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## Note

Note that using 'hold on' may affect the initial view of the 3D data!

Examples of both views are included. Either is correct and will be give full credit.

## Sample Assignment 6, Protein Data Bank

Your published report should print the source code to the three functions required by the assignment:

```
readPDBfile()
drawAtoms()
potentialHbonds()
```

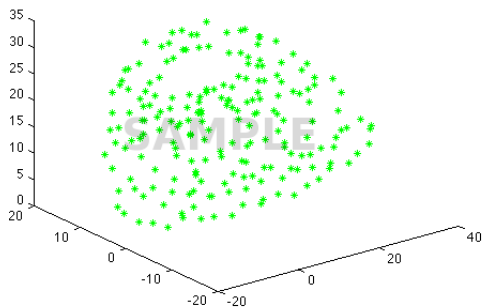
It should also include the code to the main function that computes the results for the two proteins.

## 7hvp protein

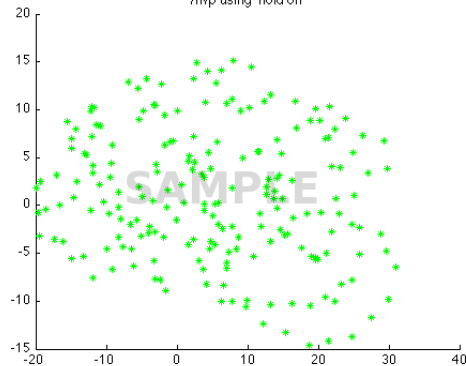
```
numOfHbondPairs =
```

```
286
```

7hvp without using 'hold on'



7hvp using 'hold on'

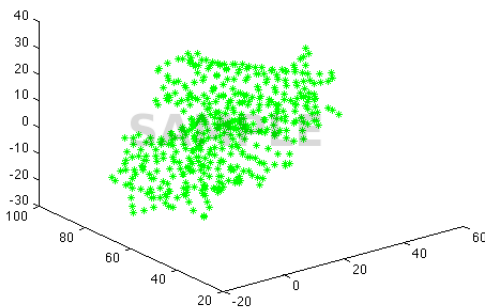


## 1gfl protein

```
numOfHbondPairs =
```

```
775
```

1gfl without using 'hold on'



1gfl using 'hold on'

