tgirtABRF

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tgirtABRF provides functions for labeling files from the work, RNA-Seq of human reference RNA samples using a thermostable group II intron reverse transcriptase.

Installing:

```
library(devtools)
install_github('wckdouglas/tgirtABRF')
```

To use the library:

```
library(tgirtABRF)
library(dplyr)
```

The naming system that was used in the project is as follow:

```
df <- read.csv('names.csv',header=F)
colnames(df) = 'names'
df</pre>
```

```
##
                   names
## 1
         ABRF-ILMN-L-A-1
## 2
         ABRF-ILMN-L-A-2
## 3
         ABRF-ILMN-L-A-3
## 4
         ABRF-ILMN-L-B-1
## 5
         ABRF-ILMN-L-B-2
## 6
         ABRF-ILMN-L-B-3
## 7
         ABRF-ILMN-R-A-1
## 8
         ABRF-ILMN-R-A-2
## 9
         ABRF-ILMN-R-A-3
## 10
         ABRF-ILMN-R-B-1
## 11
         ABRF-ILMN-R-B-2
         ABRF-ILMN-R-B-3
## 12
## 13 ABRF-ILMN-RIBO-A-1
## 14 ABRF-ILMN-RIBO-A-2
## 15 ABRF-ILMN-RIBO-A-3
## 16 ABRF-ILMN-RIBO-A-4
## 17 ABRF-ILMN-RIBO-B-1
## 18 ABRF-ILMN-RIBO-B-2
## 19 ABRF-ILMN-RIBO-B-3
## 20 ABRF-ILMN-RIBO-B-4
## 21 ABRF-ILMN-RIBO-C-1
## 22 ABRF-ILMN-RIBO-C-2
## 23 ABRF-ILMN-RIBO-C-3
## 24 ABRF-ILMN-RIBO-C-4
## 25 ABRF-ILMN-RIBO-D-1
## 26 ABRF-ILMN-RIBO-D-2
```

```
## 27 ABRF-ILMN-RIBO-D-3
## 28 ABRF-ILMN-RIBO-D-4
## 29
        ABRF-ILMN-V-A-1
## 30
         ABRF-ILMN-V-A-2
         ABRF-ILMN-V-A-3
## 31
## 32
      ABRF-ILMN-V-B-1
## 33
      ABRF-ILMN-V-B-2
## 34
         ABRF-ILMN-V-B-3
## 35
                   refA1
## 36
                   refA2
## 37
                   refA3
## 38
                   refB1
## 39
                   refB2
## 40
                   refB3
## 41
                   refC1
## 42
                   refC2
## 43
                   refC3
## 44
                   refD1
## 45
                   refD2
## 46
                   refD3
```

To extract the annotations from the sample names:

```
df %>%
    mutate(lab = getLab(names)) %>%
    mutate(mixTemplate = getTemplate(names)) %>%
    mutate(replicate = getReplicate(names)) %>%
    mutate(prep = getPrep(names)) %>%
    data.frame()
```

##		names	lab	mixTemplate	replicate	prep
##	1	ABRF-ILMN-L-A-1	L	A	1	TruSeq v2
##	2	ABRF-ILMN-L-A-2	L	A	2	TruSeq v2
##	3	ABRF-ILMN-L-A-3	L	A	3	TruSeq v2
##	4	ABRF-ILMN-L-B-1	L	В	1	TruSeq v2
##	5	ABRF-ILMN-L-B-2	L	В	2	TruSeq v2
##	6	ABRF-ILMN-L-B-3	L	В	3	TruSeq v2
##	7	ABRF-ILMN-R-A-1	R	A	1	TruSeq v2
##	8	ABRF-ILMN-R-A-2	R	A	2	TruSeq v2
##	9	ABRF-ILMN-R-A-3	R	A	3	TruSeq v2
##	10	ABRF-ILMN-R-B-1	R	В	1	TruSeq v2
##	11	ABRF-ILMN-R-B-2	R	В	2	TruSeq v2
##	12	ABRF-ILMN-R-B-3	R	В	3	TruSeq v2
##	13	ABRF-ILMN-RIBO-A-1	W	A	1	TruSeq v3
##	14	ABRF-ILMN-RIBO-A-2	W	A	2	TruSeq v3
##	15	ABRF-ILMN-RIBO-A-3	W	Α	3	TruSeq v3
##	16	ABRF-ILMN-RIBO-A-4	W	A	4	TruSeq v3
##	17	ABRF-ILMN-RIBO-B-1	W	В	1	TruSeq v3
##	18	ABRF-ILMN-RIBO-B-2	W	В	2	TruSeq v3
##	19	ABRF-ILMN-RIBO-B-3	W	В	3	TruSeq v3
##	20	ABRF-ILMN-RIBO-B-4	W	В	4	TruSeq v3
##	21	ABRF-ILMN-RIBO-C-1	W	C	1	TruSeq v3
##	22	ABRF-ILMN-RIBO-C-2	W	C	2	TruSeq v3

##	23	ABRF-ILMN-RIBO-C-3	W	C	3	TruSeq v3
##	24	ABRF-ILMN-RIBO-C-4	W	C	4	TruSeq v3
##	25	ABRF-ILMN-RIBO-D-1	W	D	1	TruSeq v3
##	26	ABRF-ILMN-RIBO-D-2	W	D	2	TruSeq v3
##	27	ABRF-ILMN-RIBO-D-3	W	D	3	TruSeq v3
##	28	ABRF-ILMN-RIBO-D-4	W	D	4	TruSeq v3
##	29	ABRF-ILMN-V-A-1	V	A	1	TruSeq v2
##	30	ABRF-ILMN-V-A-2	V	A	2	TruSeq v2
##	31	ABRF-ILMN-V-A-3	V	A	3	TruSeq v2
##	32	ABRF-ILMN-V-B-1	V	В	1	TruSeq v2
##	33	ABRF-ILMN-V-B-2	V	В	2	TruSeq v2
##	34	ABRF-ILMN-V-B-3	V	В	3	TruSeq v2
##	35	refA1	${\tt Lambowitz}$	A	1	TGIRT-seq
##	36	refA2	${\tt Lambowitz}$	A	2	TGIRT-seq
##	37	refA3	${\tt Lambowitz}$	A	3	TGIRT-seq
##	38	refB1	${\tt Lambowitz}$	В	1	TGIRT-seq
##	39	refB2	${\tt Lambowitz}$	В	2	TGIRT-seq
##	40	refB3	${\tt Lambowitz}$	В	3	TGIRT-seq
##	41	refC1	${\tt Lambowitz}$	C	1	TGIRT-seq
##	42	refC2	${\tt Lambowitz}$	C	2	TGIRT-seq
##	43	refC3	${\tt Lambowitz}$	C	3	TGIRT-seq
##	44	refD1	${\tt Lambowitz}$	D	1	TGIRT-seq
##	45	refD2	${\tt Lambowitz}$	D	2	TGIRT-seq
##	46	refD3	${\tt Lambowitz}$	D	3	TGIRT-seq