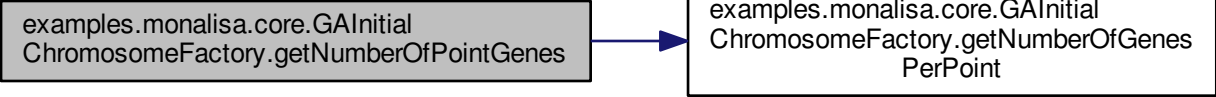


examples.monalisa.core.GAInitial  
ChromosomeFactory.getNumberOfPointGenes



```
graph LR; A["examples.monalisa.core.GAInitial  
ChromosomeFactory.getNumberOfPointGenes"] --> B["examples.monalisa.core.GAInitial  
ChromosomeFactory.getNumberOfGenes  
PerPoint"]
```

The diagram consists of two rectangular boxes connected by a horizontal arrow pointing from left to right. The left box has a light gray background and contains the text 'examples.monalisa.core.GAInitial' followed by 'ChromosomeFactory.getNumberOfPointGenes' on the next line. The right box has a white background and contains the text 'examples.monalisa.core.GAInitial' followed by 'ChromosomeFactory.getNumberOfGenes' and 'PerPoint' on the next two lines. A blue arrow points from the right side of the left box to the left side of the right box.

examples.monalisa.core.GAInitial  
ChromosomeFactory.getNumberOfGenes  
PerPoint