

```
In [22]: %time chrom_vec_size = len(chrom_vector)
```

```
CPU times: user 4 µs, sys: 1 µs, total: 5 µs
```

```
Wall time: 7.87 µs
```

```
In [23]: %time genome_matrix = np.zeros(shape=(chrom_vec_size, chrom_vec_size), dtype=np.int)
```

```
CPU times: user 17 µs, sys: 53 µs, total: 70 µs
```

```
Wall time: 73.9 µs
```