check50

cs50/problems/2023/x/dna

:) dna.py exists

Log

```
checking that dna.py exists...
```

:) correctly identifies sequences/1.txt

Log

```
running python3 dna.py databases/small.csv sequences/1.txt...
checking for output "Bob\n"...
```

:) correctly identifies sequences/2.txt

Log

```
running python3 dna.py databases/small.csv sequences/2.txt...
checking for output "No match\n"...
```

:) correctly identifies sequences/3.txt

Log

```
running python3 dna.py databases/small.csv sequences/3.txt...
checking for output "No match\n"...
```

:) correctly identifies sequences/4.txt

Log

```
running python3 dna.py databases/small.csv sequences/4.txt...
checking for output "Alice\n"...
```

:) correctly identifies sequences/5.txt

Log

```
running python3 dna.py databases/large.csv sequences/5.txt...
checking for output "Lavender\n"...
```

:) correctly identifies sequences/6.txt

Log

```
running python3 dna.py databases/large.csv sequences/6.txt...
checking for output "Luna\n"...
```

:) correctly identifies sequences/7.txt

Log

```
running python3 dna.py databases/large.csv sequences/7.txt...
checking for output "Ron\n"...
```

:) correctly identifies sequences/8.txt

Log

```
running python3 dna.py databases/large.csv sequences/8.txt...
checking for output "Ginny\n"...
```

:) correctly identifies sequences/9.txt

Log

```
running python3 dna.py databases/large.csv sequences/9.txt...
checking for output "Draco\n"...
```

:) correctly identifies sequences/10.txt

Log

```
running python3 dna.py databases/large.csv sequences/10.txt...
checking for output "Albus\n"...
```

:) correctly identifies sequences/11.txt

Log

```
running python3 dna.py databases/large.csv sequences/11.txt...
checking for output "Hermione\n"...
```

:) correctly identifies sequences/12.txt

Log
```
running python3 dna.py databases/large.csv sequences/12.txt...
checking for output "Lily\n"...
```

:) correctly identifies sequences/13.txt

Log
```
running python3 dna.py databases/large.csv sequences/13.txt...
checking for output "No match\n"...
```

:) correctly identifies sequences/14.txt

Log
```
running python3 dna.py databases/large.csv sequences/14.txt...
checking for output "Severus\n"...
```

:) correctly identifies sequences/15.txt

Log
```
running python3 dna.py databases/large.csv sequences/15.txt...
checking for output "Sirius\n"...
```

:) correctly identifies sequences/16.txt

Log
```
running python3 dna.py databases/large.csv sequences/16.txt...
checking for output "No match\n"...
```

:) correctly identifies sequences/17.txt

Log
```
running python3 dna.py databases/large.csv sequences/17.txt...
checking for output "Harry\n"...
```

:) correctly identifies sequences/18.txt

Log
```
running python3 dna.py databases/large.csv sequences/18.txt...
checking for output "No match\n"...
```

:) correctly identifies sequences/19.txt

Log
```
running python3 dna.py databases/large.csv sequences/19.txt...
checking for output "Fred\n"...
```

:) correctly identifies sequences/20.txt

Log
```
running python3 dna.py databases/large.csv sequences/20.txt...
checking for output "No match\n"...
```