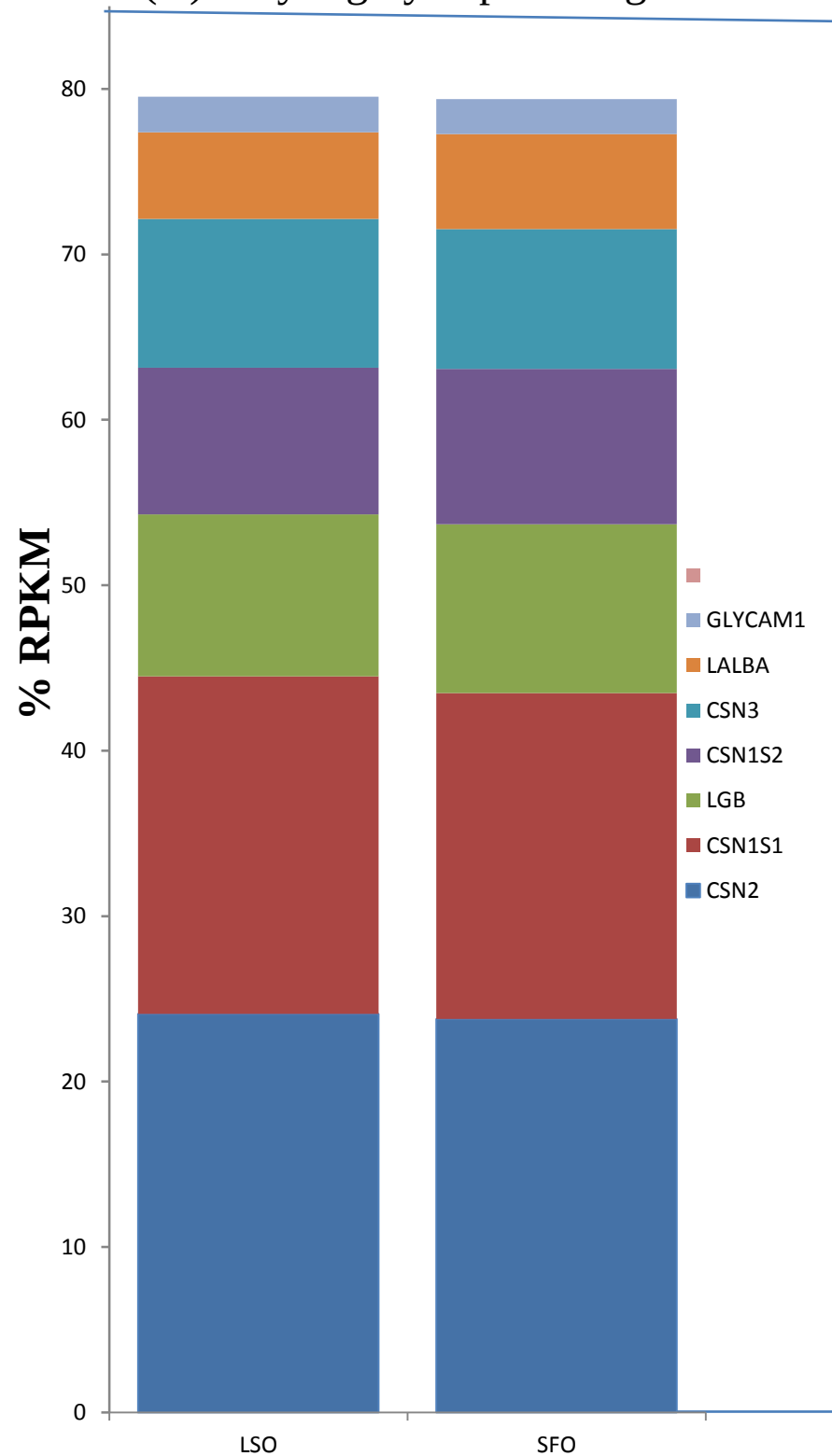
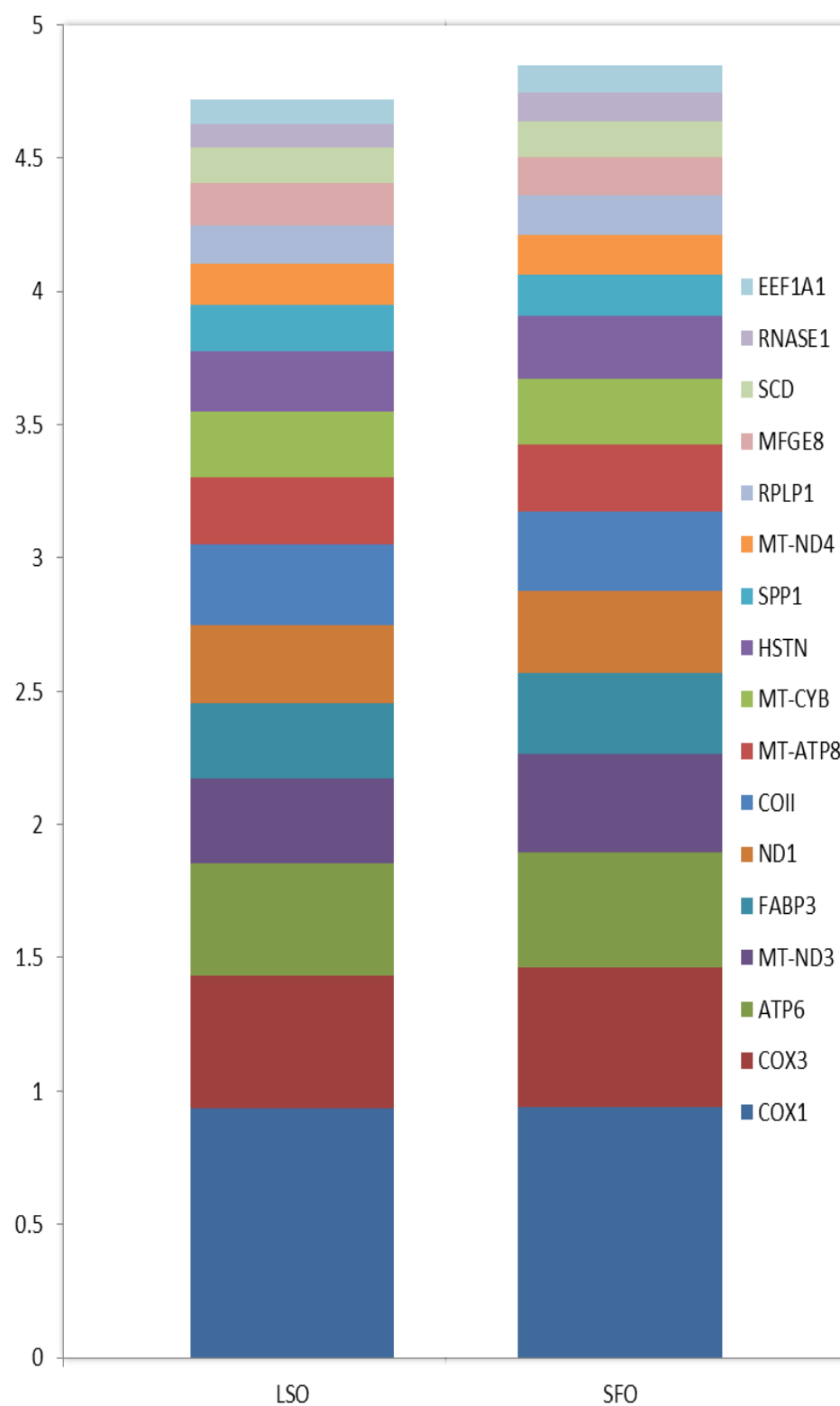


(A) Very highly expressed genes



(B) Highly expressed genes



Additional file 4: Top expressed genes according to treatment. The combined reads per kilo base per million mapped reads (RPKM) of seven very highly expressed genes constituted 79.45% of total reads while RPKM of 24 highly expressed genes constituted 4.79% of total reads.