## Fengli Fu

## List of Publications by Year

 in descending orderSource: https:|/exaly.com/author-pdf/12029698/publications.pdf
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The Role of Deleterious Substitutions in Crop Genomes. Molecular Biology and Evolution, 2016, 33, 2307-2317.

The Medicago sativa gene index 1.2: a web-accessible gene expression atlas for investigating expression differences between Medicago sativa subspecies. BMC Genomics, 2015, 16, 502.

A Roadmap for Functional Structural Variants in the Soybean Genome. G3: Genes, Genomes, Genetics, 2014, 4, 1307-1318.

An RNA-Seq based gene expression atlas of the common bean. BMC Genomics, 2014, 15, 866.
$1.2 \quad 142$

Zinc Finger Database (ZiFDB) v2.0: a comprehensive database of C 2 H 2 zinc fingers and engineered zinc finger arrays. Nucleic Acids Research, 2012, 41, D452-D455.
6.5

Predicting success of oligomerized pool engineering (OPEN) for zinc finger target site sequences.
BMC Bioinformatics, 2010, 11, 543.

Zinc Finger Database (ZiFDB): a repository for information on C 2 H 2 zinc fingers and engineered zinc-finger arrays. Nucleic Acids Research, 2009, 37, D279-D283.

High-frequency modification of plant genes using engineered zinc-finger nucleases. Nature, 2009, 459,
442-445.

Unexpected failure rates for modular assembly of engineered zinc fingers. Nature Methods, 2008, 5, 374-375.
9.0

385

Rapid â€œOpen-Sourceâ€•Engineering of Customized Zinc-Finger Nucleases for Highly Efficient Gene Modification. Molecular Cell, 2008, 31, 294-301.

Standardized reagents and protocols for engineering zinc finger nucleases by modular assembly. Nature Protocols, 2006, 1, 1637-1652.

