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· Identification of novel and conserved microRNAs related to drought stress in potato by deep sequencing	2016/10/27
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· Comparative proteomic analysis of cold-induced sweetening	2016/10/27
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· Overexpression of ERF1 gene activates stress-responsive genes and confers enhanced tolerance to drought stress in transgenic potato	2016/10/27
· Study on transcriptional expression of key genes involved in steroidal glycoalkaloid biosynthesis in potato micro-tuber	2016/10/27
· Improving farming practices reduce the carbon footprint of spring wheat production	2016/10/27
· Water-Saving Innovations in Chinese Agriculture	2016/10/27
· Microbial community diversities and taxa abundances in soils along a seven-year gradient of potato monoculture using high throughput pyrosequencing approach	2016/10/27
· Soil water status and root distribution across the rooting zone in maize with plastic film mulching	2016/10/26
· Evidence that the C-terminal region is involved in the stability and functionality of OprM in E. coli	2016/10/26
· Identification of QTLs for seedling vigor in winter wheat	2016/10/26
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