

**DATA ACCESS DISCLAIMER:**

These data are in open access. While scientists may freely publish using the IWGSC data, IWGSC does request that the source of the data be properly acknowledged.

For questions regarding data access and publication policy please contact IWGSC (communications@wheatgenome.org).

**PLEASE NOTE:**

MD5 checksums are provided to ensure complete data transfer of files, please make sure that the MD5 checksum of your copy corresponds to the one provided with the files.

**QUESTIONS**

In case of questions please contact Michael Alaux from URGI in the first instance at: [urgi-support@inra.fr](mailto:urgi-support@inra.fr)

**CREDITS:**

This data set has been generated as a collaborative effort between INRA-GDEC Clermont Ferrand, France (INRA) (Frederic Choulet, Philippe Leroy, Helene Rimbart), Plant Genome and Systems Biology, Helmholtz Zentrum, Munich, Germany (PGSB) (Klaus Mayer, Manuel Spannagl, Sven Twardziok, Heidrun Gundlach) and Earlham Institute, Norwich, UK (EI) (David Swarbreck, Luca Venturini, Gemy Kaithakottil) under the coordination of IWGSC (International Wheat Genome Sequencing Consortium) (Kellye Eversole, Jane Rogers).

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*Unplaced scaffolds*

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The unplaced scaffolds can be found on the following files:

- [iwgsc\\_refseqv1.0\\_chrUn.fsa.zip](#)

It is a “fake” chromosome with all the unplaced scaffolds randomly ordered and separated by 100 Ns.

- [iwgsc\\_refseqv1.0\\_unplaced\\_scaffolds.fsa.zip](#)

It is a multi fasta file with all unanchored scaffolds as individual sequences.