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## Assessment: Platform oriented annotation

### Finding probe sets for a gene

1/1 point (graded)

Use the `hgu133plus2.db` object loaded from the **hgu133plus2.db** package.

How many probe sets are mapped to the gene with the symbol EGFR?

Remember to use the `select()` function to search.



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You have used 1 of 5 attempts

### Probe sets for a GO category

1/1 point (graded)

Use `GO.db` to determine the tag for the term 'glial cell proliferation'. That would involve:

```
library(GO.db)
select(GO.db, key= ...)
```

Use `hgu133plus2.db` to determine how many probe sets are annotated to this biological process.



You have used 1 of 5 attempts

## Discussion

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