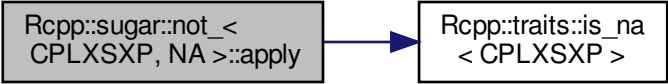


```
Rcpp::sugar::not_<  
CPLXSXP, NA >::apply
```



```
graph LR; A["Rcpp::sugar::not_<CPLXSXP, NA >::apply"] --> B["Rcpp::traits::is_na<CPLXSXP>"]
```

A diagram illustrating a transformation or relationship between two Rcpp expressions. On the left, a grey rectangular box contains the expression `Rcpp::sugar::not_<CPLXSXP, NA >::apply`. A blue arrow points from this box to a white rectangular box on the right, which contains the expression `Rcpp::traits::is_na<CPLXSXP>`. This suggests that the sugar wrapper on the left is being replaced or simplified into the traits-based function on the right.

```
Rcpp::traits::is_na  
<CPLXSXP>
```