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RAB10: an Alzheimer's disease resilience locus and potential drug target

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Abstract: Alzheimer's disease (AD) is mainly a late-onset neurodegenerative disorder. Substantial efforts have been made to solve the complex genetic architecture of AD as a means to identify therapeutic targets. Unfortunately, to date, no disease-altering therapeutics have been developed. As therapeutics are likely to be most effective in the early stages of disease (ie, before the onset of symptoms), a recent focus of AD research has been the identification of protective factors that prevent disease. One example is the discovery of a rare variant in the 3'-UTR of *RAB10* that is protective for AD. Here, we review the possible genetic, molecular, and functional role of RAB10 in AD and potential therapeutic approaches to target RAB10.

Keywords: Alzheimer's disease, RAB10, retromer, APP, resilience, GTPase

Alzheimer's disease: overview

Alzheimer's disease (AD) is mainly a late-onset neurodegenerative disease defined clinically by progressive memory decline.¹ AD currently affects nearly 50 million people worldwide, and this number is expected to rapidly increase as the population ages (<http://www.alz.co.uk/>). Neuropathologically, AD is characterized by neuronal loss and the accumulation of amyloid plaques and neurofibrillary tangles in the brain, much of which occurs many years before the onset of clinical symptoms.² While clinical diagnosis of AD is challenging because it is dependent primarily on observations of family members and caregivers, changes in protein analytes in plasma and cerebrospinal fluid (CSF) and brain imaging, although not widely used in clinical practice, may facilitate AD diagnosis.² AD can broadly be defined either as early-onset, occurring prior to age 65, or the much more common late-onset, occurring after age 65 (>96.5% of total cases).

Genetic architecture of AD

The genetics of early-onset AD (EOAD) are relatively well understood. Autosomal dominant mutations occurring in *APP*, *PSEN1*, or *PSEN2* give rise to a subset of EOAD cases. More than 200 pathogenic mutations have been reported worldwide (<http://www.molgen.ua.ac.be/admutations/>).³ Clinically, EOAD mutation carriers exhibit disease onset prior to 65 years of age.⁴ Neuropathologically, EOAD and late-onset AD (LOAD) are indistinguishable.⁵ Unfortunately, the genetics of LOAD is more complex.

Among individuals with LOAD, there is strong evidence for genetic heritability of disease.⁶⁻¹⁰ The single largest genetic risk factor for LOAD is *APOE*.¹¹ APOE plays critical roles in cholesterol transport, neuroplasticity, and inflammation.¹² APOE regulates A β metabolism directly through binding and clearance of A β ¹³ and indirectly via the LRP1 receptor.¹⁴ Moreover, APOE influences tau-mediated neurodegeneration.¹⁵

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Two different *APOE* alleles, $\epsilon 4$ and $\epsilon 2$, affect risk for AD. Risk for developing AD is 3-fold higher in individuals carrying one copy of the $\epsilon 4$ allele and 12-fold higher in individuals carrying 2 copies of *APOE* $\epsilon 4$.¹⁶ Conversely, *APOE* $\epsilon 2$ confers resilience to AD, while the $\epsilon 3$ allele is considered neutral risk.¹⁶

Genetic variants that influence risk for AD can be broadly classified into two groups: common and rare. The majority of common variants (minor allele frequency [MAF] >5%) were identified using genome-wide association studies (GWAS).¹⁷ With the exception of variants in *APOE*, all other common variants have small effect sizes on AD risk (ORs = 1.08–1.22, 0.73–0.94).¹⁷ While GWAS variants provide insights into disease processes, none of the common risk variants for AD that have been identified by GWAS have clear functional effects and efforts to identify functional variants in the regions of GWAS variants have largely been unsuccessful.

The advent of next-generation sequencing has enabled the sequencing of whole exomes and genomes, resulting in progressively larger AD datasets and providing insights into the contribution of low-frequency (MAF 1%–5%) and rare variants (MAF <1%) to the genetic architecture of AD. Using a variety of study designs, multiple rare variants have been identified that affect risk for AD.^{18–22} Rare variants typically have much larger effect sizes on disease risk than common variants identified by GWAS. Rare variants are also more likely to occur in coding or regulatory regions and, thus, are more likely to be functional. Thus, rare variants represent effective therapeutic targets.

Finally, alternative study designs can provide additional insights into the genetics of AD. For example, the use of quantitative endophenotypes (eg, CSF levels of protein analytes such as A β and tau;^{23–25} mitochondrial copy number;²⁶ metabolic efficiency;^{27,28} measurements of progressive brain atrophy;²⁹ among others) and the analysis of mitochondrial genomic variation³⁰ have expanded our understanding of the genetic architecture underlying disease onset and disease course.

Using the genetics of AD to identify therapeutic targets

Genes and pathways that regulate AD pathogenesis may represent viable targets for treating disease. For example, therapies targeting the removal of A β or inhibition of A β production are currently in clinical trials. These strategies are based on our understanding of the mechanisms by which rare mutations in *APP*, *PSEN1*, and *PSEN2* drive AD pathogenesis in EOAD and focus on inhibition of the deposition

of A β in the brain. Similarly, targeting APOE function has been investigated for its therapeutic capacity.^{31–33} AD risk genes implicated in immune function, endocytosis, and lipid biology represent novel avenues of therapeutic development. Unfortunately, to date, more than 90% of all clinical trials have failed. Given the expected increase in disease incidence, developing effective therapeutics is imperative. For the remainder of this review, we discuss the potential role of *RAB10* as a therapeutic target.

Protective factors in AD: RAB10

Common and rare variants have been identified that reduce the risk for AD.³ Those with the strongest protective effect include *APOE* $\epsilon 2$ and *APP*-A673T.^{34,35} Thus, genes and pathways involved in increasing AD risk may also confer resilience to disease. To identify additional genetic variants that confer resilience to AD, there is a need to develop novel study designs that focus on those individuals carrying protective factors for AD. In one such case, family pedigrees were identified with a statistical excess of AD mortality (ie, families with a higher number of AD deaths than expected relative to similarly sized pedigrees; termed high-risk pedigrees). Within these high-risk pedigrees, elderly individuals with genetic risk for AD (ie, *APOE* $\epsilon 4$ carriers) but who had escaped AD were evaluated using whole-genome sequencing. Ridge et al identified a rare variant (rs142787485) in the 3'-UTR of *RAB10* that confers resilience to AD.¹⁰ In a cell model of *APP* metabolism, silencing of *Rab10* expression leads to a significant decrease in A β 42 and the A β 42/40 ratio, consistent with prior reports.^{10,36} Interestingly, independent of the *RAB10* SNP, individuals with a neuropathologic diagnosis of AD exhibited significantly higher levels of *RAB10* compared to neuropathology-free controls.¹⁰ Thus, genetic and molecular evidences support a role for *RAB10* in AD pathogenesis.

RAB10 in AD pathology

RAB10 is expressed in all cell types in the brain.³⁷ *RAB10* is activated by phosphorylation at Thr73, which is mediated by *LRRK2*, a protein kinase associated with Parkinson's disease.^{38,39} This phosphorylation event may represent a pathologic feature in the brains of AD patients.³⁸ pRAB10-Thr73 was observed to colocalize with neurofibrillary tangles in AD brains as well as with granulovascular degeneration, neuropil threads, and dystrophic neurons.³⁸ Double immunofluorescence staining of these structures further revealed positive staining for phosphorylated tau. However, this colocalization of pRAB10-Thr73 and phosphorylated tau was

not complete, with some instances showing neuropil threads or dystrophic neurons containing only phosphorylated tau or only pRAB10-Thr73. Interestingly, dystrophic neurons near amyloid plaques stained positive for pRAB10-Thr73, while the amyloid plaques themselves were negative for pRAB10.³⁸ Total RAB10 staining, however, did not differ between AD and control brains. Thus, while pRAB10-Thr73 is associated with AD pathology, it remains unclear whether this represents excess activation or aberrant function of RAB10.

RAB10: function in development

RAB10 is a member of the RAB family of small GTPases, which are key regulators of vesicular trafficking. RAB proteins are cyclically controlled, requiring a GDP–GTP exchange that is facilitated by RAB guanine nucleotide exchange factors.⁴⁰ However, unlike most of the other RAB proteins, which have relatively specific roles within intracellular transport, RAB10 is relatively unique because it carries out a wide variety of functions and has multiple subcellular localizations.⁴¹

Rab10 plays essential functional roles in development: mice are embryonic lethal when both alleles are deleted.⁴² In addition, RAB10 plays a role in maintenance and regulation of endoplasmic reticulum (ER) morphology.^{43–45} When RAB10 is mutated or constitutively inactive (eg, GDP-locked), the ER contains more cisternae, suggesting that RAB10 plays a critical role in maintaining and/or generating tubule extension and fusion in the ER.⁴³ RAB10 function has

also been linked to neuronal morphology and polarization, playing a critical role in axonal development and dendrite arborization.^{46,47} During neuronal development, RAB10 associates with plasmalemmal precursor vesicles, which are linked to kinesin 1 via c-Jun N-terminal protein kinase-interacting protein 1.⁴⁸ Together this complex mediates anterograde transport of RAB10-positive vesicles to axonal tips, promoting axonal growth.⁴⁸ Thus, RAB10 function is critical for proper neuronal function.

A role for RAB10 and APP in the retromer pathway

RAB10 has been linked to many roles involved in intracellular transport (Figure 1), including but not limited to a role in endocytic recycling. In *Caenorhabditis elegans*, RAB10 is required for α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid-type glutamate receptor recycling in postsynaptic membranes.⁴⁹ RAB10 is likely functional upstream of RME-1/EHD, a protein involved in recycling endosome tubulization and endosome function.^{50–52} RAB10 is expressed at the interface of early endosomes and recycling endosomes. These structures are morphologically defective in *rab-10* mutant *C. elegans* models.^{50,51} RAB10 is required for endosomal recruitment of CNT-1.⁵³ This likely contributes to biogenesis and/or maintenance of recycling endosome compartments. RAB10 could also play a role in promoting the maturation of early endosomes to recycling endosome. RAB10 is also involved in protein degradation.

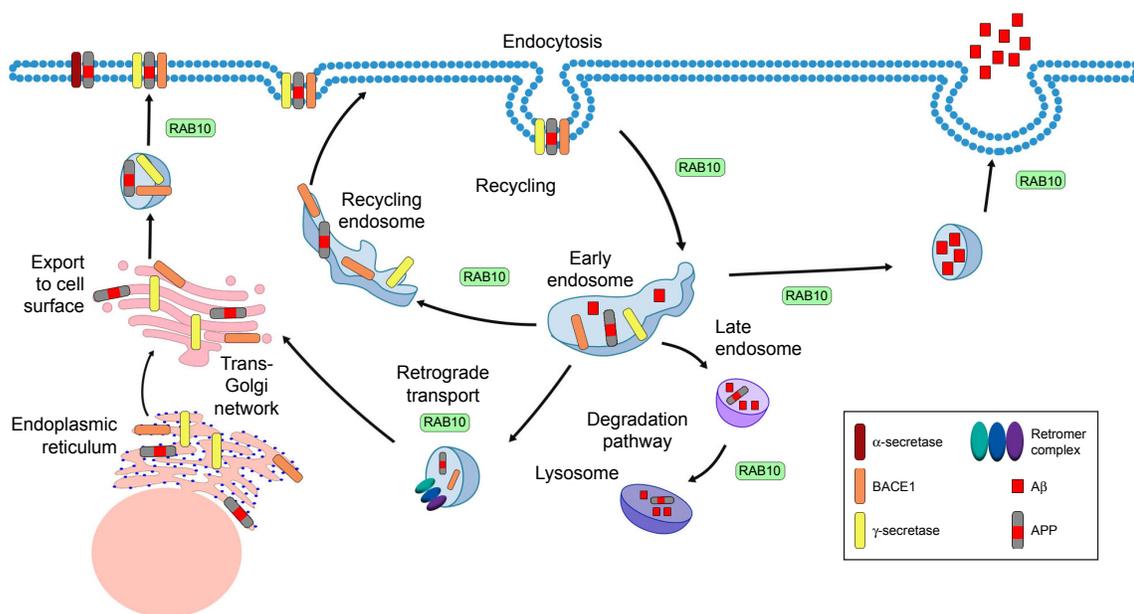


Figure 1 Diagram of APP trafficking and the role of RAB10 in these intracellular transport pathways.

RAB10 and RAB3A are essential for lysosomal exocytosis and plasma membrane repair.^{54,55} In addition, RAB10 associates with lipid droplets.⁵⁶ Complexes of RAB10–EHBP1–EHD2 promote autophagic engulfment and degradation of lipid droplets.⁵⁷ Thus, these functional roles for RAB10 in recycling and degradation make it an interesting candidate protein in AD pathogenesis.

RAB10 is involved in transport of proteins from the early endosome to the trans-Golgi network (TGN), where it may also function in the retromer pathway.⁵⁸ The retromer complex consists of Vps26, Vps35, Vps29, SNX1, SNX2, SNX5, and SNX6.⁵⁹ The retromer complex regulates transport of protein cargo from endosomes to the TGN by the retrograde pathway or to the cell surface through the recycling pathway.⁶⁰

To form the amyloid plaques that define a primary component of AD pathology, the transmembrane APP must first be processed into A β peptides. APP is thought to be retrieved from the cell surface and trafficked to the early endosomes where it is initially cleaved by β -site APP-cleaving enzyme 1 (BACE1) to generate a C-terminal fragment (β -CTF).^{61,62} The β -CTF fragments are then trafficked via retromer-dependent transport to the TGN, which facilitates further cleavage by γ -secretase to produce A β 40.⁶³ The retromer complex and its receptors can transport APP and BACE1 from endosomes to the TGN, ultimately regulating production of A β .⁶⁰ The retromer complex sorts APP and BACE1, either through direct or indirect binding to the transmembrane adaptor protein SORLA.⁶⁴ Interestingly, common variants in the gene that encodes SORLA have been associated with AD risk.⁶⁵ Thus, genes that are involved in retromer function may influence AD pathogenesis and represent promising therapeutic targets.

Retromer complex dysfunction promotes APP accumulation in neurons and A β production. In addition, the retromer also appears to have a differential effect on A β isoforms, with evidence that retromer dysfunction alters A β 40 secretion, which results in increasing the A β 42/40 ratio.⁶⁶ In mouse models, reduction of a single component within the retromer complex, Vps26, is sufficient to increase soluble A β and APP.⁶⁷ Stabilization of the retromer in human-induced pluripotent stem cell-derived neurons through the use of the pharmacologic chaperone R33 has been shown to significantly reduce the phosphorylation of Tau in a manner that correlated with, but ultimately independent of, A β production.⁶⁸ An RNAi screen of RABs that modify APP processing revealed that the silencing of *RAB10* reduces A β without altering sAPP β levels.³⁶ The LRRK2-RAB10 signaling pathway also causes an overproduction of A β .³⁸

Therapeutic implications for RAB10

Novel genes and pathways that contribute to the pathogenesis of AD provide insights into strategies to prevent or treat the disease. The amyloid hypothesis, which proposes that the accumulation of A β triggers the cascade of events that lead to AD, has been the focus of AD therapeutic development for decades. The development of anti-A β therapeutics remains the primary approach to treating AD based on our current understanding of the earliest features of this disease.⁶⁹ In this review, we emphasize the importance of RAB10 in the processing of APP and the production of A β . The function of RAB10 in intracellular trafficking and evidence that a reduction of *Rab10* results in a reduction of the A β 42/40 ratio makes inhibition of RAB10 a potential therapeutic avenue.

There are several existing strategies for therapeutic inhibition of a gene product. Potential technologies for targeting of RAB10 are small molecules, antibodies, and antisense oligodeoxynucleotide technologies. For each of these approaches the challenge of delivery across the blood–brain barrier (BBB) is a significant one.⁷⁰ A molecule with high lipid solubility and molecular mass <400 Da generally can diffuse across the BBB. These criteria limit the classes of small molecules that can be considered as potential therapeutic molecules for AD. Antibodies do not cross the BBB at therapeutic concentrations. Several approaches have been developed to deliver antibodies to the brain. Bispecific antibodies have been developed that leverage the transferrin receptor to successfully increase BBB penetration.^{71,72} In addition, antibody analogs that can cross the BBB are in development (Patent #EP3309171). Antisense oligonucleotides do not readily cross the BBB.⁷³ However, at present, intrathecal bolus injection is an effective method for ensuring distribution in neurons and glial cells in the brain.⁷⁴

Approaches to the development and delivery of small-molecule therapeutics are mature and the path and pitfalls are well established. Small molecules are effective enzyme inhibitors, receptor ligands, or allosteric modulators. The nature of small-molecule activity can make it difficult to find the balance between efficacy and side effects when treating humans.⁷⁰ Despite these challenges, small-molecule drug development has seen many successes and >90% of existing therapeutics use small molecules. Small GTPases similar to RAB10 have been successfully targeted using small molecules in the past. One important use is the development of molecules to block oncogenic properties of the RAS protein to treat cancer.⁷⁵ Recent discoveries about the structure of small GTPases and advances in targeting strategies have further increased efforts to use these important

proteins as therapeutic targets.⁷⁶ Targeting approaches, including interference with nucleotide binding, inactivation by irreversible covalent modification, inhibition of GTPase–GEF interactions, inhibition of GTPase–effector interactions, and stabilization of GTPase–protein complexes have been successful and are reviewed by Cromm et al.⁷⁶ While challenges regarding specificity and crossing the BBB remain, these advances create some optimism about the use of small molecules to inhibit RAB10 and impact AD pathology.

Antibodies function in extra- and intracellular immunity to recognize foreign or abnormal agents and target them for destruction with very high specificity. The production of monoclonal antibodies (mAbs) has resulted in their use for the treatment of a wide range of human diseases.⁷¹ There are currently >60 approved mAbs for human therapy and >50 in late-stage clinical trials.⁷⁷

RNase H-dependent antisense oligonucleotides technology is a popular method for knockdown in cell culture. It offers specific and efficient knockdown and is a powerful tool for functional studies of genes with unknown function.⁷⁸ Antisense technology has challenges such as site specificity, toxicity at high concentrations, and the difficulty of targeting specific cell types. However, it is a mature technology and antisense oligonucleotides offer a means to manipulate specific steps in mRNA processing, for example, splicing.⁶⁹ As discussed previously, in vitro reduction of *Rab10/RAB10* expression using shRNA and RNAi has been successful.³⁶ Expansion of antisense oligodeoxynucleotide from in vitro to in vivo represents a promising therapeutic avenue given recent successes with spinal muscular atrophy and Huntington's disease.^{79,80}

Conclusion

Here we have outlined the potential role of RAB10 in AD pathology and the case for RAB10 inhibition as a therapeutic intervention for AD. While small GTPases like RAB10 present significant challenges for therapeutic development, recent work in small molecules, antibodies, and antisense oligonucleotides suggests that success can be achieved.

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Disclosure

The authors report no conflicts of interest in this work.

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