A Big Earth Data Platform for Three Poles

**Genome resequencing data of main domestic animals in the Tibetan Plateau (2019)**

1、Description

In order to describe the distribution pattern of the genetic diversity of the main domesticated animals in the Qinghai Tibet Plateau and its surrounding areas, and to clarify the related genetic background. In 2019, we extracted total DNA from 21 local chicken tissue samples collected in Pakistan and Thailand, built a database and re sequenced the genome. Sequencing produced a batch of 140g genome re sequencing raw data. To provide basic data for the study of the adaptation of domestic animals to the extreme environment of the Qinghai Tibet Plateau, to explore the historical events of domestication, migration and expansion of the main domesticated animals in the region, and to further explore the adaptation mechanism of domesticated animals to the poor environment such as hypoxia, high cold and dry.

2、Keywords

Theme：Forest
Discipline：Terrestrial Surface
Places：Pan-Third Pole
Time：2019

3、Data details

1.Scale：None

2.Projection：

3.Filesize：140000.0MB

4.Data format：None

4、Space scope

|  |  |  |
| --- | --- | --- |
| - | north：36.945572 | - |
| west：63.153851 | - | east：75.259272 |
| - | south：32.293902 | - |

5、Time frame:2019-01-06 16:00:00+00:00--2020-01-06 03:59:59+00:00

6、Reference method

References to data:

LI Yan. Genome resequencing data of main domestic animals in the Tibetan Plateau (2019). A Big Earth Data Platform for Three Poles, doi:10.11888/Ecolo.tpdc.2704342020

References to articles:

7、Supporting project information

Pan-Third Pole Environment Study for a Green Silk Road-A CAS Strategic Priority A Program
Second Tibetan Plateau Scientific Expedition Program

8、Data resource provider

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