# TCGA CRC 450 dataset

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## 1 Introduction

The package contain data objects of primary colorectal cancer for a total of 450 miRNA gene expression profile

The *ExpressionSet* include samples information (phenoData) and probe (genes) information The data were provided by TCGA consortium as level3 data [1]. data were generated by HiSeq and GenomeAnalyzer platforms.

> library(TCGAcrcmiRNA)

### 2 Introduction

Data can be loaded as follow.

> data(TCGAmirna)

### 3 References

### References

[1] Network, C.G.A. Network, C.G.A. Comprehensive molecular characterization of human colon and rectal cancer Nature 487, 330-7 (2012).