

Mechanisms of Vitamin E Deficient Neurodegeneration in Small and Large Animal Models



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Overview

1. Vitamin E : Role in neurologic disease
2. Animal models of Ataxia with VitE Deficiency (AVED)
 - Equine neuroaxonal dystrophy (eNAD)
 - *Ttpa*^{-/-} mouse
3. Transcriptional profiling of CNS tissue
4. Functional electrophysiological evaluation
5. Proposed mechanisms



Vitamin E (VitE)

- Major lipid soluble antioxidant
- 8 forms

Tocopherol (saturated)	Tocotrienol (unsaturated)
α -tocopherol	α -tocotrienol
β -tocopherol	β -tocotrienol
γ -tocopherol	γ -tocotrienol
δ -tocopherol	δ -tocotrienol

- Common absorption and delivery pathways as cholesterol
- Critical in maintaining neurological health?



VitE Deficiency

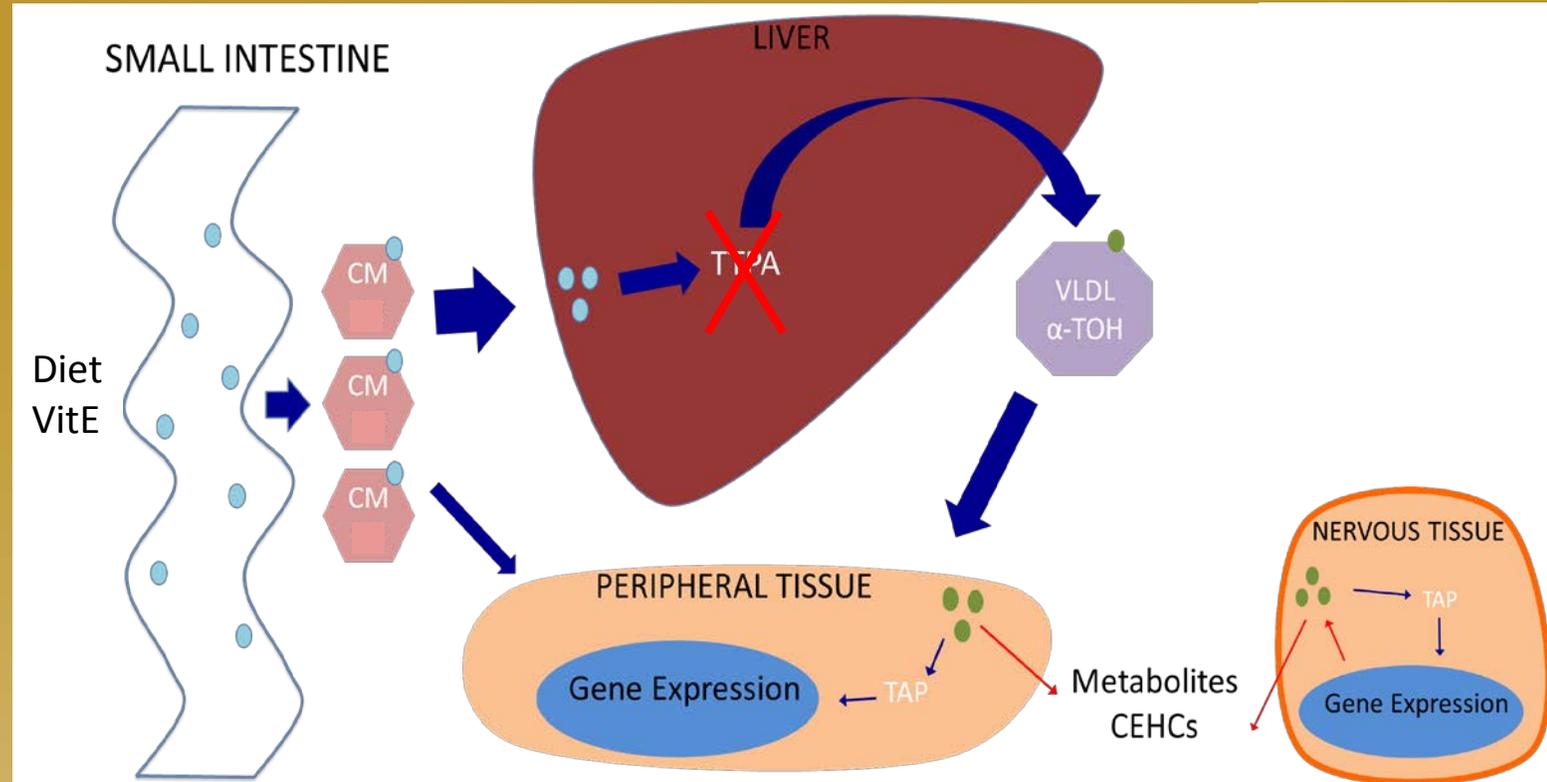
- CNS + PNS + retina + skeletal muscle
 - Areflexia
 - Cerebellar ataxia
 - Distal loss of proprioception*
 - Pigmentary retinopathy
 - Generalized muscle weakness





VitE Deficiency

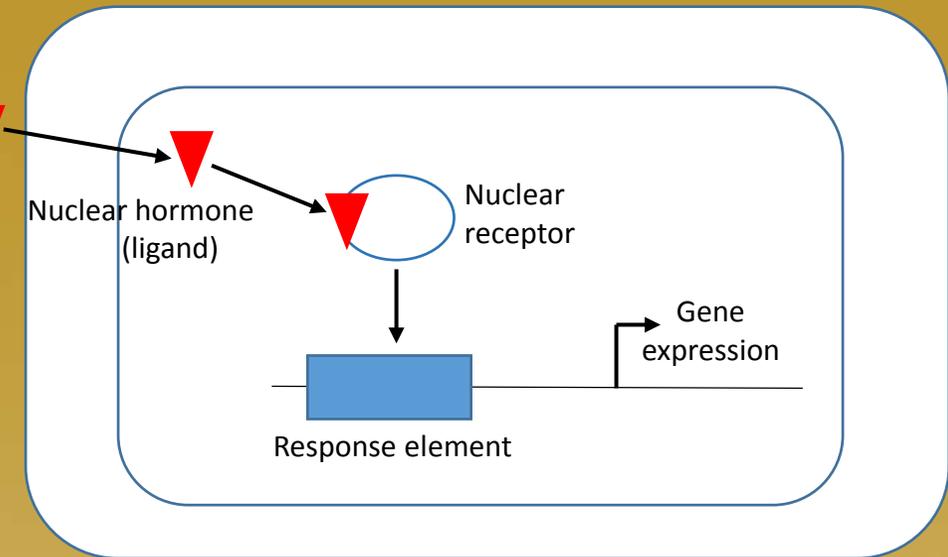
1. Abetalipoproteinemia
2. Other fat malabsorptive conditions
3. Ataxia with VitE Deficiency (AVED)
 - Low serum, CSF, tissue [α -TOH]
 - Rescued with vitE supplementation





GAPS in knowledge

1. What is the origin of the lesion?
2. Is vitE acting as an antioxidant OR modifying gene transcription?
3. How does vitE deficiency lead to sensory deprivation?

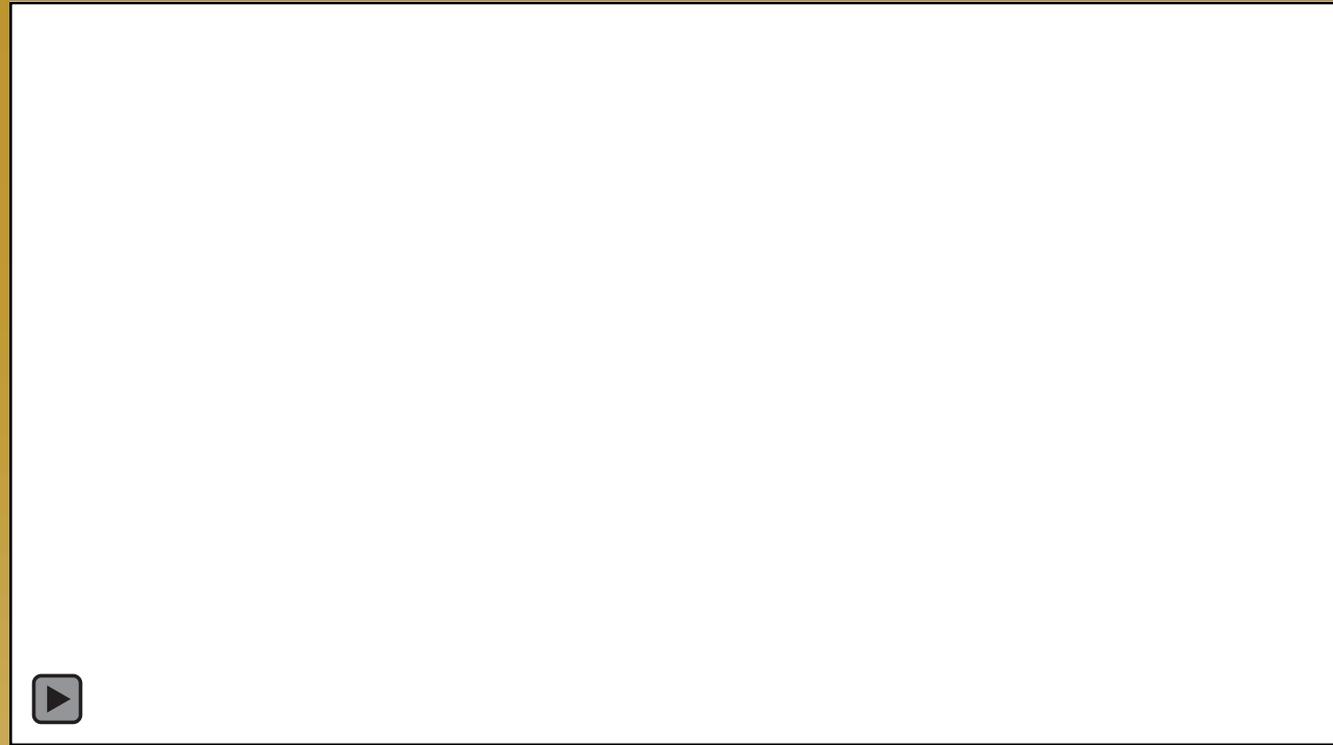




Equine Neuroaxonal Dystrophy (eNAD)



Normal Horse



Equine AVED



eNAD



J Vet Intern Med 2011;25:1439–1446

Equine Degenerative Myeloencephalopathy in Lusitano Horses

C.J. Finno, R.J. Higgins, M. Aleman, R. Ofri, S.R. Hollingsworth, D.L. Bannasch, C.M. Reilly, and J.E. Madigan

Veterinary Ophthalmology (2012) 15, Supplement 2, 3–7

DOI:10.1111/j.1463-5224.2012.00997.x

BRIEF COMMUNICATION

Electrophysiological studies in American Quarter horses with neuroaxonal dystrophy

Carrie J. Finno,* Monica Aleman,† Ron Ofri,‡ Steven R. Hollingsworth,§ John E. Madigan,** Laramie Winfield,† Danika L. Bannasch,*

J Vet Intern Med 2013;27:177–185

Pedigree Analysis and Exclusion of Alpha-Tocopherol Transfer Protein (*TTPA*) as a Candidate Gene for Neuroaxonal Dystrophy in the American Quarter Horse

C.J. Finno, T. Famula, M. Aleman, R.J. Higgins, J.E. Madigan, and D.L. Bannasch

The Veterinary Journal 202 (2014) 543–549

Contents lists available at ScienceDirect



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The Veterinary Journal

journal homepage: www.elsevier.com/locate/tvjl



J Vet Intern Med 2015;29:1667–1675

Blood and Cerebrospinal Fluid α -Tocopherol and Selenium Concentrations in Neonatal Foals with Neuroaxonal Dystrophy

C.J. Finno, K.E. Estell, S. Katzman, L. Winfield, A. Rendahl, J. Textor, D.L. Bannasch, and B. Puschner

Risk of false positive genetic associations in complex traits with underlying population structure: A case study

Carrie J. Finno ^{a,*}, Monica Aleman ^b, Robert J. Higgins ^c, John E. Madigan ^b, Danika L. Bannasch ^a





eNAD

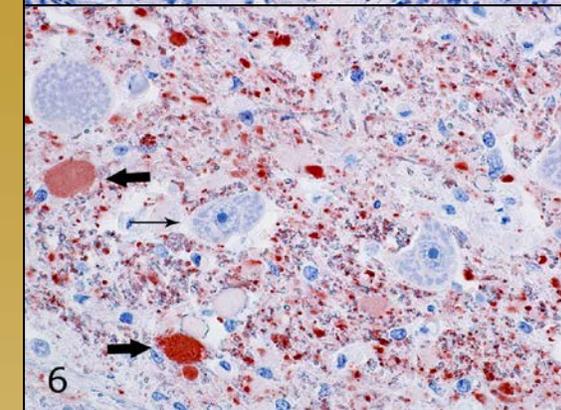
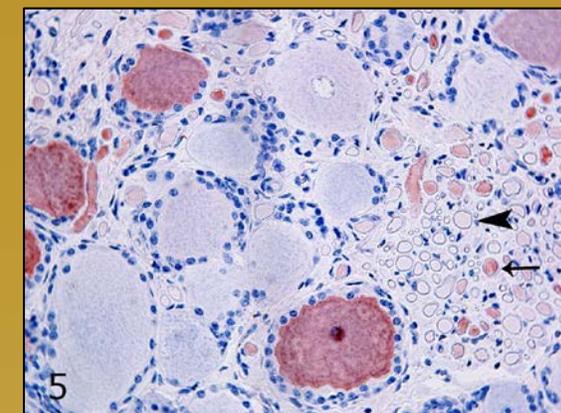
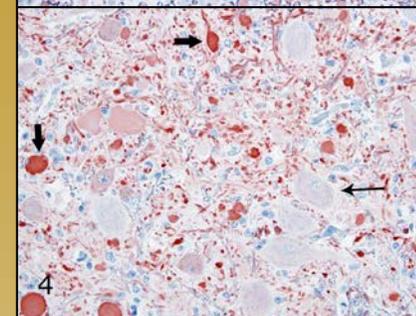
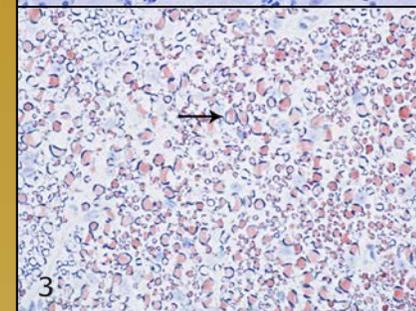
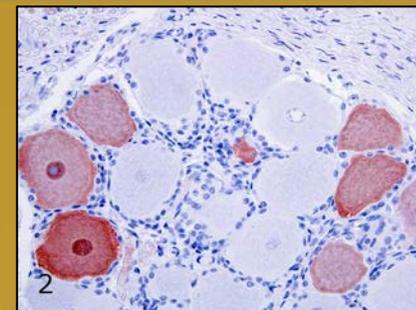


Degenerative and Immune-mediated Disease—Original Article

Evidence of the Primary Afferent Tracts Undergoing Neurodegeneration in Horses With Equine Degenerative Myeloencephalopathy Based on Calretinin Immunohistochemical Localization

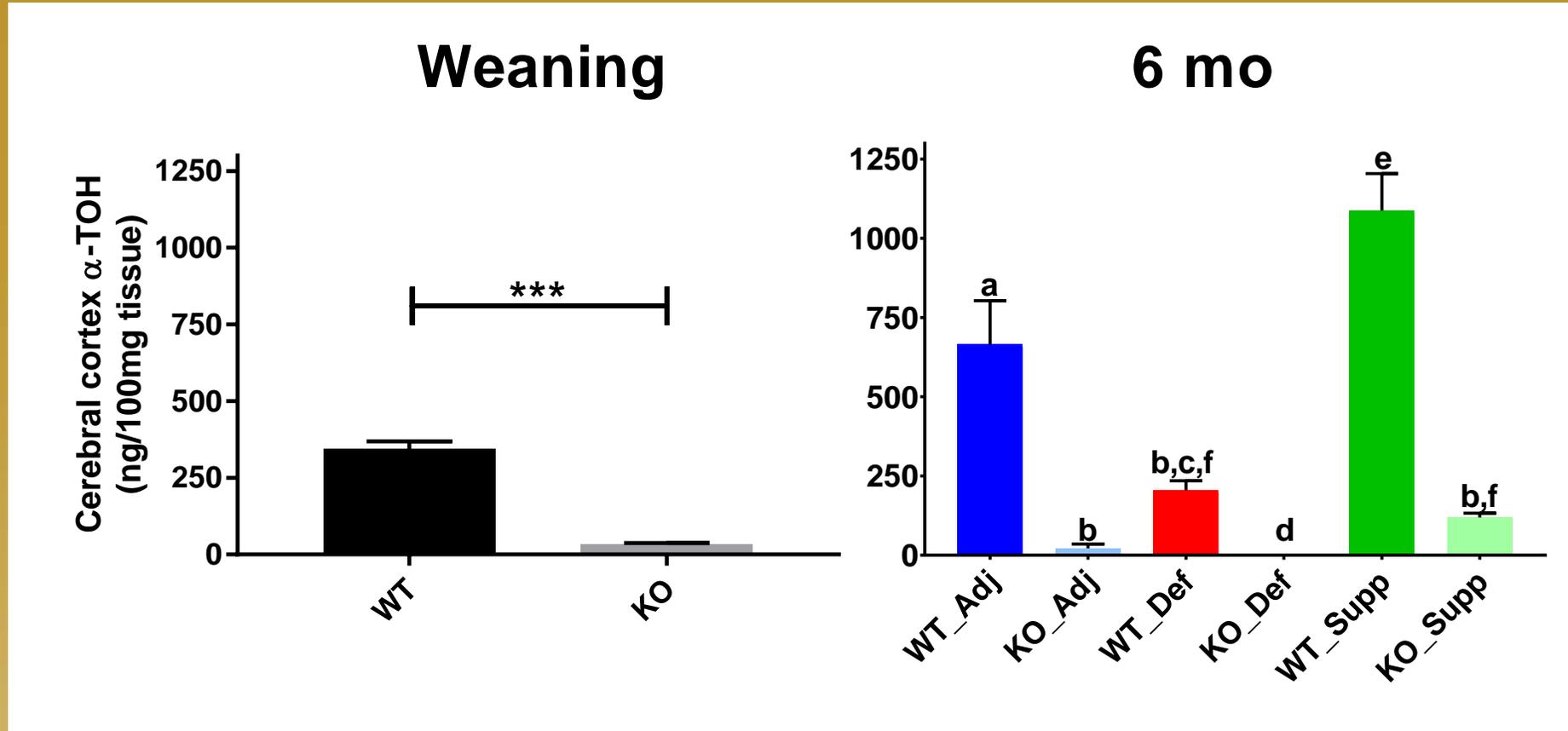
C. J. Finno¹, S. J. Valberg¹, J. Shivers², E. D'Almeida², and A. G. Armien^{1,2}

Veterinary Pathology
2016, Vol. 53(1) 77-86
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DOI: 10.1177/0300985815598787
vet.sagepub.com





Ttpa^{-/-} mouse model

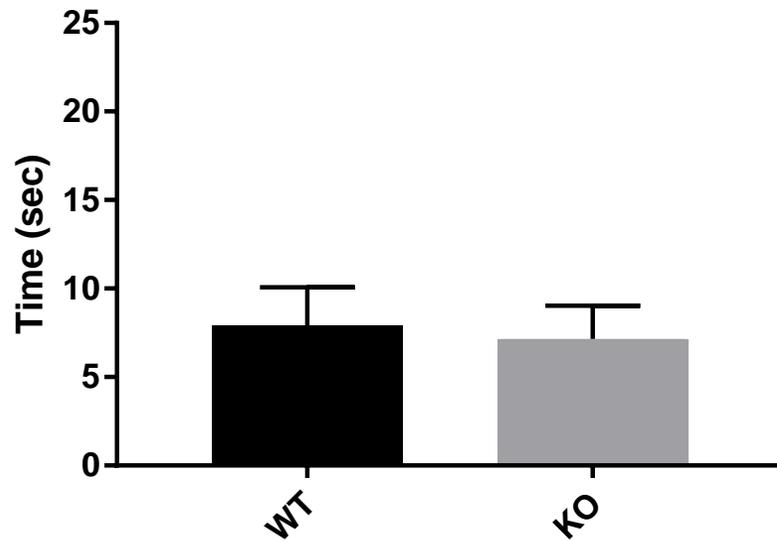




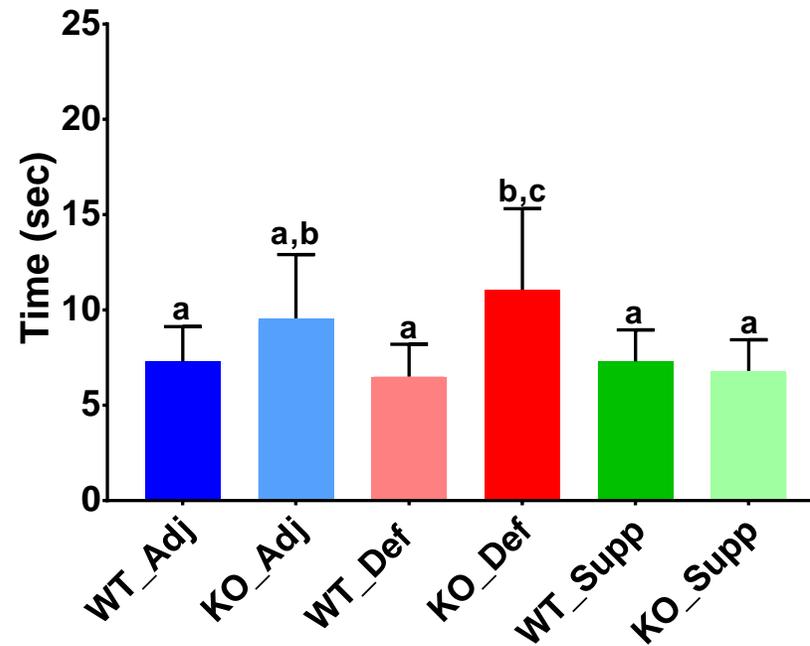
Ttpa^{-/-} mouse model



Weaning



6 mo

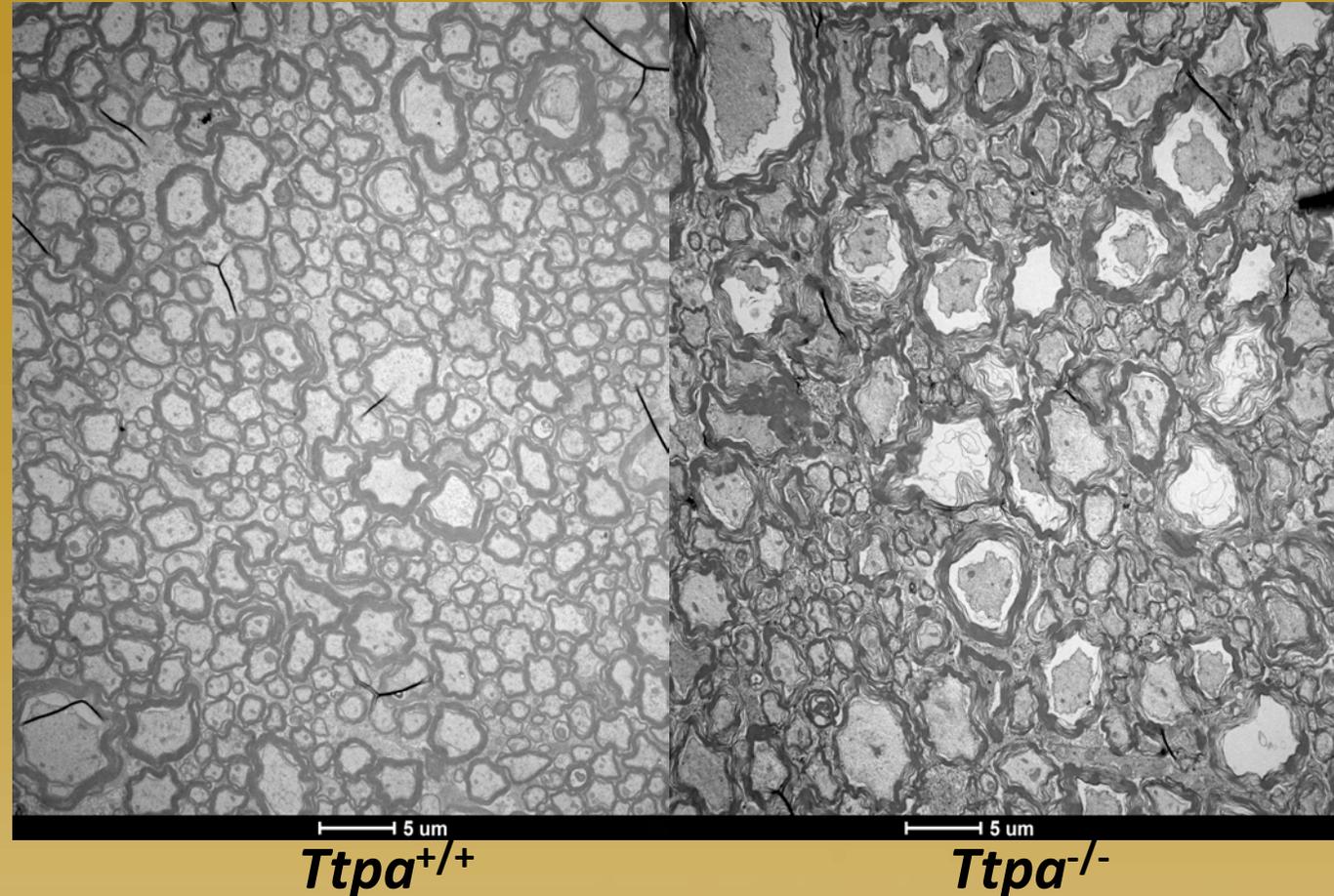




Ttpa^{-/-} mouse model



- Mimics phenotype of AVED
 - 6 months of age
 - Overt proprioceptive deficits
 - 6-12 months: histologic changes
 - Dorsal column degeneration: medulla oblongata and spinal cord
 - 17-20 months
 - Purkinje cell loss
- Rescued with early α -TOH supplementation





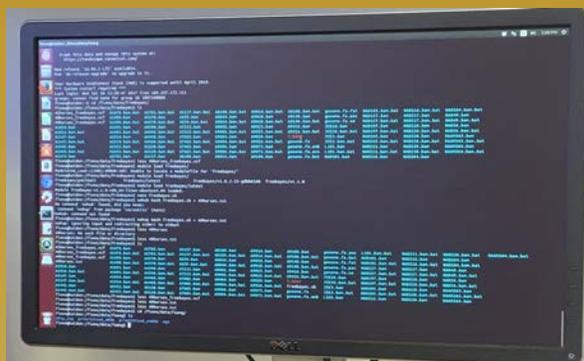
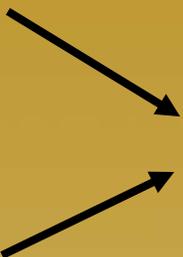
Hypothesis

Transcriptional profiling in eNAD and *Ttpa*^{-/-} mice will demonstrate similar dysregulated pathways associated with α -TOH deficiency





Methods



Identify dysregulated genes and pathways

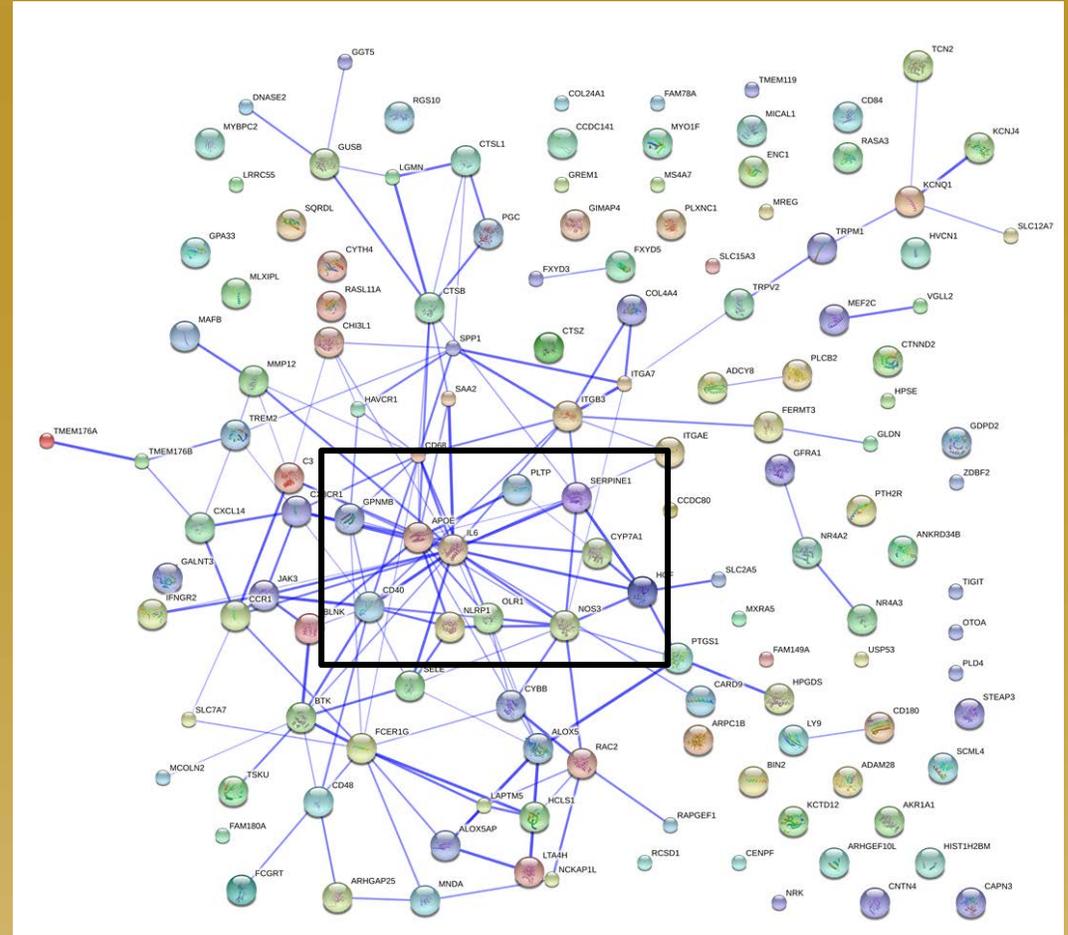


Test protein function



Results: Equine Spinal Cord

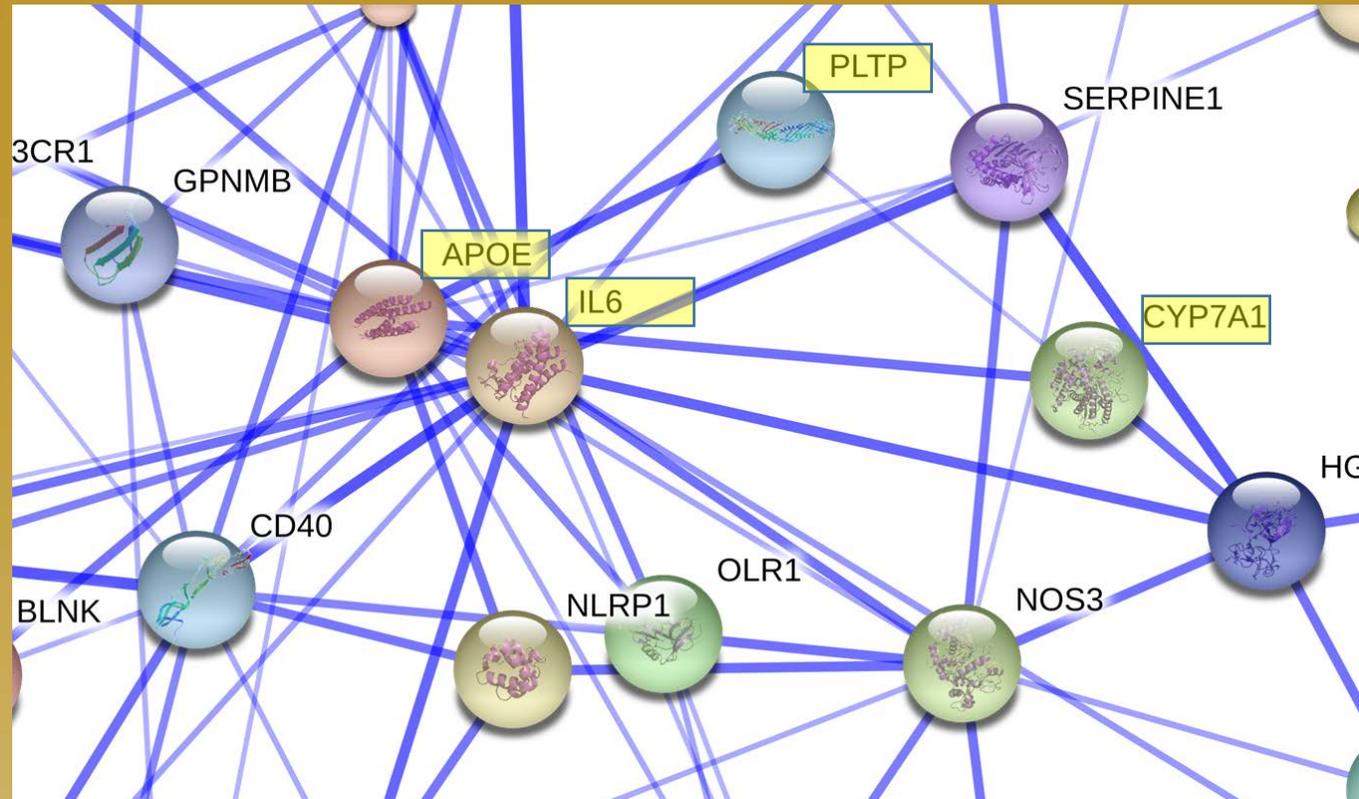
- 157 DETs (FDR=0.05)
 - 19 downregulated
 - 138 upregulated



Upregulated DETs



Results: Equine



Liver X Receptor Targeted Genes



Liver X Receptor (LXR)

- Nuclear receptor : reverse cholesterol transport
 - Cellular response to oxysterols = LXR activation and decrease cholesterol biosynthesis
- Upregulated oxidized lipoprotein receptor (*OLR1*)

Pathway	Up or down-regulated	$P_{\text{Bonferroni}}$
Ionotropic glutamate receptor pathway	-	7.7×10^{-5}
Synaptic vesicle trafficking	-	0.002
Metabotropic glutamate receptor group III pathway	-	0.009
Cholesterol biosynthesis	-	0.02

Free Radical Biology and Medicine 101 (2016) 261–271

Contents lists available at ScienceDirect

Free Radical Biology and Medicine

journal homepage: www.elsevier.com/locate/freeradbiomed

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FREE RADICAL BIOLOGY & MEDICINE

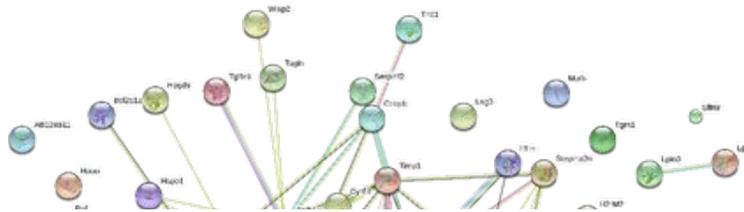
Transcriptome profiling of equine vitamin E deficient neuroaxonal dystrophy identifies upregulation of liver X receptor target genes

Carrie J. Finno^{a,*}, Matthew H. Bordbari^a, Stephanie J. Valberg^b, David Lee^c, Josi Herron^c, Kelly Hines^c, Tamer Monsour^d, Erica Scott^b, Danika L. Bannasch^a, James Mickelson^d, Libin Xu^c

CrossMark

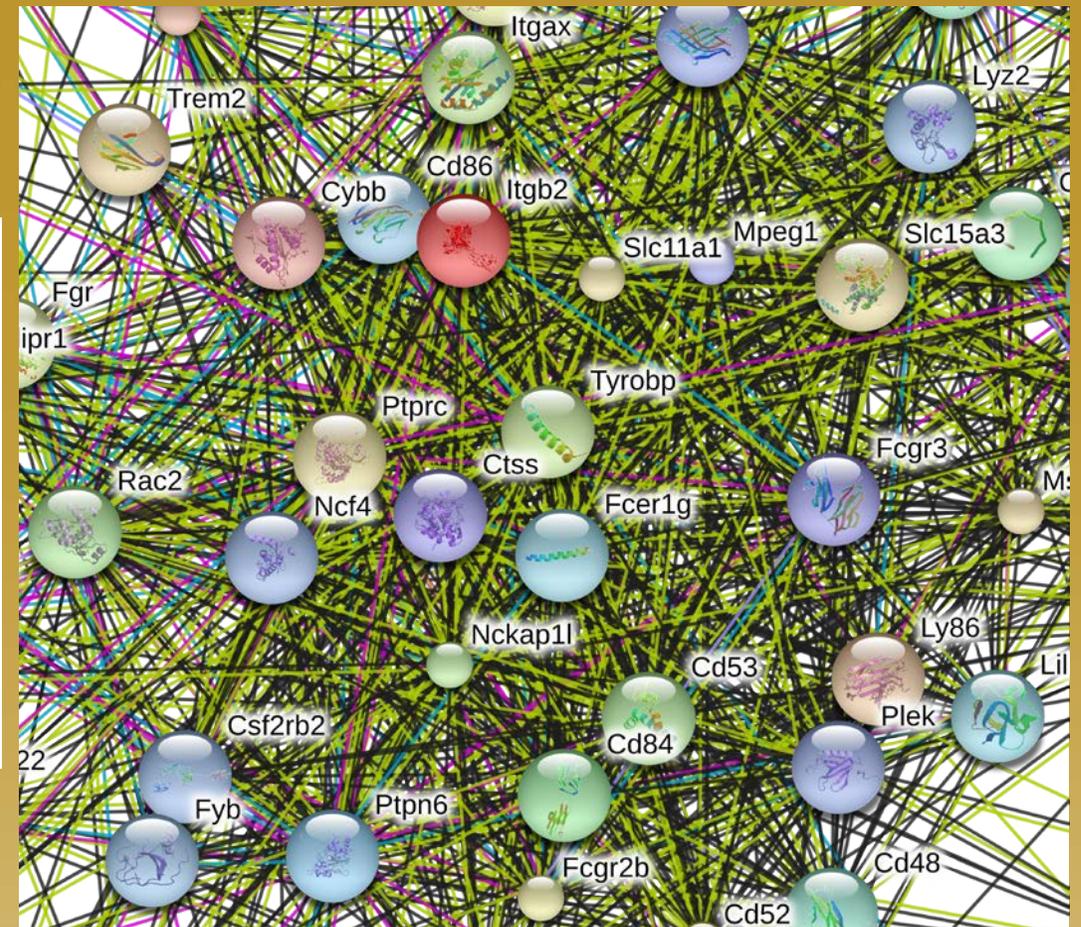
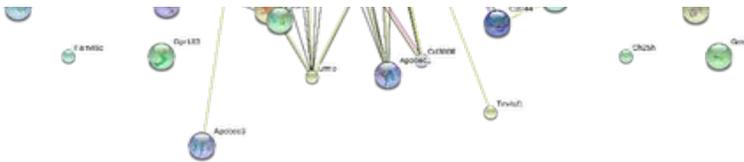


Results: 6 mo VitE- : KO vs WT



21 pathways dysregulated including:

- Innate immune response
 - (*Trem2, Cybb, Cd86, Slc11a1, Tyrobp, Fcer1g*)
- Inflammatory response
 - (*Itgb2, Ctss, Ccl3, Clec7a*)
- G-protein coupled receptor signaling
 - (*Rac2, Plek, Rgs1*)

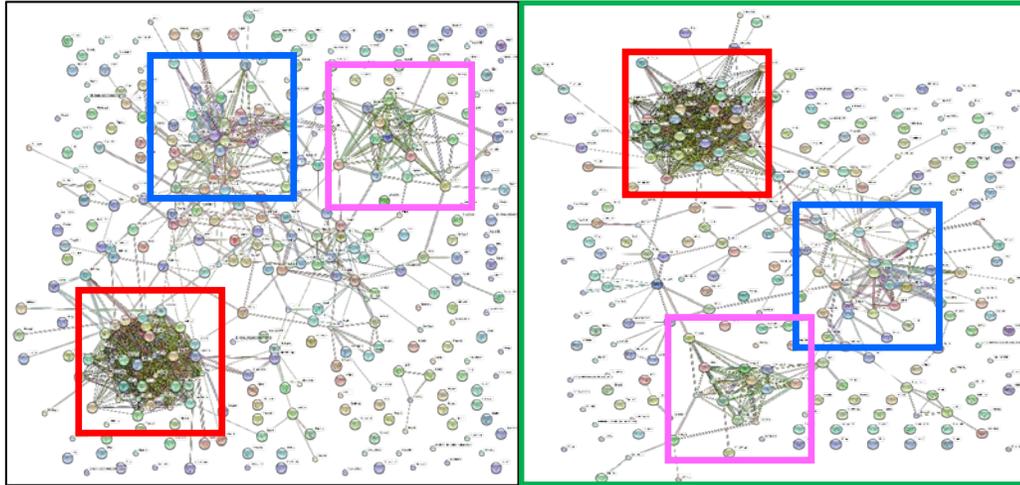




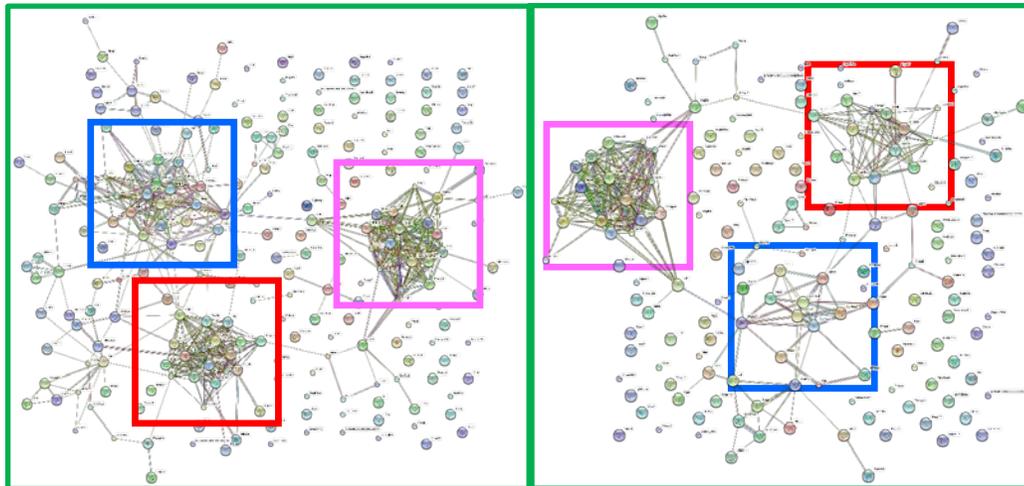
WT

vitE+

vitE-



KO



DOWNREGULATED Weaning → 6 mo

Cell cycle / Mitosis
(Aurkb, Ube2c, Cenpa)

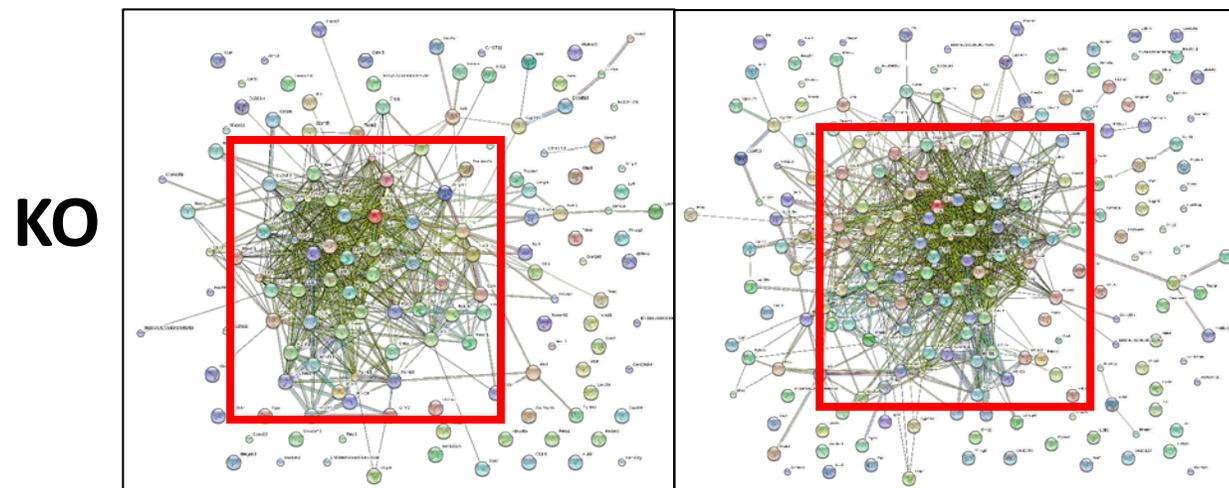
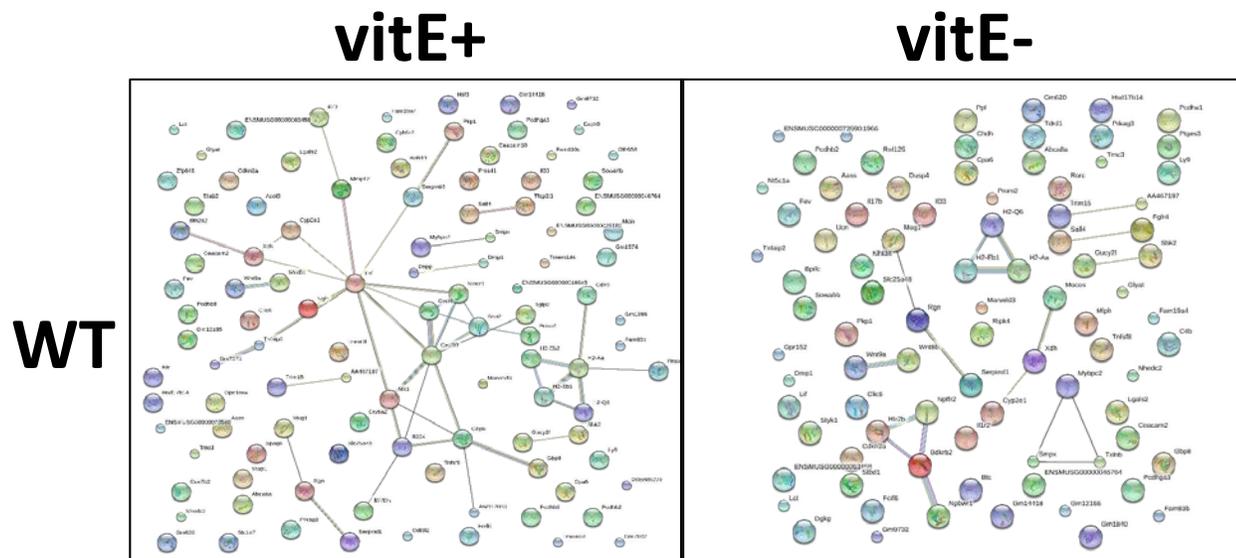
Collagen *(Col3a1, Col1a1, Col9a3)*

Cholesterol Biosynthesis
(Sqle, Dhcr7, Mvk)

Myelination
(Cntnap1, Mbp, Plp1)



UPREGULATED
Weaning → 6 mo



Inflammatory Response
(Itgb2, Ctss, Ccl3, Clec7a)



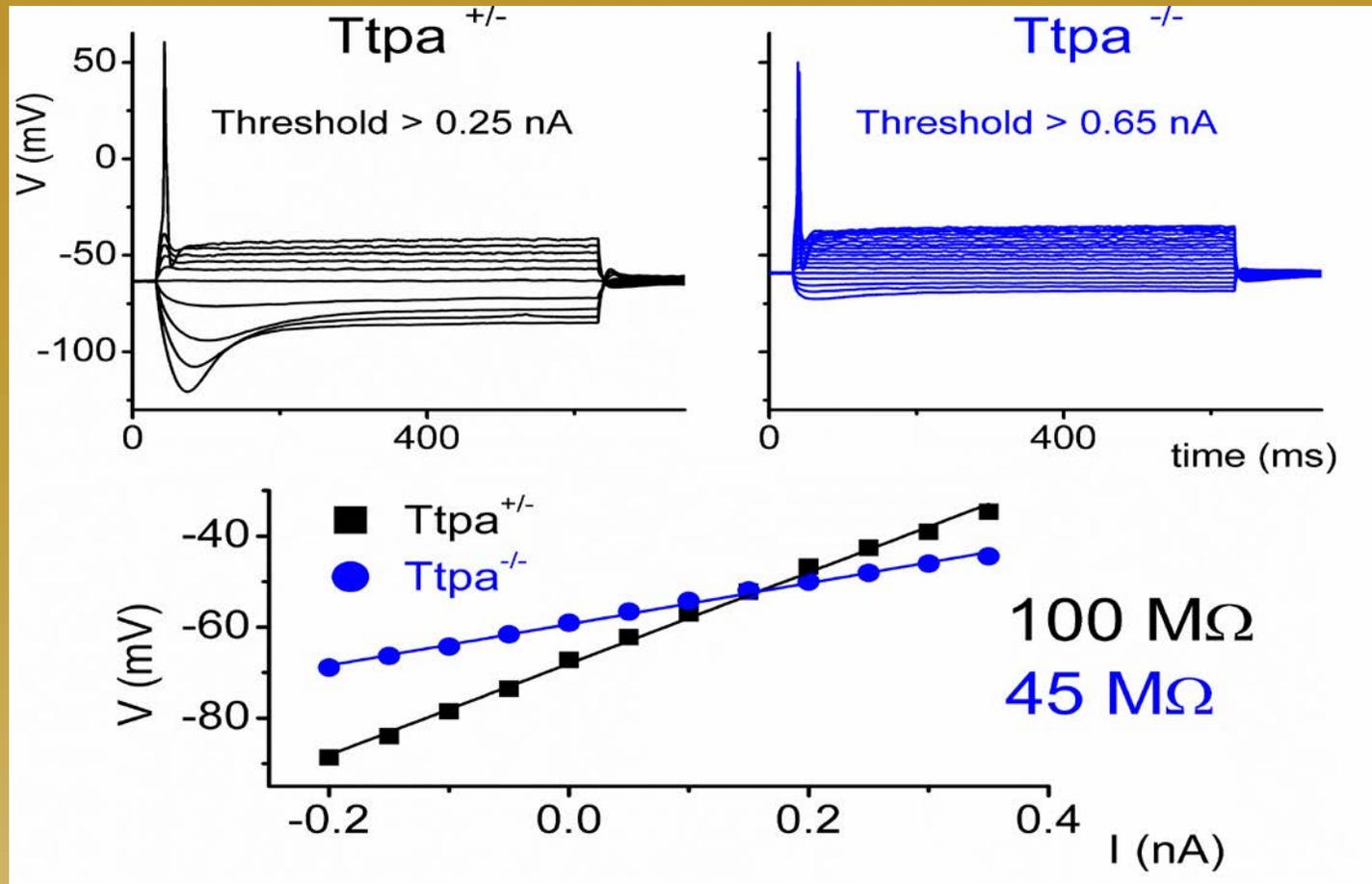
VitE : Nuclear Receptor Activation (Wean → 6 mo)



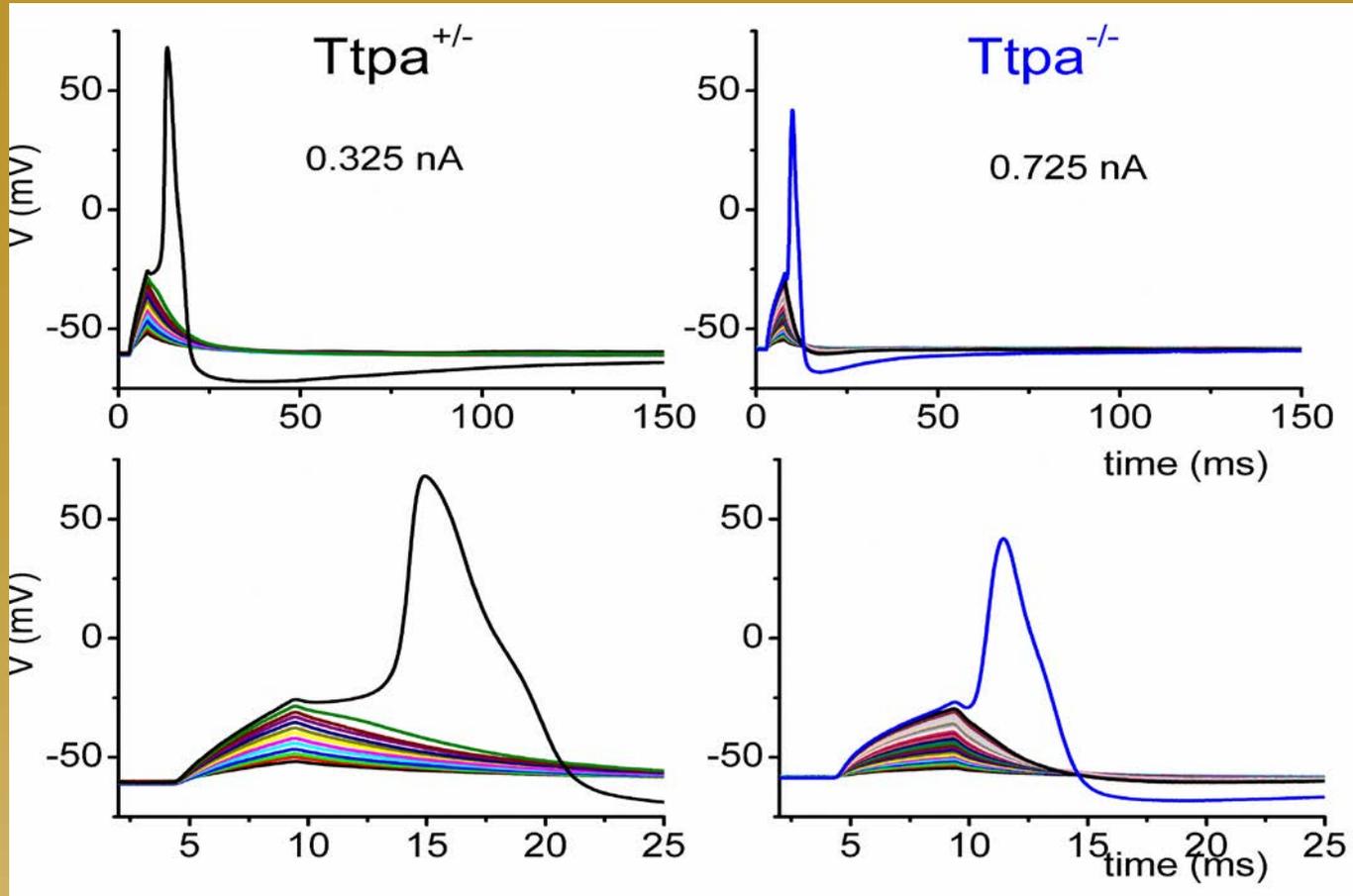
- Sufficient vitE: Retinoid orphan related receptor (*Rora*) ACTIVATED
 - Ligand= Cholesterol
 - Synaptic maintenance (*ITPR1*)
 - Antioxidant proteins (glutathione peroxidase 1, peroxiredoxin 6)
- Insufficient vitE: Liver X receptor ACTIVATED
 - Ligand= Oxidized cholesterol
 - Lipid homeostasis
 - INACTIVATES *Rora*



Input resistance of $Ttpa^{-/-}$ DRG neurons plummets by ~ 2 fold



Membrane excitability of $Ttpa^{-/-}$ DRG neurons is reduced

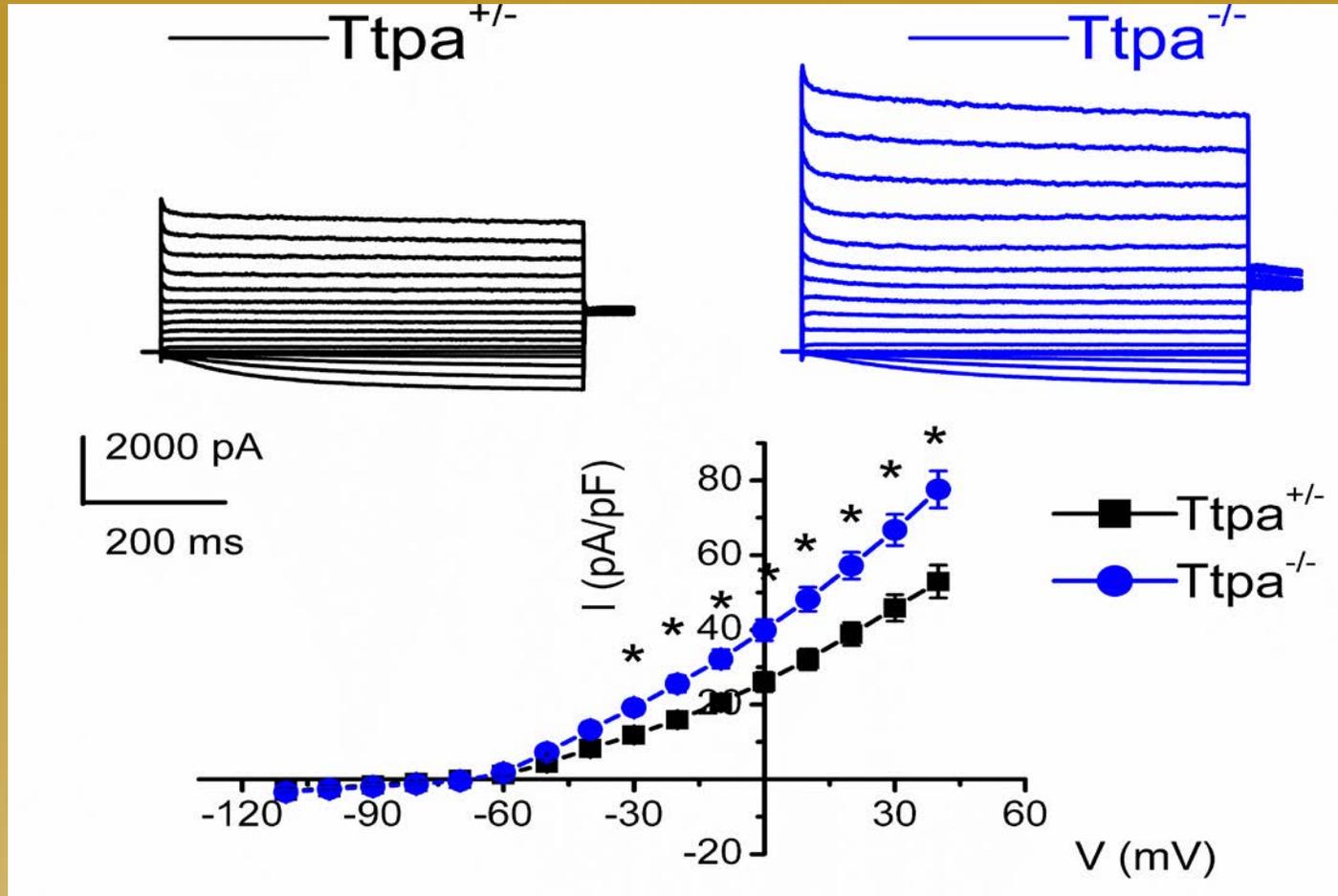


Membrane excitability of $Ttpa^{-/-}$ DRG neurons is reduced



	<i>Ttpa</i> ^{-/+} (n=9)		<i>Ttpa</i> ^{-/-} (n=9)		<i>P</i>
	Mean	SD	Mean	Sd	
RMP	-61.34	3.21	-62.17	3.89	0.75
Threshold	-33.28	2.87	-35.46	3.13	0.34
Amplitude	112.97	6.77	94.59	7.56	0.01
AP₅₀	4.51	0.34	2.45	0.21	0.0001
AHP	13.67	2.14	9.5	1.66	0.02

Enhanced outward K^+ currents in $Ttpa^{-/-}$ neurons

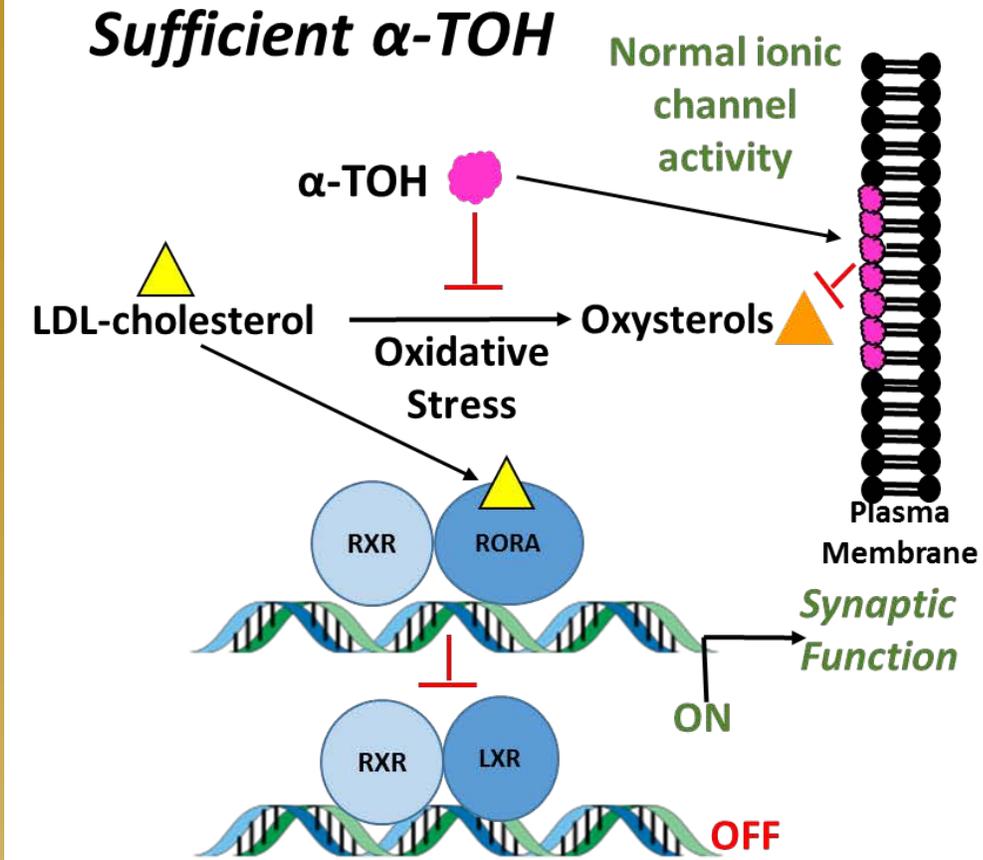


Future Goals

1. Determine identity of enhanced K^+ channels
2. Determine mechanism of enhancement

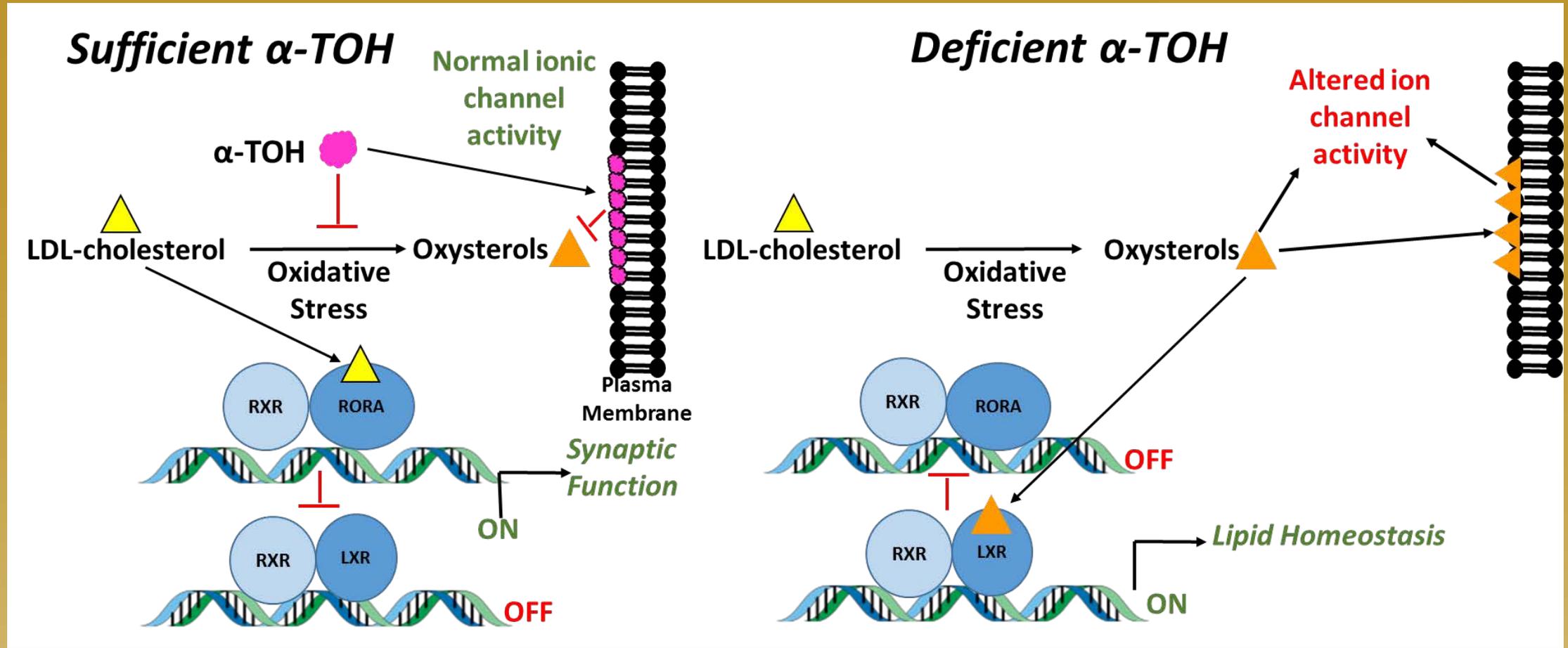


Proposed Molecular Mechanism of VitE

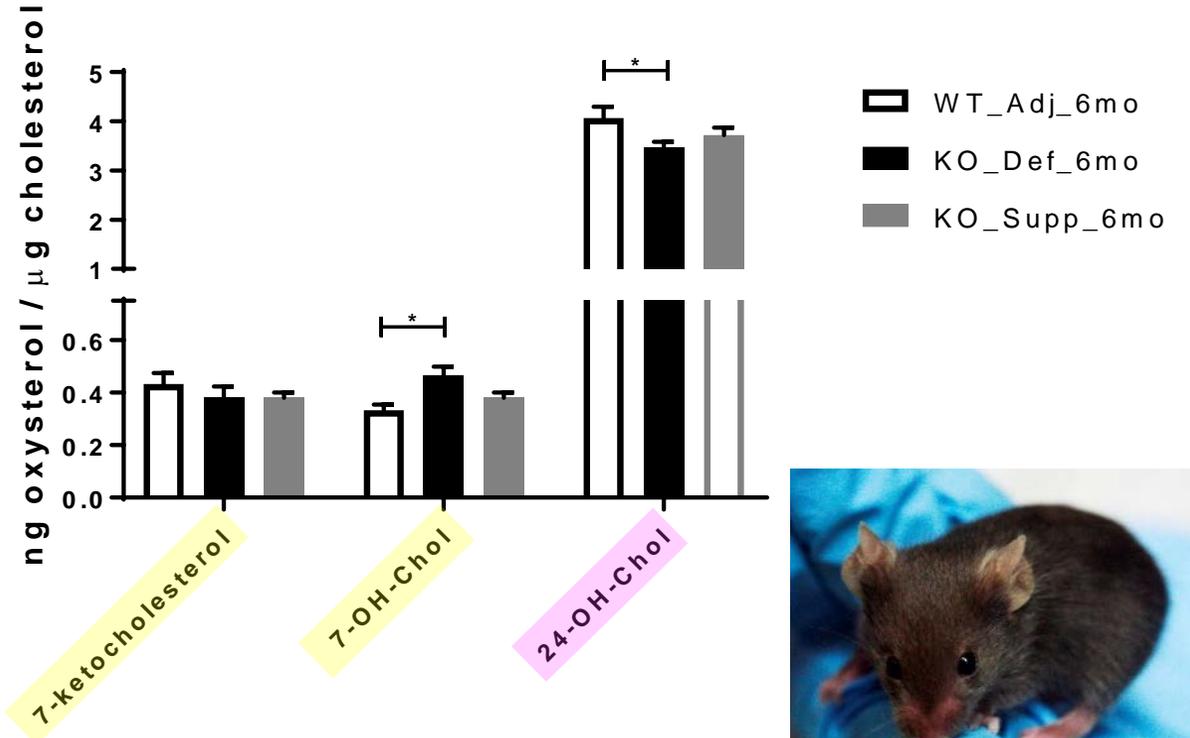
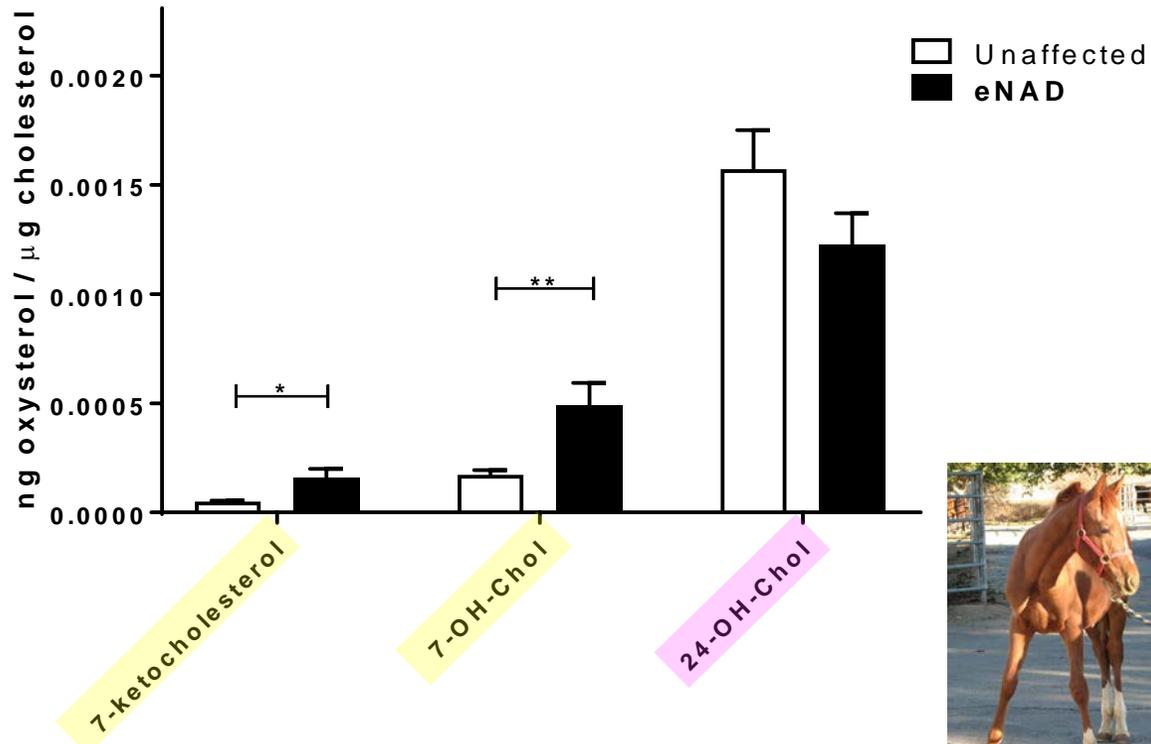




Proposed Molecular Mechanism of VitE



Spinal Cord Oxysterols





Conclusions



- Postnatal development
 - Role of α -TOH in modulating the inflammatory response
 - Decreased myelination with vitE insufficiency
 - Nuclear receptor activation affected by α -TOH concentrations
 - Sufficient vitE : RORA
 - Insufficient vitE: LXR
 - Reduced DRG membrane activity with vitE deficiency
- Molecular dysregulation may lead to neurologic phenotype

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ORIP K01 Support (2013-2018)

- 18 (of 37) publications
- 55 research abstracts
- 20 invited presentations
- Additional grants (as PI)
 - 3 Extramural
 - 6 Intramural
- 4 graduate students trained

Scott et al. *BMC Genomics* (2017) 18:511
DOI 10.1186/s12864-017-3884-2

BMC Genomics

RESEARCH ARTICLE Open Access

Identification of long non-coding RNA in the horse transcriptome CrossMark

E. Y. Scott^{1†}, T. Mansour^{2,3†}, R. R. Bellone^{2,4}, C. T. Brown², M. J. Mienaltowski¹, M. C. Penedo⁴, P. J. Ross¹, S. J. Valberg⁵, J. D. Murray^{1,2} and C. J. Finno^{2*}

ANIMAL GENETICS Immunogenetics, Molecular Genetics and Functional Genomics 

doi: 10.1111/age.12531

Deletion of 2.7 kb near *HOXD3* in an Arabian horse with occipitoatlantoaxial malformation

M. H. Bordbari^{*}, M. C. T. Penedo[†], M. Aleman[†], S. J. Valberg[§], J. Mickelson[¶] and C. J. Finno^{*}

Cerebellum
DOI 10.1007/s12311-016-0823-8

ORIGINAL PAPER

Defining Trends in Global Gene Expression in Arabian Horses with Cerebellar Abiotrophy

E. Y. Scott¹ · M. C. T. Penedo^{2,3} · J. D. Murray^{1,3} · C. J. Finno³



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- UCD Center for Equine Health

