## **PUBLICATIONS**

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- 19. Kehoe SI<sup>†</sup>, **Dill-McFarland KA**, Breaker JD, Suen G. 2019. Effects of corn silage inclusion in pre-weaned calf diets. <u>J Dairy Sci</u> *In press*
- 18. Carroll C, Olsen KD, Ricks NJ, **Dill-McFarland KA**, Suen G, Robinson TF, Chaston JM. 2018. Bacterial communities in the alpaca gastrointestinal tract vary with diet and body site. Front Microbiol. 9: 3334. doi: 10.3389/fmicb.2018.03334
- 17. **Dill-McFarland KA**\*, Tang Z\*, Kemis JH, Kerby RL, Chen G, Palloni A, Sorenson T, Rey FE<sup>†</sup>, Herd P<sup>†</sup>. Close social relationships correlate with human gut microbiota composition. Sci Rep *In press*. Preprint doi: <a href="https://doi.org/10.1101/428938">https://doi.org/10.1101/428938</a>
- 16. **Dill-McFarland KA**, Weimer PJ, Breaker JD, Suen G. 2018. Diet influences early microbiota development in dairy calves without long-term impacts on milk production. <u>Appl Environ Microbiol</u> Epub ahead of print. doi: 10.1128/AEM.02141-18
- 15. De Wolfe TJ, Eggers S, Barker AK, Kates A, Dill-McFarland KA, Suen G, Safdar N. 2018. Oral probiotic combination of *Lactobacillus* and *Bifidobacterium* alters the gastrointestinal microbiota during antibiotic treatment for *Clostridium difficile infection*. PLoS One 13(9): e0204253. doi: 10.1371/journal.pone.0204253
- 14. Cunha CS, Marcondes MI, Veloso CM, Mantovani HC, Pereira LGR, Tomich TR, **Dill-McFarland KA**<sup>†</sup>, Suen G<sup>†</sup>. 2018. Compositional and structural dynamics of the ruminal microbiota in dairy heifers and its relationship to methane production. <u>J Sci Food Agric 99(1)</u>: 210-18. doi: 10.1002/jsfa.9162
- 13 Romano KA, **Dill-McFarland KA**, Kasahara K, Kerby RL, Vivas El, Amador-Noguez D, Herd P, Rey FE. 2018. Fecal Aliquot Straw Technique (FAST) allows for easy and reproducible subsampling: Assessing interpersonal variation in trimethylamine-*N*-oxide (TMAO) accumulation. <u>Microbiome</u> 6(1): 91. doi: 10.1186/s40168-018-0458-8
- 12. Dias J, Marcondes MI, de Souza SM, da Mata BC, Noronha MF, Resende RT, Machado FS, Mantovani HC, **Dill-McFarland KA**†, Suen G†. 2018. Bacterial community dynamics across the gastrointestinal tracts of dairy calves during preweaning development. <u>Appl Environ Microbiol</u> 84(9): e02675-17. doi: 10.1128/AEM.02675-17
- 11. Williams CL, **Dill-McFarland KA**, Sparks DL, Kouba AJ, Willard ST, Suen G, Brown AE. 2018. Dietary changes during weaning shape the gut microbiota of red pandas (*Ailurus fulgens*). Conserv Physiol 6(1): cox075. doi: 10.1093/conphys/cox075
- 10. Cunha CS, Veloso CM, Marcondes MI, Mantovani HC, Tomich TR, Periera LGR, Ferreira MF, **Dill-McFarland KA**†, Suen G†. 2017. Assessing the impact of rumen microbial communities on methane emissions and production traits in Holstein cows in a tropical climate. Syst Appl Microbiol 40(8): 492-99. doi: 10.1016/j.syapm.2017.07.008
- 9. Dai X, Weimer PJ, **Dill-McFarland KA**, Brandao VL, Suen G, Faciola AP. 2017. Camelina seed supplementation at two dietary fat levels changes ruminal bacterial community composition in a dual-flow continuous culture system. <u>Front Microbiol</u> 8: 2147. doi: 10.3389/fmicb.2017.02147
- 8. Vogt NM, Kerby RL, **Dill-McFarland KA**, Harding SJ, Merluzzi AP, Johnson SC, Carlsson CM, Asthana S, Zetterberg H, Blennow K, Bendlina BB<sup>†</sup>, Rey FE<sup>†</sup>. 2017. Gut microbiome alterations in Alzheimer's disease. <u>Sci Rep</u> 7(1): 13537. doi: <u>10.1038/s41598-017-13601-y</u>
- 7. Dias J, Marcondes MI, Noronha MF, Resende RT, Machado FS, Mantovani HC, **Dill-McFarland KA**†, Suen G†. 2017. Effect of pre-weaning diet on the ruminal archaeal, bacterial, and fungal diversity of dairy calves. <u>Front Microbiol</u> 8: 1553. doi: <u>10.3389/fmicb.</u> 2017.01553

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- 3. **Dill-McFarland KA**, Weimer PJ, Pauli JN, Peery MZ, and Suen G. 2016. Diet specialization selects for an unusual and simplified gut microbiota in two- and three-toed sloths. <u>Environ Microbiol</u> 18(5): 1391-402. doi: 10.1111/1462-2920.13022
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