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Residual feed intake in late-laying hens: immune function, metabolic efficiency, and feed utilization dynamics

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The selection of high feed-efficiency animals is essential to address the increasing global demand for animal-derived products while ensuring sustainability. Residual Feed Intake (RFI), a crucial metric in poultry production, enhances feed utilization, optimizes management strategies, and promotes economic viability and environmental stewardship. However, the mechanisms underlying RFI variation remain inadequately understood. This study examined the regulatory pathways of RFI in 70-week-old Rhode Island Red laying hens through comparative analysis of phenotypic parameters, organ characteristics, and serum biochemical profiles between low-RFI (LRFI) and high-RFI (HRFI) groups. The findings demonstrate that RFI functions as a reliable indicator for feed efficiency selection, with LRFI hens demonstrating enhanced immune modulation and metabolic homeostasis while maintaining equivalent egg production performance. These results establish fundamental insights for understanding RFI regulatory mechanisms and developing precision breeding strategies for feed-efficient laying hens.

KEYWORDS

residual feed intake, late laying hens, immune function, metabolic function, feed efficiency

1 Introduction

In livestock and poultry production, feed costs constitute 60–70% of total operational expenditures, with rising feed prices and environmental concerns creating significant challenges to industry sustainability (1). Although China has achieved substantial progress in systematic breeding programs for high-yield laying hens – producing over 300 eggs per hen by 72 weeks of age – significant challenges remain in feed utilization efficiency. The average feed-to-egg ratio of 2.2–2.8:1 remains suboptimal compared to other poultry sectors (broilers, meat ducks), particularly during the late laying phase when metabolic demands increase, resulting in excessive feed waste and reduced profitability (2, 3). Residual Feed Intake (RFI), an internationally recognized metric for individual animal feed efficiency evaluation, has emerged as a vital parameter for optimizing resource allocation (4). Improving RFI-based selection strategies represents a key priority for enhancing feed conversion efficiency and promoting sustainable growth in the poultry industry.

RFI is quantitatively defined as the difference between actual feed consumption and predicted intake based on metabolic body weight, body weight gain, and egg production output, functioning as a heritable biomarker of metabolic divergence in poultry (5). This metric effectively measures species-specific energy requirements for maintenance and growth

while remaining phenotypically independent of economically valuable traits. Studies in Korat chickens demonstrate minimal correlations between RFI and meat quality parameters (6, 7). In layer breeding systems, selection for low-RFI hens improves feed conversion efficiency without compromising egg quality or growth performance, while reducing environmental nitrogen excretion by 8–12% and operational costs by 15–18% per production cycle (8). These characteristics establish RFI as an effective selection criterion for sustainable genetic improvement in feed utilization efficiency.

The regulation of RFI represents a complex biological process controlled by coordinated interactions among feeding behavior, nutrient assimilation efficiency, and metabolic pathway dynamics (9, 10). Quantitative genetic analyses indicate moderate to high heritability estimates ($h^2 = 0.25 - 0.45$ across poultry species), with genome-wide association studies identifying 12-18 candidate genes associated with RFI variation in layers (11). Selection programs utilizing RFI optimization demonstrate substantial reductions in broiler feed intake and abdominal fat deposition while maintaining carcass yield and intramuscular fat content (12). Comparative organometrics reveal lower liver mass in low-RFI hens, indicating enhanced basal metabolic rates and energy partitioning efficiency (13). Negative selection for RFI has been successfully implemented across multiple livestock species, consistently achieving feed cost reductions while preserving growth trajectory integrity (14). These multispecies validations confirm RFI's effectiveness as a selection index for improving feed efficiency in late-laying phase hens through metabolic optimization rather than production trait compromises.

This study aims to examine differences in growth performance, organ characteristics, and serum biochemical parameters between hens with high-RFI (HRFI) and low-RFI (LRFI) during the late laying phase, thereby establishing reference values for comprehensive understanding of RFI regulatory mechanisms and the breeding selection of high feed efficiency laying hen varieties.

2 Materials and methods

2.1 Birds, housing, diets and experiment groups

The experimental hens were derived from a Rhode Island Red pure-line population provided by China Beinongda Technology Co., Ltd. All subjects were individually housed in separate cages under standardized environmental conditions with ad libitum access to feed and water. The basal diet was formulated according to National Research Council (NRC) guidelines (Table 1). All hens maintained optimal health status throughout the trial, which was conducted in strict compliance with protocols approved by the Animal Welfare Committee of China Agricultural University (Approval AW62801202-1-1). After sorting in ascending order of RFI values, the top 10%/bottom 10% of individuals were selected and categorized into HRFI and LRFI groups (n = 6 per group). At trial termination (70-week age), hens were humanely euthanized for biological sampling and subsequent physico-chemical analyses.

TABLE 1 Diet's composition and nutrient contents (%).

Items	Content			
Ingredient				
Corn, %	61.06			
Soybean meal, %	23.80			
Soybean oil, %	0.80			
Wheat bran, %	3.00			
Dicalcium phosphate, %	0.90			
Stone powder, %	9.00			
Salt, %	0.40			
Antioxidant, %	0.04			
Antifungal agent, %	0.05			
Premix ¹ , %	0.05			
Total	100			
Nutrient composition				
AME (MJ/kg)	11.13			
Crude protein, %	16.13			
Methionine, %	0.36			
Lysine, %	0.87			
Calcium, %	3.71			
Phosphorus, %	0.58			

¹Premix: Vitamin premix provided/kg of diet: vitamin A, 10,000 IU; vitamin D₃, 3,000 IU; vitamin E, 30 IU; vitamin K₃, 4 mg; vitamin B₁, 3 mg; vitamin B₂, 8 mg; vitamin B₆, 6 mg, vitamin B₁₂, 0.03 mg, niacin, 40 mg, folic acid, 2 mg, biotin, 0.3 mg, pantothenic acid, 18 mg. Mineral premix provided/kg of diet: Mn 100 mg, zinc 100 mg, Fe 65 mg, Cu 9 mg, I 1 mg, Se 0.3 mg.

2.2 Sample collection and analyses

2.2.1 Measurement of production and feed efficiency traits

During weeks 67–69 of age, RFI and production traits were measured in this investigation. The evaluated parameters included body weight (BW), body weight gain (BWG), egg number (EN), egg mass (EM), and feed intake (FI). The following measurement protocols were implemented:

BW: Initial and final BW measurements were obtained using a digital platform scale (0.1 g precision), with the average value calculated as metabolic body weight (MBW).

BWG: This parameter was determined by subtracting initial BW from final BW, with daily BWG calculated by dividing the total gain by the 21-day experimental period.

Egg Production: Individual EN was systematically documented throughout the experimental duration.

EM: Daily measurements for each hen were recorded using a precision electronic balance (0.01 g accuracy), with values averaged to establish mean egg weight (EW).

FI: Individual intake monitoring utilized partitioned feeding troughs with metal baffles. Daily feed provision was weighed, with daily feed intake (DFI) derived from 21-day cumulative consumption.

 $RFI = TFI - (b_0 + b_1MBW + b_2DEM + b_3DWG).$

where TFI represents intake from 69 to 71 weeks of age, b_0 denotes the intercept, and b_1 , b_2 and b_3 represent partial regression coefficients.

2.2.2 Organ indices quantification

After euthanasia through jugular exsanguination, hepatic and splenic tissues were carefully extracted. Adherent adipose tissue was removed from organ surfaces, and excess blood was eliminated through filter paper blotting before gravimetric measurement. Organ indices were determined using the standardized formula:

Organ index (%) = (fresh tissue weight / live body weight $) \times 100$

2.2.3 Blood sampling collection

Blood samples were collected from the brachial vein at 70 weeks of age using sterile vacutainers without anticoagulants. The samples were centrifuged at 3,000 rpm for 10 min (4°C) to obtain serum separation. The separated serum was transferred to pre-chilled aseptic cryovials and stored at -20° C for subsequent biochemical analysis.

2.2.4 Serum biochemical profiling

Beijing Jinhai Keyu Biotechnology Co., Ltd. conducted serum analysis using an automated biochemistry analyzer (Kehua ZY KHB-1280) to measure seven metabolic indicators: total protein (TP), albumin (ALB), globulin (GLO), blood urea nitrogen (BUN), creatinine (CRE), uric acid (UA), and lactate dehydrogenase (LDH).

2.2.5 Serum immunological profiling

ELISA kits (Beijing Jinhai Keyu Biotechnology Co., Ltd.) were utilized to quantify serum immune parameters including: Pro-inflammatory cytokines: Interferon- γ (IFN- γ), Tumor Necrosis Factor- α (TNF- α), Interleukin-1 β (IL-1 β), Anti-inflammatory cytokines: IL-4, IL-6, IL-10, Endotoxin: Lipopolysaccharide (LPS), Immunoglobulin isotypes (IgA, IgG, IgM) were analyzed.

2.2.6 Serum Metabolic profiling

Circulating concentrations of insulin (INS), adiponectin (ADP), and insulin-like growth factor-1 (IGF-1) were measured using commercial reagent kits (Beijing Jinhai Keyu Biotechnology Co., Ltd.).

2.3 Data analysis

Data analysis was performed using Microsoft Excel 2019, and statistical evaluation was conducted through Student's t-test using SPSS 23.0 (IBM Corp.). Results are presented as mean \pm standard error of the mean (SEM). Statistical significance levels were defined as follows: non-significant (p > 0.05), significant (p < 0.05), and highly significant (p < 0.01).

3 Results

3.1 Production performance differences between LRFI and HRFI groups

As shown in Table 2, total egg production (310.17 vs. 310.33 eggs) and terminal body weight (3046.83 vs. 3010.17 g) showed no significant differences (p > 0.05) between LRFI and HRFI groups at 70 weeks of age. However, RFI values demonstrated significant divergence (-105.00 vs. 99.83 g/d, p < 0.001), establishing RFI as the primary differentiation factor while maintaining comparable egg production and terminal body weight conditions.

3.2 Visceral organ index variations between LRFI and HRFI groups

As demonstrated in Table 3, HRFI hens displayed significantly higher hepatic indices compared to LRFI hens (1.26% vs. 1.05%, p < 0.05), whereas splenic indices showed no statistical difference between the groups (0.08% vs. 0.08%, p > 0.05).

3.3 Serum biochemical profile comparisons between LRFI and HRFI groups

Table 4 shows no significant differences between groups (p > 0.05) in serum biochemical parameters, including TP, ALB, GLO, BUN, CRE, UA, and LDH.

3.4 Differential serum immune profiles between LRFI and HRFI groups

As illustrated in Table 5, LRFI hens exhibited significantly higher IL-4 (171.67 vs. 140.69 pg./mL) and IgG concentrations (10.17 vs. 7.88 g/L), while displaying lower LPS (34.34 vs. 41.55 U/L) and IL-1 β

TABLE 2 Comparison of phenotypic data of hens in late egg production with RFI.

Traits ¹	LRFI	HRFI	SEM	<i>p</i> -value
EN ₇₀	310.17	310.33	1.462	0.912
BW ₇₀	3046.83	3010.17	27.130	0.224
RFI ₆₇₋₆₉ (g/d)	-105.00 ^b	99.83ª	5.449	< 0.001

¹EN₇₀, Total egg number at 70 weeks of age; BW₇₀, Body weight at 70 weeks of age; RFI₆₇₋₆₉, Residual feed intake from 67 to 69 weeks of age. Values are presented as the mean and SEM (n = 6, 70 weeks of age), and different letters indicate significant differences.

TABLE 3 Comparison of organ index of hens in late egg-laying period with RFI.

Traits ¹	LRFI	HRFI	SEM	<i>p</i> -value
Liver index, %	1.05 ^b	1.26 ^a	0.032	0.001
Spleen, %	0.08	0.08	0.006	0.479

¹Values are presented as the mean and SEM (n = 6, 70 weeks of age), and different letters indicate significant differences.

TABLE 4 Comparison of serum biochemical indices in late egg-laying period with RFI.

Traits ¹	LRFI	HRFI	SEM	<i>p</i> -value
TP, g/L	51.08	51.24	1.736	0.926
ALB, g/L	36.45	35.55	3.018	0.771
GLO, g/L	24.95	24.08	2.060	0.682
BUN, mg/L	3.13	3.11	0.339	0.946
CRE, µmol/L	20.27	19.47	0.874	0.379
UA, μmol/L	360.19	364.01	28.591	0.896
LDH, U/L	820.81	803.67	35.112	0.641

¹TP, Total Protein; ALB, Albumin; GLO, Globulin; BUN, Blood Urea Nitrogen; CRE, Creatinine; UA, Uric Acid; LDH, Lactate Dehydrogenase. Values are presented as the mean and SEM (n = 6, 70 weeks of age).

TABLE 5 Comparison of serum immunity indices of hens in late egg production with RFI.

Traits ¹	LRFI	HRFI	SEM	<i>p</i> -value
IFN-γ, pg./mL	101.84	102.48	5.403	0.908
TNF-α, pg./ mL	74.28	78.30	3.435	0.270
LPS, U/L	34.34 ^b	41.55ª	2.114	0.007
IL-4, pg./mL	171.67ª	140.69 ^b	10.006	0.011
IL-6, pg./mL	44.19	44.37	4.102	0.966
IL-10, pg./mL	69.89	72.04	4.195	0.620
IL-1β, pg./mL	12.69 ^b	16.68ª	1.508	0.025
IgA, g/L	1.04	1.05	0.068	0.887
IgG, g/L	10.17ª	7.88 ^b	0.809	0.017
IgM, g/L	0.69	0.74	0.055	0.386

¹Values are presented as the mean and SEM (n = 6, 70 weeks of age), and different letters indicate substantial variations.

levels (12.69 vs. 16.68 pg./mL) compared to their HRFI counterparts (p < 0.05). No significant differences (p > 0.05) were detected in IFN- γ , TNF- α , IL-6, IL-10, IgA, and IgM concentrations between the groups.

3.5 Serum endocrinometabolic divergences between LRFI and HRFI groups

Table 6 demonstrates significantly higher serum glucose levels in HRFI hens compared to their LRFI counterparts, whereas ADP and IGF-1 concentrations exhibited no significant differences between groups.

4 Discussion

4.1 Impacts of RFI on growth performance, organ indices, and feed efficiency traits in late-laying hens

Feed costs constitute the primary economic factor in poultry production, with RFI functioning as a linear breeding metric whose

TABLE 6 Comparison of serum endocrine metabolism indexes in late laying hens with RFI.

Traits ¹	LRFI	HRFI	SEM	<i>p</i> -value
GLU, mmol/L	10.16	11.64	0.514	0.016
ADP, µg/L	18.03	17.09	1.478	0.543
IGF-1, ng/ml	118.16	120.49	6.856	0.741

¹GLU, Glucose; ADP, Adiponectin; IGF-1, Insulin-like Growth Factor 1. Values are presented as the mean and SEM (n = 6, 70 weeks of age), and different letters indicate substantial variations.

component trait weights are determined by biological variance (15). Evidence across livestock species [cattle (16), pig (17), poultry (5), ducks (18)] demonstrates that RFI-based selection effectively reduces feed consumption while maintaining production performance and egg/meat quality parameters, indicating its principal influence on FI and feed conversion ratio (FCR). In layer breeding programs where genetic antagonism exists between key selection traits - particularly the negative correlation between egg weight and production quantity -RFI emerges as an effective solution (19). Yuan et al. documented 8-12% FI reduction in low-RFI layers without altered egg production at 37-40 weeks of age (20), while Zhang et al. (21) reported negligible correlation between RFI and daily weight gain in Pekin ducks. The present findings align with these studies, demonstrating equivalent cumulative egg production between high and low RFI groups despite significant RFI divergence. These results validate low-RFI selection as an effective strategy for enhancing feed efficiency in late-laying hens without compromising egg production or growth performance.

The liver functions as the central hub for nutrient metabolism and storage, with hepatic index (liver-to-body weight ratio) variations reflecting hens' nutritional utilization efficiency and directly influencing laying performance (22, 23). Elevated hepatic indices typically indicate pathological lipid infiltration and metabolic dysregulation, predisposing to conditions like fatty liver hemorrhagic syndrome (24). The spleen's immunological functions-including pathogen filtration and leukocyte production-make splenic indices essential biomarkers of disease resistance capacity (25). The data revealed significantly lower hepatic indices in LRFI hens compared to HRFI counterparts, demonstrating improved hepatic metabolic efficiency through reduced lipid deposition and enhanced β -oxidation activity.

4.2 Immunological impacts of RFI in late-laying hens

Genetic and metabolic determinants of feed efficiency primarily localize to pathways governing carbohydrate/lipid/protein metabolism and ion transport, with substantial immunological cross-talk (26, 27). Enhanced immune competence enables hens to combat pathogens, maintain nutritional homeostasis, and regulate feed intake under environmental stressors (28). Serum analysis revealed distinct immunological profiles between RFI phenotypes: LRFI hens exhibited elevated IL-4 and IgG alongside reduced LPS and IL-1 β . Specifically, IL-4 potentiates B-cell differentiation for pathogen-specific antibody production (29), while IgG mediates humoral immunity through bacterial toxin neutralization and immunological memory (30). Reduced LPS levels indicate decreased bacterial endotoxin exposure (31), and lower IL-1 β reflects attenuated inflammatory responses via NF- κ B pathway regulation (32). This coordinated immunomodulation in LRFI hens - characterized by enhanced adaptive immunity and controlled inflammation-optimizes disease resistance while maintaining metabolic resources for egg production, demonstrating RFI's dual role in feed efficiency and immunological fitness in aging layers.

4.3 Metabolic implications of RFI in late-laying hens

Systemic concentrations of blood metabolites serve as physiological biomarkers for feed efficiency, as demonstrated by the investigation into serum metabolic profiles of RFI-divergent hens (33, 34). The analysis revealed significantly lower circulating glucose levels in LRFI hens compared to HRFI counterparts, paralleling previous observations in ducks where low-RFI individuals exhibited reduced triglycerides (TG) and GLU concentrations (35). Significant positive correlations between RFI and both TG and GLU indicate intrinsic linkages between energy metabolism regulation and feed efficiency (36). This metabolic advantage in LRFI hens likely derives from decreased insulin sensitivity, upregulated hepatic glucokinase expression, and optimized gut microbiota composition, collectively maintaining postprandial glucose homeostasis while minimizing energy dissipation through futile cycling pathways.

5 Conclusion

This investigation confirms RFI as an effective selection criterion for improving feed conversion efficiency in laying hens, achieving reduced feed consumption while maintaining egg production and growth performance. Serum biomarker analysis revealed enhanced immunological and metabolic profiles in low-RFI hens. These findings establish a theoretical foundation for breeding feed-efficient layer strains through RFI optimization, supporting sustainable poultry production through improved metabolic homeostasis and prolonged productive longevity.

Data availability statement

The original contributions presented in the study are included in the article/supplementary material, further inquiries can be directed to the corresponding author.

Ethics statement

The animal studies were approved by Animal Welfare Committee of China Agricultural University (ApprovalAW62801202-1-1). The

References

1. Makkar H. Feed demand landscape and implications of food-not feed strategy for food security and climate change. *Animal.* (2018) 12:1744–54. doi: 10.1017/S175173111700324X

2. Quintana GA, Alfaro MC, Oviedo EO, Ruiz JR, Bernal LC, Martinez GD. Data analytics of broiler growth dynamics and feed conversion ratio of broilers raised to 35 d

studies were conducted in accordance with the local legislation and institutional requirements. Written informed consent was obtained from the owners for the participation of their animals in this study.

Author contributions

ZG: Writing – original draft, Writing – review & editing. ZM: Supervision, Writing – review & editing. LX: Supervision, Writing – review & editing. GM: Investigation, Writing – review & editing. YW: Formal analysis, Writing – review & editing. GX: Project administration, Validation, Writing – review & editing.

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Conflict of interest

ZM, GM, and YW were employed by Beinongda Technology Co., Ltd.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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under commercial tropical conditions. *Animals.* (2023) 13:2447. doi: 10.3390/ani13152447

3. Walk CL, Addo-Chidie EK, Bedford MR, Adeola O. Evaluation of a highly soluble calcium source and phytase in the diets of broiler chickens. *Poult Sci.* (2012) 91:2255–63. doi: 10.3382/ps.2012-02224

4. Prakash A, Saxena VK, Singh MK. Genetic analysis of residual feed intake, feed conversion ratio and related growth parameters in broiler chicken: a review. *World Poultry Sci J.* (2020) 76:304–17. doi: 10.1080/00439339.2020.1735978

5. Ramankevich A, Danko S, Banaszkiewicz R, Kasperek K, Zięba G. Residual feed intake as a behavioral, nutritional and economic criterion in poultry production. *Animals.* (2025) 15:1115. doi: 10.3390/ani15081115

6. Poompramun C, Mignon-Grasteau S, Juanchich A, Hennequet-Antier C, Thumanu K, Bihan-Duval EL, et al. Phenotypic correlations between feed efficiency, growth, and meat quality of slow-growing chickens. *Br Poult Sci.* (2022) 63:742–6. doi: 10.1080/00071668.2022.2094218

7. Poompramun C, Hennequet-Antier C, Thumanu K, Sinpru P, Pengsanthia S, Molee W, et al. Revealing pathways associated with feed efficiency and meat quality traits in slow-growing chickens. *Animals*. (2021) 11:2977. doi: 10.3390/ani11102977

8. Zerjal T, Härtle S, Gourichon D, Guillory V, Bruneau N, Laloë D, et al. Assessment of trade-offs between feed efficiency, growth-related traits, and immune activity in experimental lines of layer chickens. *Genet Sel Evol.* (2021) 53:44. doi: 10.1186/s12711-021-00636-z

9. Taussat S, Boussaha M, Ramayo-Caldas Y, Martin P, Venot E, Cantalapiedra-Hijar G, et al. Gene networks for three feed efficiency criteria reveal shared and specific biological processes. *Genet Sel Evol.* (2020) 52:1–14. doi: 10.1186/s12711-020-00585-z

10. Yang C, Ding Y, Dan X, Shi Y, Kang X. Multi-transcriptomics reveals RLMF axismediated signaling molecules associated with bovine feed efficiency. *Front Vet Sci.* (2023) 10:1090517. doi: 10.3389/fvets.2023.1090517

11. Yuan J, Wang K, Yi G, Ma M, Dou T, Sun C, et al. Genome-wide association studies for feed intake and efficiency in two laying periods of chickens. *Genet Sel Evol.* (2015) 47:1–13. doi: 10.1186/s12711-015-0161-1

12. Abebe BK, Wang J, Guo J, Wang H, Li A, Zan L. A review of emerging technologies, nutritional practices, and management strategies to improve intramuscular fat composition in beef cattle. *Anim Biotechnol.* (2024) 35:2388704. doi: 10.1080/10495398.2024.2388704

13. Metzler-Zebeli BU, Siegerstetter S, Magowan E, Lawlor PG, O Connell NE, Zebeli Q. Feed restriction reveals distinct serum metabolome profiles in chickens divergent in feed efficiency traits. *Meta*. (2019) 9:38. doi: 10.3390/metabo9020038

14. Datt C, Sharma VK, Dudi K, Baban BN, Sharma PS, Negesse T, et al. Residual feed intake as a tool for selecting more efficient animals: a review. *Indian J Anim Nutr.* (2017) 34:238–55. doi: 10.5958/2231-6744.2017.00041.X

15. Fathi MM, Galal A, Al-Homidan I, Abou-Emera OK, Rayan GN. Residual feed intake: a limiting economic factor for selection in poultry breeding programs. *Ann Agric Sci.* (2021) 66:53–7. doi: 10.1016/j.aoas.2021.03.001

16. Belous AA, Sermyagin AA, Zinovieva NA. Beef cattle evaluation by feeding efficiency and growth energy indicators based on bioinformatic and genomic technologies. *Agric Biol.* (2022) 57:1055–70. doi: 10.15389/agrobiology.2022.6.1055eng

17. Sánchez JP, Ragab M, Quintanilla R, Rothschild MF, Piles M. Genetic parameters and expected responses to selection for components of feed efficiency in a Duroc pig line. *Genet Sel Evol.* (2017) 49:86. doi: 10.1186/s12711-017-0362-x

18. Zeng T, Zhang H, Liu J, Chen L, Tian Y, Shen J, et al. Genetic parameters of feed efficiency traits and their relationships with egg quality traits in laying period of ducks. *Poult Sci.* (2018) 97:758–63. doi: 10.3382/ps/pex337

19. Thiruvenkadan AK, Panneerselvam S, Prabakaran R. Layer breeding strategies: an overview. *World Poultry Sci J.* (2010) 66:477–502. doi: 10.1017/S0043933910000553

20. Yuan J, Chen S, Shi F, Wu G, Liu A, Yang N, et al. Genome-wide association study reveals putative role of gga-miR-15a in controlling feed conversion ratio in layer chickens. *BMC Genomics*. (2017) 18:1–11. doi: 10.1186/s12864-017-4092-9

21. Zhang Y, Guo ZB, Xie M, Zhang Z, Hou S. Genetic parameters for residual feed intake in a random population of Pekin duck. *Asian Australas J Anim Sci.* (2016) 30:167. doi: 10.5713/ajas.15.0577

22. Mohajan HK. A study on functions of liver to sustain a healthy liver. *Innov Sci Technol.* (2025) 4:77–87. doi: 10.56397/IST.2025.01.08

23. Navarro-Villa A, Mica JH, De Los Mozos J, Den Hartog LA, García-Ruiz AI. Nutritional dietary supplements to reduce the incidence of fatty liver syndrome in laying hens and the use of spectrophotometry to predict liver fat content. *J Appl Poult Res.* (2019) 28:435–46. doi: 10.3382/japr/pfz005

24. Kaneva AM, Bojko ER. Fatty liver index (FLI): more than a marker of hepatic steatosis. *J Physiol Biochem.* (2024) 80:11–26. doi: 10.1007/s13105-023-00991-z

25. Li G, Zhao Y, Guo S, Liu B, Chen Y, Sun X, et al. Comparative analysis of spleen transcriptome detects differences in evolutionary adaptation of immune defense functions in bighead carp and silver carp. *Fish Shellfish Immunol.* (2019) 84:148–57. doi: 10.1016/j.fsi.2018.09.077

26. Sutton J, Habibi M, Shili CN, Beker A, Salak-Johnson JL, Foote A, et al. Lowprotein diets differentially regulate energy balance during Thermoneutral and heat stress in cobb broiler chicken (Gallus domesticus). *Int J Mol Sci.* (2024) 25:4369. doi: 10.3390/ijms25084369

27. Gao Z, Zheng C, Mao Z, Zheng J, Xu G, Liu D. A comprehensive study of liver-gut microbiota and antioxidant enzyme activity mediated regulation of late-laying hens by high and low residual feed intake. *Int J Biol Macromol.* (2025) 298:139938. doi: 10.1016/j.ijbiomac.2025.139938

28. Właźłak S, Pietrzak E, Biesek J, Dunisławska A. Modulation of the immune system of chickens a key factor in maintaining poultry production—a review. *Poult Sci.* (2023) 102:102785. doi: 10.1016/j.psj.2023.102785

29. Chakraborty S, Khamaru P, Bhattacharyya A. Regulation of immune cell metabolism in health and disease: special focus on T and B cell subsets. *Cell Biol Int.* (2022) 46:1729–46. doi: 10.1002/cbin.11867

30. Castro-Dopico T, Clatworthy MR. IgG and Fcy receptors in intestinal immunity and inflammation. Front Immunol. (2019) 10:805. doi: 10.3389/fimmu.2019.00805

31. Kroon S, Malcic D, Weidert L, Bircher L, Boldt L, Christen P, et al. Sublethal systemic LPS in mice enables gut-luminal pathogens to bloom through oxygen speciesmediated microbiota inhibition. *Nat Commun.* (2025) 16:1–18. doi: 10.1038/s41467-025-57979-0

32. Wang P, Qian H, Xiao M, Lv J. Role of signal transduction pathways in IL-1 β -induced apoptosis: pathological and therapeutic aspects. *Immun Inflamm Dis.* (2023) 11:e762. doi: 10.1002/iid3.762

33. Li J, Mukiibi R, Wang Y, Plastow GS, Li C. Identification of candidate genes and enriched biological functions for feed efficiency traits by integrating plasma metabolites and imputed whole genome sequence variants in beef cattle. *BMC Genomics*. (2021) 22:1–12. doi: 10.1186/s12864-021-08064-5

34. Guo X, Li J, Li X, Sun J, Zou X, Ji J, et al. Synergy of genetics and lipid metabolism driving feed utilization efficiency in chickens. *Poult Sci.* (2025) 104:104885. doi: 10.1016/j.psj.2025.104885

35. Bai H, Guo Q, Yang B, Dong Z, Li X, Song Q, et al. Effects of residual feed intake divergence on growth performance, carcass traits, meat quality, and blood biochemical parameters in small-sized meat ducks. *Poult Sci.* (2022) 101:101990. doi: 10.1016/j.psj.2022.101990

36. Mota LF, Santos SW, Júnior GAF, Bresolin T, Mercadante ME, Silva JA, et al. Metaanalysis across Nellore cattle populations identifies common metabolic mechanisms that regulate feed efficiency-related traits. *BMC Genomics*. (2022) 23:424. doi: 10.1186/s12864-022-08671-w