fastx-utils之根据标识符去除重复序列: dedup

一、fastx-utils dedup 介绍

功能描述:

fastx-utils dedup 根据序列去除重复。

命令行接口:

\$ fastx-utils dedup

Usage: fastx-utils dedup <in.fa/fq>

二、使用场景实例及其用法

示例演示:

示例文件: protein.faa

\$ cat protein.faa | head -n 10

>AGE80385.1

MKKIIIISATTIVIGITSFAYFGSKTPLHNEAKAVESQKHNNHKKEEIPAFPKADHNAKKIDNDFSVVTNPKSNLVLINK HRKLPDGYIPEDLTRPNVPFISPKDKEKTLLRKDAAEALENMFKAAKKEGLDLTAVSGYRSYKRQKSLHDTYVRRQGKAE ANSVSAIPGTSEHQTGLAMDISSKSAKFQLEPIFGETAEGKWVAEHAHEFGFVIRYLEDKTDTTEYAYEPWHLRYVGNPY ATYLYKHHLTLEEAMEDKK

>AGE79558.1

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>AGE80385.1

 ${\tt MQNGKVKWFNSEKGFGFIEVEGGEDVFVHFSAIQGEGFKTLEEGQEVTFEVEQGNRGPQATNVNKK}$

>AGE81522.1

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运行命令: 去除 protein. faa 文件中的重复序列.

\$ fastx-utils dedup protein.faa | head -n 10

>AGE80385.1

MKKIIIISATTIVIGITSFAYFGSKTPLHNEAKAVESQKHNNHKKEEIPAFPKADHNAKKIDNDFSVVTNPKSNLVLINK HRKLPDGYIPEDLTRPNVPFISPKDKEKTLLRKDAAEALENMFKAAKKEGLDLTAVSGYRSYKRQKSLHDTYVRRQGKAE ANSVSAIPGTSEHQTGLAMDISSKSAKFQLEPIFGETAEGKWVAEHAHEFGFVIRYLEDKTDTTEYAYEPWHLRYVGNPY ATYLYKHHLTLEEAMEDKK

>AGE79558.1

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>AGE81522.1

MKKTILTTTAALTMMGTGMGINVDHIKPAEVKADTISFYDVPNNHWATKAITNLANRNIVVGYGNGQFGFGDNVTRGQVA RMIYNYLKPADAGNFKNPFSDIKGHMFEKEILALAKVGIIKGYGEGKFGPDDILTREQMAQVLTNAFKFEGTKKTSFVDV DKNSWSYKAIGALEEKGVTIGTGGNMYSPTSVVTREQYSQFLFNSINVIEKETKPEEKPNTGGEVKPEEKPNTGGEVKPEEKPNTGGETKPVNIPEWLETSLATNDFTFTQAWYDGSEAINKAASTNAQQIVKNINSKYGTNLKYSEVGAIVQLVDGAREQLWLAGMNVNDFRVTFRVSNNAMIELTKELVTLVNSDLNLDQEIQEIPSAPMKIKNVEKGDYKIRISPAMADQMITIIIE

>AGE80137.1

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