

Exposure to a Virtual Environment Induces Biological and Microbiota Changes in Onset-of-Lay Hens

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Redweik GAJ, Millman ST, Parsons RL, Hurtado Terminel AN, Radkowski R, Daniels K, Lyte M, Oliver J and Mellata M (2022) Exposure to a Virtual Environment Induces Biological and Microbiota Changes in Onset-of-Lay Hens. Front. Virtual Real. 3:891584. doi: 10.3389/frvir.2022.891584 Increasing demand for cage-free eggs arises from goals to provide hens with better welfare, particularly in terms of natural behavior. However, most laying hens are kept in conventional cages, and cage-free systems can present challenges, such as injuries, floor eggs, and bacterial infections. We proposed using virtual reality (VR) as a feasible means for combining the positive attributes of natural environments while mitigating health risks. To our knowledge, no animal study has provided evidence that VR can trigger biological changes to improve animal health and well-being nor whether VR can affect the gut microbiota. In this study, we used VR technology to simulate a natural environment in laying hen housing. Early-lay White Leghorn hens were placed in pens with (VR) or without (CON) video projections displaying free-range chickens interacting with indoor and outdoor environmental features over 5 days. Using in vitro blood bactericidal assays, VR hens exhibited higher resistance against avian pathogenic Escherichia coli versus CON (p < p0.05), which was positively associated with corticosterone levels (p < 0.01). Analyzing intestinal neurochemicals via ultra-high pressure liquid chromatography, salsolinol was the only neurochemical metabolite affected by VR, being greater in CON ileal content (p < p0.0001), in VR ileal mucus (p < 0.01), and in VR ceca tissue (p < 0.05). Using 16S rRNA sequencing and QIIME2 analyses, no differences in alpha nor beta diversity were determined between groups. Although several genera (Megamonas, Ruminococcus, Slackia) were reduced in VR hens versus CON, Mucispirillum schaedleri (member of Deferribacteres Phylum) was the only taxon increased in VR hens, being elevated in ileal mucus (p < 0.05). Lastly, using the QIIME2 plugin mmvec to map microbe-metabolite cooccurrences, we identified several positive relationships between bacterial phyla and neurochemical metabolites, notably finding dopamine and salsolinol levels were related to Deferribacteres and Tenericutes levels. In conclusion, we found that several biological parameters were influenced by VR treatment in hens, suggesting that VR can be used to improve host resistance to pathogens and gut health in poultry.

Keywords: virtual reality, chickens, corticosterone, neurochemicals, gut microbiota, APEC, salsolinol

INTRODUCTION

Commercial poultry are housed in high-density environments to maximize output of products meat and eggs. However, in many caged and cage-free facilities, chickens lack complex natural environmental features with biological relevance, drawing ethical concerns from the public. Although public interests have pushed commercial egg farms to shift towards cage-free housing, cage-free animals are at higher risk for infections, and undesirable behaviors (e.g. cannibalism), as well as higher rates of injury and mortality (Heerkens et al., 2015). Although free-range operations are also used to raise chickens, these conditions present greater risks of predation, parasites/pathogens (Bestman and Bikker-Ouwejan, 2020), and exposure to inclement weather (Singh et al., 2017). Thus, technologies that allow the combination of naturalistic stimuli of free-range housing with the productivity and husbandry benefits in poultry facilities are drastically needed.

Virtual reality (VR) has recently garnered much attention in human medicine given its benefits on post-traumatic stress (Maples-Keller et al., 2017), sports training (Stinson and Bowman, 2014), autism spectrum disorder (Yang et al., 2018), and stroke rehabilitation (Maier et al., 2019). In addition to humans, VR modulates social behaviors in fish, mice, and flies (Stowers et al., 2017). To our knowledge, no study has demonstrated if VR can be protective against infectious disease or whether it can induce changes in the intestinal environment (microbiota, neurochemical production, *etc.*).

Compared to humans, birds' vision is tetrachromic with poor depth perception and they better recognize objects when they are moving than stationary (Broom, 1969; Dawkins and Woodington, 2000). Environment factors, such as color, light quality, duration, and intensity affect feeding behaviors of poultry (Lien et al., 2008). Additionally, video images reduces fear response, as birds quickly approach the video image to which they are exposed (Clarke and Jones, 1999). Furthermore, when looking at a video of chicks feeding, birds will imitate these behaviors and approach their feed more quickly (Clarke and Jones, 2001). Thus, chickens are highly receptive to visual stimuli, and VR design should be adapted to accommodate the sensory preferences of chickens. However, the extent to which biological functions aside from behavior are affected by VR is unclear.

Overall, we hypothesized that VR could be designed to impact biological factors, and thus improve laying hen welfare and health. The objectives of the study were to 1) create a virtual free-range environment from video recordings of freerange chickens performing behaviors associated with positive affect (*e.g.*, nesting, dustbathing, preening, perching) to display on the walls of an enclosed pen, 2) test the effect of VR on overall chicken health and welfare, 3) characterize stress and immunological responses to VR, and 4) investigate how VR exposure may affect neurochemical-microbial interactions in the chicken intestine.

MATERIALS AND METHODS

Ethics Statement

Animal experiments were approved by Iowa State University Institutional Animal Care and Use Committee, $\log \#19-211$. Animal distress was minimized during experimental procedures via enrichments (ex: perches). Open-floor pens were implemented to enable social interactions between chickens, and chickens were fed with no-medicated feed and water *ad libitum*. Upon experiment completion at 5 days post pen-placement, chickens were euthanized via CO₂ asphyxiation in accordance with American Veterinary Medical Association Guidelines (2020).

Chickens, Housing, and Husbandry

Chickens used in video recordings for the creation of VR were enrolled from and maintained on three privately owned backvard flocks in Iowa during summer 2019. These flocks consisted of mature hens, roosters, pullets, and chicks, including a diversity of fancier-type egg-laying, meat and dual-purpose breeds. Housing included one or more indoor facilities, equipped with perches, nest boxes, feeders, drinkers and litter areas, with access to an outdoor fenced scratch area or unfenced open prairie with grasses, shrubs and flowers. Chickens used in the laboratory testing of VR consisted of 34 White Leghorn hens (Dekalb Whites) at early-lay stage (15 weeks post-hatch), randomly split into two groups, one exposed to virtual reality (VR) and the other unexposed (CON) (n = 15) via staff blinded to the experimental groups. Birds were individually identified using food dye on plumage and colored leg bands. Each group was housed in a 9 ft \times 9 ft \times 7 ft pen with steel spindle walls and pine shavings litter, and pens were maintained at 74°F. In both VR and CON rooms, pen walls were wrapped with vinyl projector screen material on all four sides. Each pen was equipped with a feed hopper, drinker, perches, nest boxes, and a dustbathing area. The photoperiod was 16L:8D. Husbandry consisted of daily inspection of the chickens, replenishing water and a commercial layer diet was provided ad libitum.

Virtual Reality Design

Hens from the VR and control groups were housed in 9 ft \times 9 ft \times 7 ft steel cages wrapped with vinyl projector screen material on all four sides. In the VR pen (Figure 1), video-recorded scenes of free-range chickens (see Supplemental Materials for example videos) were projected on each of the four walls during daylight hours. Each wall depicted a free-range flock interacting with one of the following scenes: 1, perching area; 2, nesting area; 3, outdoor dirt yard scratch area; 4, outdoor vegetative area. The storyboard for wall was constructed with hourly segments depicting scenes intended to provoke desirable behaviors (e.g., nesting, dustbathing, preening, perching) appropriate to the known ethogram and time budget of the laying hen. For example, VR projected behind the nest boxes displayed hens actively performing nesting behavior during the morning hours, when oviposition typically occurs, whereas empty nests with no hens present were displayed during

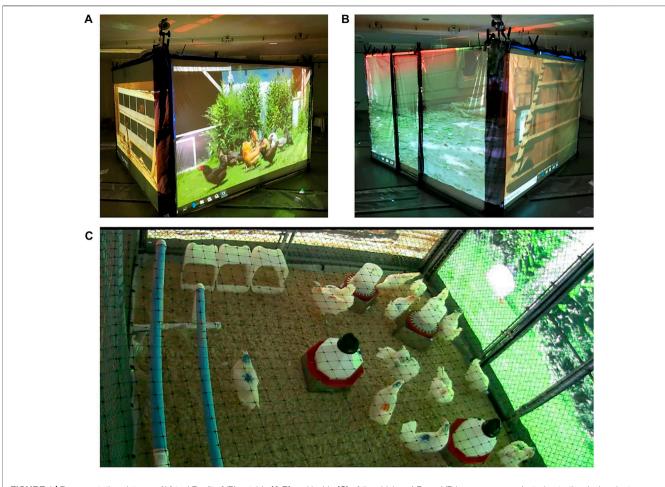


FIGURE 1 | Representative pictures of Virtual Reality (VR) outside (A,B) and inside (C) of the chickens' Pens. VR images were projected onto the vinyl projector screens placed against each pen wall in the VR group only.

afternoon and evening hours when oviposition is unlikely. The perching area VR storyboard switched hourly between still images of perches with no hens present, followed by video images of hens actively perching during the daylight hours, and concluding with a still image of perches with motionless hens perching immediately prior to and during the dimming lights and dark periods.

General Welfare Assessment

Prior to necropsy, all hens were weighted and evaluated according to welfare characteristics, including plumage damage, cleanliness, keel bone deformation, comb pecking wounds and comb abnormality as previously described (Oliveira et al., 2019).

Serum Corticosterone

Prior to euthanasia at day 5 of VR or control treatment, fresh blood (100 μ L) was collected via wing vein via filter-sterilized heparin (1,000 U/ml)-coated needles and immediately placed in citrate-coated tubes at room temperature for 4 h. Blood was centrifuged at 4,000×g for 20 min, and non-hemolyzed serum was individually stored at -80°C. For quantification of corticosterone, serum samples were run in triplicate using a

commercial corticosterone ELISA kit (Enzo Life Sciences) in accordance to the manufacturer directions.

Whole Blood Bactericidal Assay

Prior to euthanasia at day 5 of VR or control treatment, fresh blood (200 µL) was collected from the right-wing vein of hens via filter-sterilized heparin-coated needles and placed on ice. Blood was evenly split into 2 pools per treatment group (n = 7hens per pool), then diluted 1:8 in CO₂-independent media with 2 mM L-glutamine (Gibco). Cell viability was determined to be greater than 98% via trypan blue staining for all pools. Prior to experiments, reference E. coli strains (Table 1) were stored in peptone-glycerol solution at -80°C. Strains were grown in Luria-Bertani (LB) agar (0.1% glucose) overnight at 37°C, colonies were mixed into phosphate-buffered saline (PBS) until OD₆₀₀ reached 0.1, and solutions were diluted to reach 10² CFU/20 µL. Individual wells in 96-well plates were filled with whole blood solution and bacterial inoculum (9:1, respectively) and incubated for 30 min at 40°C. After brief resuspension, samples were serially diluted, plated on MacConkey agar, and incubated overnight at 37°C. Assays were performed in triplicate.

TABLE 1 | Summary of Escherichia coli isolates used for in vitro resistance assays.

Escherichia coli	Relevant phenotype and characteristics	References
APEC O1	APEC isolate, O1 serotype	Johnson et al. (2007
APEC O2	APEC isolate, O2 serotype	Johnson et al. (2006
χ7122	APEC isolate, O78 serotype	Stacy et al. (2014)
MG1655	Non-pathogenic laboratory strain, K-12, OR:K-:H48	Barbieri et al. (2013)

Heterophil to Lymphocyte Ratio

To calculate H:L ratios, a biomarker for stress (Gross and Siegel, 1983), approximately 3 μ L of fresh blood (acquired prior to euthanasia as described earlier) were aliquoted and smeared onto a standard microscope slide (one slide per hen, n = 15 per group), briefly air dried, fixed and stained using a Neat Stain Hematology Staining Kit (Polysciences, Inc.). For each slide, approximately 200 leukocytes were enumerated via light microscopy based on phenotypic characteristics (de Macchi et al., 2013). Thereafter, H:L ratios were then calculated by averaging heterophils and lymphocytes identified within each slide.

Neurochemical and Short Chain Fatty Acid Analyses

Post-euthanasia at day 5 of VR or control treatment, intestinal segments (i.e., ileum and ceca) were aseptically removed from birds in both groups (n = 15 per group). Ileal and cecal contents were squeezed from the intestinal segments into respective tubes. Mucus from each section was collected by longitudinally dissecting intestinal tissues to expose the intestinal lumen and then gently scraping the mucosa with fresh, autoclaved glass microscope slides, which were then directly transferred into respective cryotubes. The remaining tissues were then placed into respective cryotubes. All sample cryotubes were briefly homogenized with 0.2 M perchloric acid (1:10 sample-acid ratio) and flash-frozen via liquid nitrogen prior to -80°C storage. For neurochemical and short chain fatty acid analyses, samples were homogenized using bead mill homogenization (Omni Bead Ruptor Elite) and centrifuged at 2,950 \times g at 4°C. Thereafter, supernatant was transferred to 2-3 kDa spin filter and centrifuged again at $2,950 \times g$ at 4°C. Flow-through was collected and then analyzed via ultra-high-performance liquid chromatography with electrochemical detection (UHPLC-ED) using the 99 Dionex UltiMate 3000 with MD-TM Mobile Phase Solution as sample diluent (Fisher Scientific) as described previously (Villageliú et al., 2018b).

Microbiome High-Throughput Sequencing and Analysis

Total DNA was isolated from 0.25 g intestinal samples (ileal mucus, ceca mucus, ceca content) via bead-beating using the DNeasy PowerSoil Kit (Qiagen). Extracted DNAs were assessed for quality using a NanoDrop 2000 spectrophotometer 260–280 nm ratios. Concentrations were determined using a

Qubit fluorometer with the double-stranded DNA broad range kit (ThermoFisher Scientific) and adjusted to 50 ng/µL in nuclease-free water. Ileal content was initially-sampled, but insufficient DNA yield prevented further analyses. Library preparation was performed using the MiSeq and HiSeq2500 kit (Illumina) following all manufacturer's instructions with 151 × 151 paired-end MiSeq sequencing (Illumina). DNAs were then sequenced via Illumina MiSeq (v3) at the Iowa State DNA facility. For 16S rRNA analysis, QIIME2 (version 2019.10) was used to analyze the 16S rRNA data between all sequenced groups. Sequences were demultiplexed using the QIIME2 demux emp-paired function and denoised using the QIIME2 plugin DADA2. The number of good quality reads for taxonomic assignment ranged from 37,495 to 114,115 reads. GreenGenes database (version 13.8) at the 99% operational taxonomic units (OTUs) spanning the V4 and V5 regions (515F, GTGYCAGCMGCCGCGGTAA; 926R. CCGYCAATTYMTTTRAGTTT) was used to classify each of the reads using QIIME2's feature-classifier function. Alpha and beta diversity analyses were calculated using QIIME2's built-in functions. Mmvec plugin (Morton et al., 2019) was used to explore taxonomic balances and taxonomic group differences between treatment groups. T tests were used to determine statistical significance between groups (see Section 2.10 for further details).

Statistical Analyses

GraphPad Prism software version 6.0 (San Diego, CA) was used to calculate significance and construct figures for all analyses. Parametric, unpaired t-tests (two-tailed; $\alpha = 0.05$) were used to compare differences between CON and VR groups. For bactericidal assays, where samples were pooled, assays were performed in triplicate per pool (n = 6 per treatment group). For 16S rRNA bacterial abundances and neurochemical data, differences were compared within sample site (ex: content, mucus, or tissue from ceca or ileum). For all statistical tests, data sets for each group were assumed to follow a normal distribution. *p* values <0.05 were considered significant.

RESULTS

Welfare Conditions

General welfare assessment has shown no significant differences in body weight between control ($1.15 \text{ kg} \pm 0.066$) and VR ($1.15 \text{ kg} \pm 0.063$) groups (**Supplementary**

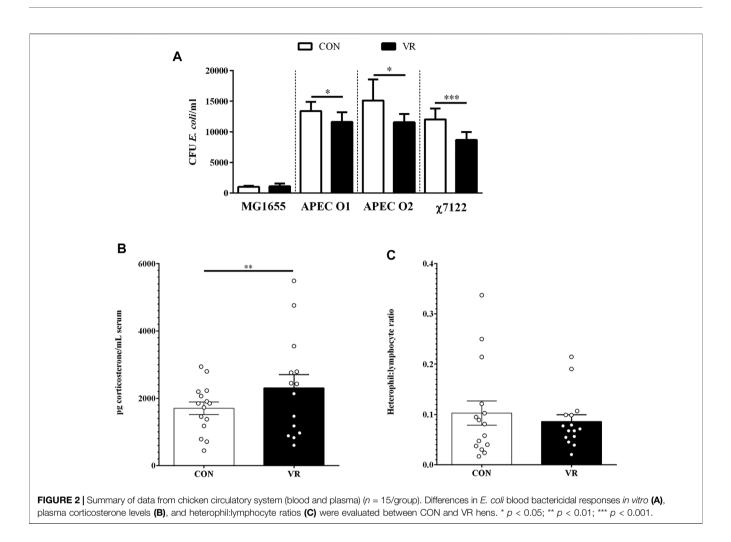


Figure S1). Additionally, birds in both treatment groups exhibited no signs of injury from negative behaviors like pecking.

Blood *E. coli* Killing Was Increased in VR Hens and Was Correlated With Corticosterone Level

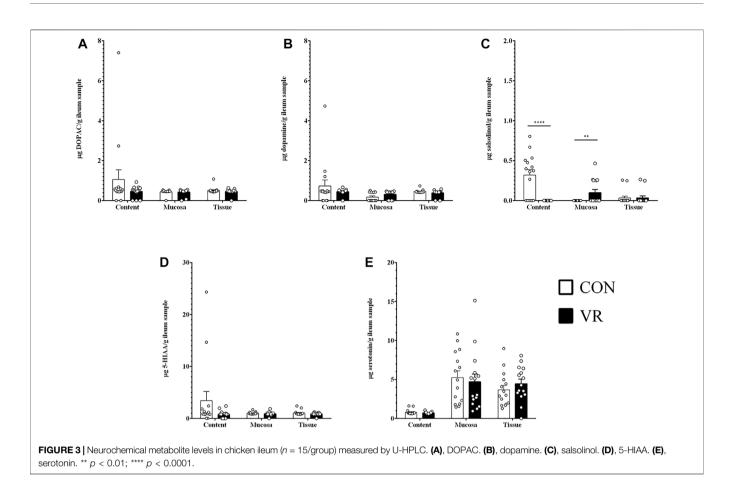
Using a blood bactericidal assay to test resistance to avian pathogenic *E. coli* (APEC), we found that blood from VR hens exhibited higher APEC killing to the three APEC serotypes O1, O2, and O78 tested compared to CON hens (p < 0.05; **Figure 2A**). This phenotype was positively-associated with an increase in serum corticosterone levels (p < 0.01; **Figure 2B**). However, H:L ratio was not significantly different between treatment groups (p > 0.05; **Figure 2C**).

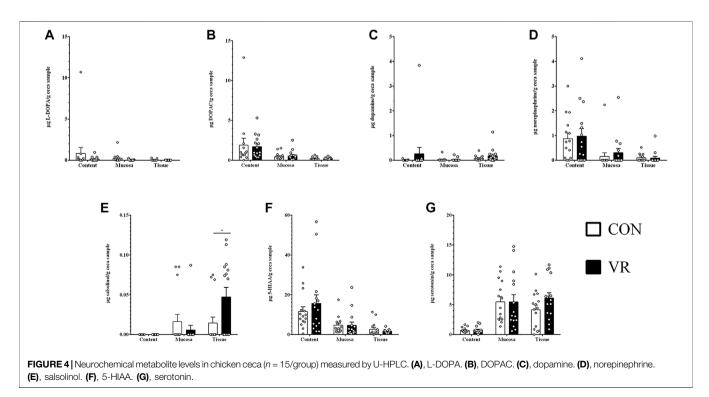
Salsolinol Was the Only Neurochemical Altered by VR Exposure

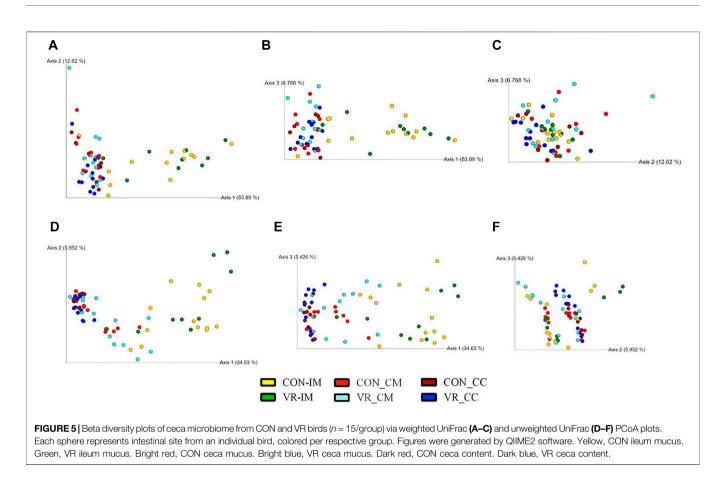
Ultra-high pressure liquid chromatography (U-HPLC) data for neurochemicals and related metabolites in the content, mucus, and tissue of the chicken gut are summarized in **Figure 3** (ileum) and **Figure 4** (ceca). In the ileum, all detected neurochemicals were not significantly different in each site except for salsolinol (**Figure 3C**), which was higher in the content of CON hens (p < 0.0001) yet higher in mucus of VR hens (p < 0.01). Similarly, only salsolinol levels changed in the ceca between groups (**Figure 4E**), although differences were only seen in the tissue with VR hens having greater salsolinol levels (p < 0.05).

VR Exposure Induced Shifts in Abundances of Gut Microbes

Using 16S rRNA sequencing and QIIME2 analysis, microbiome alpha diversity (Faith PD, Pielou's Evenness) was similar between groups (**Supplementary Figure S2**). Similarly, microbiome beta diversity was highly similar between groups at each intestinal site, although samples from the ileum mucus clustered distinctly along Axis 1 away from other intestinal samples in every beta diversity measure (**Figures 5A-C**, Weighted Unifrac; 5D-F, Unweighted UniFrac; 6A-C, Jaccard; 6D-E, Bray-Curtis). Looking at bacterial abundances, several taxa were consistently lower in VR hens compared to CON. For







example, looking at ceca genera, *Megamonas* were lower in the mucus and content (p < 0.01), whereas *Ruminococcus* and *Slackia* were lower in content alone (p < 0.05; **Figure 6A**). Looking at the species level, *Barnesiella viscericola* and *Blautia producta* were reduced in the ceca mucus of VR hens (p < 0.05; **Figure 6B**). Uniquely, *Mucispirillum schaedleri* was the only taxon higher in VR hens, being greater in ileum mucus versus CON hens (p < 0.05; **Figure 7B**). Notably, these shifts did not have any impacts on short chain fatty acid (SCFA) production (**Supplementary Figure S3**).

Several Correlations Between Microbial Taxa and Neurochemicals Were Detected

To explore the possibility that specific bacterial taxa may be related to intestinal neurochemical synthesis, we used the QIIME2 plugin mmvec (Morton et al., 2019) to draw relationships between neurochemical and microbiome data from the ileum mucus, ceca mucus, and ceca content. Using an Emperor plot to display microbial or metabolite co-occurrences, two phyla co-occurrences (1: *Firmicutes* and *Cyanobacteria*; 2: *Tenericutes*, *Deferribacteres*, and *Euryarchaeota*) appear frequently along Axis 1 (**Supplementary Figure S4A**). These co-occurrences are amplified when plotted against Axis 3 (**Supplementary Figure S4B**) versus Axis 2 (**Supplementary Figure S4A**). Similarly, norepinephrine and homovillanic acid appear to frequently co-occur regardless of axes (**Supplementary Figure S4**). Lastly, to observe microbial-metabolite

co-occurrences, inferred log conditional probabilities (LCPs) of microbial correlations with neurochemicals are displayed as a heatmap in Figure 8 and numerically summarized in Supplementary Table S1, in which positive probabilities are distinguished as red whereas low and negative values (i.e., light red to dark blue) indicate no relationship, not necessarily a negative relationship (see Github mmvec page: https://github.com/ biocore/mmvec). L-DOPA and norepinephrine are positively related to Cyanobacteria (LCP = 1.52) and Euryarchaeota (LCP = 1.78), respectively. Dopamine and its waste metabolite salsolinol are positively related with Deferribacteres (LCP = 1.51 and 1.73, respectively) and Tenericutes (LCP = 1.69 and 1.71, respectively). Serotonin (5-HT) and its waste metabolite 5-hydroxyindoleacetic acid (5-HIAA) are positively related with Bacteroidetes (LCP = 3.19 and 2. 05, respectively), Cyanobacteria (LCP = 3.43 and 2.33, respectively), Firmicutes (LCP = 3.30 and 2.63, respectively), and Proteobacteria (LCP = 2.74 and 2.07, respectively), with serotonin being specifically related with Spirochaetes (LCP = 1.71) and Tenericutes (LCP = 1.72). Uniquely, homovillanic acid and DOPAC are not related to any bacterial taxa at the phylum level.

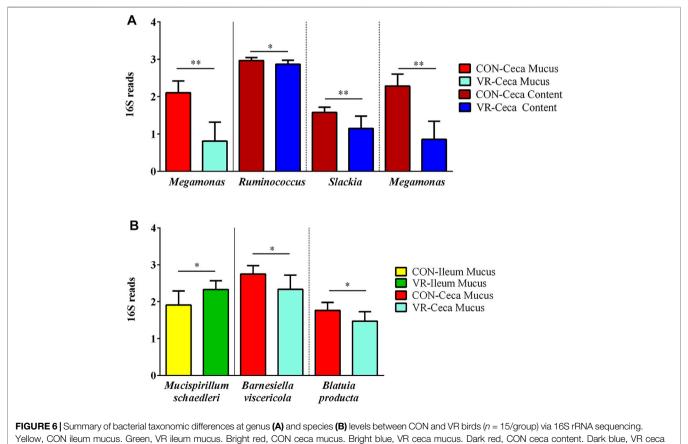
DISCUSSION

In this study, we tested whether VR video in chickens' pens could improve hens' welfare and health. Five-day-exposure to a VR video of free-range hens induced changes in several

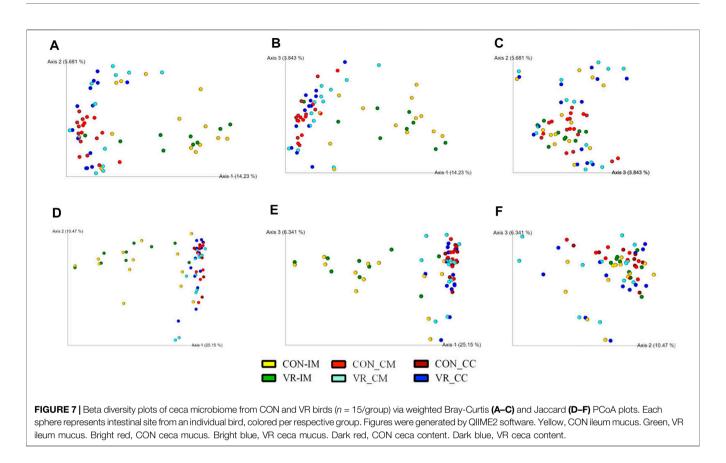
biological parameters in onset-lay hens. Given that layer hens at this stage (15 weeks post-hatch) are moved from pullet to egg-laying facilities and are plausibly at greater risk for stress (Janczak and Riber, 2015), we used this period to model how VR could be implemented in a commercial practice. Previous studies have demonstrated a positive correlation between an exposure to a video image of conspecific animals with increased food intake and positive social behaviors in chickens (Keeling and Hurnik, 1993). However, our study did not find an effect of a short exposure period to VR on chickens' body weight, as both VR and control had the same body weight (~1.15 kg) and similar levels of stress-related catecholamines like norepinephrine and epinephrine. Furthermore, although corticosterone level was higher in VR hens, corticosterone concentrations from both groups, ranging from <1 to nearly 6 ng corticosterone/ml serum, were in-line for what has been observed in unstressed hens, which have been reported to exceed over 20 ng corticosterone/ml serum (Scanes, 2016). Thus, it appears our pen design itself may have minimized stress in both animal groups. The little effect of VR on chickens could be due to the short duration of exposure (5 days) to VR videos. Previous work has shown that the interest to the video image by experimental birds is gradual. At Day 1, hens avoid looking at the screen,

whereas hens demonstrate an increased interest by Day 8 (Clark and Jones, 2001). Future studies should test longer exposure time to VR and consider the incorporation of our VR construct with hens from different stages (*i.e.*, pullet stage) to determine if VR exposure is more influential at certain ages.

Reducing stress could also affect immunological responses against bacterial pathogens like APEC (Mellata, 2013). Given the interconnectedness between the animal brain, gut, and its gut microbiota (Clarke et al., 2013; Yano et al., 2015; Valles-Colomer et al., 2019), we hypothesized that VR stimuli may modulate neurochemical metabolism and microbiota composition in the gut. Most VR studies focus on behavioral and neuroscience changes (Bohil et al., 2011); therefore, we focused on how VR affected the immune system and microbiota of hens. One notable finding was that on the fifth 5 day of VR exposure, blood from VR hens had increased killing of APEC in vitro compared to that of control birds. Furthermore, this observation was positivelyassociated with an increase in serum corticosterone. Although high glucocorticoid concentrations (i.e., > 9 ng/ml) induce immunosuppression (Davison et al., 1988; El-Lethey et al., 2003; Shini et al., 2010), acute exposure to moderate levels of cortisol (5 ng/ml) in bovine endometrial epithelial cells



content. * p < 0.05; ** p < 0.01.



increase inflammatory responses against *E. coli* when compared to cells not conditioned with cortisol (Cui et al., 2020). Thus, the moderate increase in corticosterone could have primed immune cells to allow a greater APEC-bactericidal response. Since our VR model only lasted for 5 days, it is not clear if continued exposure to these visual stimuli would result in chronically-elevated corticosterone, which might have deleterious effects on immune function (Coutinho and Chapman, 2011).

To our knowledge, no study has investigated whether exposure to VR can alter the gut microbiota. Here, almost every taxon with significantly-changed relative abundances were higher in CON versus VR hens, regardless of intestinal site. Many of these taxa (ex: Megamonas, Blautia, and Ruminococcus) are major producers of SCFAs (Takahashi et al., 2016; Louis and Flint, 2017; Zhao et al., 2017), which confer numerous benefits such as pathogen exclusion and immune activation (Parada Venegas et al., 2019). However, despite these microbiome changes, we did not identify any changes in SCFAs in any intestinal site. Thus, it appears that changes in these SCFA-producing populations did not have a functional impact on VR hens in terms of SCFA production at the time point tested. Additionally, a single taxon, M. schaedleri, was higher in the ileal mucus of VR hens relative to CON. M. schaedleri, the sole member of the Deferribacteres phylum detected in this study, is an obligate anaerobic bacterium, which colonizes (but does not degrades) host

mucus (Loy et al., 2017) and is associated with several inflammatory pathologies in mammals (Loy et al., 2017) and fat deposition in poultry (Wen et al., 2019). At this time, it is unclear how VR treatment increases M. schaedleri abundances in the ileum. However, we did observe a strong, positive relationship between Deferribacteres (i.e., M. schaedleri) and salsolinol-dopamine in the intestine. Salsolinol is a dopaminergic waste metabolite (Yamanaka et al., 1970), which has gained recent attention given its role in Parkinson's disease in mammals (Naoi et al., 2002). Altogether, these findings suggest that M. schaedleri may either produce salsolinol directly [as demonstrated by other microbes (Villageliú et al., 2018a)] or positively-responds to its presence in the intestine. However, the changes to both M. schaedleri and salsolinol production are driven by VR exposure, and future studies will investigate the mechanisms in which these visual stimuli drive such changes in the chicken intestine.

In addition to the relationship between *Deferribacteres* and salsolinol-dopamine, we identified several other positive relationships between microbial phyla and neurochemicals. We measured these relationships using co-occurrence probabilities (*i.e.*, the QIIME2 mmvec plugin) versus correlative measures like Pearson's, which are too simplistic for compositional data due to high false-discovery rates (Morton et al., 2019). Similar to *Deferribacteres, Tenericutes* abundances exhibited high relatedness with salsolinol and dopamine, suggesting that

members of these two phyla may collaborate to directly or indirectly facilitate dopamine-metabolism to salsolinol. Notably, we also identified several phyla (*Bacteriodetes*, *Cyanobacteria*, *Firmicutes*, *Proteobacteria*, *Spirochaetes*, and *Tenericutes*) positively-related to serotonin in the chicken intestine. In mice, spore-forming gut microbes are crucial for host serotonin synthesis in the gut (Yano et al., 2015), the source of roughly 95% of the body's total serotonin (Gershon and Tack, 2007). Given that several *Firmicutes* form endospores (Galperin, 2013) and the positive relationship between *Firmicutes* and serotonin in our study, we speculate a similar relationship may exist in chickens. However, it is unclear whether these microbes are directly producing these neurochemicals (Lyte, 2014), influencing host neurochemical synthesis (Yano et al., 2015), or are responding to these neurochemicals in the gut via activities like virulence gene expression (Freestone et al., 2008).

In conclusion, we found host resistance to *E. coli*, corsticosterone levels, neurochemical metabolism, and gut microbiota composition were all affected by VR exposure in laying hens. Importantly, this study is the first to demonstrate that VR exposure can modulate the gut microbiota, which was related to the production of several neurochemicals and associated metabolites in the gut. Current work is underway to look at whether more-complex animal behaviors were either affected by VR in general or if birds were particularly-responsive to certain scenes in these VR videos.

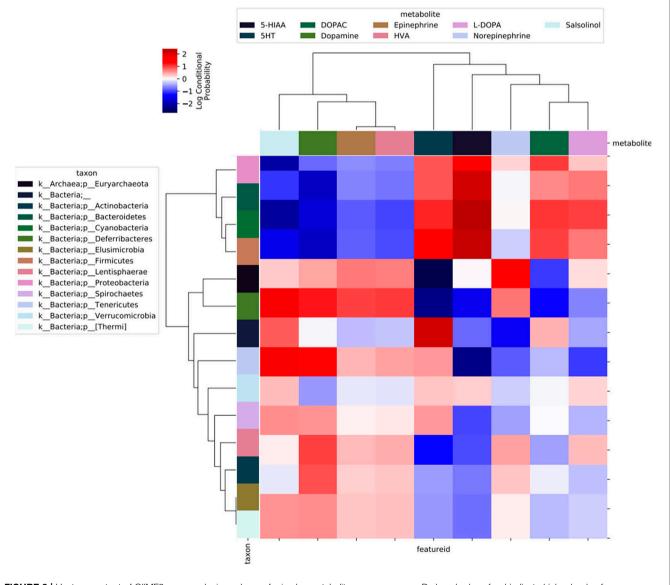


FIGURE 8 | Heatmap output of QIIME2 mmvec plugin analyses of microbe-metabolite co-occurrences. Darker shades of red indicate higher levels of cooccurrences between bacterial phyla and neurochemical metabolites.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found below: https://www.ncbi. nlm.nih.gov/, PRJNA638088.

ETHICS STATEMENT

The animal study was reviewed and approved by Iowa State University Institutional Animal Care and Use Committee, log #19-211. Written informed consent was obtained from the owners for the participation of their animals in this study.

AUTHOR CONTRIBUTIONS

MM and JO conceived the project; GR, SM, RR, ML, JO, and MM designed the experiments. GR, RP, AH, KD, and MM performed the experiments and analyzed the data. GR and MM wrote the manuscript. SM, ML, and JO revised the manuscript. MM, SM, RR, and ML contributed reagents, materials, and analysis tools. All authors read and approved the final manuscript.

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SUPPLEMENTARY MATERIAL

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Conflict of Interest: GR, SM, RR, ML, JO, and MM are listed as inventors on a pending patent covering research presented in this paper.

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