

```
In [54]: %time out_file = input_file.rstrip(".reg.gz") + (".dat")
CPU times: user 7 µs, sys: 7 µs, total: 14 µs
Wall time: 11.9 µs
```

```
In [55]: import time
```

```
In [56]: def procedure():
.....:     np.savetxt(out_file, genome_matrix, fmt="%d", delimiter=' ')
.....:
```

```
In [57]: t0 = time.clock()
```

```
In [58]: procedure()
```

```
In [59]: print time.clock() - t0, "seconds process time"
4.370414 seconds process time
```

```
In [60]: █
```