Regeneration *Drosophila melanogaster* new RNAseq samples sequenced at the CRG

Outline

Overview of RNA-seq samples

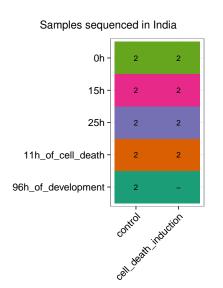
Mapping statistics

Correlation between replicates

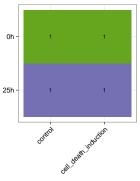
Comparison between ERC wing disc and 96h contro

PCA and Clustering

RNAseq samples



Samples sequenced at the CRG



Outline

Overview of RNA-seq samples

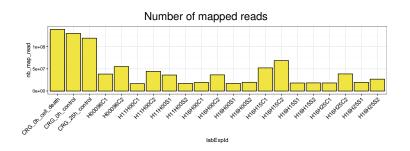
Mapping statistics

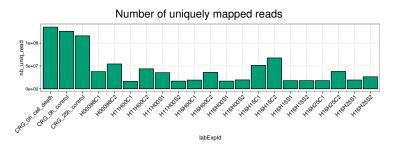
Correlation between replicates

Comparison between ERC wing disc and 96h contro

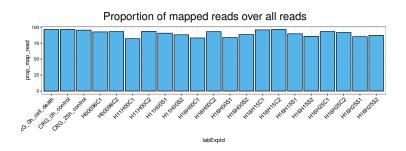
PCA and Clustering

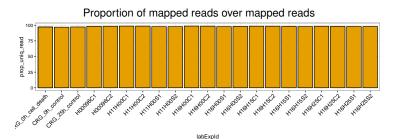
Mapping statistiques



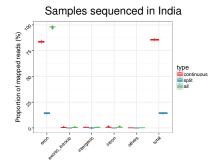


Mapping statistiques

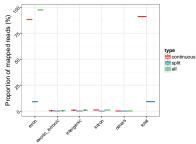




Genomic regions

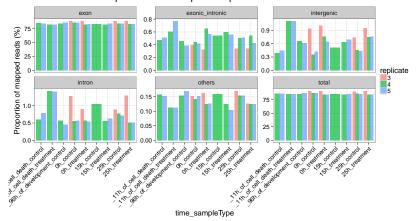






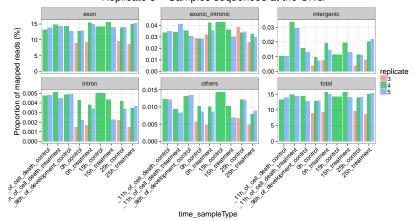
Genomic regions - continuous mapping

Replicates 4 and 5 = Samples sequenced in India
Replicate 3 = Samples sequenced at the CRG



Genomic regions - split mapping

Replicates 4 and 5 = Samples sequenced in India Replicate 3 = Samples sequenced at the CRG



Outline

Overview of RNA-seq samples

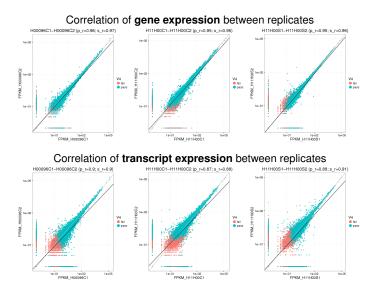
Mapping statistics

Correlation between replicates

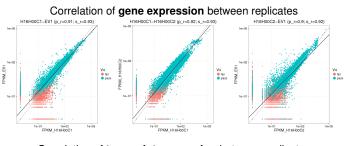
Comparison between ERC wing disc and 96h contro

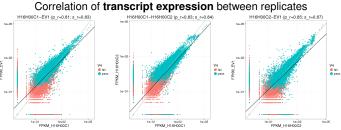
PCA and Clustering

96h control and 11h cell death

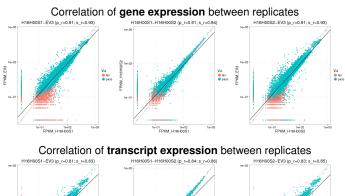


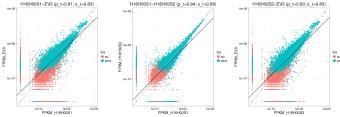
Oh control



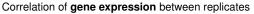


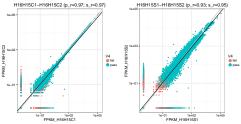
Oh cell death



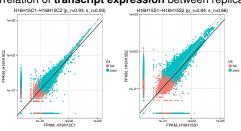


15h control and cell death - not sequenced in CRG

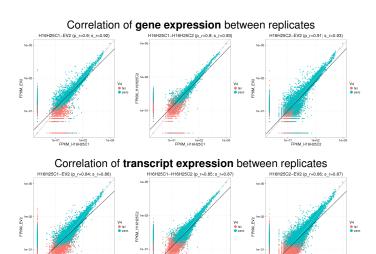




Correlation of transcript expression between replicates



25h control



FPKM H16H25C1

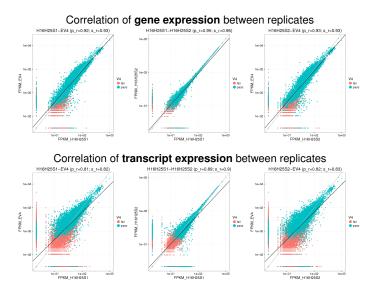
10+05

FPKM H16H25C2

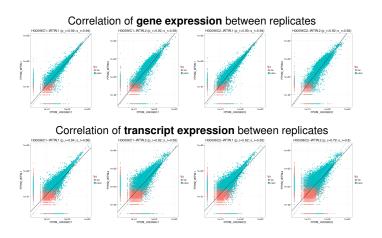
10+05

FPKM H16H25C1

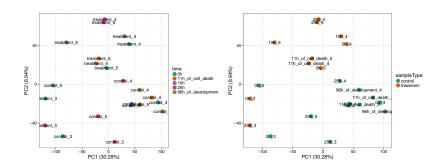
25h cell death



Comparison between ERC wing disc and 96h control



PCA and Clustering



PCA and Clustering

