

Sequence::canBeNucleic

Sequence::getSequenceSymbol

Sequence::canBeRna

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graph LR; A[Sequence::canBeNucleic] --> C[Sequence::canBeRna]; B[Sequence::getSequenceSymbol] --> C;
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The diagram illustrates a relationship between three methods. On the left, two white rectangular boxes with black borders contain the text 'Sequence::canBeNucleic' (top) and 'Sequence::getSequenceSymbol' (bottom). On the right, a gray rectangular box with a black border contains the text 'Sequence::canBeRna'. Two blue arrows originate from the right side of the left boxes and point towards the left side of the gray box, indicating that both 'canBeNucleic' and 'getSequenceSymbol' are associated with or point to 'canBeRna'.