

## Addendum: Resequencing of 31 wild and cultivated soybean genomes identifies patterns of genetic diversity and selection

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Raw data of short reads for each soybean accession can be downloaded as individual files from the NCBI website. The corresponding file name and soybean accession are listed in the addendum. Further information on the soybean accessions can be found in the supplementary information online.

The full dataset can also be downloaded from the following ftp site: [ftp://public.genomics.org.cn/BGI/soybean\\_resequencing/](ftp://public.genomics.org.cn/BGI/soybean_resequencing/).

### Corresponding table for NCBI data file and soybean accession number<sup>a</sup>

NCBI sample code <sup>b</sup>	Corresponding soybean accession number
SRS079322	C08
SRS079323	C16
SRS079324	C17
SRS079325	C24
SRS079326	C30
SRS079327	C33
SRS079328	C34
SRS079329	W12
SRS079330	W02
SRS079331	W03
SRS079332	W04
SRS079333	W07
SRS079334	W08
SRS079335	W10
SRS079336	C35
SRS079337	C14
SRS079338	W01
SRS079339	C27
SRS079340	C19
SRS079341	W13
SRS079342	W15
SRS079343	C01
SRS079344	C02
SRS079345	C12
SRS079346	W11
SRS079347	W16
SRS079348	W06
SRS079349	W09
SRS079350	W14
SRS079351	W17
SRS079352	W05

<sup>a</sup>Each file contains the complete set of raw data (short reads) of each soybean accession. Further information on the soybean accessions can be found in **Supplementary Figure 1** and **Supplementary Table 1**. <sup>b</sup>The full dataset can also be downloaded from the following ftp site: [ftp://public.genomics.org.cn/BGI/soybean\\_resequencing/](ftp://public.genomics.org.cn/BGI/soybean_resequencing/).