

Comprehensive analyses of bone and cartilage transcriptomes evince ion transport, inflammation and cartilage development-related genes involved in chickens' femoral head separation

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Additional file S2.

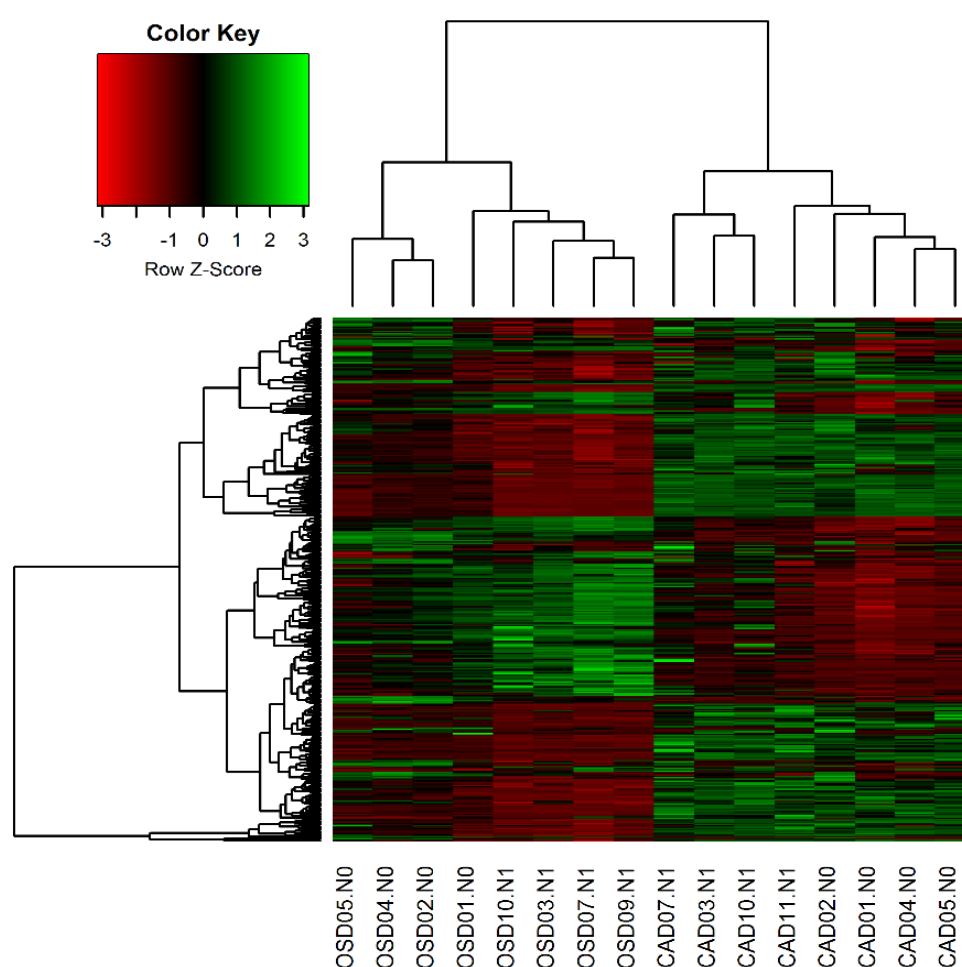


Fig S1. Heatmap gene cluster classification for AC and GP normal and FHS-affected samples. In the heatmap, the expression for each gene is presented in the rows and sample is visualized

in the columns, showing a hierarchical clustering of genes and samples. In red, DE genes were downregulated and in green, upregulate in the affected samples.