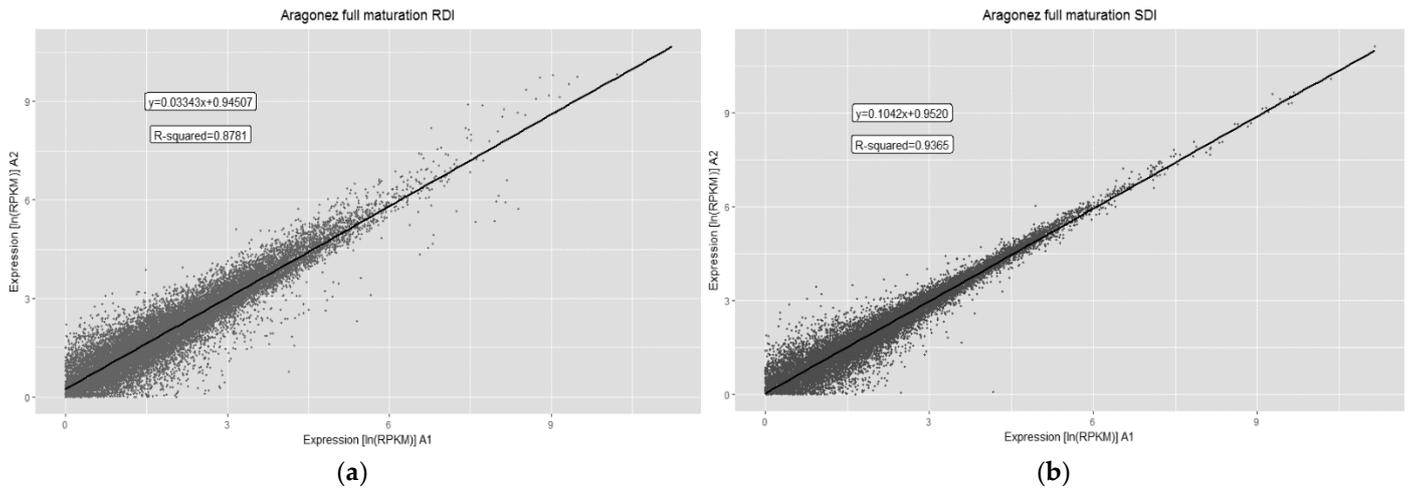


Supplementary Figure. Correlation between biological replicates in RNA-seq.



Supplementary Figure S1. Correlation between the logarithmized Reads per Kilobase transcript per Million reads (RPKM) of two biological replicates (in the X and Y axes). Dots represent the transcripts. Pearson Method was used to calculate linear regression (black line). Coefficient of determination (R-squared) and the linear regression equation are provided; (a) full maturation SDI; (b) full maturation RDI.