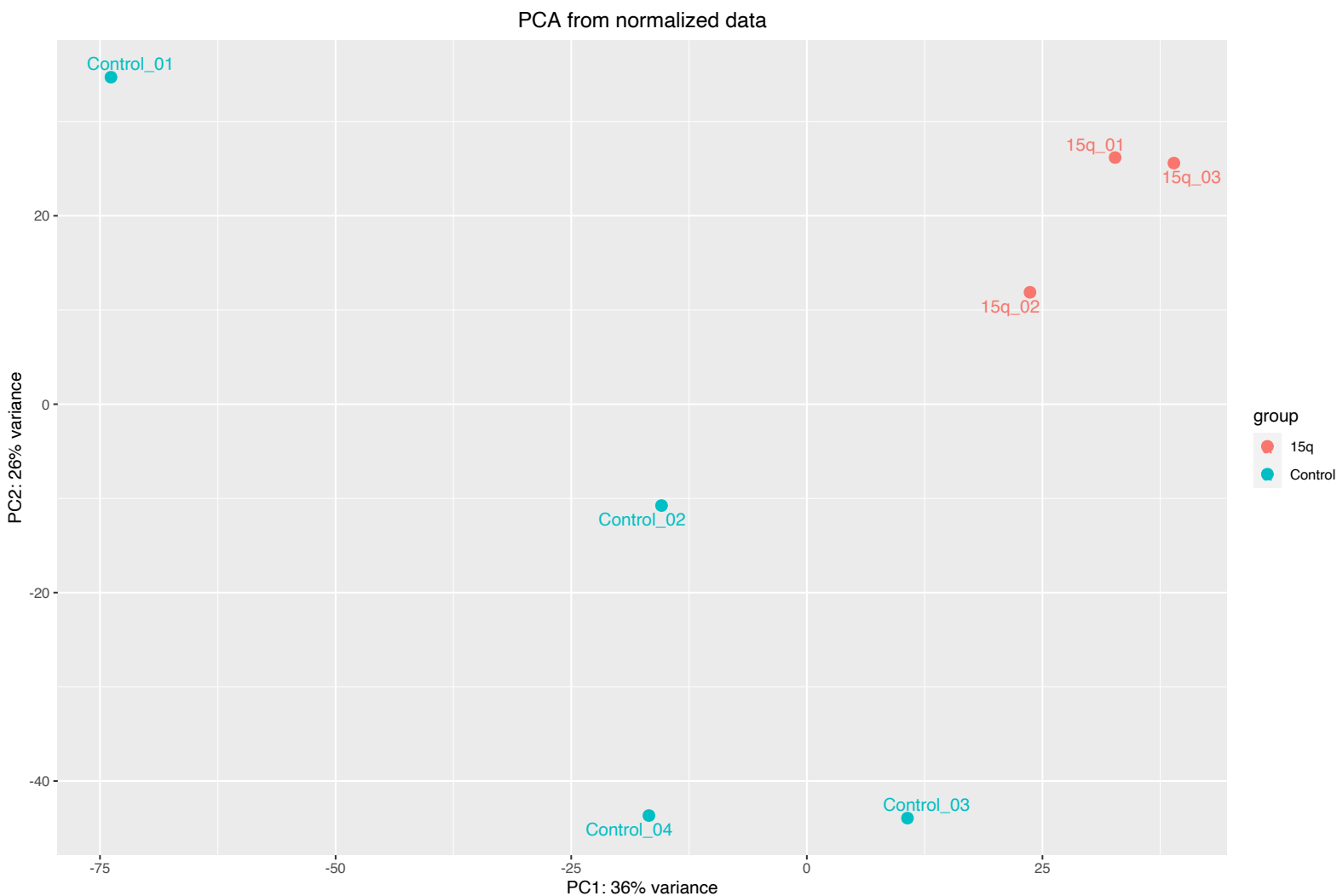




Additional file 1. Principal component analysis (PCA) plot of human RNA-seq samples. The first two principal components (PCs) are plotted and coloured according to group (15q – individuals with 15q13.3 microdeletion syndrome and Control – age- and gender-matched controls). PCA was performed using all expressed genes. Percentage of variation accounted for by each principal component is shown with the axis label.