ANKS1B

Disease/Syndrome Features:

In 2017, chromosome microarray analyses identified a ~370 kb deletion in the ANKS1B gene in individuals from a single family presenting with speech apraxia, motor dyspraxia, developmental delays, and autism spectrum disorders (ASD). This deletion was paternally inherited and present in 3 out of 4 children [1]. Since then, 11 additional individuals in the USA, Europe, and the Middle East displaying speech and motor deficits, autism, and developmental delays were found to have related deletions [1]. The ANKS1B gene is only found in vertebrates, and with highest homology orthologs present in mammals. The gene spans ~1.2 mb across chromosome 12 band 23.1 (12q23.1) and contains approximately 30 exons, with the exact number depending on the genome repository. Deletions identified in probands are roughly 100 to 400 kb in size and result in the loss of 3-7 exons in the 5' region of ANKS1B. No deletions in the 3' region have been identified, and all probands identified to date are heterozygous. Normal transcription of ANKS1B results in multiple variants that can be broadly classified as large ANKS1B variants containing the 5' exons, or shorter variants lacking the exons deleted in probands [2]. This suggests that the deletions identified in humans so far would only affect expression of larger ANKS1B transcripts.

Almost all subjects with *ANKS1B* deletions show speech impairments, including delayed achievement of developmental milestones for speech. In some cases, a formal diagnosis of speech apraxia or developmental articulation disorder has been made. Motor phenotypes are also observed, including delays in developing fine motor and/or gross motor skills. Developmental coordination disorder and Tourette's are among the motor disorders diagnosed. Several subjects have also been diagnosed with autism spectrum disorder. Cognitive impairment is a variable finding, with subjects ranging from normal IQ to intellectual disability. On neuroimaging, three subjects show thinning of the corpus callosum by anatomical MRI, two show enlargement of brain ventricles. In most cases, heterozygous deletions were inherited from a parent with a normal to mild phenotype.

Protein/Pathway:

ANKS1B encodes the protein AIDA-1 (amyloid precursor protein intracellular domain associated-1 protein). AIDA-1 is an adaptor protein implicated in synaptic and nuclear function¹ that is enriched in the cerebral cortex, hippocampus, and cerebellum². AIDA-1 is principally expressed in neurons where it is one of the most abundant proteins at postsynaptic densities (PSDs)[3-6]. At the PSD it binds to NMDARs and the adaptor protein PSD95 [3] through the first two PDZ domains on PSD95. Synaptic activity causes AIDA-1 to shuttle into the nucleus and regulate Cajal body stability and nucleolar morphology [3,7]. Recent work shows that AIDA-1 is required for the transport of GluN2B subunits of NMDARs into hippocampal synapses [8]. Loss of AIDA-1 leads to GluN2B accumulation in the endoplasmic reticulum (ER), reduces levels of GluN2B at synapses, decreases GluN2B-mediated currents, and impairs NMDAR-mediated long-term potentiation (LTP) and long-term depression (LTD) in the hippocampus [8]. A growing number of studies implicate *ANKS1B* in disease. As with NMDARs and their

associated proteins [9-11], recent studies link *ANKS1B* to neuropsychiatric disorders including schizophrenia [12-15], and autism spectrum disorders (ASD) [16-19]. Analysis of the AIDA-1 interactome using quantitative proteomic techniques has uncovered roles in protein networks involved in synaptic function and the etiology of neurodevelopmental disorders [1], which provides potential causal links between disruption of this gene and the associated disorders.

Mice lacking ANKS1B in the nervous system (using a Nestin-Cre driver) have been developed, but only heterozygotes are viable. Homozygotes die late in embryogenesis or perinatally for unknown reasons, which may explain the lack of probands identified with homozygous deletions. Mice with homozygous deletion specifically in neurons of the forebrain (using a CaMKIIa-cre driver) are viable and show impaired synaptic function and reduced NMDA-dependent plasticity in the hippocampus [8]. Synapses in stratum radiatum of mutant mice show increased conductance of the NMDA receptor subunit GluN2A with a corresponding decrease in GluN2B-mediated currents. In the absence of AIDA-1, GluN2B subunits become enriched in the endoplasmic reticulum, suggesting AIDA-1 regulates the intracellular transport or maturation of NMDA receptor subunits [8]. Heterozygous mice display behavioral correlates of neurodevelopmental disorders. Mutant mice display deficits in sociability as measured using the three chambered sociability test, mild hyperactivity, and show deficits both in acoustic startle and pre-pulse inhibition [1]. Tests of fine motor skills using tape removal show deficits in the absence of AIDA-1. Results suggest heterozygotes may represent a viable model of a novel ANKS1B-related neurodevelopmental syndrome.

REFERENCES:

[1] Carbonell, A.U., Cho, C.H., Tindi, J.O., Counts, P.A., Bates, J.C., Erdjument-Bromage, H., Cvejic, S., Iaboni, A., Kvint, I., Rosensaft, J., Banne, E., Anagnostou, E., Neubert, T.A., Scherer, S.W., Molholm, S., Jordan, B.A. Haploinsufficiency in the ANKS1B Gene Encoding AIDA-1 Leads to a Neurodevelopmental Syndrome *Nat Commun* 10(1) doi: 10.1038/s41467-019-11437-2 (2019)

[2] Ghersi, E., Vito, P., Lopez, P., Abdallah, M., D'Adamio, L. The intracellular localization of amyloid beta protein precursor (AbetaPP) intracellular domain associated protein-1 (AIDA-1) is regulated by AbetaPP and alternative splicing. *J Alzheimers Dis* **6**, 67-68, doi: 10.3233/jad-2004-6108 (2004)

[3] Jordan, B.A., Fernholz, B.D., Khatri, L. & Ziff, E.B. Activity-dependent AIDA-1 nuclear signaling regulates nucleolar numbers and protein synthesis in neurons. *Nat Neurosci* **10**, 427-435, doi: 10.1038/nn1867 (2007)

[4] Jacob, A. L., Jordan, B. A. & Weinberg, R. J. Organization of amyloid-beta protein precursor intracellular domain-associated protein-1 in the rat brain. *J Comp Neurol* **518**, 3221-3236, doi:10.1002/cne.22394 (2010).PMC2894292

[5] Dosemeci, A. *et al.* Composition of the synaptic PSD-95 complex. *Mol Cell Proteomics* **6**, 1749-1760, doi:10.1074/mcp.M700040-MCP200 (2007).PMC2096750

[6] Jordan, B. A. *et al.* Identification and verification of novel rodent postsynaptic density proteins. *Mol Cell Proteomics* **3**, 857-871, doi:10.1074/mcp.M400045-MCP200 (2004)

[7] Xu, H. & Hebert, M. D. A novel EB-1/AIDA-1 isoform, AIDA-1c, interacts with the Cajal body protein coilin. *BMC Cell Biol* **6**, 23, doi:10.1186/1471-2121-6-23 (2005).PMC1097723

[8] Tindi, J. O. *et al.* ANKS1B Gene Product AIDA-1 Controls Hippocampal Synaptic Transmission by Regulating GluN2B Subunit Localization. *J Neurosci* **35**, 8986-8996, doi:10.1523/JNEUROSCI.4029-14.2015 (2015).PMC4469732

[9] Lau, C. G. & Zukin, R. S. NMDA receptor trafficking in synaptic plasticity and neuropsychiatric disorders. *Nat Rev Neurosci* **8**, 413-426, doi:10.1038/nrn2153 (2007)

[10] Paoletti, P., Bellone, C. & Zhou, Q. NMDA receptor subunit diversity: impact on receptor properties, synaptic plasticity and disease. *Nat Rev Neurosci* **14**, 383-400, doi:10.1038/nrn3504 (2013)

[11] Tarabeux, J. *et al.* Rare mutations in N-methyl-D-aspartate glutamate receptors in autism spectrum disorders and schizophrenia. *Transl Psychiatry* **1**, e55, doi:10.1038/tp.2011.52 (2011).PMC3309470

[12] Fromer, M. *et al.* De novo mutations in schizophrenia implicate synaptic networks. *Nature* **506**, 179-184, doi:10.1038/nature12929 (2014).PMC4237002

[13] McClay, J. L. *et al.* Genome-wide pharmacogenomic study of neurocognition as an indicator of antipsychotic treatment response in schizophrenia. *Neuropsychopharmacology* **36**, 616-626, doi:10.1038/npp.2010.193 (2011).PMC3055694

[14] McClay, J. L. *et al.* Genome-wide pharmacogenomic analysis of response to treatment with antipsychotics. *Mol Psychiatry* **16**, 76-85, doi:10.1038/mp.2009.89 (2011).PMC2888895

[15] Purcell, S. M. *et al.* A polygenic burden of rare disruptive mutations in schizophrenia. *Nature* **506**, 185-190, doi:10.1038/nature12975 (2014).PMC4136494

[16] Hu, V. W., Addington, A. & Hyman, A. Novel autism subtype-dependent genetic variants are revealed by quantitative trait and subphenotype association analyses of published GWAS data. *PLoS One* **6**, e19067, doi:10.1371/journal.pone.0019067 (2011).PMC3083416

[17] Meehan, T. F. *et al.* Autism candidate genes via mouse phenomics. *J Biomed Inform* **44 Suppl 1**, S5-11, doi:10.1016/j.jbi.2011.03.003 (2011).PMC3263820

[18] Pinto, D. *et al.* Functional impact of global rare copy number variation in autism spectrum disorders. *Nature* **466**, 368-372, doi:10.1038/nature09146 (2010).PMC3021798

[19] Uddin, M. *et al.* Brain-expressed exons under purifying selection are enriched for de novo mutations in autism spectrum disorder. *Nat Genet* **46**, 742-747, doi:10.1038/ng.2980 (2014)

Support Groups and Information:

Anks1b Deletion (private Facebook group)

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