

Low base quality



AGACTTGGGC**T**CCCCCATTTC
AGACT**T**GGGC**T**CCCC**C**CCATTTC**C**A
AGACT**A**GGGCC**C**CCCC**A**CCCCATTTC**A**AGG
ACTTGGGC**T****C**CTCCCCATTTC**C**AGGTC
TTGGGC**T**CCCCCATTTC**C**AGGTCCTT
GC**C**CCCC**A**CCCC**A**ATTTC**A**AGGTCCTT**C**
CCCC**A**CCCCATTTC**C**AGGTCCTT**C**
TCCCCATTTC**C**AGGTC**C**TC

Reference seq

A G A C T T G G C C C C C T C C C C A T T C A A G G T C T T C

Allelic counts

R : 2 3 4 4 5 2 5 6 6 2 7 6 7 5 8 7 8 7 8 8 8 8 2 7 6 6 5 5 3 3 3
A : 0 0 0 0 0 1 0 0 0 4 0 0 0 3 0 0 0 1 0 0 0 0 5 0 0 0 0 0 0 0 0

Predicted dosage

0 0 0 0 0 **1** 0 0 0 **1** 0 0 0 **1** 0 0 0 0 0 0 0 0 0 **1** 0 0 0 0 0 0 0 0 0