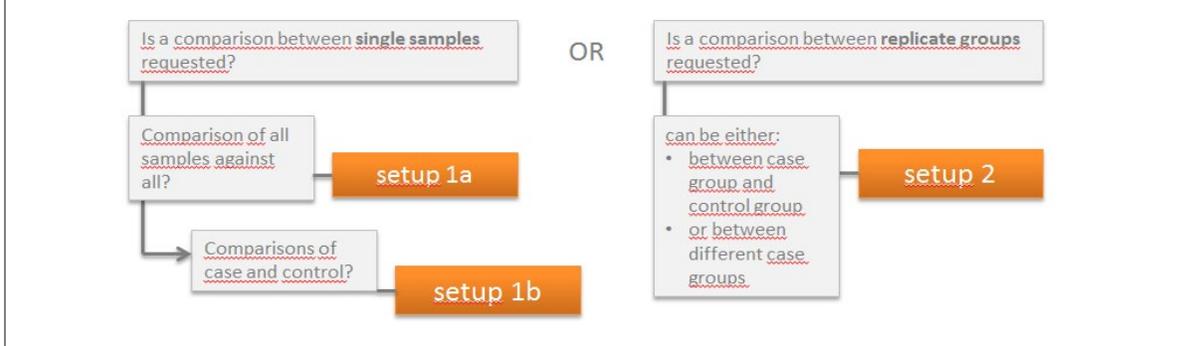


# INFORMATION REQUEST FORM

This information is mandatory for performing the Transcriptome or Methylation Analysis.  
Please fill in this form electronically and send it back to GATC Biotech by e-mail

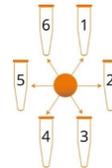
## Information about the experimental design

### Find the right setup for your analysis



### Setup 1a: sample wise comparison, all against all

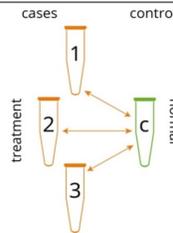
Samples will be compared singlewise, all against all.



### Setup 1b: sample wise case / control setup

#### Sample assignment to case / control

sample name	case	control
normal		x
treatment_1	x	
treatment_2	x	
treatment_3	x	

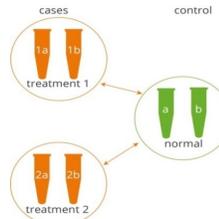


### Setup 2: Group wise comparison for replicates

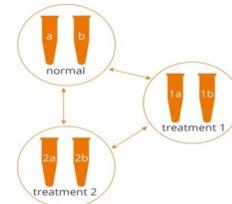
#### Sample assignment to groups

sample name	group name
sample_a	normal
sample_b	normal
sample_1_a	treatment_1
sample_1_b	treatment_1
sample_2_a	treatment_2
sample_2_b	treatment_2

- valid characters for group names: 0 to 9, Aa-Zz \_
- invalid characters will be converted to "\_"



OR



#### Groups to be compared

case group	vs.	control group or alt. case group
normal		treatment_1
normal		treatment_2
treatment_1		treatment_2

### Setup 2: Group wise comparison for replicates

### Groups to be compared



