Explain ceRNA effect from genomic muatations

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December 9, 2013

1 Background and Questions

Based on Hua-sheng's defense topic which excluded mutation, cnv for ceRNA effect Question 1: SNPs/CNVs/Methylations/Somatic mutations regulate gene expressions, the expression of genes have effect on ceRNA effect, but how much does each of them contribute to the final ceRNA effect?

Question 2:

Goal:

Find snps that are associated with DEGs, and explian it's effect using network approach

2 Methods

1.SNPs association with Gene Expression * SNP filtering process 1: -SNP quality, -SNP MAF - HWE test *association study : - Cohen's Kappa http://en.wikipedia.org/wiki/Cohen's_kappa feature selection methodsgroup lasso hierarchary structure: 2. TSS 3. 3' UTR

3 Data

1. TCGA level
2 SNP array 2. TCGA level
3 CNV (SNP array) 3. TCGA level
3 RNAseq \dots

Workflow

- 3.1
- 3.2

4 Validation

UK10K 1000 Genome Encode data 3C, 5C, H-G

5 Conclusion

Acknowledgement

6 Figures