

FASTAQuery

```
graph BT; SVData[SVData] -.->|ref_genome| FASTAQuery[FASTAQuery];
```

The diagram illustrates a dependency between two classes. At the top is a white box labeled 'FASTAQuery'. At the bottom is a gray box labeled 'SVData'. A dashed purple arrow points from 'SVData' up to 'FASTAQuery'. To the right of the arrow is the text 'ref_genome' in a light gray font.

ref_genome

SVData