

Aligned reads

AGACTTGGCTCCCTCCCAATTTC
AGACTTGGCTCCCTCCCAATTTC
AGACTAGGCTCCCTCCCAATTTC
ACTTGGCTCCCTCCCAATTTC
TTGGCTCCCTCCCAATTTC
GCCTCCCTCCCAATTTC
CCCCTCCCAATTTC
TCCCTCCCAATTTC
CAGGTCTTC
CAGGTCTTC

Reference seq

AGACTTGGCCCCCTCCCCATTCAAGGTCTTC

Allelic counts

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

Predicted dosage

0 0 0 0 0 0 0 0 0 0 1 0 0 0 1 0 0 0 0 0 0 0 0 0 0 1 0 0 0 0 0 0 1 0 0