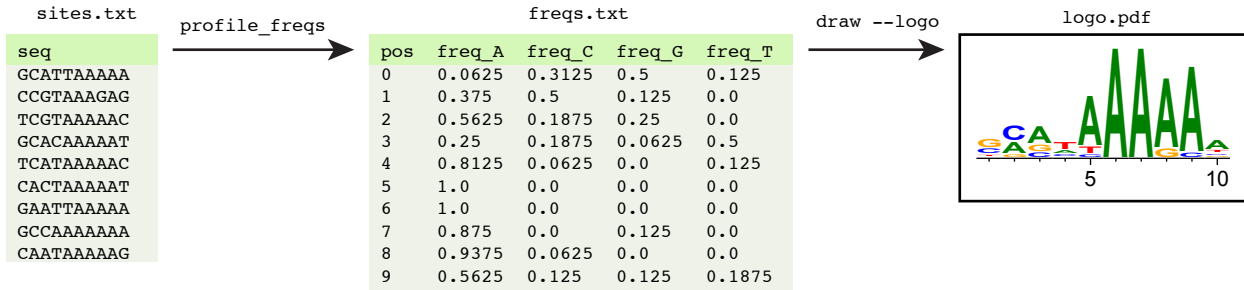


A



B

```
$ sortseq profile_freqs -i sites.txt -o freqs.txt
$ sortseq draw -i freqs.txt -o logo.pdf --logo
```

C

```
$ echo sites.txt | sortseq profile_freqs | sortseq draw --logo > logo.pdf
```

D

```
import sortseq
data = sortseq.load("data.txt")
freqs = sortseq.profile_freqs(data)
fig = sortseq.draw(freqs, "logo")
fig.save("logo.pdf")
```

Figure 3