickson^{1,14} & Wolfdieter Springer^{1,14,*}

Abstract

Mutations in PINK1 and PARKIN cause recessive, early-onset Parkinson's disease (PD). Together, these two proteins orchestrate a protective mitophagic response that ensures the safe disposal of damaged mitochondria. The kinase PINK1 phosphorylates ubiquitin (Ub) at the conserved residue S65, in addition to modifying the E3 ubiquitin ligase Parkin. The structural and functional consequences of Ub phosphorylation (pS65-Ub) have already been suggested from in vitro experiments, but its (patho-)physiological significance remains unknown. We have generated novel antibodies and assessed pS65-Ub signals in vitro and in cells, including primary neurons, under endogenous conditions. pS65-Ub is dependent on PINK1 kinase activity as confirmed in patient fibroblasts and postmortem brain samples harboring pathogenic mutations. We show that pS65-Ub is reversible and barely detectable under basal conditions, but rapidly induced upon mitochondrial stress in cells and amplified in the presence of functional Parkin. pS65-Ub accumulates in human brain during aging and disease in the form of cytoplasmic granules that partially overlap with mitochondrial, lysosomal, and total Ub markers. Additional studies are now warranted to further elucidate pS65-Ub functions and fully explore its potential for biomarker or therapeutic development.

Keywords early-onset Parkinson's disease; mitophagy; Parkin; phosphorylated ubiquitin; PINK1

Subject Categories Neuroscience; Post-translational Modifications, Proteolysis & Proteomics; Molecular Biology of Disease DOI 10.15252/embr.201540514 | Received 8 April 2015 | Revised 24 June 2015 | Accepted 25 June 2015 | Published online 10 July 2015 EMBO Reports (2015) 16: 1114-1130

Introduction

Mutations in PINK1 and PARKIN are the most common cause of recessive early-onset Parkinson's disease (PD). Together, they coordinate a mitochondrial quality control pathway that ensures safe disposal of defective (mitophagy) and maintenance of healthy mitochondria [1]. This stress-induced pathway is tightly controlled and underlies complex regulation at multiple steps of a sequential process [2]. Upon mitochondrial damage, the protein kinase PINK1 is stabilized on the outer membrane and recruits the E3 ubiquitin (Ub) ligase Parkin from the cytosol [3]. PINK1 has been shown to phosphorylate Parkin [4-6] in its Nterminal Ub-like (UBL) domain, which is required for Parkin's structural [7] and functional activation [8]. Parkin is "charged" with Ub by E2 co-enzymes that modulate its mitochondrial translocation and enzymatic functions, both of which are linked [9,10]. Parkin then labels mitochondrial substrate proteins with poly-Ub chains of distinct topologies to mediate their sequestration and/or degradation. Parkin and generated Ub conjugates are also subject to regulation by specific de-ubiquitinating enzymes

Department of Neuroscience, Mayo Clinic, Jacksonville, FL, USA 1

Institut de Neuropatologia, Servei d'Anatomia Patològica, Hospital Universitari de Bellvitge, Hospitalet del Llobregat, Spain 2

³ CIBERNED, Centro de Investigación Biomédica en Red sobre Enfermedades Neurodegenerativas, Instituto de Salud Carlos III, Barcelona, Spain

⁴ Department of Neurology, Parkinson's and Movement Disorders Unit, Hospital Universitario Insular de Gran Canaria, Las Palmas de Gran Canaria, Spain

Department of Neurology, School of Medicine in Katowice, Medical University of Silesia, Katowice, Poland

⁶ Neuroregeneration and Stem Cell Programs, Institute for Cell Engineering, Johns Hopkins University School of Medicine, Baltimore, MD, USA

⁷ Department of Neurology, Johns Hopkins University School of Medicine, Baltimore, MD, USA

Department of Physiology, Johns Hopkins University School of Medicine, Baltimore, MD, USA 8

⁹ Adrienne Helis Malvin Medical Research Foundation, New Orleans, LA, USA

¹⁰ Diana Helis Henry Medical Research Foundation, New Orleans, LA, USA

Solomon H. Snyder Department of Neuroscience, Johns Hopkins University School of Medicine, Baltimore, MD, USA 11

¹² Department of Pharmacology and Molecular Sciences, Johns Hopkins University School of Medicine, Baltimore, MD, USA

Department of Neurology, Mayo Clinic, Jacksonville, FL, USA 13

¹⁴

Neurobiology of Disease, Mayo Graduate School, Jacksonville, FL, USA

^{*}Corresponding author. Tel: +1 904 953 6129; Fax: +1 904 953 7117; E-mail: springer.wolfdieter@mayo.edu

(DUBs) [11]. Removal of individual Ub moieties or chains from substrates modulates downstream functions that are decoded by Ub-binding adaptors.

PINK1 has just recently been identified to phosphorylate Ub, in addition to the Ub ligase Parkin, at a conserved serine 65 (S65) residue [12–14]. Both phosphorylation events are required for full activation of Parkin by feed-forward mechanisms during mitophagy [15–17]. While phosphorylation of the modifier protein further increases complexity, it also provides more selectivity and specificity for a seemingly universal ubiquitination process. In addition to activation of Parkin, consequences of pS65-Ub on structure, chain assembly, hydrolysis, and recognition have been reported *in vitro* [18]. During preparation of this manuscript, another study suggested pS65-Ub as the Parkin receptor on damaged mitochondria [19]. However, the (patho-)physiological significance of this post-translational modification in particular in neurons and in brain remains unclear.

Here, we developed and carefully characterized two phosphospecific antibodies as tools to demonstrate the (patho-)physiological relevance of pS65-Ub. While one of the antibodies was specific to pS65-Ub, the other antibody recognized both pS65-Ub and pS65-Parkin. We confirmed that the obtained signals were: (i) specific to phosphorylated S65, (ii) induced by mitochondrial stress, (iii) dependent on PINK1 kinase, and (iv) reversible by and sensitive to phosphatase activity. For the first time, we corroborated the presence of pS65-Ub under endogenous conditions in stressed primary neurons and *in vivo* in human postmortem brains. Importantly, primary cells and brain tissue from PD patients carrying *PINK1* mutations were largely devoid of pS65-Ub signal. Our findings suggest that pS65-Ub accumulates with stress, disease, or age, and highlight its significance and potential for future biomarker and/or therapeutic development.

Results

Validation of pS65-Ub antibodies in vitro

We sought to develop antibodies specific to Ub phosphorylated at Ser65 (pS65-Ub) to investigate its significance in primary neurons, in human brain, and in PD patient samples. Affinity purification yielded two selective and sensitive rabbit polyclonal antibodies (hereafter referred to as pS65-Ub#1 and pS65-Ub#2) as shown by dot blot with the immunogenic and an unmodified control peptide (Fig 1A). Western blots (WBs) of synthetic or PINK1-phosphorylated pS65-Ub confirmed their selectivity for modified Ub only (Fig 1B and C). Similar to monomeric pS65-Ub, antibodies also detected poly-Ub chains that had been phosphorylated by PINK1 wild-type (WT), but not kinase-dead (KD) mutant (Figs 1D and EV1A). The slight preference for K48 over K63 linkage might be explained by the proximity of S65 to K63 (Fig 1E) and distinct topologies of the respective chains (Fig EV1B). K63 linkage could equally affect PINK1 phosphorylation of S65 or binding of the pS65-Ub antibodies.

To determine a potential cross-reactivity of the pS65-Ub antibodies with Parkin, the other substrate of PINK1, we performed further experiments. Direct comparison of S65-phosphorylated Ub and Parkin peptides showed a minor detection of pS65-Parkin in addition to pS65-Ub, with one of two antibodies (pS65-Ub#2, Fig EV2A). *In vitro* phosphorylation of Parkin with PINK1 confirmed some cross-reactivity of pS65-Ub#2 with phosphorylated full-length Parkin. However, compared to pS65-Ub (Fig 1C), pS65-Parkin was detected only after longer kinase reactions (60 min) and longer exposures (Fig EV2B). As this could be consistent with the idea that Ub is the preferred substrate for PINK1 over Parkin, we generated equimolar amounts of both phosphorylated proteins (in a 2-day kinase reaction to ensure complete modification of all Ub and Parkin molecules). In this setting, pS65-Ub#2 showed a stronger signal for pS65-Parkin compared to pS65-Ub (Fig EV2C).

Cellular pS65-Ub signal is induced by stress and amplified by functional Parkin

Next, we tested pS65-Ub antibodies on samples from human HeLa cells. Parental cells, which lack detectable amounts of endogenous Parkin, or cells stably overexpressing native, untagged Parkin, were treated with the mitochondrial uncoupler carbonyl cyanide m-chlorophenyl hydrazone (CCCP) (Fig 2A). WB of lysates revealed almost no signal in untreated cells, but a robust increase in pS65-Ub signal with mitochondrial damage over time. Presence of functional Parkin WT amplified the pS65-Ub levels, likely through enhanced formation of poly-Ub chains that in turn serve as substrates for PINK1. While pS65-Ub signal steadily increased over longer times CCCP treatment in cells without Parkin, it never reached levels observed in the presence of functional Parkin. Here, a peak was reached already around 4 h upon CCCP treatment, after which pS65-Ub levels decreased possibly due to substrate degradation. In addition to the prominent high molecular weight (HMW) smear, we also detected monomers and dimers of pS65-modified Ub. Interestingly, monomeric pS65-Ub accumulated more in HeLa cells lacking Parkin and appeared to be utilized in Parkin overexpressing cells for the formation of HMW conjugates. Immunofluorescence (IF) staining of HeLa cells expressing GFP-Parkin corroborated lack of pS65-Ub signal at basal conditions and robust induction upon CCCP treatment. Moreover, IF revealed an exclusive co-localization of pS65-Ub with Parkin on mitochondria as expected (Fig 2B and C).

pS65-Ub#2 recognizes both PINK1 kinase products, pS65-Ub and pS65-Parkin

To further investigate the PINK1- and Parkin-dependent amplification of pS65-Ub signal, we stably expressed a catalytically dead Parkin C431S mutant in HeLa cells. Similar to parental cells that lack Parkin, expression of this inactive variant did not result in amplification of pS65-Ub signal and appearance of HMW species as compared to cells expressing Parkin WT (Fig EV3A). The Parkin C431S mutant, which traps the Ub moiety on the catalytic center in a stable oxyester, but not WT, was recognized by the antibody pS65-Ub#2 as a discrete doublet band upon CCCP treatment. The upper band of these was indeed sensitive to NaOH cleavage, suggesting the accumulation of Ub-charged pS65-Parkin C431S over time (Fig EV3B). Of note, pS65-Ub#2 did not recognize the Parkin S65A mutant, suggesting its specificity for the S65-phosphorylated form of Parkin (Fig EV3C). To exclude a major contribution of pS65-Parkin detection to the overall cellular signal observed with pS65-Ub#2, we studied GFP-Parkin WT and C431S HeLa cells. We





Figure 1. Novel antibodies selectively detect phosphorylated Ub monomers and poly-Ub chains in vitro.

- A Dot blots were performed with the immunogen (pSer65, amino acids 59–71 of Ub: YNIQKE[pS]TLHLVL) and a corresponding, non-phosphorylated peptide (Ser65) and probed with sera and affinity-purified antibodies pS65-Ub#1 and #2.
- B Western blots (WBs) with increasing amounts of Ub and synthetic pS65-Ub (both N-terminally biotinylated, N-biotin) were probed with anti-pS65-Ub antibodies or streptavidin coupled to horseradish peroxidase (SA-HRP) as indicated.
- C Monomeric N-biotin-Ub was incubated with recombinant MBP-tagged PINK1 kinase wild-type (WT) or kinase-dead (KD) mutant for the indicated times *in vitro* prior to WB.
- D N-biotin-tagged K48- and K63-linked poly-Ub chains (*n* = 2–7) were incubated with or without MBP-tagged PINK1 WT *in vitro* prior to WB. Membranes were also probed with SA-HRP and K48 or K63 linkage-specific anti-Ub antibodies as controls, which did not discriminate between phosphorylated or non-phosphorylated Ub conjugates.
- E Ribbon diagram of the pS65-Ub monomer. pSer65 is shown in VdW representation, while all seven lysine (Lys) residues are shown in stick/licorice, both colored by atom type.



Figure 2. Cellular pS65-Ub signal is stress-induced and amplified by functional Parkin on mitochondria.

- A Parental HeLa cells and cells stably overexpressing untagged, native human Parkin WT were treated with CCCP for the indicated times, and lysates were analyzed by WB as indicated. HeLa cells without or with functional Parkin both showed an increase in pS65-Ub signal over time of CCCP treatment; however, the signal was strongly amplified in the presence of Parkin. Inlay WB (16%) for the stronger pS65-Ub#2 antibody shows phosphorylated mono- and di-Ub (as indicated by one or two black circles). Note the increase of both species over time in parental cells. In the presence of Parkin, these appear to be utilized for conjugation of higher molecular weight (HMW) phosphorylated poly-Ub chains as indicated by enhanced overall total Ub levels.
- B, C HeLa cells stably overexpressing GFP-tagged Parkin (green) were treated with CCCP as indicated, fixed and stained with anti-TOM20 (mitochondria, cyan) and antipS65-Ub#1 (B) or anti-pS65-Ub#2 (C) (red) as well as Hoechst (nucleus, blue). Upon CCCP treatment, pS65-Ub signal forms on mitochondria in the presence of functional Parkin. Scale bars, 10 μm.

confirmed that the pS65-Ub signal is amplified only in the presence of functional Parkin (Appendix Fig S1). Yet, after longer times of CCCP incubation (8 h), we noticed a very weak signal in the presence of non-functional C431S Parkin with the antibody pS65-Ub#1 and, to a stronger extent, with pS65-Ub#2. However, Parkin C431S did not translocate to damaged mitochondria and remained evenly distributed throughout the cell (Appendix Fig S1). The signal obtained with pS65-Ub#2 rather resembled the intracellular localization of mitochondria.

To finally confirm the nature of the cellular signal obtained with the pS65-Ub antibodies, we performed immunoprecipitation (IP) of denatured lysates. Both pS65-Ub antibodies pulled down increasing amounts of Ub conjugates over time following CCCP treatment from HeLa cells overexpressing Parkin WT (Appendix Fig S2A). Yet, Parkin was not detectable from either pS65-Ub IP (not shown). Reciprocal anti-FLAG IP, this time under non-denaturing, but stringent RIPA buffer conditions, pulled down substantial amounts of phosphorylated poly-Ub conjugates with 3× FLAG-tagged Parkin WT, but not C431S mutant (Appendix Fig S2B). Using an anti-FLAG antibody, Parkin protein did not show a major shift into HMW species (Appendix Fig S2B). This suggests that Parkin strongly binds to pS65-Ub chains rather than it is overtly modified by them.

pS65-Ub is specific to mitochondrial damage in all cells

Given that our novel tools allowed detection of a mitochondrial stress-induced signal, we aimed to corroborate the mitochondrial co-localization of pS65-Ub with Parkin. Subcellular fractionations from HeLa cells stably expressing 3× FLAG-Parkin WT or C431S identified stabilized PINK1 protein and functional Parkin in mitochondrial samples upon CCCP treatment (Fig 3A). In contrast to WT, Parkin C431S remained in the cytosol and was detected only with pS65-Ub#2 as the characteristic doublet band. In both cells, polymeric HMW pS65-Ub species accumulated in the mitochondrial fraction, at much higher levels in the presence of functional Parkin as seen with a total Ub antibody. However, monomeric pS65-Ub and dimeric pS65-Ub were exclusively detected in the cytoplasmic fraction and were much more abundant in the absence of functional Parkin.

To further show that pS65-Ub is specific to mitochondrial damage, we challenged cells by ER (tunicamycin) or DNA (etoposide) stress (Appendix Fig S3A). Only CCCP treatment robustly induced the PINK1-/Parkin-dependent formation of phosphorylated poly-Ub chains, whereas unfolded protein response or DNA damage response did not. This increase of pS65-Ub upon mitochondrial damage was not cell type specific since it occurred in several non-neuronal and neuronal-like cells (Appendix Fig S3B).

pS65-Ub depends on PINK1 kinase activity and is reversible

Lysates from CCCP-treated HeLa cells that were incubated with alkaline phosphatase had greatly reduced anti-pS65-Ub signal, while total Ub levels remained unchanged (Fig EV4A), showing phosphorylation dependence. The swift reversion of the pS65-Ub signal became apparent in CCCP washout experiments (Fig EV4A). HeLa cells stably expressing $3 \times$ FLAG-Parkin WT were stressed for 2 h and then washed with fresh medium lacking the mitochondrial depolarizer to allow their recovery over time. pS65-Ub signals started to fade already after 30 min of washout and were followed by deconjugation of Ub polymers as judged by the total Ub signal. While levels of pS65-Ub obviously declined shortly after CCCP washout, quantification of the de-phosphorylation events from WB experiments reached significance only after 3 h recovery (Fig EV4C). Comparable kinetics upon CCCP washout were observed for de-phosphorylation of Parkin using the C431S mutation (Appendix Fig S4). In addition, we also measured (de-)phosphorylation of pS65-Ub over time using high content imaging (HCI) (Fig EV4D). HeLa cells stably expressing native, untagged Parkin showed significant increases in pS65-Ub levels over 2 or 4 h CCCP treatment. Given higher sensitivity of imaging vs. WB, de-phosphorylation of pS65-Ub was significant already after 15 min of recovery from CCCP.

Consistent with the PINK1-dependent phosphorylation of Ub, siRNA-mediated loss of the kinase activity almost entirely suppressed the induction of pS65-Ub upon CCCP treatment (Fig 3B). This was observed for phosphorylated HMW as well as mono- and di-Ub species. Knockdown of PINK1 also suppressed formation of poly-Ub chains (see total Ub antibody) and degradation of the Parkin substrate mitofusin 2 (Mfn2). Next, we analyzed human dermal fibroblasts from healthy control subjects or PD patients harboring a homozygous PINK1 Q456X mutation [20]. pS65-Ub was strongly induced by CCCP over time in control human fibroblasts and was absent in PINK1 mutant cells or in cells prior to mitochondrial damage (Fig 4A; for valinomycin treatment, see Fig EV5A). In line with these findings, control cells showed a strong increase in pS65-Ub and almost exclusive co-localization with clustered mitochondria when analyzed by IF (Fig 4B). However, in PINK1 mutant fibroblasts, pS65-Ub antibodies showed only faint background signal under both non-stress conditions and CCCP treatment. Similar to HeLa cells (Fig EV4B), primary human fibroblasts showed swift reversion of pS65-Ub signal in response upon CCCP washout (Fig EV5B). PINK1 was again de-stabilized as early as 10 min after CCCP washout. pS65-Ub levels strongly declined within 30 min of recovery from mitochondrial stress reaching statistical significance after 1 h as measured from WB experiments (Fig EV5C). De-phosphorylation of pS65-Ub was accompanied by re-establishment of mitofusin 1 protein levels shortly after.

pS65-Ub in human patients' fibroblast-derived and primary mouse neurons

To demonstrate the relevance of pS65-Ub in neuronal cells, primary human fibroblasts were directly converted into induced neurons (iNeurons) using short hairpin RNA targeting polypyrimidine-tractbinding protein (shPTBP) [21,22]. iNeurons differentiated from control or PINK1 Q456X fibroblasts were treated with valinomycin, and cell lysates were analyzed by WB. Similar to other cell types, mitochondrial stress robustly induced pS65-Ub in iNeurons from controls, concomitant with stabilization of PINK1 (Fig 5A). Induction of pS65-Ub signals was not evident from PINK1 mutant iNeurons. IF analysis of iNeurons confirmed significantly elevated pS65-Ub levels upon valinomycin treatment specifically in control cells (Fig 5B and C). Again, PINK1 mutant iNeurons showed only background pS65-Ub antibody signal that was not enhanced upon mitochondrial damage.



Figure 3. Levels of cellular pS65-Ub species are dependent on PINK1 kinase activity.

- A HeLa cells stably expressing 3× FLAG-tagged Parkin WT or C431S were treated for 2.5 h with CCCP and subjected to subcellular fractionation. Post-nuclear supernatant (PNS, as the total lysate) as well as mitochondrial and cytosolic fractions are shown. Functional Parkin showed enhanced levels of phosphorylated poly-Ub chains in the mitochondrial fraction and reduced levels of phosphorylated mono- and di-Ub levels in the cytoplasmic fraction (indicated by one or two black circles), compared to the inactive C431S mutant. Upon CCCP, PINK1 and Parkin WT accumulate in the mitochondrial fraction. Note: PINK1 has generally higher levels in the absence of functional Parkin and vice versa, consistent with the literature. pS65-Parkin (open arrowhead) and "Ub-charged" Parkin (filled arrowhead) are only detectable for C431S mutant, but not for WT. Both species are prominently detected by pS65-Ub#2, but not the Ub-specific pS65-Ub#1. Anti-VDAC1 and p38 MAPK antibodies served as controls for fractionation into mitochondria and cytosol, respectively.
- B HeLa Parkin cells were transfected with control or PINK1 siRNA and treated with CCCP as indicated. Phosphorylated mono-, di-, and poly-Ub species were almost undetectable upon knockdown of PINK1. Loss of kinase activity also suppressed the amplification of total poly-Ub signal otherwise observed in the presence of functional Parkin. In agreement, ubiquitination/degradation of the control mitophagy substrate mitofusin 2 was not observed upon silencing of PINK1.



Figure 4. pS65-Ub signal is absent in PD patients' fibroblasts with a PINK1 mutation.

Primary skin fibroblasts derived from healthy controls or PD patients with a homozygous PINK1 Q456X mutation were incubated with CCCP as indicated.

- A Cells were lysed and analyzed by WB with antibodies against pS65-Ub and total Ub. Mitofusin 1 serves as a control substrate that is ubiquitinated and degraded in response to CCCP in the presence of functional PINK1. For treatment of fibroblasts with valinomycin, see Fig EV5A.
- B Fibroblasts were fixed and stained with anti-TOM20 (mitochondria, red), anti-pS65-Ub (green), and Hoechst (nuclei, blue). The right panels represent enlarged views of the boxed areas. Upper set: anti-pS65-Ub#1 and lower set: anti-pS65-Ub#2. pS65-Ub signal was induced in CCCP-treated control fibroblasts with clear co-localization to clustered mitochondria. As expected, inducible pS65-Ub signal was absent in PD patients' cells lacking PINK1 kinase. Scale bars, 10 μm.



Figure 5.

Figure 5. pS65-Ub is induced in human iNeurons from controls but not from PINK1 patients.

Primary human fibroblasts were directly converted into iNeurons by viral transduction with shPTBP [21,22].

- A iNeurons were treated with valinomycin as indicated, lysed, and analyzed by WB. Inducible pS65-Ub was detectable in controls, but not in iNeurons from PINK1 patients. Loss of PTBP and appearance of several neuronal markers (TUJ1, MAP2A/B, and MAP2D) show the successful conversion of fibroblasts (-) to iNeurons (iN).
- B iNeurons were treated with valinomycin, fixed and stained with anti-TOM20 (mitochondria, green), anti-pS65-Ub#2 (red), anti-TUJ1 (neuronal marker, cyan), and Hoechst (nuclei, blue). Scale bars, 20 μm. Bottom images represent enlarged view of the boxed areas.
- C Quantification of pS65-Ub signal in iNeurons. TUJ1 staining was used to generate a mask outlining the total area of the iNeurons. Mean pS65-Ub signal intensity was measured in four images per condition from three independent experiments (one-way ANOVA with Tukey *post hoc*; **P < 0.005, ***P < 0.0005). Shown are mean values \pm SD.

Since Ub and in particular S65 are highly conserved across all species, we tested the ability of the antibodies to recognize pS65-Ub in primary mouse neurons. IF of untreated vs. valinomycin-treated mouse embryonic cortical cultures demonstrated strongly induced levels of pS65-Ub upon mitochondrial damage (Fig 6A). Indeed, valinomycin treatment significantly enhanced pS65-Ub levels in primary mouse neurons (see mask in Fig 6A and quantification of intensity in Fig 6B). Of note, most of the pS65-Ub signal was derived from cytosolic granules (see Fig 6A and quantification of spots in Fig 6C) that were largely co-localized with fragmented mitochondria. In summary, we show the induction of pS65-Ub upon mitochondrial damage in primary and neuronal cells and absence in cells from PINK1 mutation carriers.

pS65-Ub in human postmortem brain

To further demonstrate the pathophysiological relevance of pS65-Ub, we stained human postmortem brain sections of the substantia nigra (SN) from neurologically normal, young (< 40 years, n = 4), and old (> 60 years, n = 6) individuals as well as from PD patients (n = 6) (Fig 7; for details on brain samples, see Appendix Table S1). Very few cytoplasmic granules were recognized in human SN from younger neurologically normal individuals. In contrast, SN from older healthy individuals showed much more cytoplasmic granules, here often within optically clear vacuoles (Fig 7A). These granules were exclusively detectable with bleeds upon immunization and were absent from sections stained with pre-immune sera (Fig 7B). Importantly, brain sections from a compound heterozygous PINK1 mutation carrier [23] showed only SN characteristic neuromelanin pigment, but were in large parts devoid of specific pS65-Ub signal (indicated by arrows, Fig 7C).

Interestingly, in PD patient brain, pS65-Ub signals were detected in granules adjacent to, but not within PD characteristic Lewy bodies (LBs) and Lewy neurites (Fig 7D). Double-label IHC and IF of PD brains with pS65-Ub and different markers (Fig 7E) showed a strong overlap with lysosomes (Fig 7E, upper panel) and a partial co-localization with mitochondria (Fig 7E, middle panel), as would be expected for a mitophagy label. While pS65-Ub staining also overlapped with total Ub signal (Fig 7E, lower panel), both patterns were largely distinct from each other. Taken together, pS65-Ub-positive mitophagy granules accumulate with age and disease in human brain.

Discussion

Here, for the first time, we show that pS65-Ub is present in primary neurons under endogenous conditions and is inducible by and specific to mitochondrial damage. Using two novel rabbit antibodies, we confirmed that pS65-Ub is dependent on PINK1 kinase activity as it was absent from primary cells and brain tissue of PD patients with PINK1 mutations. We demonstrated the (patho-)physiological relevance for pS65-Ub as a specific label for mitochondrial stress, as it accumulates with organelle damage, age, and disease. Mitochondrial stress-induced phosphorylation of Ub was observed in all analyzed cell types including mouse primary neurons and human iNeurons. Monomeric pS65-Ub and dimeric pS65-Ub were identified in the cytoplasm, whereas most of the pS65-Ub signal was derived from higher molecular weight species conjugates to mitochondrial proteins. In human brain, we found pS65-Ub within cytoplasmic structures that partially co-localized with ubiquitin, mitochondria, and particularly with lysosomes. Though adjacently localized, LBs characteristic for PD were not stained with pS65-Ub antibodies. Rather pS65-Ub-positive granules were reminiscent of pathology in granulovacuolar degeneration. Enhanced, persistent levels of the pS65-Ub mitophagy label are indicative of increased mitochondrial damage and/or decreased flux through the degradation systems. However, the nature and composition of the pS65-Ub-positive structures must now be exactly determined from larger case series.

Together with PINK1, functional Parkin amplified the pS65-Ub signal in cells in response to mitochondrial damage as would be expected from the involved feed-forward and feedback loop mechanisms [24]. Parkin itself is likely an (pS65-)Ub substrate to some extent, but strongly binds to phosphorylated poly-Ub chains, consistent with their role as the Parkin receptor [19]. Further functional effects of pS65 on Ub structure, chain assembly, and hydrolysis

Mouse primary neuronal cultures were left untreated or challenged with valinomycin as indicated.

- B Quantification of the mean intensity of the pS65-Ub signal in (A).
- C The pS65-Ub mask shown in black/white in (A) is a binary image that was generated to quantify the area of the pS65-Ub spots.

Figure 6. pS65-Ub is induced in mouse primary neurons upon mitochondrial damage.

A Two representative images are shown for each condition. Cells were fixed and stained with anti-cyclophilin F (mitochondria, red), anti-pS65-Ub#2 (green), anti-TUJ1 (cyan), and Hoechst (blue). Scale bars, 10 μm.

Data information: Analyses were performed on five images for each condition for three independent experiments (two-sided unpaired t-test; *P < 0.05). Shown are mean values \pm SD.



Figure 6.



Figure 7.

have already been shown *in vitro* [18]. Of note, the reversibility of both posttranslational modifications allows complex control over mitophagy and modulation over time. Upon washout of CCCP,

pS65-Ub quickly disappeared and was followed by disassembly of earlier formed poly-Ub conjugates, consistent with the idea that phosphorylation of poly-Ub blocks their cleavage by DUBs [18].

Figure 7. pS65-Ub is detected in human brain and increases with age and disease.

Human postmortem brains were stained with pS65-Ub antibodies. Representative images of pS65-Ub#2 are shown. Similar results were obtained for both antibodies. A Representations of human substantia nigra (SN) sections from young and old brain stained with pS65-Ub antibodies. Only few cytoplasmic granules are present in a 37-year-old man, while many granules, often within optically clear vacuoles, are present in a 93-year-old man.

- B Test of (pre-)immune sera and affinity-purified pS65-Ub antibodies in human brains. Pre-immune serum and affinity-purified pS65-Ub antibody were tested on brain sections from a 67-year-old woman with ALS. Basal nucleus neurons are shown with only non-specific lipofuscin staining of the pre-immune serum, but many cytoplasmic granules detected with anti pS65-Ub. Scale bar, 6 μm.
- C SN sections from a PD patient with compound heterozygous mutations in PINK1 (c.1252_1488del and c.1488+1G>A) and age-matched control brain. Both PINK1 mutations should lead to in-frame loss of exon 7 (which encodes parts of the kinase domain) either by direct deletion or splicing-mediated exon skipping, respectively [23]. Control brain: 37-year-old man (neurologically normal, type 1 diabetes, end-stage renal disease); note sparse cytoplasmic granules and granules in cell processes (arrows). PINK1 mutant brain: 39-year-old man; note only non-specific neuromelanin staining, characteristic for SN, is observed. The asterisk indicates an unstained Lewy body.
- D, E Double staining of pS65-Ub with different antibody markers by classical IHC and by IF labeling. (D) Representative image of many cytoplasmic granules adjacent to Lewy body (asterisk) in a 84-year-old man with PD. Double IF staining with anti-α synuclein corroborates non-overlap, but localization of pS65-Ub adjacent to Lewy neurites and bodies. (E) Upper panel: Double stain (cathepsin D for lysosomes, blue; pS65-Ub, brown) in 78-year-old man with PD (pS65-Ub granules next to Lewy body (asterisk)). IF corroborates partial co-localization of pS65-Ub with a lysosomal marker (cathepsin D). Middle panel: Double stain (M117 for mitochondria, blue; pS65-Ub, brown) in 78-year-old man with PD. IF corroborates partial co-localization of pS65-Ub with a mitochondrial marker (cyclophilin F). Lower panel: Double stain (total Ub, blue; pS65-Ub, brown) in a granular axonal spheroid in 78-year-old man with PD. IF corroborates partial co-localization of pS65-Ub with total Ub. Scale bar, 6 µm for IHC and 10 µm for IF images.

Several studies indicate a role of the mitochondrial phosphatase PGAM5 for PD. In mice, loss of PGAM5 was found to result in PD-like phenotypes, potentially by dysregulating PINK1 during mitophagy [25]. In flies, however, PGAM5 deficiency rescued loss of PINK1, but not Parkin, suggesting a functional role downstream of PINK1 either between PINK1 and Parkin or independently of Parkin [26]. Going forward it will be important to determine the exact function(s) of PGAM5 and if these are mediated through de-phosphorylation of the PINK1 substrates Ub and/or Parkin.

Both of our pS65-Ub antibodies recognized phosphorylated forms of Ub, but not unmodified, across several applications. One of the antibodies (pS65-Ub#2) additionally cross-reacted with pS65-Parkin, which was only detectable by using recombinant proteins or in cells expressing the inactive Parkin C431S mutant, but not WT. In vitro, PINK1 phosphorylated much more Ub moieties over shorter time, compared to Parkin as a substrate. The preference of PINK1 for Ub as a substrate, vast cellular levels of Ub and relatively low expression of Parkin, together with an amplification of the pS65-Ub signal by PINK1 and Parkin cooperation, highly suggests a much greater cellular abundance of pS65-Ub over pS65-Parkin. Consistently, the staining pattern of both pS65-Ub antibodies was identical in all applications and from all samples tested. Although not noticeable, we cannot formally exclude some minor contribution of pS65-Parkin cross-signal to the staining obtained with pS65-Ub#2. Regardless, Ub and Parkin are both PINK1 substrates that accumulate on damaged mitochondria and jointly act as important regulators of the mitochondrial quality control pathway.

Taken together, we have investigated the pathophysiological significance of pS65-Ub signaling in neurons and in disease pathogenesis. Our findings show that pS65-Ub antibodies can be used as a tool to monitor mitochondrial damage in various applications from cell models and brain tissue. Given that the epitope is conserved across all species, further analyses of pS65-Ub in different genetic or environmental animal models of PD are now warranted. Moreover, PINK1-/Parkin-dependent mitochondrial quality control has recently been linked to other human diseases besides PD [27]. Since mitochondrial and autophagic dysfunctions emerge as a common leitmotif during aging and disease, pS65-Ub might have far-reaching implications for biomarker and therapeutic development.

Materials and Methods

pS65-Ub peptide and antibody generation

Polyclonal anti-phospho serine 65 ubiquitin (pS65-Ub) antibodies were generated by immunization of rabbits with two phosphorylated Ub peptides C-Ahx-YNIQKE[pS]TLHLVL-amide and Ac-YNIQ KE[pS]TLHLVL-Ahx-C-amide (pSer65-Ub) corresponding to aa 59–71 of Ub (21st Century Biochemicals, Inc., Marlborough, MA, USA). Affinity purification (AP) and immunodepletion (ID) were performed using phosphorylated and non-phosphorylated peptides (C-Ahx-YNIQKESTLVLHL, Ser65-Ub), respectively. A phosphorylated Parkin-specific peptide (C-Ahx-DLDQQ[pS]IVHIVQ-amide pSer65-Parkin) corresponding to aa 60–71 or 61–73 was used to analyze cross-reactivity of the antibodies.

N-terminally biotinylated, synthetic pS65-Ub (Biotin-AHX MQIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQQRLIFAGK QLEDGRTLSDYNIQKE[pS]TLHLVLRLRGG) was generated by Thinkpeptides (Oxford, UK).

Western blots

Protein was subjected to SDS–PAGE using 14, 16, or 8–16% Trisglycine gels (Invitrogen, Carlsbad, CA, USA) and transferred onto polyvinylidene fluoride (PVDF) or nitrocellulose (NC) membranes (both Millipore, Billerica, MA, USA). Membranes were incubated with primary antibodies overnight at 4°C followed by HRPconjugated secondary antibodies (1:10,000; Jackson Immuno-Research Laboratories, West Grove, PA, USA). Bands were visualized with Immobilon Western Chemiluminescent HRP Substrate (Millipore) on Blue Devil Lite X-ray films (Genesee Scientific, San Diego, CA, USA) or on a Fuji LAS-3000 (Fujifilm Life Science USA, Stamford, CT, USA) system. For densitometric analysis, ImageJ software version 1.46R was used. Total ubiquitin blots were either prepared on NC that was microwaved 10 min in ddH₂O or on PVDF washed 5 min in 4% paraformaldehyde (PFA) before incubation with primary antibody for antigen retrieval.

In vitro phosphorylation efficiency of Parkin and Ub was determined by sample separation on 14% (Ub) or 10% (Parkin) Tris–glycine gels containing 50 μ M phos-tag acrylamide AAL-107 (Wako

Chemicals, Richmond, VA, USA) and 100 μ M ZnCl₂. Prior transfer onto PVDF membranes, gels were incubated for 20 min in transfer buffer containing 0.5 mM EDTA and 0.01% of SDS and twice in the absence of EDTA.

Dot blots

pUb immunization, Ub immunodepletion, and pParkin peptides were dissolved in ddH_2O and spotted onto 0.02-µm NC membranes that were dried overnight and incubated with primary antibodies for 2 h at room temperature followed by secondary antibody incubation. Dot blots were imaged on a FUJI LAS-3000 system.

Antibodies

Following antibodies were used for Western blot analysis (WB), dot blots (DBs), immunofluorescence (IF), or immunohistochemistry (IHC): mouse anti-cathepsin D (IHC/IF, 1:5,000, IM03, Millipore), rabbit anti-mouse anti-cyclophilin F (IF, 1:1,000, ab110324, Abcam, Cambridge, UK), mouse anti-FLAG (WB, 1:250,000, F3165, M2, Sigma), mouse anti-FLAG-HRP (WB, 1:200,000, A8592, M2, Sigma), mouse anti-GAPDH (WB, 1:150,000, H86504M, Meridian Life science, Memphis, TN, USA), mouse anti-human mitochondria (IHC, 1:15, M117, Leinco Technologies, Saint-Louis, MO, USA), rabbit anti-maltose-binding protein (MBP, WB, 1:800,000, E8030S, New England Biolabs Inc., Ipswich, MA, USA), rabbit anti-MAP2 (WB, 1:1,000, M3696, Sigma), mouse anti-mitofusin 1 (WB, 1:5,000, ab57602, Abcam), mouse anti-mitofusin 2 (WB, 1:5,000, ab56889, Abcam), mouse anti-Parkin (WB, 1:1,000-1:800,000, #4211, Prk8, Cell Signaling, Beverly, MA, USA), rabbit anti-PINK1 (WB, 1:500-2,000, BC100-494, Novus Biologicals, Littleton, CO, USA), goat anti-PTBP1 (WB, 1:500, ab5642, Abcam), mouse anti-α synuclein (IF, 1:50, 18-0215, LB509, Invitrogen), rabbit anti-TUJ1 (WB, 1:1,000, β3-tubulin, D71G9, CST), chicken anti-TUJ1 (IF, 1:500, β3-tubulin, AB9354, Millipore), rabbit anti-Ub K48-linked (WB, 1:50,000, #05-1307, Apu2, Millipore), rabbit anti-Ub K63-linked (WB, 1:50,000, #05-1308, Apu3, Millipore), mouse anti-Ub (WB, 1:2,000-1:500,000, IHC/IF, 1:1,000-30,000, MAB1510, ubi-1, Millipore), mouse anti-TOM20 (IF, 1:1,000, sc-17764, Santa Cruz Biotechnology, Dallas, TX, USA), mouse anti-vinculin (WB, 1:100,000-1:500,000, V9131, Sigma). pS65-Ub antibodies were used 1:1,000 for WB and DB, 1:200 for IF, and 1:100-1:1,000 for IHC. Streptavidin-HRP (Jackson ImmunoResearch 016-030-084) was diluted 1:5,000,000.

In vitro kinase assays

MBP-tagged insect (*Tribolium castaneum*) wild-type (WT, 66-0043-050) and kinase-dead (KD, D359A, 66-0044-050) PINK1 as well as untagged human Parkin (63-0048-025) were purchased from Ubiquigent (Dundee, UK). Untagged Ub chains (UC-240, UC-340), N-terminally biotinylated Ub (UB-560), and Ub chains (UCB-230, UCB-330) were from Boston Biochem (Cambridge, MA, USA). Reactions were performed in kinase buffer (20 mM Hepes pH 7.4, 10 mM DTT, 0.1 mM EGTA, 100 μ M ATP and 10 mM MgCl₂) and incubated at 30°C for the indicated times. For standard reactions, 1 μ g PINK1 protein was used per μ g of substrate. To achieve complete phosphorylation of equimolar amounts of ubiquitin and Parkin, reactions containing 1 μ g of MBP-PINK1 per 1 μ g of Parkin or 0.17 μ g of

N-terminally biotinylated Ub (both 19 pmol) were incubated for 2 days at 30°C. All reactions were stopped by addition of SDS sample buffer and denaturing at 95°C. 15–300 ng was loaded per lane.

Cell culture

All cells were grown at 37°C under humidified conditions in 5% CO₂/air. Human epithelial cancer (HeLa), human neuroglioma H4, and human neuroblastoma BE(2)-M17 (M17) cells were obtained from the ATCC (American Type Culture Collection, Manassas, VA, USA) and human embryonic kidney (Hek293E) cells from Invitrogen. Parental HeLa cells or clonal cells stably expressing untagged Parkin (HeLa Parkin), EGFP-Myc Parkin (HeLa GFP-Parkin), or 3× FLAG-Parkin WT or C431S [9] were grown in Dulbecco's modified Eagle medium (DMEM, Invitrogen) supplemented with 10% fetal bovine serum (FBS, Gemini Bio-Products, West Sacramento, CA, USA). Human neuroglioma and M17 cells were grown in Opti-MEM (Invitrogen) supplemented with 10% FBS. Control fibroblasts were obtained from Cell Applications (San Diego, CA, USA). PINK1 Q456X fibroblasts have been described previously [20]. Fibroblasts were grown in DMEM supplemented with 10% FBS, 1% PenStrep, and 1% non-essential amino acids (both Invitrogen). Fibroblasts were collected under approved Mayo Clinic IRB protocols.

Chemical treatments, DNA, and siRNA transfection of cells

Carbonyl cyanide m-chlorophenyl hydrazone (CCCP) and tunicamycin were purchased from Sigma-Aldrich (Sigma-Aldrich, St. Louis, MO, USA) and valinomycin and etoposide from Axxora (Axxora, Farmingdale, NY, USA). If not stated otherwise, CCCP and valinomycin were used in a final concentration of 10 or 1 μ M, respectively. Washout of CCCP was performed by three washes in 1× PBS. Cells were incubated further in medium lacking CCCP.

siRNA transfections were performed on two consecutive days using HiPerfect with 20 nM control (all stars negative control) or PINK1-specific siRNA (5'-GACGCTGTTCCTCGTTATGAA-3', both from Qiagen, Valencia, CA, USA).

HeLa cells were transfected with either FLAG-Parkin C431S or the double mutant C431S/S65A using Lipofectamine 2000 according to manufacturer's protocol, and medium was replaced 4 h later.

Cell lysates and subcellular fractionation

For regular Western blot analysis, cells were lysed in RIPA buffer (50 mM Tris pH 8.0, 150 mM NaCl, 1% NP-40, 0.5% deoxycholate, 0.1% SDS) plus Complete protease and PhosSTOP phosphatase inhibitor (Roche Applied Science, Penzberg, DE). Protein concentration was determined by the use of bicinchoninic acid (Pierce Biotechnology, Rockford, IL, USA). For phosphatase treatment, lysates prepared in RIPA buffer without PhosSTOP and were incubated in the presence of FastAP thermosensitive alkaline phosphatase ($1U/\mu g$ protein, Thermo Scientific, Waltham, MA, USA) in 1× enzyme buffer for 24 h at 37°C.

For Oxyester analysis, cells were harvested in preheated (95°C) SDS lysis buffer (50 mM Tris pH 7.6, 150 mM NaCl, 1% SDS). Lysates were homogenized by 10 strokes through a 20-G needle. To verify the band shift by oxyester formation, aliquots of lysates were

treated with or without NaOH (final concentration: 100 mM) for 1 h at 37°C before loading onto SDS gels.

For subcellular fractionation, cells were harvested in fractionation buffer containing 20 mM HEPES pH 7.4, 1 mM EDTA, 250 mM sucrose, and cocktail of protease and phosphatase inhibitors (Roche). After twenty strokes through a 22-G needle, nuclei and cell debris were removed by centrifugation at 1,500 g for 5 min. The post-nuclear supernatant was transferred into a new tube and centrifuged at 12,000 g for 10 min. The mitochondria-enriched pellet was resuspended in cell fractionation buffer. Cytoplasmic fractions were isolated as supernatant after ultracentrifugation at 100,000 g for 30 min.

Immunoprecipitation

For pS65-Ub IP, cells were lysed in SDS lysis buffer. DNA was sheared by 22-G needle, and samples were boiled at 95°C. Lysates were 10× diluted with NP-40-containing buffer (50 mM Tris pH 7.6, 150 mM NaCl, and 0.5% NP-40). Five hundred microgram of total cell lysate was incubated overnight at 4°C with 5 μ g of control rabbit IgG (Jackson ImmunoReseach) or pS65-Ub antibodies. Formed immunocomplex was captured by addition of 40 μ l Protein G Agarose Fast Flow (Millipore 16-266) for 2 h at 4°C and eluted with 2× SDS sample buffer.

For FLAG IP, HeLa cells stably expressing $3 \times$ FLAG-Parkin WT or C431S were left untreated or treated with 10 μ M CCCP for 2 or 4 h and lysed in RIPA buffer plus protease and phosphatase inhibitors. One thousand microgram protein lysate was immunoprecipitated using 20 μ l of anti-FLAG EZview beads (Sigma F2426) at 4°C for 4 h. Beads were washed three times in cold RIPA buffer and eluted with 150 μ l 2× SDS sample buffer and boiled at 95°C. Ten microgram of input cell lysates and 15 μ l of immunoprecipitates were analyzed by Western blot.

Differentiation of iNeurons

iNeurons were generated as described previously with modifications [21,22]. Briefly, human PTBP shRNA (shPTBP) cloned into lentiviral vector pLKO.1 (Sigma), a kind gift from Dr. Fu (University of California, San Diego), was packaged in HEK293FT cells using Virapower (Invitrogen). Viral particles were collected 48 and 72 h post-transfection. Fibroblasts were seeded on a poly-D-lysine (PDL)coated surface and transduced with lentiviral particles containing shPTBP for 16–18 h in the presence of 5 µg/ml Polybrene (Sigma). Transduced cells were selected with 1.5 µg/ml puromycin (Invitrogen) starting 2 days after transduction. Two days later, 10 ng/ml basic fibroblast growth factor (bFGF, GenScript Z02734, Piscataway, NJ, USA) was added to the fibroblast media and cultivation continued for additional two days. From day 6 on, cells were maintained in differentiation medium containing DMEM/F12 (Invitrogen), 5% FBS (further reduced to 2% after 2 days), 25 μ g/ml insulin, 100 nM putrescine, 50 µg/ml transferrin, 30 nM sodium selenite (all from Sigma), and 15 ng/ml bFGF. After 4 days, B27 supplement without antioxidants (Invitrogen) and 10 ng/ml each of brain-derived neurotrophic factor (BDNF, R&D Systems 248-BD-025, Minneapolis, MN, USA), glial cell-derived neurotrophic factor (GDNF, R&D Systems 212-GD-050), neurotensin-3 (NT3, Peprotech AF-450-03, Rocky Hill, NJ, USA), and ciliary neurotrophic factor (CNTF, Peprotech 450-13) were added to the differentiation medium. The cells were used for experiments 2–4 days later. Neuronal differentiation was confirmed by expression of the neuronal marker TUJ1 and MAP2.

Preparation of mouse cortical cultures

Primary cortical cultures were prepared from embryonic day 18 (E18) C57BL6/J mouse pups as previously described [28] with slight modifications. Briefly, fetal cortices were dissected in Hibernate A media without calcium (BrainBits, Springfield, IL, USA) and subsequently transferred into 10 ml of growth media consisting of Neurobasal A media supplemented with antioxidant-free B27, GMAX, gentamicin, and bFGF (all from Invitrogen). Cells were dissociated in growth media and strained through a 40-µm Falcon cell strainer (Becton Dickinson Biosciences, San Jose, CA, USA) to remove cellular debris and to obtain a uniform, single-cell suspension. For IF studies, cells were plated onto PDL-coated coverslips in 24-well tissue culture plates at a density of 3×10^4 cells/well. Media was partially replaced every 2-3 days until experiments were performed at 7 days in vitro. All procedures using animals were approved by the Institutional Animal Care and Use Committees at Mayo Clinic. Cells were mock-treated or treated with 1 µM valinomycin in the presence of 50 µM z-VAD-FMK (Apex Bio, Houston, TX, USA) as previously described [29].

Immunofluorescence staining of cells

Cells were grown on glass coverslips coated with PDL (Sigma P0899). Cells were fixed with 4% (w/v) PFA and permeabilized with 1% Triton X-100. Cells were incubated with primary antibodies followed by incubation with secondary antibodies (anti-mouse IgG Alexa Fluor 488, 568, or 647 and anti-rabbit Alexa Fluor 488 or 568 and anti-chicken Alexa Fluor 647—all Invitrogen) diluted 1:1,000. Nuclei were stained with Hoechst 33342 (Invitrogen) diluted 1:5,000. Coverslips were mounted onto microscope slides using fluorescent mounting medium (Dako, Carpinteria, CA, USA). Confocal fluorescence images were taken with an AxioObserver microscope equipped with an ApoTome Imaging System or a laser scanning microscope 510 META equipped with 405-, 488-, 543-, and 633-nm lasers (Zeiss, Jena, DE).

Immunohistochemistry (IHC) and IF of brain tissue

IHC has been described before [30]. In brief, sections from paraffinembedded tissue were cut at a thickness of 5 μ m, mounted on onto positively charged glass slides, and allowed to dry overnight in a 60°C oven. Following deparaffinization and rehydration, target retrieval was performed by steaming the sections for 30 min in deionized water. Endogenous peroxidase was blocked for 5 min with 0.03% hydrogen peroxide. Sections were then treated with 5% normal goat serum for 20 min, followed by incubation in the primary antibodies for 45 min at room temperature. Immunostaining was performed on the Dako Autostainer using Envision Plus or Envision G/2 Doublestain kit (Dako, Carpinteria, CA). Horseradish peroxidase (HRP) labeling of immune-depleted pS65-Ub was visualized using 3,3'-diaminobenzidine (DAB+) as the chromogen. Alkaline phosphatase labeling of mouse monoclonal antibodies M117, Ubi-1, and cathepsin D was visualized using Vector Blue Alkaline Phosphatase Substrate kit III (Vector Laboratories, Burlingame, CA). After dehydration, sections were coverslipped with or without hematoxylin counterstaining. For IF, sections were blocked with serum-free protein block and incubated overnight with primary antibodies in background-reducing antibody diluent (both Dako) after deparaffinization and steaming. Sections were washed and incubated with secondary antibodies plus Hoechst for 2 h. Signal was quenched in 3% Sudan black in 70% MeOH solution for 2 min before sections were washed and coverslipped.

High content imaging (HCI)

HeLa Parkin cells were seeded with 7,000 cells/100 µl per well in 96-well imaging plates (BD Biosciences). Twenty-four hours after plating, cells were treated with 10 µM CCCP for 0, 2, or 4 h. For CCCP washout, cells were washed three times with media lacking the uncoupler and further incubated for the indicated times. Cells were then fixed for 20 min in 4% PFA and stained with pS65-Ub antibodies and Hoechst 33342 (1:5,000, Invitrogen). Plates were imaged on a BD Pathway 855 system (BD Biosciences) with a $20\times$ objective using a 3×3 montage (no gaps) with laser autofocus every second frame. Raw images were processed using the built-in AttoVision V1.6 software. Regions of interest (ROIs) was defined as cytoplasm using the built-in "RING 2 outputs" segmentation for the Hoechst channel after applying a shading algorithm. Resulting mean intensity values were normalized.

Image quantification and statistical analysis

Images were analyzed using ImageJ (1.48v). The mean intensity of pS65-Ub signal was analyzed over the whole image (primary neurons) or using a binary mask created from the corresponding TUJ signal (iNeurons). For area analysis of pS65-Ub spots, ImageJ built-in "analyze particles" was used after applying a uniform threshold over all images. Spots with < 4 pixels were excluded from the analysis. For quantification of pS65-Ub signal on WB, the mean value of the entire lane (25–250 kDa) was used and normalized to the corresponding loading control (GAPDH). Statistical analysis of three to five independent experiments was performed with one-way ANOVA or two-sided *t*-test, as indicated.

Modeling ubiquitin structures with interpolation and minimization

The sequence of human ubiquitin proteins (PDB codes: 2KDF and 3H7S) [31,32], a cellular protein encoded by the polyubiquitin-C gene, was taken from the NCBI Reference Sequence NP_066289.3. The modeling was built using the above PDB structures and cleaned using protein preparation wizard within Maestro console of Schrodinger [33,34]. The connectors between K48- and K63-linked chains were built using Maestro, and then, superposition was applied on duplicate structures to generate transformations in X, Y, Z Cartesian coordinate space for rotation and translation to generate octa-mers lysine-linked chains. Structures were inspected using Yasara program WHAT-IF [35] and have valid conformation consistent with good phi-psi space [36–38], which includes packing (1D and 3D), dihedrals, phi-psi, Backbone, and Packing3 (WHAT-IF implementation), plus force field considerations. The superposition and

subsequent minimization of each structure was completed to generate didactic models for various linked forms of pS65-Ub protein. The final model was subjected to energy optimization with PR conjugate gradient with an R-dependent dielectric. Model manipulation was done with Maestro (Macromodel, version 9.8, Schrödinger, LLC, New York, NY, 2010). Previous models have been successfully generated using these techniques [7,28,39].

Modifications modeling

Modifications for amino acids, such as phosphorylation of serine 65 (pSer65), was achieved using the 2D sketcher and importing as an new amino acid type which can be parameterized using the Schrödinger force fields. Additionally, ubiquitin linkages were also obtainable in this manner. Using OPLS2005 or YASARA2, one can parameterize these modification and import into existing molecular dynamics integration engines, as the parameters for the modification are well documented [34,40–43].

Expanded View for this article is available online: http://embor.embopress.org

Acknowledgements

We thank Dr. Fu for the lentiviral shPTBP plasmid. We are grateful to the patients and their families for providing tissues. This study was supported by NIH/NINDS R01NS085070, the Michael J. Fox Foundation for Parkinson's Research and the Foundation for Mitochondrial Medicine, Mayo Clinic Foundation and the Center for Individualized Medicine, the Marriott Family Foundation, and a Gerstner Family Career Development Award to WS. Mayo Clinic Florida is a Morris K. Udall Parkinson's Disease Research Center of Excellence (NINDS P50 #NS072187; DWD, ZKW, and OAR). LC, VLD, and TMD are supported by grants from the NIH/NINDS Udall Center P50 NS38377, MDSCRF 2007-MSCRFI-0420-00, 2009-MSCRFII-0125-00, MDSCRF 2012-MSCRFII-0268-00, MDSCRF 2013-MSCRFII-0105-00, and the JPB Foundation. TMD and VLD acknowledge the joint participation by the Adrienne Helis Malvin and the Diana Helis Henry Medical Research Foundations through their direct engagement in the continuous active conduct of medical research in conjunction with the Johns Hopkins Hospital and the Johns Hopkins University School of Medicine and the Foundation's Parkinson's Disease Programs, M-1 and H-2014. TMD is the Leonard and Madlyn Abramson Professor in Neurodegenerative Diseases. Further support: The Uehara Memorial Foundation (MA), Mayo Clinic Center for Regenerative Medicine (POB), ALS Association (POB), NIH/NINDS R01NS078086 (OAR).

Author contributions

FCF and WS conceived the study. FCF, DWD, and WS designed the experiments. FCF, MA, RH, ARH, ELML, TRC, and MCC performed experiments. FCF, DWD, and WS analyzed the data. FCF and WS wrote the manuscript. POB, JNS, ZKW, OAR, IF, JMA, OLB, JS, LC, VLD, and TMD provided reagents and discussion.

Conflict of interest

The authors declare that they have no conflict of interest.

References

 Scarffe LA, Stevens DA, Dawson VL, Dawson TM (2014) Parkin and PINK1: much more than mitophagy. *Trends Neurosci* 37: 315–324

- Pickrell AM, Youle RJ (2015) The Roles of PINK1, Parkin, and mitochondrial fidelity in Parkinson's disease. *Neuron* 85: 257–273
- Geisler S, Holmstrom KM, Skujat D, Fiesel FC, Rothfuss OC, Kahle PJ, Springer W (2010) PINK1/Parkin-mediated mitophagy is dependent on VDAC1 and p62/SQSTM1. *Nat Cell Biol* 12: 119–131
- Iguchi M, Kujuro Y, Okatsu K, Koyano F, Kosako H, Kimura M, Suzuki N, Uchiyama S, Tanaka K, Matsuda N (2013) Parkin-catalyzed ubiquitinester transfer is triggered by PINK1-dependent phosphorylation. *J Biol Chem* 288: 22019–22032
- Kondapalli C, Kazlauskaite A, Zhang N, Woodroof HI, Campbell DG, Gourlay R, Burchell L, Walden H, Macartney TJ, Deak M *et al* (2012) PINK1 is activated by mitochondrial membrane potential depolarization and stimulates Parkin E3 ligase activity by phosphorylating Serine 65. Open Biol 2: 120080
- Shiba-Fukushima K, Imai Y, Yoshida S, Ishihama Y, Kanao T, Sato S, Hattori N (2012) PINK1-mediated phosphorylation of the Parkin ubiquitin-like domain primes mitochondrial translocation of Parkin and regulates mitophagy. *Sci Rep* 2: 1002
- Caulfield TR, Fiesel FC, Moussaud-Lamodiere EL, Dourado DF, Flores SC, Springer W (2014) Phosphorylation by PINK1 releases the UBL domain and initializes the conformational opening of the E3 ubiquitin ligase Parkin. *PLoS Comput Biol* 10: e1003935
- Shiba-Fukushima K, Inoshita T, Hattori N, Imai Y (2014b) PINK1-mediated phosphorylation of Parkin boosts Parkin activity in Drosophila. *PLoS Genet* 10: e1004391
- Fiesel FC, Moussaud-Lamodiere EL, Ando M, Springer W (2014) A specific subset of E2 ubiquitin-conjugating enzymes regulate Parkin activation and mitophagy differently. J Cell Sci 127: 3488–3504
- Kazlauskaite A, Kelly V, Johnson C, Baillie C, Hastie CJ, Peggie M, Macartney T, Woodroof HI, Alessi DR, Pedrioli PG *et al* (2014a) Phosphorylation of Parkin at Serine65 is essential for activation: elaboration of a Miro1 substrate-based assay of Parkin E3 ligase activity. *Open Biol* 4: 130213
- 11. Dikic I, Bremm A (2014) DUBs counteract parkin for efficient mitophagy. *EMBO J* 33: 2442–2443
- Kane LA, Lazarou M, Fogel Al, Li Y, Yamano K, Sarraf SA, Banerjee S, Youle RJ (2014) PINK1 phosphorylates ubiquitin to activate Parkin E3 ubiquitin ligase activity. *J Cell Biol* 205: 143–153
- Kazlauskaite A, Kondapalli C, Gourlay R, Campbell DG, Ritorto MS, Hofmann K, Alessi DR, Knebel A, Trost M, Muqit MM (2014b) Parkin is activated by PINK1-dependent phosphorylation of ubiquitin at Ser65. *Biochem J* 460: 127–139
- Koyano F, Okatsu K, Kosako H, Tamura Y, Go E, Kimura M, Kimura Y, Tsuchiya H, Yoshihara H, Hirokawa T *et al* (2014) Ubiquitin is phosphorylated by PINK1 to activate parkin. *Nature* 510: 162–166
- Ordureau A, Sarraf SA, Duda DM, Heo JM, Jedrychowski MP, Sviderskiy VO, Olszewski JL, Koerber JT, Xie T, Beausoleil SA *et al* (2014) Quantitative proteomics reveal a feedforward mechanism for mitochondrial PARKIN translocation and ubiquitin chain synthesis. *Mol Cell* 56: 360–375
- Shiba-Fukushima K, Arano T, Matsumoto G, Inoshita T, Yoshida S, Ishihama Y, Ryu KY, Nukina N, Hattori N, Imai Y (2014a) Phosphorylation of mitochondrial polyubiquitin by PINK1 promotes Parkin mitochondrial tethering. *PLoS Genet* 10: e1004861
- Zhang C, Lee S, Peng Y, Bunker E, Giaime E, Shen J, Zhou Z, Liu X (2014) PINK1 triggers autocatalytic activation of Parkin to specify cell fate decisions. *Curr Biol* 24: 1854–1865
- Wauer T, Swatek KN, Wagstaff JL, Gladkova C, Pruneda JN, Michel MA, Gersch M, Johnson CM, Freund SM, Komander D (2014) Ubiquitin Ser65

phosphorylation affects ubiquitin structure, chain assembly and hydrolysis. *EMBO J* 34: 307-325

- Okatsu K, Koyano F, Kimura M, Kosako H, Saeki Y, Tanaka K, Matsuda N (2015) Phosphorylated ubiquitin chain is the genuine Parkin receptor. J Cell Biol 209: 111–128
- Siuda J, Jasinska-Myga B, Boczarska-Jedynak M, Opala G, Fiesel FC, Moussaud-Lamodiere EL, Scarffe LA, Dawson VL, Ross OA, Springer W et al (2014) Early-onset Parkinson's disease due to PINK1 p. Q456X mutation–clinical and functional study. *Parkinsonism Relat Disord* 20: 1274–1278
- Su Z, Zhang Y, Gendron TF, Bauer PO, Chew J, Yang WY, Fostvedt E, Jansen-West K, Belzil VV, Desaro P et al (2014) Discovery of a biomarker and lead small molecules to target r(GGGGCC)-associated defects in c9FTD/ALS. *Neuron* 83: 1043–1050
- Xue Y, Ouyang K, Huang J, Zhou Y, Ouyang H, Li H, Wang G, Wu Q, Wei C, Bi Y *et al* (2013) Direct conversion of fibroblasts to neurons by reprogramming PTB-regulated microRNA circuits. *Cell* 152: 82–96
- Samaranch L, Lorenzo-Betancor O, Arbelo JM, Ferrer I, Lorenzo E, Irigoyen J, Pastor MA, Marrero C, Isla C, Herrera-Henriquez J et al (2010) PINK1-linked parkinsonism is associated with Lewy body pathology. Brain 133: 1128–1142
- 24. Riley BE, Olzmann JA (2015) A polyubiquitin chain reaction: parkin recruitment to damaged mitochondria. *PLoS Genet* 11: e1004952
- Lu W, Karuppagounder SS, Springer DA, Allen MD, Zheng L, Chao B, Zhang Y, Dawson VL, Dawson TM, Lenardo M (2014) Genetic deficiency of the mitochondrial protein PGAM5 causes a Parkinson's-like movement disorder. *Nat Commun* 5: 4930
- Imai Y, Kanao T, Sawada T, Kobayashi Y, Moriwaki Y, Ishida Y, Takeda K, Ichijo H, Lu B, Takahashi R (2010) The loss of PGAM5 suppresses the mitochondrial degeneration caused by inactivation of PINK1 in Drosophila. *PLoS Genet* 6: e1001229
- 27. Palikaras K, Tavernarakis N (2012) Mitophagy in neurodegeneration and aging. *Front Genet* 3: 297
- Zhang YJ, Caulfield T, Xu YF, Gendron TF, Hubbard J, Stetler C, Sasaguri H, Whitelaw EC, Cai S, Lee WC *et al* (2013) The dual functions of the extreme N-terminus of TDP-43 in regulating its biological activity and inclusion formation. *Hum Mol Genet* 22: 3112–3122
- 29. Cai Q, Zakaria HM, Simone A, Sheng ZH (2012) Spatial parkin translocation and degradation of damaged mitochondria via mitophagy in live cortical neurons. *Curr Biol* 22: 545–552
- Lin WL, Castanedes-Casey M, Dickson DW (2009) Transactivation response DNA-binding protein 43 microvasculopathy in frontotemporal degeneration and familial Lewy body disease. J Neuropathol Exp Neurol 68: 1167–1176
- Weeks SD, Grasty KC, Hernandez-Cuebas L, Loll PJ (2009) Crystal structures of Lys-63-linked tri- and di-ubiquitin reveal a highly extended chain architecture. *Proteins* 77: 753–759
- Zhang N, Wang Q, Ehlinger A, Randles L, Lary JW, Kang Y, Haririnia A, Storaska AJ, Cole JL, Fushman D *et al* (2009) Structure of the s5a:k48linked diubiquitin complex and its interactions with rpn13. *Mol Cell* 35: 280–290
- Jacobson MP, Friesner RA, Xiang Z, Honig B (2002) On the role of the crystal environment in determining protein side-chain conformations. J Mol Biol 320: 597–608
- Krieger E, Joo K, Lee J, Raman S, Thompson J, Tyka M, Baker D, Karplus K (2009) Improving physical realism, stereochemistry, and side-chain accuracy in homology modeling: four approaches that performed well in CASP8. *Proteins* 77(Suppl 9): 114–122

- Laskowski RA, MacArthur MW, Moss DS, Thornton JM (1993) PROCHECK: a program to check the stereochemical quality of protein structures. J Appl Cryst 26: 283–291
- Joosten RP, Salzemann J, Bloch V, Stockinger H, Berglund AC, Blanchet C, Bongcam-Rudloff E, Combet C, Da Costa AL, Deleage G et al (2009)
 PDB_REDO: automated re-refinement of X-ray structure models in the PDB. J Appl Crystallogr 42: 376–384
- Krieger E, Nabuurs SB, Vriend G (2003) Homology modeling. Methods Biochem Anal 44: 509-523
- Vriend G (1990) WHAT IF: a molecular modeling and drug design program. J Mol Graph 8: 52–56, 29

- Caulfield TR, Devkota B, Rollins GC (2011) Examinations of tRNA Range of Motion Using Simulations of Cryo-EM Microscopy and X-Ray Data. *J Biophys* 2011: 219515
- Krieger E, Dunbrack RL Jr, Hooft RW, Krieger B (2012) Assignment of protonation states in proteins and ligands: combining pKa prediction with hydrogen bonding network optimization. *Methods Mol Biol* 819: 405-421
- 41. Schrödinger (2010) Maestro, version 9.1. New York, NY: Schrödinger, LLC
- 42. Schrödinger (2012) Suite 2012. New York, NY: Schrödinger, LLC
- Schrödinger (2013) Biologics Suite: BioLuminate, version 1.1. New York, NY: Schrödinger, LLC