

How to quantify multiple bam files

Dear Sammeth,

Is there any support to quantify multiple bam files using flux capacitor. I can quantify one file using following command:

```
flux-capacitor -p param.txt -i myMapping1.bam -a myAnnotation.gtf -o myout.flux --threads 2
```

I mean, is there any support to provide multiple bam files like following:

```
flux-capacitor -p param.txt -i myMapping1.bam,myMapping2.bam,myMapping3.bam, -a myAnnotation.gtf -o myout.flux --threads 2
```