**Table S1.** Repetitive sequences identified using RepeatMasker in the Plum contigs with low ALCR and high SDR.

|  |  |  |  |
| --- | --- | --- | --- |
|  | Total length | GC level | Repetitive sequence length |
| contig\_741 | 3193 bp | 55.28% | 2273 bp (71.19 %) |
| contig\_2322 | 8911 bp | 43.87% | 6280 bp (70.47 %) |
| contig\_829 | 9400 bp | 47.70% | 6477 bp (68.90 %) |
| contig\_710 | 22313 bp | 43.08% | 14894 bp (66.75 %) |
| contig\_2344 | 14975 bp | 43.45% | 8704 bp (58.12 %) |
| contig\_1293 | 103789 bp | 41.27% | 42073 bp (40.54 %) |
| contig\_1115 | 107136 bp | 41.36% | 38336 bp (35.78 %) |