Table S2. SSR markers Allele size range, frequency of predominant alleles, number of alleles per locus and genetic diversity index of each locus for Korean and Chinese soybean populations.

<table>
<thead>
<tr>
<th>Marker</th>
<th>Allele size range (bp)</th>
<th>Predominant alleles (%)</th>
<th>Number of alleles/locus</th>
<th>Genetic diversity index (H)</th>
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<td>Korea</td>
<td>Total</td>
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</table>

† Digit in parenthesis indicates soybean chromosome number
‡ Digit in parenthesis indicates the number of alleles specific to the population
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<th>Marker</th>
<th>Allele size range (bp)</th>
<th>Predominant alleles (%)</th>
<th>Number of alleles/locus</th>
<th>Genetic diversity index (H)</th>
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\(\dagger\) Digit in parenthesis stands for soybean chromosome number

\(\ddagger\) Digit in parenthesis indicates the number of alleles specific to the population
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<td>Korea</td>
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</tbody>
</table>

**Sum** | 751 | 638 (219) | 527 (112) | 0.77  | 0.76  | 0.72  |

† Digit in parenthesis stands for soybean chromosome number

‡ Digit in parenthesis indicates the number of alleles specific to the population