

Alignment 1
sequence2
Pyrus calleryanaSRR7135497 (+)
1-160117
Criteria: 70%, 100 bp
Regions: 182

Alignment 2
sequence3
Pyrus calleryanaSRR7135498 (+)
1-159937
Criteria: 70%, 100 bp
Regions: 181

Alignment 3
sequence4
Pyrus calleryanaSRR7135500 (+)
1-159939
Criteria: 70%, 100 bp
Regions: 181

X-axis: sequence1
Resolution: 63
Window size: 100 bp

Legend:
grey arrow: contig
blue box: gene
blue box: exon
red box: UTR
green box: CNS
grey line: mRNA

Suppl. Fig. 2: Alignment of *Pyrus calleryana* chloroplasts to OM541581 using mVista Viewer. Shown are identity scores (graph) and coding vs. non-coding regions (blue vs. red) as well as the annotation of the reference *P. calleryana*. Green boxes denote locations of regions analyzed by Zheng et al. 2014.

