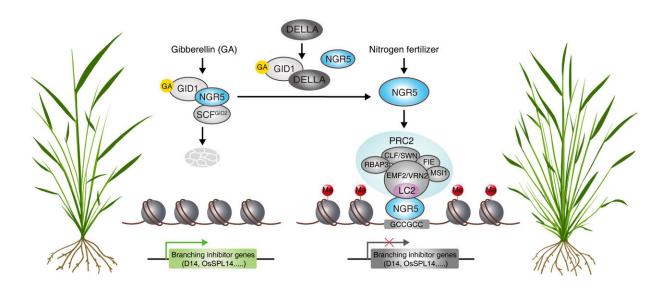


Branching out for a new green revolution

February 7 2020



GA-induced H3K27me3 modification enhances nitrogen stimulation of tillering. Credit: Xiangdong Fu, Institute of Genetics and Developmental Biology, Chinese Academy of Science, Beijing, China

Researchers at the University of Oxford and the Chinese Academy of Sciences have discovered a new gene that improves the yield and fertilizer use efficiency of rice.

The worldwide late-20th century 'Green Revolution' saw dramatic year-by-year increases in global grain yields of rice and other cereals. The Green Revolution was fueled by new high-yielding dwarfed Green Revolution Varieties (GRVs) that are still in widespread use today, and by increased fertilizer use.



The numbers of grain-bearing branches ('tillers') per plant are increased in GRVs, and further enhanced by increased nitrogen fertilizer use, thus boosting grain yield. However, fertilizers are costly to farmers and cause extensive environmental damage. Developing new GRVs combining increased tiller number and grain yield with reduced nitrogen use is thus an urgent global sustainable agriculture goal.

A major new study, published today as the cover story of journal *Science*, led by Professor Xiangdong Fu from the Chinese Academy of Sciences' Institute of Genetics and Developmental Biology, and Professor Nicholas Harberd from the Department of Plant Sciences at the University of Oxford, part-funded by the BBSRC-Newton Rice Initiative, has for the first time discovered a gene that can help reach that goal.

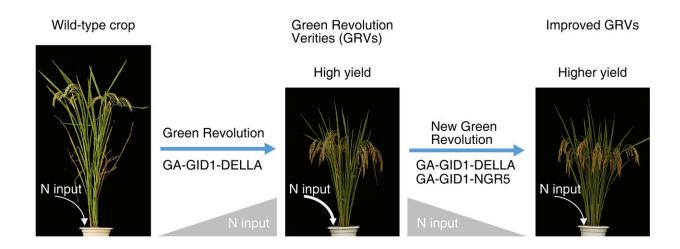
The study identified a rice gene that responds to nitrogen, and hence increases the accumulation in plant cells of a protein called NGR5. Nitrogen-stimulated NGR5 accumulation then alters the structure of genes that inhibit tiller growth, switching them off and thus increasing the numbers of yield-enhancing tillers.

Professor Harberd said: "Discovering how nitrogen stimulates tiller growth was exciting in itself. But our discovery was particularly exciting because NGR5 controls the activity (via a mechanism known as chromatin modulation) of multiple genes in the rice genome, genes likely responsible for many different rice responses to soil nitrogen in addition to tiller growth."

The increased tiller number of GRVs is also caused by accumulation of another branching-promoting protein called DELLA, an accumulation that is reduced by the plant hormone gibberellin (GA). The study found that GA also reduces NGR5 accumulation, and that tiller growth is the product of complex interactions between the NGR5 and DELLA



proteins.



Increased NGR5 activity boosts grain yield and nitrogen use efficiency. Credit: Xiangdong Fu, Institute of Genetics and Developmental Biology, Chinese Academy of Science, Beijing, China

Professor Harberd said: "We next reasoned that further increase in the accumulation of NGR5 might increase tiller number and yield with reduced fertilizer use. To our delight, we found that increasing NGR5 accumulation caused an increase in both tiller number and grain yield of a current elite <u>rice</u> GRV, especially at low fertilizer levels."

The researchers say NGR5 should now become a major target for plant breeders in enhancing crop yield and <u>fertilizer</u> use efficiency, with the aim of achieving the global grain yield increases necessary to feed a growing world population at reduced environmental cost.

Professor Harberd added: "This study is a prime example of how



pursuing fundamental plant science objectives can lead rapidly to potential solutions to global challenges. It discovers how <u>plants</u> coordinate their growth in response to soil nitrogen availability, then shows how that discovery can enable breeding strategies for sustainable food security and future new green revolutions."

More information: K. Wu el al., "Enhanced sustainable green revolution yield via nitrogen-responsive chromatin modulation in rice," *Science* (2020). science.sciencemag.org/cgi/doi ... 1126/science.aaz2046

Provided by University of Oxford

Citation: Branching out for a new green revolution (2020, February 7) retrieved 19 April 2024 from https://phys.org/news/2020-02-green-revolution.html

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