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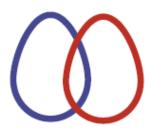
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Association of VIPR-1 Gene Polymorphism and CRBP-IV Gene Expression with Layer Traits in Rhode Island Red Chicken

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The investigation was carried out to determine polymorphism at VIPR-1 gene and mRNA expression of CRBP-IV gene in various tissues in Rhode Island Red (RIR) chicken. Hundred eleven pullets belonging to five hatches were studied. Data on BW20 and layer traits were analyzed by least squares taking sire as random and hatch as fixed effects. Average ASM and EP40 were 135.19±1.15 days and 124.55±1.94 eggs, respectively. Hatch effect was significant (P<0.05) on ASM, EW28 and EP40. BW20 revealed low, but positive genetic and phenotypic correlations with EP40. Allelic profiles of VIPR-1 at its three loci were performed using Tail, Hhal and Tagl restriction enzymes. Digestion was monomorphic for Tail and Hhal and polymorphic for Tagl with allelic frequency 0.98 (A) and 0.02 (B). Genotypic frequencies for AA and AB were 0.96 and 0.04, respectively; but did not show any significant effect on layer traits. Relative mRNA expression of CRBP-IV was studied in kidney, liver and oviduct tissues in four different egg production-body weight groups by gRT-PCR and data was analyzed using JMP of SAS, CRBP-IV expression revealed significant difference among different egg production-body weights groups only in kidney (P≤0.05). Expression of CRBP-IV was also differed significantly (P≤0.0001) among tissues and highest expression was observed in oviduct. This investigation generated information related to VIPR-1 gene polymorphism and analysis with large sample size may yield significant findings. Further analysis of CRBP-IV expression at different ages and also in various tissues would help in exploitation of selection for higher reproductive efficiency in poultry.

Keywords: Candidate genes, Genotyping, Layer traits, mRNA expression, RIR chicken

Using high throughput phenotyping of growth and feed intake to improve adaptation of chickens to sustainable diets

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To refine selection of broilers on feed efficiency, we developed an electronic feed station allowing a continuous record of body weight, feed intake and feeding behavior of animals reared on floor. Animals can express their natural feeding behavior as they are raised as a group. Using this station, we compared growth and feed efficiency traits of two genotypes (Label Rouge and standard broilers) fed with two diets (corn-soybean or an alternative diet including more sustainable feedstuffs) which were adapted to the nutritional needs of each genotype.

Diet change had moderate effect on growth and feed efficiency even if the response differed between the two genotypes. Indeed, Label Rouge chickens fed with the alternative diet grew faster (+ 3,3% at slaughter), ate more (+1,1%) and had a higher FCR (+2,6%) than those fed with the corn-soybean diet. At the opposite, standard chickens fed with the alternative diet grew less (-1,7%), ate less (-4,6%) and had a lower FCR (-2,8%) than those fed with the classic diet.

Theses first results indicate that animals can adapt to alternative feedstuffs, even those with a rapid growth rate. The variability of response to these alternative diets will now be tested within-genotype in order to check whether some feeding behavior or continuous feed efficiency traits can be used in selection.

Keywords: High throughput phenotyping, Feed efficiency, Feeding behavior, Growth, Alternative diet