BMC Neuroscience



Open Access Research article

Dynamic and redundant regulation of LRRK2 and LRRK1 expression

Saskia Biskup¹, Darren J Moore¹, Alexis Rea², Bettina Lorenz-Deperieux³, Candice E Coombes⁴, Valina L Dawson¹, Ted M Dawson^{*1} and Andrew B West*5

Address: ¹Institute for Cell Engineering and Department of Neurology, Johns Hopkins University School of Medicine, Baltimore, USA, ²Institute of Genetic Medicine, Johns Hopkins University School of Medicine, Baltimore, USA, ³Institute of Human Genetics, GSF National Research Center for Environment and Health, Munich-Neuherberg, Germany, ⁴Department of Molecular Biology and Genetics, Johns Hopkins University School of Medicine, Baltimore, USA and ⁵Center for Neurodegeneration and Experimental Therapeutics and Department of Neurology, University of Alabama at Birmingham, Birmingham, USA

Email: Saskia Biskup - saskia.biskup@arcor.de; Darren J Moore - dmoore20@jhmi.edu; Alexis Rea - area5@jhmi.edu; Bettina Lorenz-Deperieux - Lorenz-Depiereux@gsf.de; Candice E Coombes - dmoore20@jhmi.edu; Valina L Dawson - vdawson@jhmi.edu; Ted M Dawson* - tdawson@jhmi.edu; Andrew B West* - abwest@uab.edu

* Corresponding authors

Published: 28 November 2007

BMC Neuroscience 2007, 8:102 doi:10.1186/1471-2202-8-102

This article is available from: http://www.biomedcentral.com/1471-2202/8/102

© 2007 Biskup et al; licensee BioMed Central Ltd.

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/2.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Received: 30 July 2007 Accepted: 28 November 2007

Abstract

Background: Mutations within the leucine-rich repeat kinase 2 (LRRK2) gene account for a significant proportion of autosomal-dominant and some late-onset sporadic Parkinson's disease. Elucidation of LRRK2 protein function in health and disease provides an opportunity for deciphering molecular pathways important in neurodegeneration. In mammals, LRRK1 and LRRK2 protein comprise a unique family encoding a GTPase domain that controls intrinsic kinase activity. The expression profiles of the murine LRRK proteins have not been fully described and insufficiently characterized antibodies have produced conflicting results in the literature.

Results: Herein, we comprehensively evaluate twenty-one commercially available antibodies to the LRRK2 protein using mouse LRRK2 and human LRRK2 expression vectors, wild-type and LRRK2null mouse brain lysates and human brain lysates. Eleven antibodies detect over-expressed human LRRK2 while four antibodies detect endogenous human LRRK2. In contrast, two antibodies recognize over-expressed mouse LRRK2 and one antibody detected endogenous mouse LRRK2. LRRK2 protein resides in both soluble and detergent soluble protein fractions. LRRK2 and the related LRRK I genes encode low levels of expressed mRNA species corresponding to low levels of protein both during development and in adulthood with largely redundant expression profiles.

Conclusion: Despite previously published results, commercially available antibodies generally fail to recognize endogenous mouse LRRK2 protein; however, several antibodies retain the ability to detect over-expressed mouse LRRK2 protein. Over half of the commercially available antibodies tested detect over-expressed human LRRK2 protein and some have sufficient specificity to detect endogenous LRRK2 in human brain. The mammalian LRRK proteins are developmentally regulated in several tissues and coordinated expression suggest possible redundancy in the function between LRRK1 and LRRK2.

Background

Mutations in the leucine-rich repeat kinase 2 (LRRK2, OMIM 609007) gene are the most common known cause for sporadic and familial Parkinson's Disease (PD) [1-4]. The most frequent mutation in North America and Europe is a glycine to serine substitution at residue 2019 (G2019S) in the LRRK2 protein which accounts for up to 10% in familial cases and up to 2% in sporadic cases [5-9]. Frequencies of LRRK2 mutations in PD cases are reported as high as 40% in North Africans and Ashkenazi Jews [10,11], but since penetrance of the mutation in 80 year old carriers may be as low as 30% with standard diagnosis [12] the discrimination between familial and sporadic disease becomes difficult [13]. LRRK2 mutation carriers present with similar clinical and neurochemical features to sporadic disease, and with only few exceptions, present with typical Lewy body pathology [14,15].

Due to the unprecedented frequency of mutations in LRRK2 occurring in PD, a number of laboratories have embarked on the development of LRRK2-null and LRRK2 transgenic animals. In multiple-tissue Northern blots, LRRK2 is nearly constitutively expressed but the timing of expression has not been described nor the profile of the related LRRK1 gene [1,2,16]. Given the sequence similarity between LRRK1 and LRRK2, a complete expression profile may indicate whether redundancy in function could mask a phenotype in transgenic and knock-out animals. In addition, inappropriate expression in transgenic animals due to off-tissue expression or abnormal developmental expression may initiate undesirable phenotypes and reduce the utility as a PD model. To provide the complete expression profile of LRRK2 and LRRK1, we utilize quantitative PCR, Northern blot and Western blot experiments to track levels of expression.

The 2527 amino acid multi-domain protein kinase LRRK2 is associated with membranes in rodent brain and is present in tissues relevant to PD in human and rodent brain [17-20]. Some studies suggest the presence of LRRK2 protein in disease-associated aggregations such as Lewy bodies and Lewy neurites, using different antibodies against human LRRK2 [21-23]. To aid the neurodegeneration research community we present a detailed characterization of twenty-one commercially available antibodies to mouse and human LRRK2 purported as specific to the 280 kDa protein.

Results

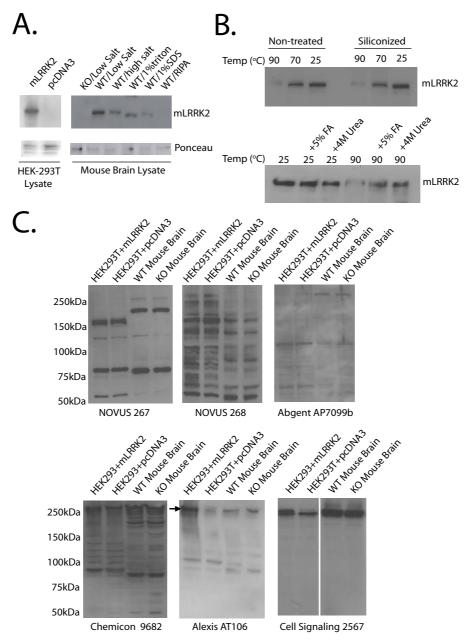
LRRK2 is present in both freely soluble and SDS-soluble fractions of mouse brain lysate as determined by antibody IH5514

Published literature to date suggests many commercially available antibodies raised against peptides derived from human LRRK2 sequence detect a protein of approximately 280 kDa in size in mouse or human tissue and/or cell lines [21-28]. To confirm whether these antibodies detect mouse LRRK2 or cross-reactive protein species unrelated to LRRK2 migrating near 280 kDa, we performed immunoblot analyses of the various antibodies in wild type mice and LRRK2 knock-out mice in which exons 39 and 40 are disrupted (S. Biskup, M. Sasaski, V.L. Dawson and T.M. Dawson, personal observation) (Figure 1). Our previously characterized antibody JH5514 recognizes human over-expressed and endogenous LRRK2 protein in mouse and humans with minimal cross-reactivity with other protein species and LRRK2 immunoreactivity is absent in LRRK2 KO mice (Figure 1A). No endogenous LRRK2 can be detected in HEK-293T cells even with long exposures and high antibody concentrations (Figure 1A and data not shown). Our validated antibody JH5514 does not detect a protein of the appropriate size, determined through alignment with recombinant LRRK2 protein, in any tested cell line lysate including human embryonic kidney (HEK-293T) cells, human neuroblastoma (SH-SY5Y) cells, human neuroblastoma (BE(2)-M17) cells and rat adrenal (PC-12) cells (data not shown). Several of these cell lines, including HEK-293T and SH-SY5Y express a low level of LRRK2 mRNA compared with brain tissue suggesting post-transcriptional regulation (data not shown).

With a series of different extraction buffers for mouse brain using LRRK2-null brain tissue as a control, we show that membrane disruption through simple mechanical homogenization of brain tissue in PBS (low salt buffer) or in high salt buffer is sufficient to extract the majority of mouse LRRK2 that can be detected by Western Blot (Figure 1A). Mouse LRRK2 protein is also detected in relatively insoluble fractions only resolved through the addition of 1% SDS buffer. Incubation of lysate samples supplemented with 1x Laemmli sample buffer for ten minutes at temperatures above room temperature led to a loss of recoverable levels of LRRK2 (Figure 1B). This phenomenon is apparently not due to an interaction with plastic disposables since siliconized tubes and tips had no effect. Supplementation of the sample buffer lysate solution with either 4 M urea or 5% Formic acid (FA) after incubation for ten minutes at 90°C resulted in a significant recovery of LRRK2 protein levels, suggesting that LRRK2 may form SDS-resistant insoluble aggregates with elevated temperatures (Figure 1B).

Commercially available antibodies that recognize overexpressed and endogenous mouse LRRK2

Using a panel of over-expressed mouse *LRRK2* (*mLRRK2*) versus empty vector transfected HEK-293T cells in combination with *LRRK2*-null versus wild-type mouse brain tissue, we tested twenty-one commercially available LRRK2 antibodies to determine specificity. Most of the antibodies cannot detect over-expressed mLRRK2 or endogenous



Biochemical characterization of LRRK2 and testing of LRRK2 antibodies on mouse tissue. A) Over-expressed mouse LRRK2 detected by JH5514 versus empty vector transfected HEK-293T cells. Whole mouse brain was mechanically homogenized in either PBS (low salt buffer), high salt buffer (PBS supplemented to 600 mM NaCl), 1% Triton X-100 (in PBS), 1% SDS (in PBS) or RIPA (1% SDS, 1% sodium deoxycholate, 1% NP-40). All buffers contained complete protease inhibitors (Roche). A knockout mouse brain is homogenized in PBS (low salt) alone and shown as control. B) Lysates derived from HEK-293T cells transfected with mLRRK2 plasmids were incubated in 1 × Laemmli sample buffer at the indicated temperatures for 10 minutes and then analyzed by SDS-PAGE. Plasticware used to process the protein samples were either siliconized or left untreated. Indicated samples were supplemented with 5% Formic acid (FA) or 4M urea after 10 minute incubation at the indicated temperature. C) LRRK2 antibodies Novus 267 and 268, Abgent AP7099b, and Chemicon (AB9682) in addition to Alexis (AT106) and Cell Signaling (2567) were tested on over-expressed mouse LRRK2 protein and wildtype and *LRRK2*-deficient mouse brain homogenized in PBS alone. All antibodies recognize cross-reactive bands near the expected size of LRRK2. An arrow denotes the position of LRRK2 in the Alexis (AT106) blot. All antibodies were tested on at least two independent membranes and lysates using optimized exposure conditions with similar results.

wild-type mLRRK2 protein (For a summary, see Table 1). For example, the antibodies Novus 267, Novus 268, Abgent AP7099b do not have the sufficient specificity to detect mLRRK2, either over-expressed or endogenous using standard blotting conditions (Figure 1C). Of note,

antibody AP7099b from Abgent was raised against a peptide identical between mouse and human. Typical for the majority of antibodies tested here, cross-reactive species near the molecular weight of LRRK2 protein confound the interpretation of blots when appropriate controls such as

Table I: Summary of antibodies tested on Western Blot

Antibody	Overexpressed mLRRK2	Overexpressed hLRRK2	Endogenous mLRRK2	Endogenous hLRRK2	Peptide Location
Abgent AP7099a	-	-	-	-	N/A
Abgent AP7099b	-	-	-	-	(LRR, AA 1246–1265, 100% identity) H: WSRVEKLHLSHNKLKEIPPE M: WSRVEKLHLSHNKLKEIPPE
Abgent AP7099c	-	-	-	-	(N-Term, around L229) H: EEIVLHVLHCLHSLAIPCNNVEVLMSGNVR M: KEIVYHVLCCLHS L AVTCSNVEVLMSGNV
Abgent AP7099d	-	-	-	-	(N-Term, around E285) H: VSCCLLHRLTLGNFFNILVLNEVHEFVVKA M: VSCSLFQKLTLGNFFNILVLNEVHVFVVKA
Abgent AP7099e	-	+	-	+	(N-Term, around E519) H: RAILHFIVPGMPEESREDTEFHHKLNMVKK M: RAILHFVVPGLLEESREDSQCRPNVLRK
Abgent AP7099f	-	+	-	-	(N-Term, around L893) H: AQSDDLDSEGSEGSFLVKKKSNSISVGEFY M: GQSDDLDSEGSESSFLVKRKSNSISVGEVY
Abgent AP7099g	-	+	-	-	(N-Term, around L899) H: AQSDDLDSEGSEGSFLVKKKSNSISVGEFY M: GQSDDLDSEGSESSFLVKRKSNSISVGEVY
Abgent AP7099h	-	+	-	-	(N-Term, around L955) H: IFDHEDLLKRKRKILSSDDSLRSSKLQSHM M: VFDHEDLLRRKRKILSSDESLRSSRLPSHM
Abgent AP7099i	-	-	-	-	(C-Term, around C2354) H: STERNVMWGGCGTKIFSFSNDFTIQKLIE M: SSERHITWGGCGTKVFSFSNDFTIQKLIET
Abgent AP7099j	-	-	-	-	(C-Term, around K2402) H: CVHFLREVMVKENKESKHKMSYSGRVKTI M: CVHFLKEVMVKLNKESKHQLSYSGRVKAI
Abgent AP7099k	-	-	-	-	(C-Term, around Y2475) H: MTAQLGSLKNVMLVLGYNRKNTEGTQKQFM: ATAQLGSLKNVMLVLGYKRKSTEGIQEQ
Abgent AM7099a	-	-	-	-	(monoclonal) N/A
Abgent AM7099b	-	-	-	-	(monoclonal) N/A
Abcam 27482	-	+	-	-	N/A
Abcam 19906	-	-	-	-	N/A
Novus Biological 267	-	+	-	+	(N-Term, AA 920–945) H: SNSISVGEFYRDAVLQRCSPNLQRHS M: SNSISVGEVYRDLALQRYSPNAQRHS
Novus Biological 268	-	+	-	+	(C-Term, AA 2500–2527) H: INLPHEVQNLEKHIEVRKELAEKMRRTSVE M: LNLPHEVQNLEKHIEVRTELADKMRKTSV
Chemicon AB9682	-	+	-	-	N/A, synthetic peptide from rat
Chemicon AB9704	-	+	-	-	N/A
Alexis (AT106)	+	+	+	+	(AA 1838–2133)
Cell Signaling 2567	+	+	-	-	(C-Term, G2090) H: FPNEFDELEIQGKLPDPVKEYGCAPWPMV M: FPNEFDELAIQGKLPDPVKEYGCAPWPMV
JH5514	+	+	+	+	(C-Term, AA 2500–2515), 100% identity H: INLPHEVQNLEKHIE M: LNLPHEVQNLEKHIE

Summary of the results of commercially available antibodies against LRRK2 tested on Western Blot. A (+) denotes antibodies that produce a discernable band specific to LRRK2 protein. A (-) denotes antibodies unable to detect a band specific to LRRK2 protein. Antibodies were designed against the human sequence unless otherwise indicated.

LRRK2-null brain lysate are not included. With very high concentrations and exposure times, many cross reactive bands appear near the expected size of LRRK2 protein but none are changed in the *LRRK2*-null brain lysate for all commercially available antibodies with the exception of the Alexis (AT106) antibody, which recognizes a faint band immediately under a prominent cross-reactive protein species near 280 kDa (Figure 1C). Other antibodies capable of detecting over-expressed *mLRRK2* in HEK 293T cells such as the Cell Signaling (2567) and Chemicon (ab9682) antibodies recognize cross-reactive protein species that presumably mask endogenous mLRRK2 protein due to the insufficient separation afforded by commercially available gradient mini-gels.

Commercially available antibodies that recognize overexpressed and endogenous human LRRK2

All antibodies were additionally tested on human overexpressed LRRK2 versus empty vector transfected HEK-293T cells (Table 1). Figure 2A shows the panel of antibodies capable of detecting over-expressed human LRRK2 (Novus 267, Novus 268, Abgent AP7099e, AP7099f, AP7099g, AP7099h, Abcam 27482 and Cell Signaling 2567). We tested antibodies capable of robustly detecting over-expressed human LRRK2 protein (Novus 267, Novus 268, Abgent AP7099e, AP7099f, AP7099g, AP7099h, Abcam 27482) using human brain lysate derived from fresh-frozen samples of human temporal lobe. In situ hybridization analyses have previously demonstrated LRRK2 mRNA is at the highest levels in the cortex and striatum and we would expect significant quantities of LRRK2 protein in the temporal lobe [16,18,29-32]. Novus 267, Novus 268 and AP7099e all have sufficient specificity to detect human LRRK2 based on their ability to resolve a band of the predicted molecular weight of LRRK2 with identical solubility profiles (Figure 2B). Novus 268 and AP7099e are shown in comparison with our in-house antibody JH5514 (Figure 2B). Using gel systems other than Tris-Glycine such as Tris-Acetate or Bis-Tris gels causes a dramatically different migration of LRRK2 protein with respect to protein markers, with migration near 240 kDa in Tris-Acetate gels (data not shown).

We visualized LRRK2 protein by Western blot in human brain lysates derived from tissue with minimal delay in processing (fresh frozen brain samples) and have been unsuccessful in detecting LRRK2 protein from the vast majority of archived frozen brain samples that correspond to typical post-mortem delays (4–24 hours) (data not shown). This may be due to a high rate of degradation or processes that otherwise render LRRK2 inaccessible to detection by immunoblotting due to typical post-mortem delay in most human tissue samples available. We have not observed significant decay in LRRK2 protein within

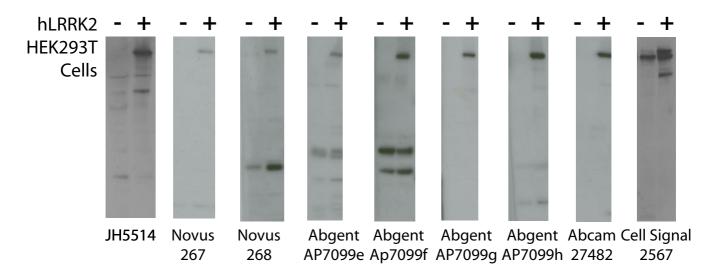
protein lysate when stored at -20 or -80 degrees or dramatic loss of LRRK2 due to freeze/thaw cycles (data not shown).

LRRK1 and LRRK2 expression in development and adulthood

Many invertebrate organisms such as Drosophila and lower vertebrate organisms such as Fugu and Zebrafish possess only a single LRRK gene, suggesting mammalian LRRK1 and LRRK2 protein diverged from a single founder [33]. LRRK-like proteins and orthologues extend well beyond complex invertebrates into even simple singlecelled eukaryotic organisms [34], suggesting a fundamental albeit unknown role for the LRRK family of proteins in mammals. Northern blot analysis has previously shown a broad distribution of LRRK2 mRNA in the brain and other organs in humans [1,2]. In order to better understand the regulation of LRRK2 expression we measured LRRK1 and LRRK2 mRNA in multiple organs in neonatal and adult mice (Figure 3A, B). We observed statistically identical levels of LRRK1 and LRRK2 mRNA in the lung, heart, skeletal muscle and lymph node (Figure 3B). LRRK2 mRNA is more abundant in kidney and brain tissue, whereas LRRK1 mRNA is more abundant in the gut.

Using Northern (whole mouse embryo) and Western (whole brain lysate) blotting we show that LRRK2 does not have an expression profile typical of a gene important during early development (Figure 4A,B). mRNA levels measured from whole mouse embryos show that LRRK2 expression increases from embryonic day 15 to day 17 (Figure 4A). Given the large increase in expression of LRRK2 during the latest stages of development, we created a panel of cDNA derived from developing mouse organs, including brain, lung, heart and liver, at various stages of development. We then went on to compare LRRK1 and LRRK2 levels using quantitative RT-PCR and observed a dynamic expression profile that distinguishes LRRK1 from LRRK2. LRRK2 most dramatically increases in lung and kidney between E15.5 and E19.5 partially accounting for the results of the whole-embryo Northern blot. Surprisingly, LRRK2 mRNA is constitutively expressed early in development of the brain and does not substantially increase upon cell maturation as might be expected by the Northern blot of whole embryo (Figure 4C). LRRK1 levels spike at E15.5 in the brain then decrease below LRRK2 levels and remain low with respect to LRRK2 throughout adulthood. In other organs such as the lung and liver, LRRK2 dramatically increases in the latest stages of development in contrast to the more constitutive expression in the heart where LRRK1 and LRRK2 are nearly identically expressed both in development and throughout adulthood.





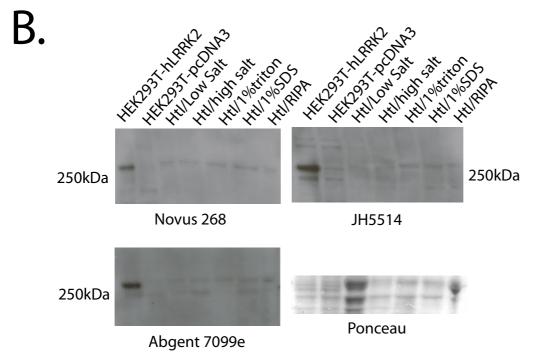


Figure 2
Testing of LRRK2 antibodies on human tissue. A) Antibodies able to detect over-expressed human LRRK2 protein. B)
Fresh frozen tissue samples from the human temporal lobe (Htl) is subjected to different extraction buffers as in figure legend IA. Novus 267, Novus 268, Abgent AP7099e and JH5514 have sufficient specificity to detect endogenous human LRRK2. All antibodies were tested on at least two independent membranes and lysates using optimized exposure conditions with similar results.

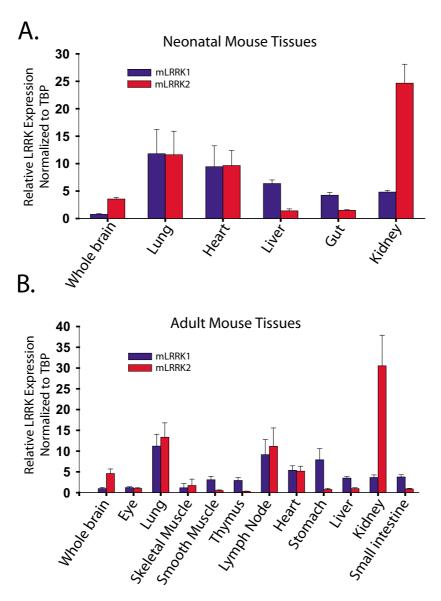


Figure 3 LRRK1 and **LRRK2** mRNA levels in development and adulthood. A) Neonatal *LRRK1* and *LRRK2* mRNA levels. Relative *LRRK* expression was calculated by normalization to *TBP* (*TATA binding protein*) within each tissue sample using the delta delta CT method. Similar results were obtained with internal normalization to *GAPDH*. Error bars represent standard error mean derived from two each independently analyzed male and female 14 day old CD-1 mice B) *LRRK1* and *LRRK2* mRNA levels in adult mouse tissues as normalized to *TBP*. Similar results were obtained with normalization to *GAPDH*. Error bars represent standard error mean independently derived from two each female and male 3 month old CD-1 mice.

Discussion

The function of LRRK proteins in mammals remains enigmatic yet strong evolutionary conservation suggests important roles. Besides obvious interest in understanding the only proteins in the mammalian proteome encoding both a kinase and GTPase domain, the striking association between *LRRK2* and PD further justifies the speedy discovery of LRRK2 protein function in cells. Within the present study we show that several commer-

cially available antibodies recognize endogenous human LRRK2 protein but no tested commercially available antibody can efficiently resolve mouse LRRK2 protein despite a number of reports that did not have the benefit of *LRRK2*-null mice. To the contrary, more abundant crossreactive species near or at the molecular weight of mLRRK2 protein are detected by most commercially available antibodies as shown for the human specific Novus antibodies and the Abgent antibody AP7099b. Cross-reac-

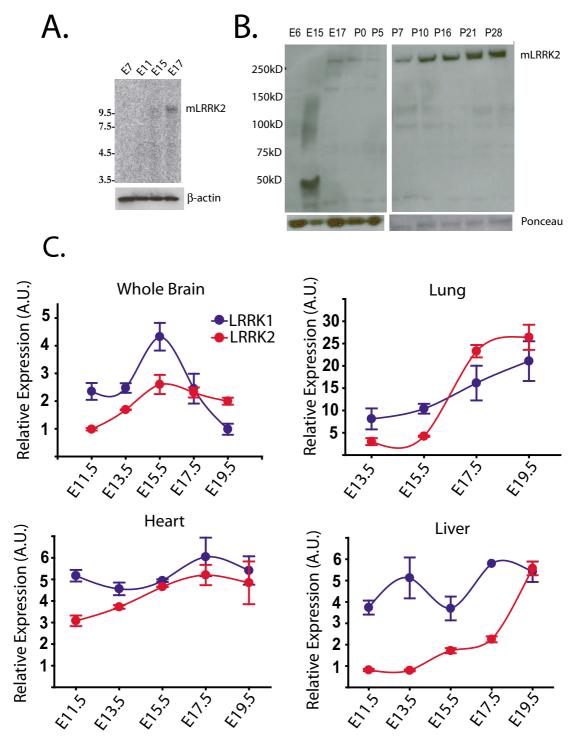


Figure 4 LRRK1 and LRRK2 mRNA levels in development and adulthood. A) Northern blot analysis of different embryonic stages (whole mouse embryo) using a *LRRK2* specific riboprobe. B) LRRK2 protein can be detected by JH5514 in whole brain lysates on E17 and throughout post-natal development C) Relative *LRRK1* and *LRRK2* mRNA levels in direct comparison calculated from whole brain, lung, heart and liver during embryonic stages E11.5 to E19.5, with internal normalization to *TBP* transcript. Error bars represent standard error mean independently derived from a total of four CD-1 embryos at the indicated age.

tive species likely mask endogenous mLRRK2 protein in the case of antibodies from Cell Signaling and Chemicon. However, both antibodies are capable of detecting over-expressed mouse protein. The Alexis antibody detects over-expressed mouse protein and endogenous mouse protein that migrates slightly below cross-reactive species. It may be possible with additional purification steps, blocking optimization, extraction optimization and improved washing and blotting conditions that some of the antibodies tested here will prove useful in the future.

In all tissues, *LRRK1* and *LRRK2* mRNA is rare compared to even nominally expressed transcripts such as *TATA* binding protein (*TBP*) and cyclophilin-A, both used here in addition to *GAPDH* as normalization controls. The combination of a poorly expressed transcript with an apparently unstable protein of above-average size creates a worst case scenario for routine detection of endogenous protein by Western blotting and immunocytochemistry despite the high solubility of LRRK2 protein.

We describe cloning of mouse *LRRK2* into expression vectors and the expression profile of LRRK1 and LRRK2 in the developing and adult mouse. Mouse LRRK2 expression increases from embryonic day 15 to day 17 and continues through adulthood. In some organs, such as the brain, LRRK2 expression apparently increases to a more nominal extent at earlier time points. We have been unable to detect LRRK2 protein in the brain before embryonic day 17 using the western blotting conditions and protein extraction protocols described here. LRRK1 and LRRK2 expression overlaps in all tissues used in this study (whole brain, eye, lung, skeletal muscle, smooth muscle, thymus, lymph node, heart, stomach, liver, kidney and gut), and in some tissues expression levels are identical. LRRK2 expression levels are particularly high in embryonic and adult lung and kidney. As kinase enzymes, LRRK protein may target tissue specific substrates for phosphorylation. It is possible that LRRK protein is so critical to cell function that the protein has diverged into two related proteins to provide a protective measure of redundancy in function.

Conclusion

Knowledge about localization, function and substrates of LRRK1 and 2 protein will help clarify *LRRK2* associated disease. We characterized twenty-one commercially available antibodies against LRRK2 protein and describe the biochemical properties of mouse and human LRRK2 proteins with regards to solubility and expression. The importance of *LRRK1* and *LRRK2* for mouse development and adult mice in different tissues is actively pursed by many research groups. *LRRK1* and *LRRK2* are comparably expressed in most tissues with stable expression in adulthood but dynamic expression during development, with

a general trend towards increased expression during organogenesis and maturation.

Methods

Antibody production and western blotting

Affinity-purified rabbit polyclonal antibodies were generated to LRRK2 protein as described previously [17]. Protein extracts from adult mouse and fresh frozen tissue samples from the human temporal lobe were prepared as described [17] and Western blot analysis (6% SDS-PAGE gels) was conducted as previously described for LRRK2 [19]. Human tissue samples were obtained from human brain temporal lobe resections snap-frozen immediately after removal and stored at -80°C until homogenization. Briefly: whole mouse brain or temporal human brain was mechanically homogenized in either PBS (low salt buffer), high salt buffer (PBS supplemented to 600 mM NaCl), 1% Triton X-100 (in PBS), 1% SDS (in PBS) or RIPA (1% SDS, 1% sodium deoxycholate, 1% NP-40). All buffers contained complete protease inhibitors (Roche). Antibodies were diluted 1:1000 in TBS-T containing 5% skimmed milk if not otherwise indicated. Incubation with primary antibodies was performed overnight. LRRK2 knockout mice were generated by disruption of exons 39 and 40 of the mouse LRRK2 gene (S. Biskup, M. Sasaki, V.L. Dawson and T.M. Dawson, personal observation). Wild-type, heterozygote and knockout LRRK2 mice were generated by crossbreeding of germline chimaeras to C57BL/6J mice. Protein extracts from mouse and fresh frozen tissue samples from the human temporal lobe were processed for Western blotting with LRRK2 antibodies as described above. LRRK2 antibodies were obtained from Abgent, Novus Biologicals, Abcam, Chemicon, Alexis and Cell Signaling.

Generation of human and mouse LRRK2 constructs

The generation of human *LRRK2* constructs has been previously described [19,35].

The full-length cDNA of mouse LRRK2 was amplified from a Marathon-Ready mouse embryo 17d cDNA library (BD Biosciences). Using nested primers based on mouse LRRK2 (GenBank accession number NM 025730) and PfuTurbo hotstart DNA polymerase (Stratagene) according to the manufacturer's protocol, two overlapping fragments from exons 1-28 (~3.9 kb) and exons 15-51 (~5.9 kb) were generated. The first part of the LRRK2 cDNA (exons 1-28) was cloned in pBluescript SK+ (Stratagene) (KpnI/NotI) and sequenced with an ABI Prism Big Dye Terminator cycle sequencing kit on an ABI Prism 3730 DNA analyzer (Applied Biosystems). To generate a full-length LRRK2 clone the second part of LRRK2 (exons 15-51) was cloned in the pBluescript plasmid already containing one part of LRRK2 (NdeI/NotI). The entire open reading frame of LRRK2 (7584 bp) was confirmed by sequencing.

Northern blotting

Whole mouse embryonic membranes were purchased from Clontech. Membranes were hybridized overnight at 65°C with a biotinylated antisense riboprobe synthesized using the MAXIscript in vitro transcription kit (Ambion) from a PCR-amplified DNA template with incorporated T7 promoter, and washed with $2 \times SSC/1\%$ SDS and then 0.1 × SSC/0.1% SDS at 65°C. Filters were visualized by incubation with streptavidin-alkaline phosphatase and chemiluminescence using the BrightStar BioDetect kit (Ambion). The riboprobe sequence for LRRK2 corresponds to nucleotide positions 4292 to 4813 of LRRK2 cDNA. PCR primers used were: forward 5'-TGAAGCCTT-GGCTCTTCAAT-3' and reverse 5'-TAATACGACTCACTAT-AGGTACAAAGCCACTTGGGTTCC-3' with bold nucleotides representing the T7 promoter priming site.

Quantitative reverse-transcription PCR

RNA was isolated from embryonic and adult CD-1 mice (Charles Rivers) via mechanical homogenization and extraction in TriZOL and both cleaned and concentrated using a commercially available kit (Qiagen RNEasy Micro; Qiagen). Reverse transcription was performed using the Applied Biosystems High Capacity cDNA Reverse Transcription Kit (Applied Biosystems); cDNA concentrations were determined with a Nanodrop spectrophotometer (Nanodrop Technologies) prior to PCR. cDNA samples underwent absolute quantitative real-time PCR on the 9700HT instrument (Applied Biosystems) under the default 40-cycle program. TaqMan gene expression assays were purchased from Applied Biosystems, including LRRK1 and LRRK2 expression assays (assay Mm0481934 and Mm00713303), TBP (assay Mm00446973_m1), (Mm02342429_g1) Aand (4352932E). For each amplicon, PCR efficiency was estimated near 1.0 by serial dilutions of cDNA. Relative quantities of mRNA were estimated using the delta delta CT method.

List of abbreviations

leucine-rich repeat kinase 2 (LRRK2), leucine-rich repeat kinase 1 (LRRK1), Parkinson's Disease (PD), polymerase chain reaction (PCR), human embryonic kidney cells (HEK cells), pheochromocytoma cells (PC12 cells), phosphate buffered saline (PBS), sodium dodecyle sulfate (SDS), SDS polyacrylamide gel electrophoresis (SDS-PAGE), formic acid (FA), real time PCR (RT-PCR), embryonic day (E), postnatal day (P), glyceraldehyde 3-phosphate dehydrogenase (GAPDH), TATA binding protein (TBP), sodium chloride/sodium citrate (SSC), Radioimmunoprecipitation (RIPA), sodium chloride (NaCl), Nonidet P-40 (NP-40), human temporal lobe (Htl), wild-type (WT), knock-out (KO).

Authors' contributions

SB, ABW, AR and CC carried out all experiments, BLD generated the mouse *LRRK2* clone, DJM developed and purified the JH5514 antibody. TMD and VLD provided reagent and infrastructure support for the experiments and critically reviewed the manuscript. All authors read and approved the final manuscript.

Acknowledgements

Supported by grants from the NIH, NINDS NS 38377, the Lee Martin Trust, the Sylvia Nachlas Trust, the National Parkinson Foundation and the American Parkinson Disease Association. T.M.D. is the Leonard and Madlyn Abramson Professor in Neurodegenerative Diseases. S.B. is supported by the German Research Foundation (Deutsche Forschungsgemeinschaft). A.B.W. is supported by a NIH/NINDS R00 NS058111. We would like to thank Dr. Laura Marsh for human brain tissue used in this study.

References

- 1. Zimprich A, Biskup S, Leitner P, Lichtner P, Farrer M, Lincoln S, Kachergus J, Hulihan M, Uitti RJ, Calne DB, et al.: Mutations in LRRK2 cause autosomal-dominant parkinsonism with pleomorphic pathology. Neuron 2004, 44(4):601-607.
- Paisan-Ruiz C, Jain S, Evans EW, Gilks WP, Simon J, van der Brug M, Lopez de Munain A, Aparicio S, Gil AM, Khan N, et al.: Cloning of the gene containing mutations that cause PARK8-linked Parkinson's disease. Neuron 2004, 44(4):595-600.
- Hardy J, Cai H, Cookson MR, Gwinn-Hardy K, Singleton A: Genetics of Parkinson's disease and parkinsonism. Ann Neurol 2006, 60(4):389-398.
- Moore DJ, West AB, Dawson VL, Dawson TM: Molecular pathophysiology of Parkinson's disease. Annu Rev Neurosci 2005, 28:57-87.
- Gasser T: Genetics of Parkinson's disease. Curr Opin Neurol 2005, 18(4):363-369.
- Bonifati V: Genetics of Parkinson's disease. Minerva Med 2005, 96(3):175-186.
- 7. Di Fonzo A, Rohe CF, Ferreira J, Chien HF, Vacca L, Stocchi F, Guedes L, Fabrizio E, Manfredi M, Vanacore N, et al.: A frequent LRRK2 gene mutation associated with autosomal dominant Parkinson's disease. Lancet 2005, 365(9457):412-415.
- Gilks WP, Abou-Sleiman PM, Gandhi S, Jain S, Singleton A, Lees AJ, Shaw K, Bhatia KP, Bonifati V, Quinn NP, et al.: A common LRRK2 mutation in idiopathic Parkinson's disease. Lancet 2005, 365(9457):415-416.
- Kachergus J, Mata IF, Hulihan M, Taylor JP, Lincoln S, Aasly J, Gibson JM, Ross OA, Lynch T, Wiley J, et al.: Identification of a novel LRRK2 mutation linked to autosomal dominant parkinsonism: evidence of a common founder across European populations. Am J Hum Genet 2005, 76(4):672-680.
- Lesage S, Ibanez P, Lohmann E, Pollak P, Tison F, Tazir M, Leutenegger AL, Guimaraes J, Bonnet AM, Agid Y, et al.: G2019S LRRK2 mutation in French and North African families with Parkinson's disease. Ann Neurol 2005, 58(5):784-787.
- Ozelius LJ, Senthil G, Saunders-Pullman R, Ohmann E, Deligtisch A, Tagliati M, Hunt AL, Klein C, Henick B, Hailpern SM, et al.: LRRK2 G2019S as a cause of Parkinson's disease in Ashkenazi Jews. N Engl J Med 2006, 354(4):424-425.
- Goldwurm S, Zini M, Mariani L, Tesei S, Miceli R, Sironi F, Clementi M, Bonifati V, Pezzoli G: Evaluation of LRRK2 G2019S penetrance: relevance for genetic counseling in Parkinson disease. Neurology 2007, 68(14):1141-1143.
- Klein C, Lohmann-Hedrich K, Rogaeva E, Schlossmacher MG, Lang AE: Deciphering the role of heterozygous mutations in genes associated with parkinsonism. Lancet Neurol 2007, 6(7):652-662.
- Ross OA, Toft M, Whittle AJ, Johnson JL, Papapetropoulos S, Mash DC, Litvan I, Gordon MF, Wszolek ZK, Farrer MJ, et al.: Lrrk2 and Lewy body disease. Ann Neurol 2006, 59(2):388-393.
- Giordana MT, D'Agostino C, Albani G, Mauro A, Di Fonzo A, Antonini A, Bonifati V: Neuropathology of Parkinson's disease associated with the LRRK2 Ile1371Val mutation. Mov Disord 2007, 22(2):275-278.

- Simon-Sanchez J, Herranz-Perez V, Olucha-Bordonau F, Perez-Tur J: LRRK2 is expressed in areas affected by Parkinson's disease in the adult mouse brain. Eur J Neurosci 2006, 23(3):659-666.
- Biskup S, Moore DJ, Celsi F, Higashi S, West AB, Andrabi SA, Kurkinen K, Yu SW, Savitt JM, Waldvogel HJ, et al.: Localization of LRRK2 to membranous and vesicular structures in mammalian brain. Ann Neurol 2006, 60(5):557-569.
- Higashi S, Moore DJ, Colebrooke ŘE, Biskup S, Dawson VL, Arai H, Dawson TM, Emson PC: Expression and localization of Parkinson's disease-associated leucine-rich repeat kinase 2 in the mouse brain. J Neurochem 2007, 100(2):368-381.
- West AB, Moore DJ, Biskup S, Bugayenko A, Smith WW, Ross CA, Dawson VL, Dawson TM: Parkinson's disease-associated mutations in leucine-rich repeat kinase 2 augment kinase activity. Proc Natl Acad Sci USA 2005, 102(46):16842-16847.
- Gloeckner CJ, Kinkl N, Schumacher A, Braun RJ, O'Neill E, Meitinger T, Kolch W, Prokisch H, Ueffing M: The Parkinson disease causing LRRK2 mutation 12020T is associated with increased kinase activity. Hum Mol Genet 2006, 15(2):223-232.
- 21. Giasson Bl, Covy JP, Bonini NM, Hurtig Hl, Farrer MJ, Trojanowski JQ, Van Deerlin VM: Biochemical and pathological characterization of Lrrk2. Ann Neurol 2006, 59(2):315-322.
- Zhu X, Siedlak SL, Smith MA, Perry G, Chen SG: LRRK2 protein is a component of Lewy bodies. Ann Neurol 2006, 60(5):617-618. author reply 618–619
- Miklossy J, Arai T, Guo JP, Klegeris A, Yu S, McGeer EG, McGeer PL: LRRK2 expression in normal and pathologic human brain and in human cell lines. J Neuropathol Exp Neurol 2006, 65(10):953-963.
- MacLeod D, Dowman J, Hammond R, Leete T, Inoue K, Abeliovich A: The familial Parkinsonism gene LRRK2 regulates neurite process morphology. Neuron 2006, 52(4):587-593.
- Greggio E, Jain S, Kingsbury A, Bandopadhyay R, Lewis P, Kaganovich A, van der Brug MP, Beilina A, Blackinton J, Thomas KJ, et al.: Kinase activity is required for the toxic effects of mutant LRRK2/ dardarin. Neurobiol Dis 2006, 23(2):329-341.
- Greggio E, Lewis PA, van der Brug MP, Ahmad R, Kaganovich A, Ding J, Beilina A, Baker AK, Cookson MR: Mutations in LRRK2/dardarin associated with Parkinson disease are more toxic than equivalent mutations in the homologous kinase LRRK1. J Neurochem 2007, 102(1):93-102.
- Hatano T, Kubo S, Imai Ś, Maeda M, Ishikawa K, Mizuno Y, Hattori N: Leucine-rich repeat kinase 2 associates with lipid rafts. Hum Mol Genet 2007, 16(6):678-690.
- Melrose HL, Kent CB, Taylor JP, Dachsel JC, Hinkle KM, Lincoln SJ, Mok SS, Culvenor JG, Masters CL, Tyndall GM, et al.: A comparative analysis of leucine-rich repeat kinase 2 (Lrrk2) expression in mouse brain and Lewy body disease. Neuroscience 2007.
- Higashi S, Biskup S, West AB, Trinkaus D, Dawson VL, Faull RL, Waldvogel HJ, Arai H, Dawson TM, Moore DJ, et al.: Localization of Parkinson's disease-associated LRRK2 in normal and pathological human brain. Brain Res 2007, 1155:208-219.
- Galter D, Westerlund M, Carmine A, Lindqvist E, Sydow O, Olson L: LRRK2 expression linked to dopamine-innervated areas. Ann Neurol 2006, 59(4):714-719.
- Melrose H, Lincoln S, Tyndall G, Dickson D, Farrer M: Anatomical localization of leucine-rich repeat kinase 2 in mouse brain. Neuroscience 2006, 139(3):791-794.
- Taymans JM, Van den Haute C, Baekelandt V: Distribution of PINKI and LRRK2 in rat and mouse brain. J Neurochem 2006, 98(3):951-961.
- 33. Marin I: The Parkinson disease gene LRRK2: evolutionary and structural insights. *Mol Biol Evol* 2006, **23(12)**:2423-2433.
- Bosgraaf L, Van Haastert PJ: Roc, a Ras/GTPase domain in complex proteins. Biochim Biophys Acta 2003, 1643(1-3):5-10.
- 35. West AB, Moore DJ, Choi C, Andrabi SA, Li X, Dikeman D, Biskup S, Zhang Z, Lim KL, Dawson VL, et al.: Parkinson's disease-associated mutations in LRRK2 link enhanced GTP-binding and kinase activities to neuronal toxicity. Hum Mol Genet 2007, 16(2):223-232.

Publish with **Bio Med Central** and every scientist can read your work free of charge

"BioMed Central will be the most significant development for disseminating the results of biomedical research in our lifetime."

Sir Paul Nurse, Cancer Research UK

Your research papers will be:

- available free of charge to the entire biomedical community
- peer reviewed and published immediately upon acceptance
- cited in PubMed and archived on PubMed Central
- yours you keep the copyright

Submit your manuscript here: http://www.biomedcentral.com/info/publishing_adv.asp

