The Power of Forage: The Importance of Feeding the Gut Microbiome in Horses









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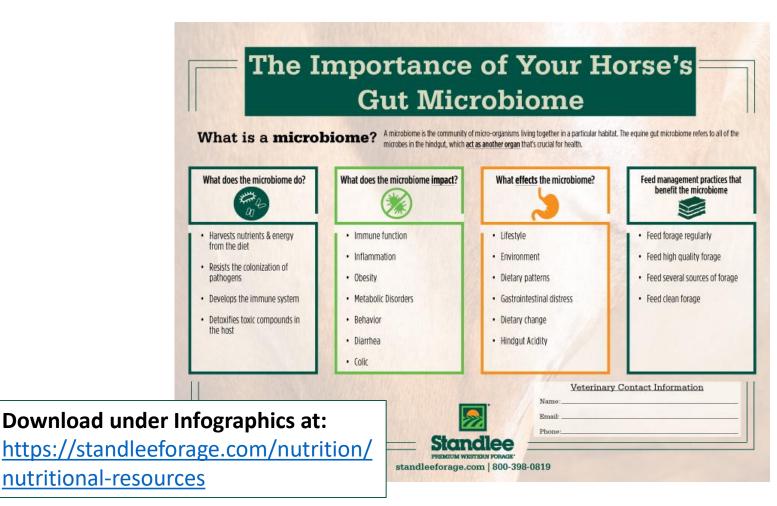
For more information, please contact kstarr@standleeforage.com.

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YOUR HORSE'S GUT MICROBIOME







NUTRITIONAL RESOURCES



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"Nutrition is the key to success"

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NUTRITIONAL RESOURCES

Standlee Premium Western Forage[®] creates and maintains a wealth of nutritional resources to help you maintain the nutrition of your horses. You can find those resources here.

Nutritional Webinars

- How to Decrease Your Horse's Risk of Colic with Nutrition Management (Watch Video Recording Now)
- Winter Feeding (Watch Video Recording Now)
- Gastric Ulcers in Horses (Watch Video Recording Now)
- Metabolic Disease Prevention and Managment (Watch Video Recording Now)
- What Do I Need to Know About Raising Chicks? (Watch Video Recording Now)
- When Quality Hay Is In Short Supply, What Can I Feed My Horse? (Watch Video Recording Now)
- Beet Pulp What Is It and Why Do Horses Need It? (Watch Video Recording Now)



The Power of Forage: The Importance of Feeding the Gut Microbiome in Horses



DR. TANIA CUBITT PERFORMANCE HORSE NUTRITION









7

"Nutrition is the key to success"

Digestion

OUTLINE

- Gut Microbiome
 - Obesity
 - Immune function
 - Inflammation
 - Behavior
- Management









Digestion







BACKGROUND

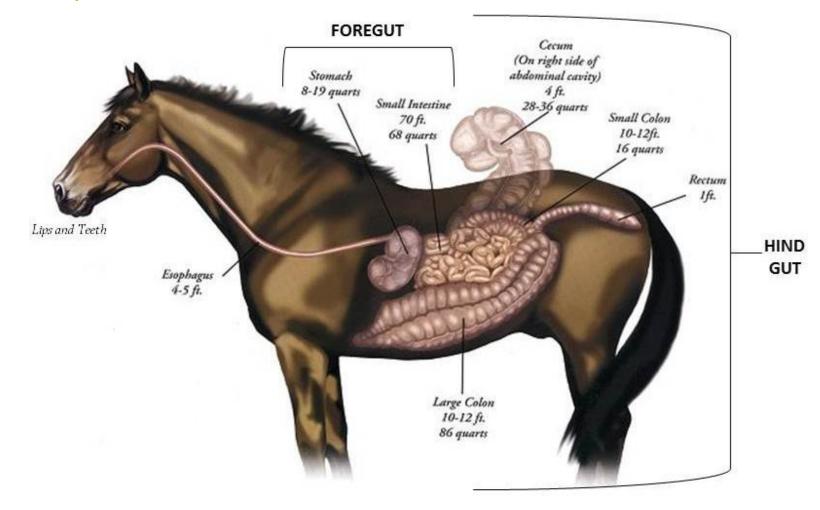
- Grazing non-ruminant herbivore
 - Hindgut fermenter capable of utilizing a wide range of plant species (Costa et al., 2012)
- Hindgut fermenters contain a complex microbial digestion uniquely adapted to:
 - Grazing on high-fiber, low-energy fodder (Dougal et al., 2012)
 - Degrade and ferment to short chain fatty acids that in turn can be absorbed and utilized as energy sources (Blackmore et al., 2013)





EQUINE DIGESTIVE TRACT









FORAGE AMOUNT

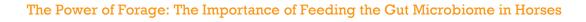
"Nutrition is the key to success"

- Absolute Minimum = 1% of B.W.
 1000 lb horse = 10 lbs forage (DM)
- Weight Loss = 1.2% of B.W. 1000 lb horse = 12 lbs forage (DM)
- Recommended Minimum = 1.5% of B.W. 1000 lb horse = 15 lbs forage (DM)
- Normal Forage Intake = 1.8 to
 2.5% of B.W. (DM)
- Maximum Intake = 3 to 3.5% of B.W. (DM)

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QUESTIONS?

"Nutrition is the key to success"

- We know the numbers
- We know the horse needs fiber

BUT WHY?

• We know gut health can effect many functions in the horse

BUT HOW?

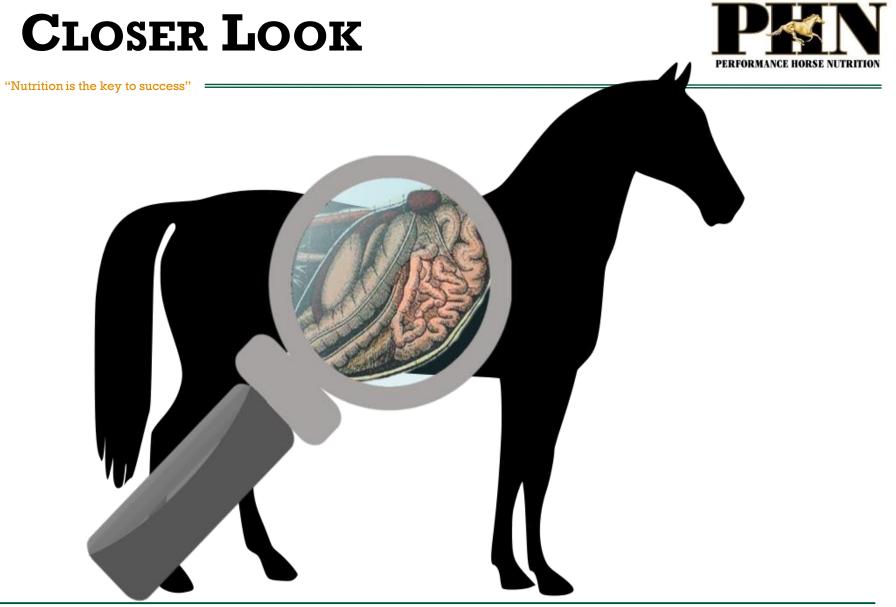




Gut Microbiome









MICROBIOME



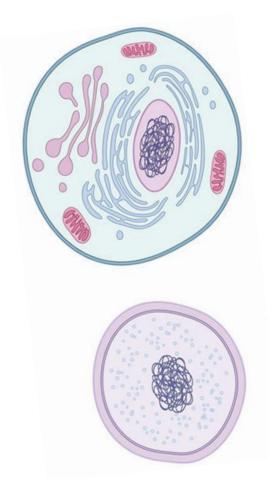
- A microbiome is the community of micro-organisms living together in a particular habitat. Humans, animals and plants have their own unique microbiomes.
- The equine gut microbiome refers to all of the microbes in hindgut, which <u>act as another organ</u> that's crucial for health.
- Gut microbiome plays important roles in:
 - Harvesting nutrients & extracting energy from the diet
 - Resisting the colonization of pathogens
 - Developing the immune system
 - Detoxifying toxic compounds in the host





MICROBIOME

- Consists of:
 - Bacteria
 - Fungi (yeast)
 - Protozoa
 - Viruses
 - Archaea
- Microbes outnumber the cells in our body by a factor of about 10 to 1







The gut microbiome can be affected by external factors such as:

• Lifestyle

"Nutrition is the key to success"

- Environment
- Dietary patterns

MICROBIOME

- Gastrointestinal disease
- Dietary change
- Fermentative acidosis
- Laminitis
- Colic











SCIENTIFIC REPORTS

Article | OPEN | Published: 14 November 2017

Evaluating the impact of domestication and captivity on the horse gut microbiome

Jessica L. Metcalf, Se Jin Song, James T. Morton, Sophie Weiss, Andaine Seguin-Orlando, Frédéric Joly, Claudia Feh, Pierre Taberlet, Eric Coissac, Amnon Amir, Eske Willerslev, Rob Knight, Valerie McKenzie & Ludovic Orlando [™]

Scientific Reports 7, Article number: 15497 (2017) | Download Citation 🚽

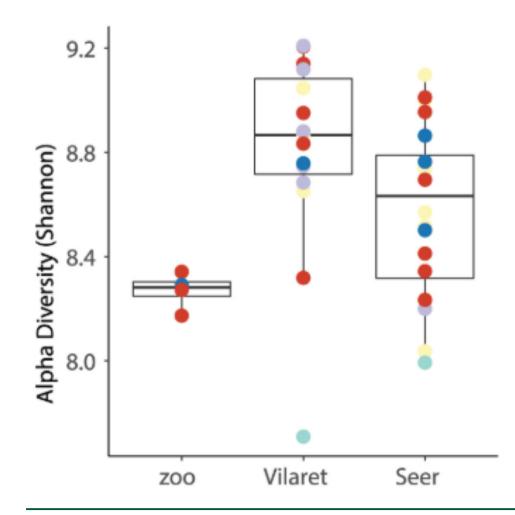


The Power of Forage: The Importance of Feeding the Gut Microbiome in Horses

DOMESTICATION



"Nutrition is the key to success"



Gut microbial diversity based on birth place and domestication:

- Zoo (domesticated)
- Vilaret (wild reserve in France not domesticated)
- Seer (wild reserve in Mongolia not domesticated)

Significantly stunted microbial diversity when born in a zoo





Article

Asian-Australasian Journal of Animal Sciences (AJAS) 2016; 29(9): 1345-1352. Published online: December 1, 2015 DOI: https://doi.org/10.5713/ajas.15.0587

Comparison of Fecal Microbiota of Mongolian and Thoroughbred Horses by High-throughput Sequencing of the V4 Region of the 16S rRNA Gene

Yiping Zhao, Bei Li, Dongyi Bai, Jinlong Huang, Wunierfu Shiraigo, Lihua Yang, Qinan Zhao, Xiujuan Ren, Jing Wu, Wuyundalai Bao, Manglai Dugarjaviin^{*}





- Mongolian horses vs.
 Thoroughbred
- All housed in Mongolia 2 different farms
- Mongolian horses roamed and grazed prairies
- Thoroughbreds fed typical hay and grain rations housed in stalls



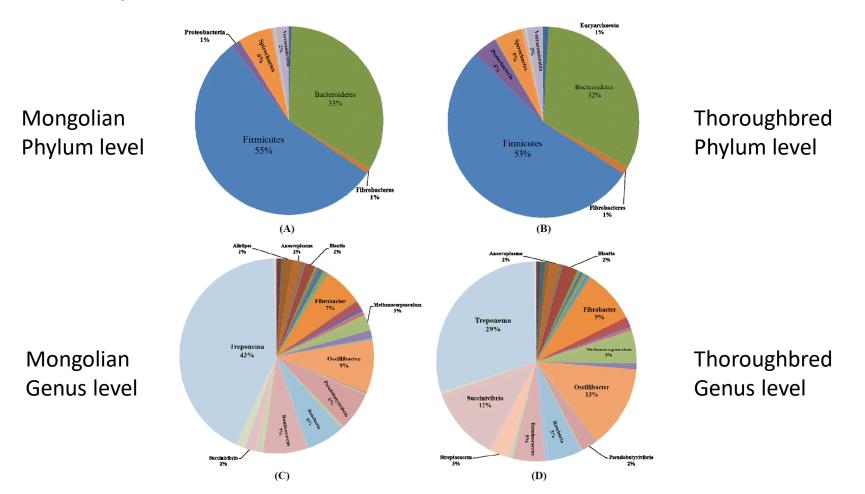








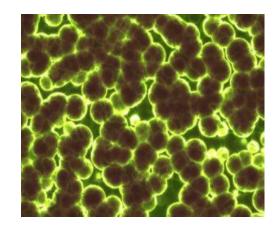








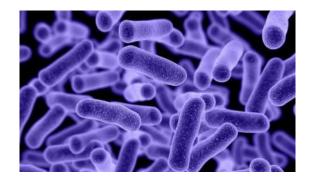
- Previous studies in horses reported:
 - Firmicutes as the predominant taxa (39.5% to 72.6%)
 - Firmicutes are the main bacterial phylum, comprising over 250 genera, including Lactobacillus, Streptococcus, Mycoplasma, and Clostridium which are able to produce several short chain fatty acids (SCFAs) like butyrate







- Previous studies in horses reported:
 - Bacteroidetes (2nd to Firmicutes) (8.9% to 21.3%) in healthy horses populations Costa et al. (2012)
 - Bacteroidetes are gramnegative bacteria that ferment polysaccharides and otherwise indigestible carbohydrates and produce short-chain fatty acids (SCFAs) that have many beneficial effects in the gut







"Nutrition is the key to success"

Current study reported the same predominance

However

- Proteobacteria was significantly different in two breeds
 - (1% and 4%, Mongolian horses and Thoroughbred horses respectively)
 - Proteobacteria included many pathogenic bacteria, such as *salmonella*, *V. cholera*, *Helicobacter pylori*, etc.
- Streptococcus was 3% for Thoroughbred horses and less than 1% for Mongolian horses
 - Streptococcus also included pathogenic bacteria (e.g. Streptococcus pneumoniae, Streptococcus pyogenes, Streptococcus agalactiae)

Link?

• Thoroughbred horses more susceptible to diseases than Mongolian horses?







ORIGINAL RESEARCH published: 20 September 2018 doi: 10.3389/fvets.2018.00225



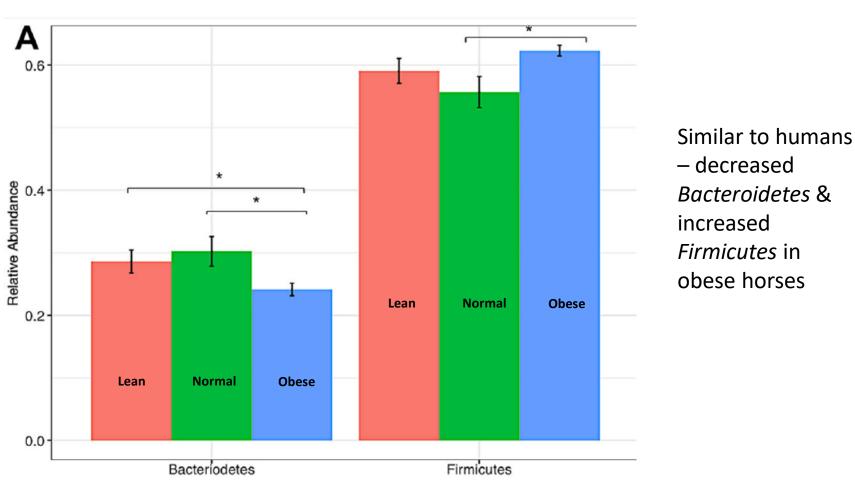
Microbiome and Blood Analyte Differences Point to Community and Metabolic Signatures in Lean and Obese Horses

Amy S. Biddle^{1*}, Jean-Francois Tomb² and Zirui Fan²

¹ Department of Animal and Food Science, University of Delaware, Newark, DE, United States, ² Department of Computer and Information Sciences, University of Delaware, Newark, DE, United States





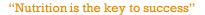














- Human obesity has been associated with reduced proportions of *Bacteroidetes*, and correspondingly increased abundances of *Firmicutes* in fecal samples (Ley et al., 2006)
- Human obesity is associated with a reduction in insulin sensitivity (Kahn and Flier, 2000), and in turn, a reduced insulin sensitivity has been associated with decreased microbial richness (Le Chatelier et al., 2013)







Journal of Equine Veterinary Science Volume 76, May 2019, Pages 40-41



Microbiome effects on metabolic efficiencies in easy and hard keepers

A.C.B. Johnson*, A.S. Biddle



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Poll Question



"Nutrition is the key to success"





The Power of Forage: The Importance of Feeding the Gut Microbiome in Horses

Metabolic Status



- Hard Keeper (HK)
- Easy Keeper (EK)
- Medium Keeper (MK)
- HK contained the least bacterial diversity followed by EK then MK respectively
- HK –less lactic acid producing bacteria, less Amino Acid utilizing bacteria
- MK are the most efficient at nutrient digestion and host absorption
- Reduced bacterial diversity in HK leads to insufficient nutrient levels to support a healthy microbiome and maintain horse condition





PLoS One. 2014; 9(2): e87424. Published online 2014 Feb 4. doi: <u>10.1371/journal.pone.0087424</u> PMCID: PMC3913607 PMID: <u>24504261</u>

Characterisation of the Faecal Bacterial Community in Adult and Elderly Horses Fed a High Fibre, High Oil or High Starch Diet Using 454 Pyrosequencing

<u>Kirsty Dougal</u>, ¹ <u>Gabriel de la Fuente</u>, ¹ <u>Patricia A. Harris</u>, ² <u>Susan E. Girdwood</u>, ¹ <u>Eric Pinloche</u>, ¹ <u>Raymond J. Geor</u>, ³ <u>Brian D. Nielsen</u>, ³ <u>Harold C. Schott</u>, II, ³ <u>Sarah Elzinga</u>, ³ and <u>C. Jamie Newbold</u> ¹, *





AGE/DIET"Nutrition is the key to success"

- 17 horses (2 different groups -Adult vs. Elderly)
 - 8 in the 5-12 year range
 - 9 in the 19-28 year range
- 3 different diets

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- Hay (NSC 11%; Fat 4%)
- Hay + CHO (NSC 42%; Fat 5%)
- Hay + Fat (NSC 13%; Fat 8.5%)





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AGE/DIET

- Core Community -
 - HAY diet 15.9%
 - OIL diet 10.3%
 - CHO 5.4%
- Lachnospiraceae being the most abundant in the core community
 - Lachnospiraceae are known butyrate producers and butyrate is known to have a protective function on colonocytes in the gut wall
- Fecal microbiome significantly reduced in elderly horses (perhaps due to reduced dentition/fiber digestion? Or is this a result?)





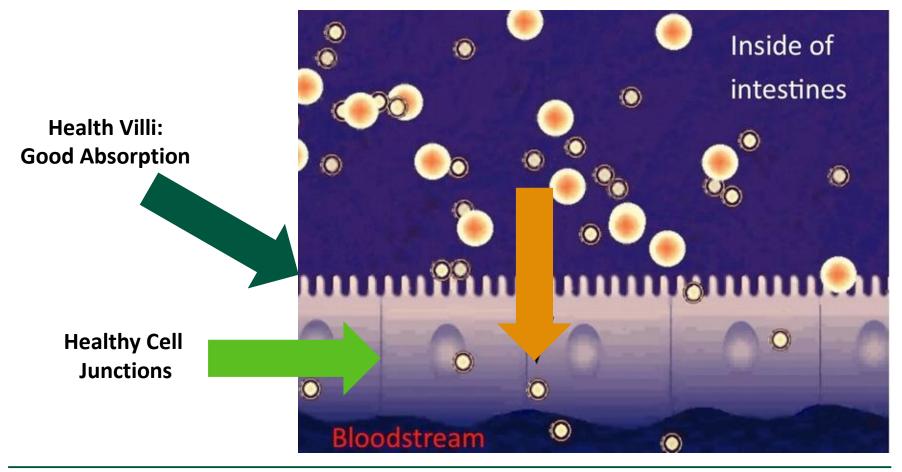


HEALTHY GUT



"Nutrition is the key to success"

Small, properly digested food particles are absorbed - everything else kept out

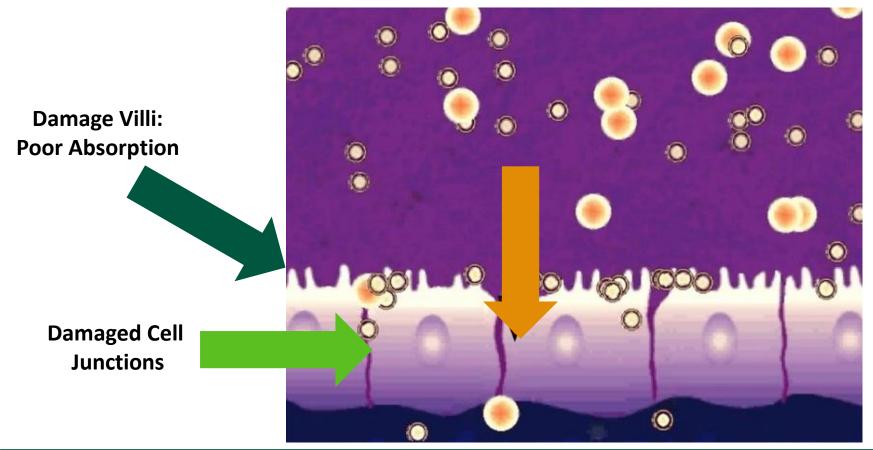








Large food particles and bacterial debris leak through the damaged intestinal barrier





Poll Question







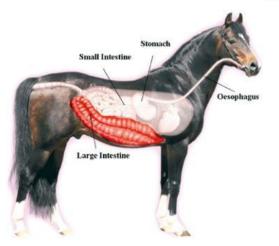


Comparison of the Fecal Microbiota of Healthy Horses and Horses with Colitis by High Throughput Sequencing of the V3-V5 Region of the 16S rRNA Gene

Marcio C. Costa¹*, Luis G. Arroyo², Emma Allen-Vercoe³, Henry R. Stämpfli², Peter T. Kim⁴, Amy Sturgeon¹, J. Scott Weese¹

1 Department of Pathobiology, Ontario Veterinary College, University of Guelph, Guelph, Ontario, Canada, 2 Department of Clinical Studies, Ontario Veterinary College, University of Guelph, Guelph, Ontario, Canada, 3 Department of Molecular and Cellular Biology, College of Biological Sciences, University of Guelph, Guelph, Guelph, Ontario, Canada, 4 Department of Mathematics and Statistics, College of Physical and Engineering Science, University of Guelph, Guelph, Ontario, Canada

- 16 horses
- 6 healthy
- 10 colitis
- Fecal samples





COLITIS

• Phylum: "Firmicutes"

• Order: Clostridiales

Domain: Bacteria

• Family: Lachnospiraceae



- 100% 90% 80% 70% Other Fusobacteria 60% Spirochaetes 50% Proteobacteria 40% Actinobacteria Bacteroidetes 30% Firmicutes 20% 10% 0% Healthy Colitis
- Clostridiales, members of the Lachnospiraceae family, were the most abundant bacteria shared between healthy horses
- Previous study showed Lachnospiraceae highest in horses eating high forage diets





Journal of Equine Veterinary Science 44 (2016) 9-16



Original Research

Comparison of the Fecal Microbiota in Horses With Equine Metabolic Syndrome and Metabolically Normal Controls Fed a Similar All-Forage Diet

Sarah E. Elzinga^a, J. Scott Weese^b, Amanda A. Adams^{a,*}

^a Department of Veterinary Science, M.H. Gluck Equine Research Center, University of Kentucky, Lexington KY ^b Ontario Veterinary College, University of Guelph, Ontario, CA





EMS

"Nutrition is the key to success"

- 20 horses
- 10 with EMS
- 10 non EMS control
- Forage only diet

Characterization of phenotypic measures in equine metabolic syndrome (EMS) versus non-EMS control horses.

Group	BCS	CNS	Basal Insulin (µIU/mL)	Post Oral Sugar Insulin (µIU/mL)
EMS	7.13 ± 0.15	3.57 ± 0.27	33.85 ± 2.66	63.26 ± 5.29
Control	5.85 ± 0.23	1.95 ± 0.18	13.03 ± 1.46	26.11 ± 3.77
P value	<.001	<.001	<.001	<.001

Average body condition score (BCS) and cresty neck score (CNS), and basal insulin and insulin 60 minutes after oral sugar administration in n = 10 EMS and n = 10 non-EMS control horses. Results are represented as the mean plus or minus the standard error of the mean.



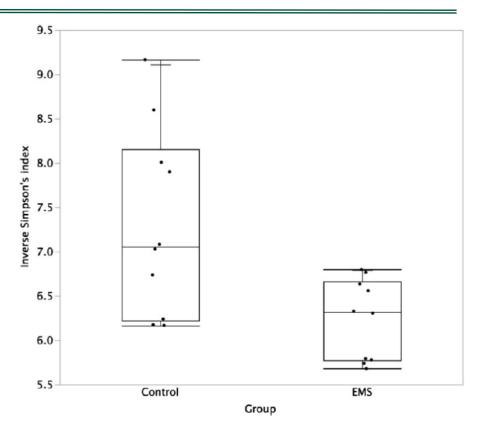


Fig. 1. Intestinal microbiota diversity in equine metabolic syndrome (EMS) versus non-EMS control horses. Intestinal microbiota diversity as represented by an inverse Simpson's index in n = 10 EMS and n = 10 non-EMS control horses.





Steelman et al. BMC Veterinary Research 2012, 8:231 http://www.biomedcentral.com/1746-6148/8/231



RESEARCH ARTICLE

Open Access

Pyrosequencing of 16S rRNA genes in fecal samples reveals high diversity of hindgut microflora in horses and potential links to chronic laminitis

Samantha M Steelman¹, Bhanu P Chowdhary¹, Scot Dowd², Jan Suchodolski³ and Jan E Janečka^{1*}



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LAMINITIS

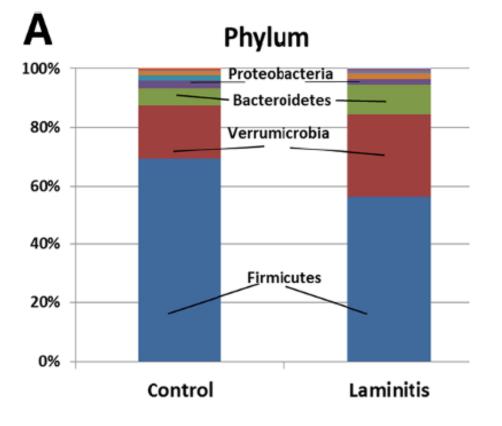
- 18 horses
 - 10 normal
 - 8 laminitis

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PREMILIM WESTERN FORAGE

- Results
 - Less Firmicutes in Laminitic horses



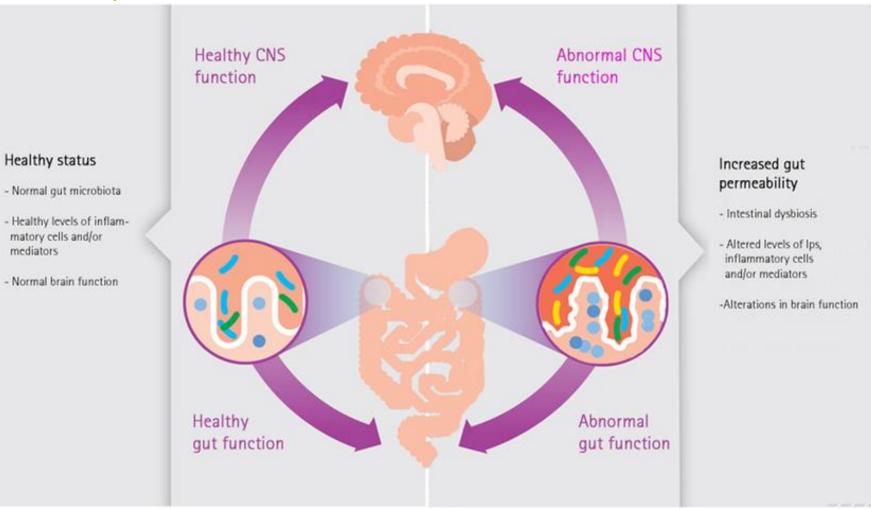




GUT - BRAIN AXIS (BEHAVIOR)









Management

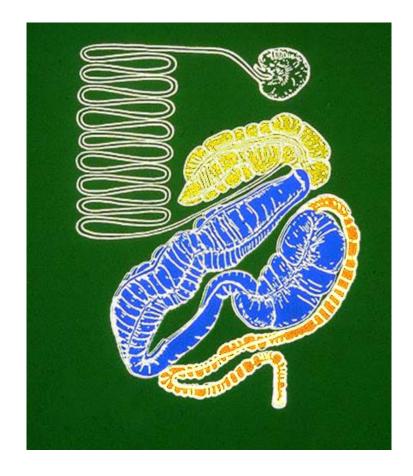






MANAGEMENT

- Focus on feeding the "Microbiome"
- Feeding strategies that:
 - Promote a healthy "microbiome"
 - Discourage pathogenic bacteria
- Gut microbiome is the link between nutrition and health







FORAGE AMOUNT REMINDER



- Absolute Minimum = 1% of B.W.
 1000 lb horse = 10 lbs forage (DM)
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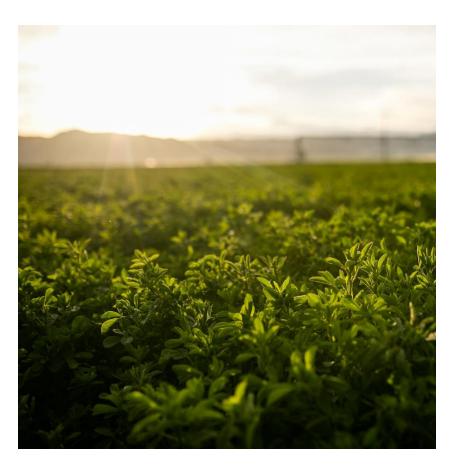








- High quality forage
- Consistently provided
- Feeding management practices that favor the microbiome
- Research ongoing





Q & A Session







Thank you for joining us!

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