## GTEx expression data

-:-- A ---

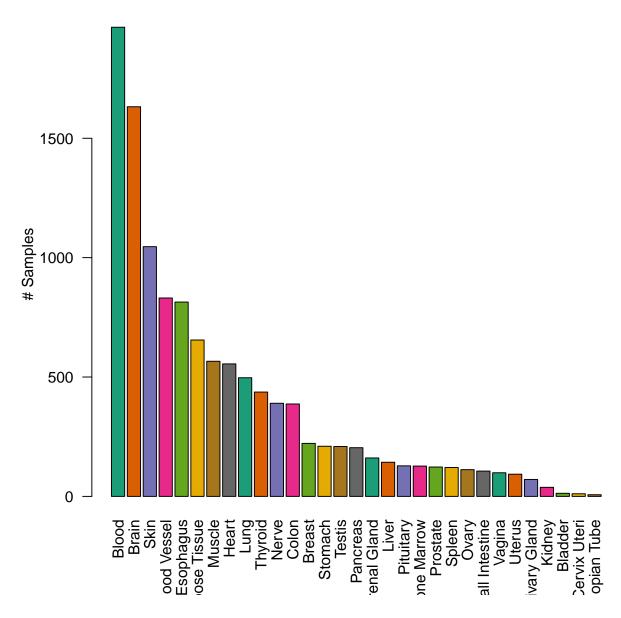
# Rainet project

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#### 1 Samples per tissue

Number of samples per tissue using the whole GTEx\_Data\_V6\_Annotations\_SampleAttributesDS.txt file with the attribute SMTS (the less-specific tissue terms).

#### Samples per tissue



```
## [1] "Total # of tissues : 31"

## [1] "Total # of samples : 11973"

## [1] "Minimum # samples per tissue: 7"

## [1] "Maximum # samples per tissue: 1965"
```

#### 2 Expression per tissue

 $\label{lem:continuous} Expression\ values\ per\ tissue\ using\ the\ whole\ GTEx\_Analysis\_v6\_RNA-seq\_Flux1.6\_transcript\_rpkm\_test.txt\ file.$ 

```
## Error in scan(file, what, nmax, sep, dec, quote, skip, nlines, na.strings, : line 1 did not
have 57124 elements
```

### 3 TO DO

In this section..