

Genome of Chinese plum provides hope of improved fruit

A Chinese research team, led by Beijing Forestry University, BGI, Beijing Lin Fu Ke Yuan Flowers Co., Ltd, and other institutes, has completed the first genomic sequence of *Prunus mume*, known as mei.

Wenbin Chen, Project Manager of BGI, said, “The *P. mume* genome lays a solid foundation for identification of important economic traits, and provides a valuable resource for *P. mume* breeding and other Rosaceae species studies. The work here also brings a new approach for further exploring the biosynthesis of floral scent and regulation mechanism of early blooming in endodormancy, and other comparative genomics studies on Rosaceae species.”

View the original article here: [First Genome Sequence of Chinese Plum Provides Important Resource for Fruit Improvement](#)