

OLC

A	C	D	E	F	G	H	I	K	L	M	N	P	Q	R	S	T	V	W	Y
81	30	61	51	77	94	26	71	58	103	10	88	62	66	42	103	95	92	12	53

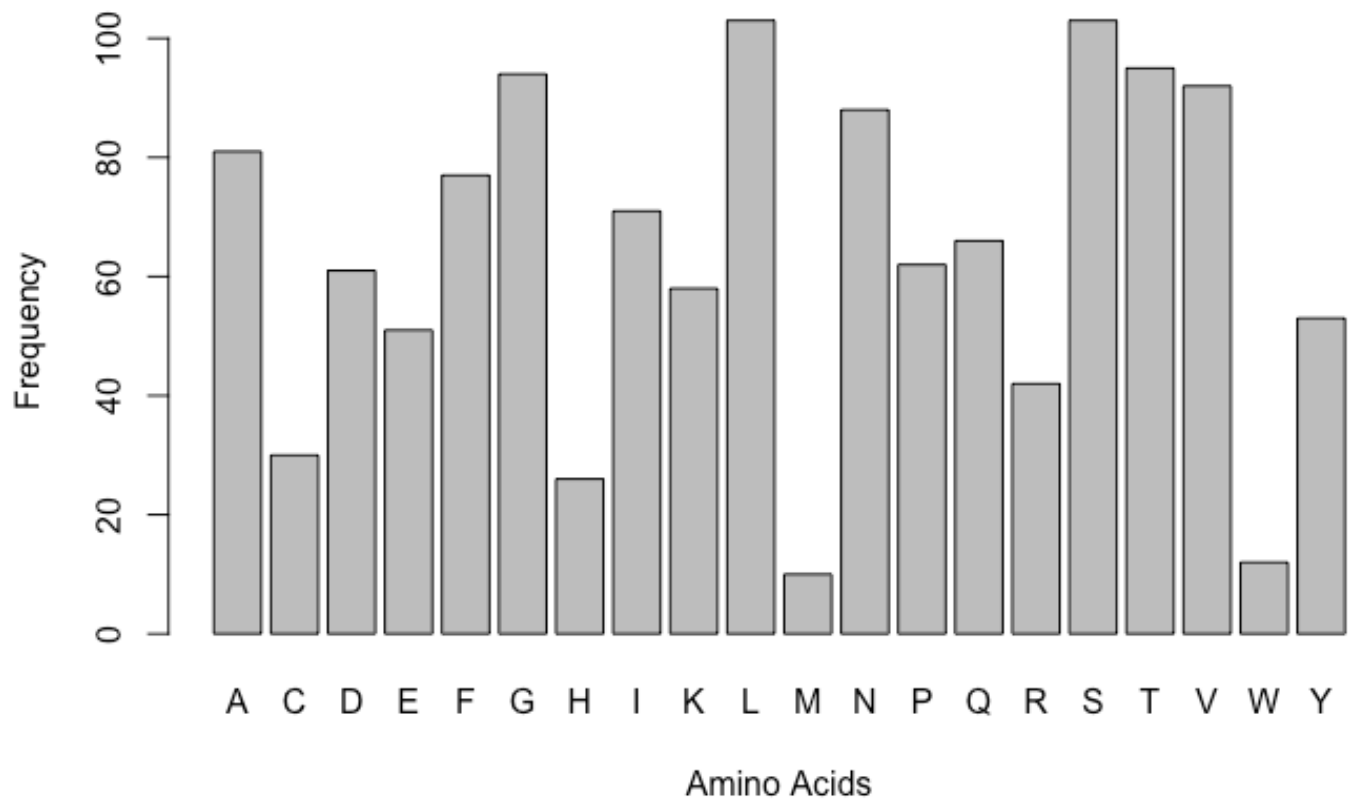
```
aminosorted <- sort((table(OLC)),decreasing=TRUE,na.last=TRUE)
```

aminosorted

OLC

L	S	T	G	V	N	A	F	I	Q	P	D	K	Y	E	R	C	H	W	M
103	103	95	94	92	88	81	77	71	66	62	61	58	53	51	42	30	26	12	10

```
plot((OLC), ylab = 'Frequency', xlab = 'Amino Acids')
```



NA

NA