

SVData::getRefGenome



```
graph LR; A[SVData::getRefGenome] --> B[FASTAQuery::getFilepath]
```

The diagram consists of two rectangular boxes connected by a horizontal blue arrow pointing from left to right. The left box is filled with a dark gray color and contains the text 'SVData::getRefGenome'. The right box is white with a black border and contains the text 'FASTAQuery::getFilepath'.

FASTAQuery::getFilepath