PloidPy Usage Guide

BAM file
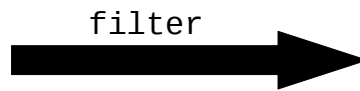
`process_bam` →
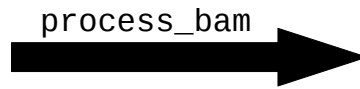
```
94 425
86 397
87 363
94 378
190 384
105 406
```

Raw count file

```
94 425
86 397
87 363
94 378
190 384
105 406
```

Raw count file

`filter` →

```
94 425
86 397
87 363
94 378
190 384
105 406
```

Filtered count file

```
94 425
86 397
87 363
94 378
190 384
105 406
```

Filtered count file

`jdist` →
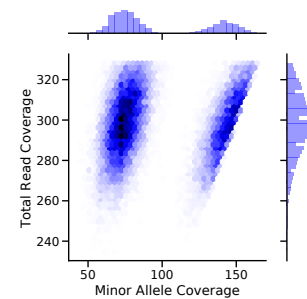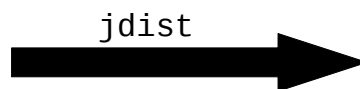
Joint distribution plot

```
94 425
86 397
87 363
94 378
190 384
105 406
```

Raw or **filtered** count file

`assess` →
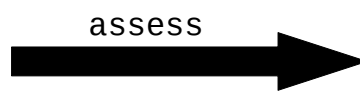
2n

vs

3n

vs

4n

...

Ploidy prediction