

You cannot specify a READ_Descriptor for BAM files

Hi,

I'm annotating some filtered, sorted bam files with their index files and when I run:

```
flux-capacitor -p <parameter_file> -i <sample.bam> -o <output.gtf>
```

with:

```
ANNOTATION_FILE gencode.v12.annotation.gtf  
READ_DESCRIPTOR {ID}/{MATE}[1,2]
```

```
ANNOTATION_MAPPING PAIRED
```

I get "You cannot specify a READ_Descriptor for BAM files".

I see this has been raised and apparently resolved in

- [BARNA-261](#)

It still seems to occur. I'm using Win 7 OS 64bit - flux-capacitor-1.5.2.

Any prompt help will be great!

Thanks.