

known samples  
correctly (Figure A).  
the 5 unknown sam  
out of 5 samples w  
correctly with refere  
genotypes obtai  
sequencing (Figure  
rs1801187 only the g  
could be called a  
genotypes AG and A  
be distinguished (Fi  
the 5 unknown samp  
5 samples were call  
(Figure D

## CONCLUS

**In this study, two SM  
templates amenable  
templates were com  
followed by nested P  
to that of genomic D  
unknown samples, a**

**Being the largest kn**