



Development of Next-Generation CO2-Fixing Plant Through the Gene Optimization, Distant Hybrid, and Microbial Symbiosis

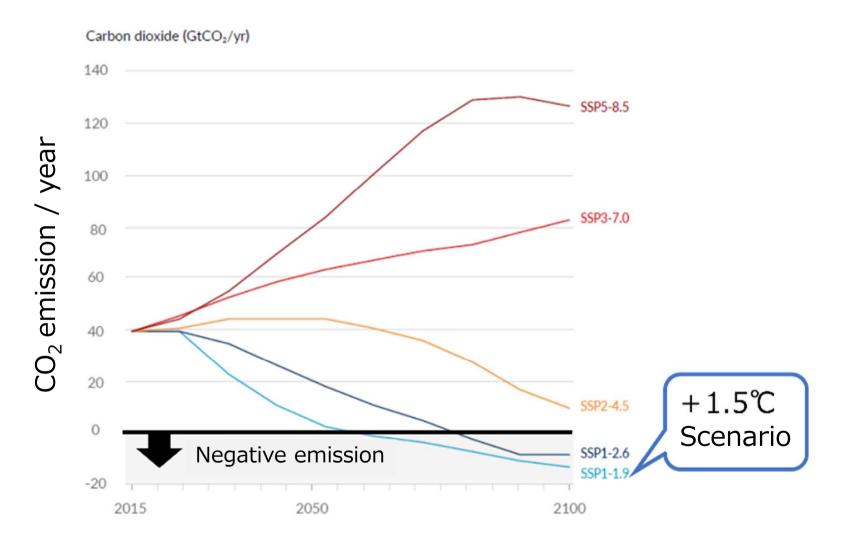


PM : Nobutaka MITSUDA Natl. Inst. Adv. Ind. Sci. Tech. (AIST)PJ member : AIST, Tokyo Metropolitan Univ., SUMITOMO FORESTRY Co., Ltd.

Background

• To stop global warming…

Five scenarios about transition of CO_2 emission (IPCC AR6)

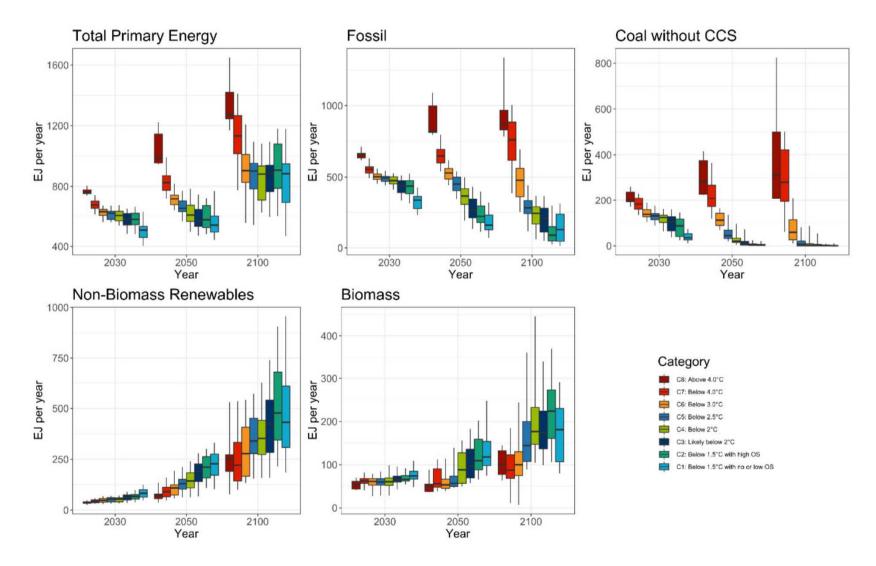


To achieve the +1.5°C scenario, CO₂ emissions in NET must be reduced even from 2023, and negative emissions in NET must be achieved around 2055

Background

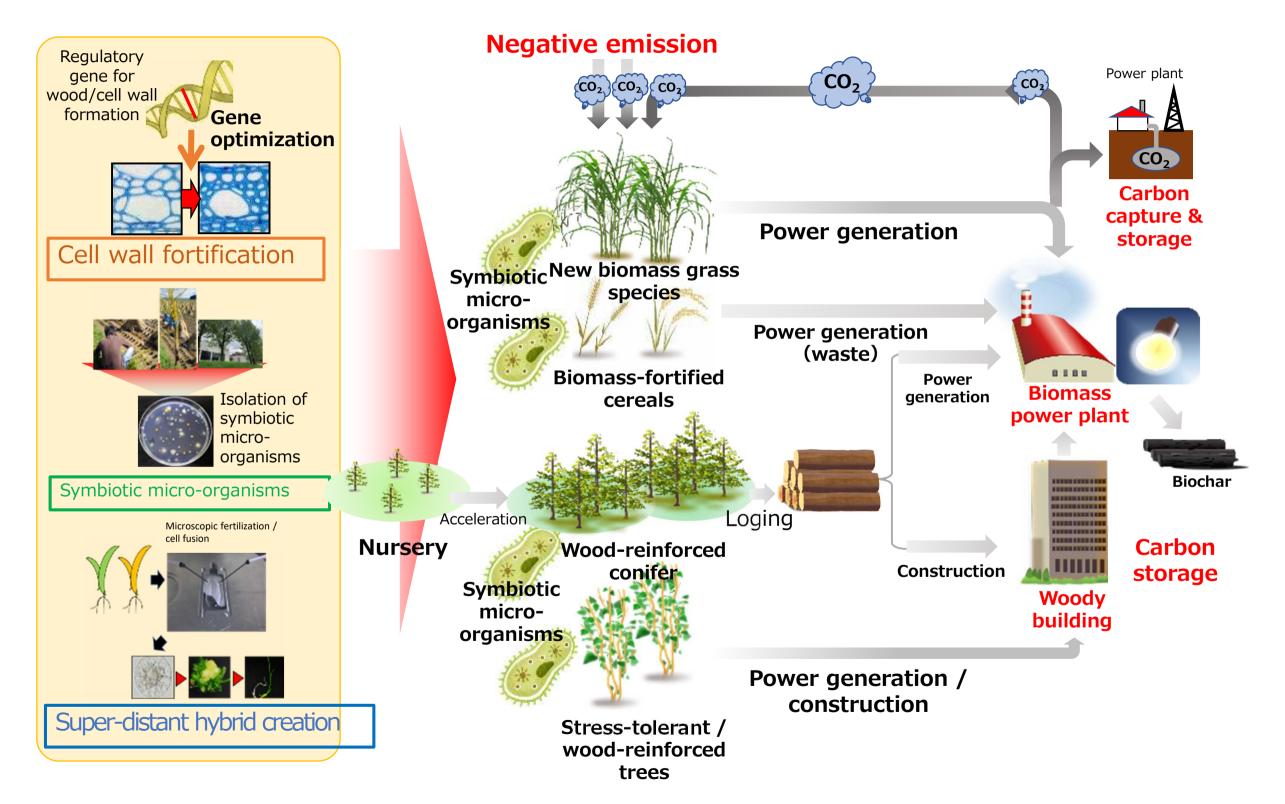
• To achieve +1.5℃ scenario …

Energy source transition in each scenario (IPCC AR6)

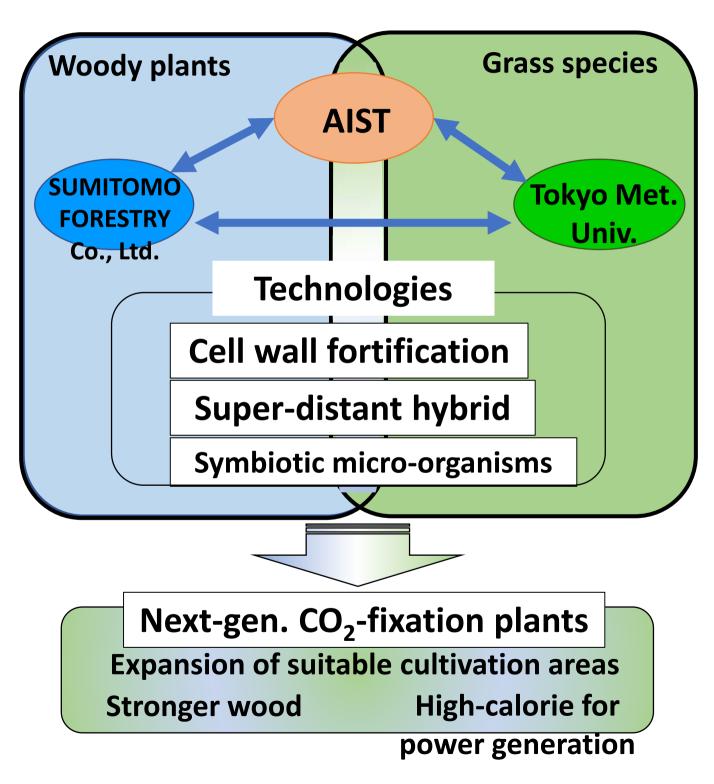


- For +1.5°C scenario, about a quarter of all energy consumption must come from biomass in 2100
- > High expectations for biomass, not only for energy

Graphical abstract



Project's structure · Goals



- Aiming to improve biomass productivity by 30% compared to conventional methods for genetic optimization, hyper distantly related hybrids, and symbiotic microorganism optimization, respectively.
 - Results can be used in combination with conventional breeding
- Using the optimization of symbiotic micro-organisms as a hub, combined with super distantly related hybrids in the short term and genetic optimization in the long term, aim to increase biomass productivity by 50% compared to conventional methods. 5

R&D schedule

	2022~2023FY	2024FY
1. Establishment of biomass enhancement strategy by gene optimization	Wood reinforcement in woody and herb Examination of primary cell wall enhance	>
2. Establishment of new biomass plant creation method by super-distant hybrid creation technology	Development of enhanced cell fusion sy Creation of new biomass plants by sup	>
3. Establishment of plant growth promotion system by symbiotic micro- organism	Exploration of symbiotic micro-organism Evaluation of the effect of symbiotic mic	

Three major technologies

Gene optimization

Super-distant hybrids

Symbiotic micro-organisms

Three major technologies

Gene optimization

Super-distant hybrids

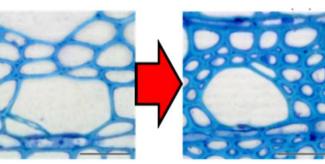
Symbiotic micro-organisms

Biomass fortification by gene optimization

Three strategies

Based on gene edit technology

1. Wood reinforcement

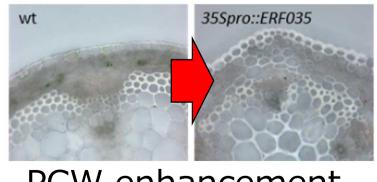


Wood reinforcement

- 2. Increased strength
 - Beneficial trait in addition to higher wood productivity



3. Primary cellwall enhancement(only for grass)



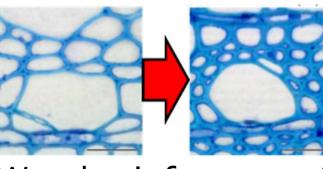
PCW enhancement

Biomass fortification by gene optimization

Three strategies

Based on gene edit technology

1. Wood reinforcement



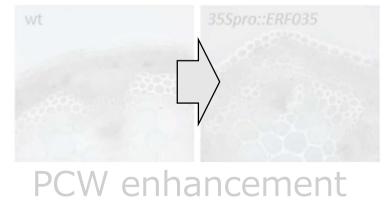
Wood reinforcement

2. Increased strength

Beneficial trait in addition to higher wood productivity

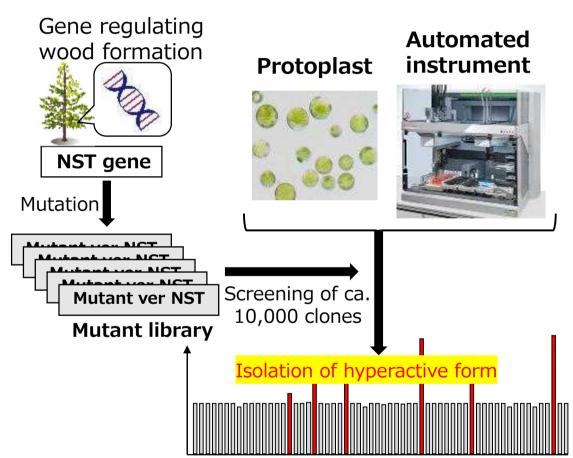


3. Primary cell wall enhancement (only for grass)



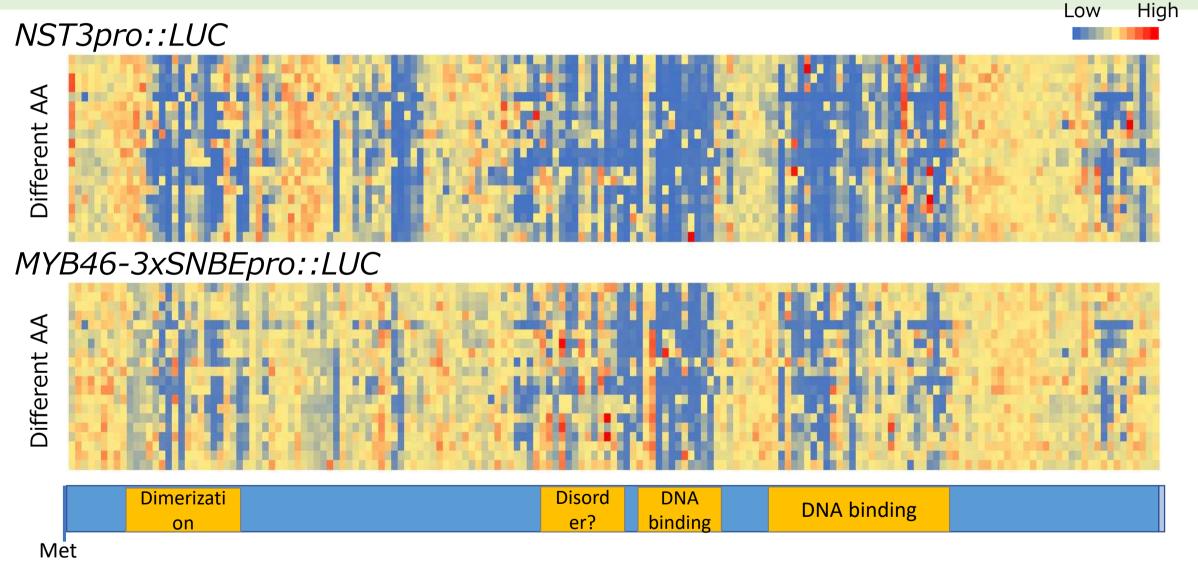
Hyperactivating NST transcription factor regulating wood formation

NST transcription factor Getting hyperactive form of NST transcription factor NST WT knockout



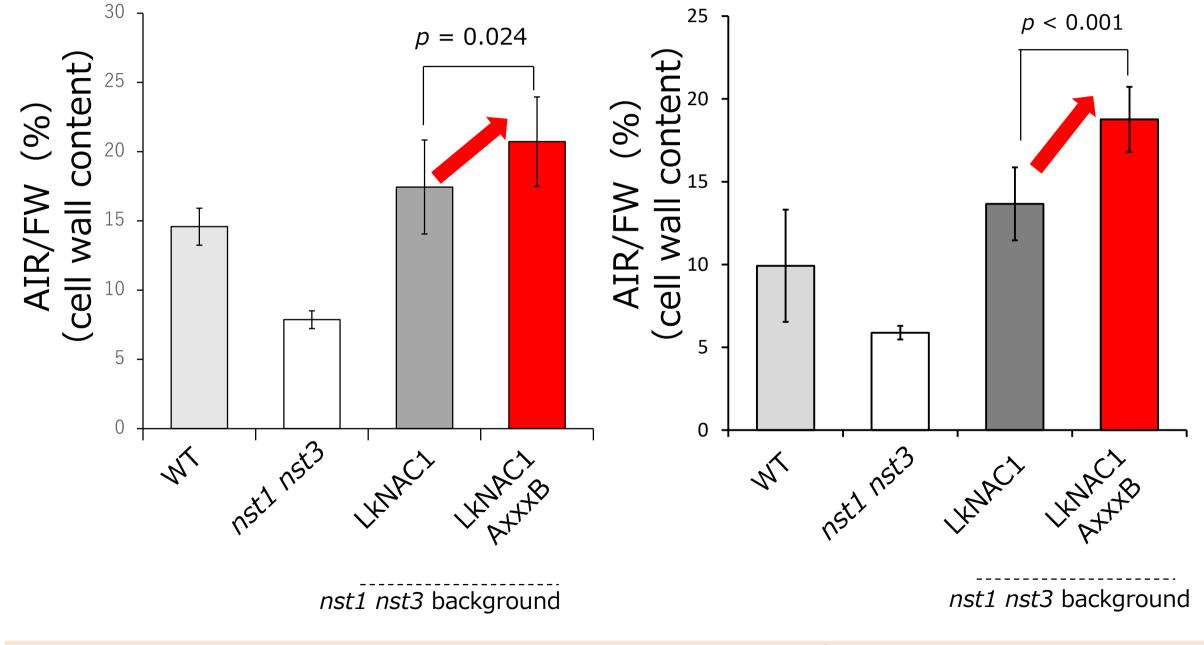
 \rightarrow Developing a technology to apply the gene edit in practical plants

 Results of NST transcription factor activity measurements utilizing two reporters.



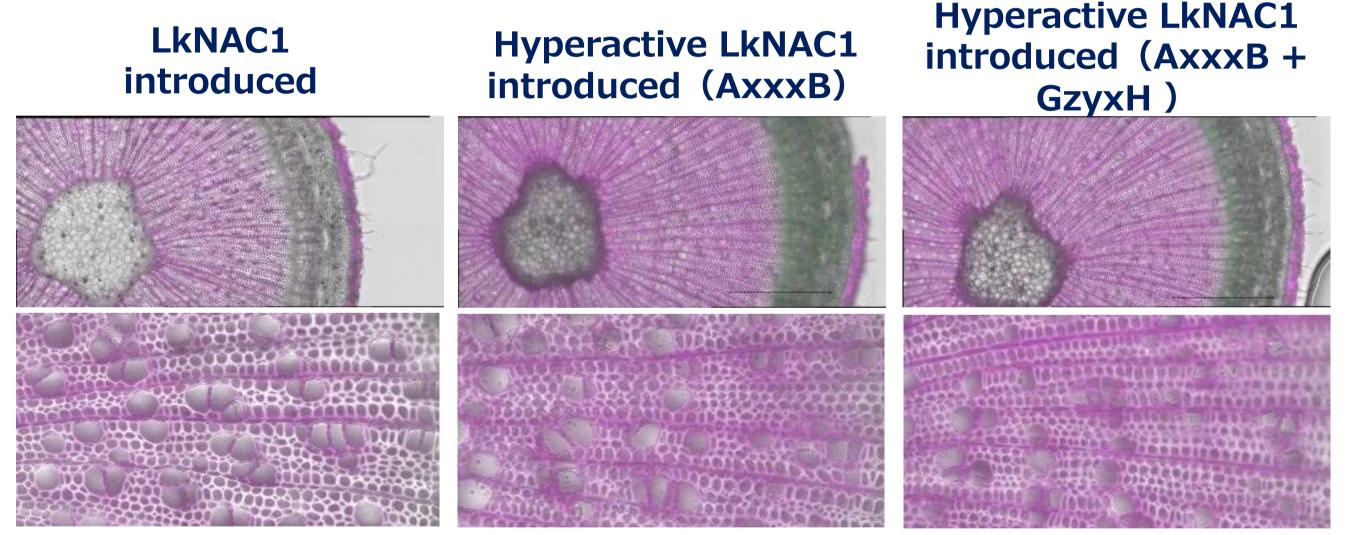
> NST転写因子を高活性化する変異を約60カ所同定(産総研ー住友林業BIP)

 Cell wall amount of transgenic Arabidopsis thaliana for NST transcription factor genes with candidate mutations.



▶ 有望変異が木質増強効果を有することをシロイヌナズナで確認(産総研-住友林業BIP)

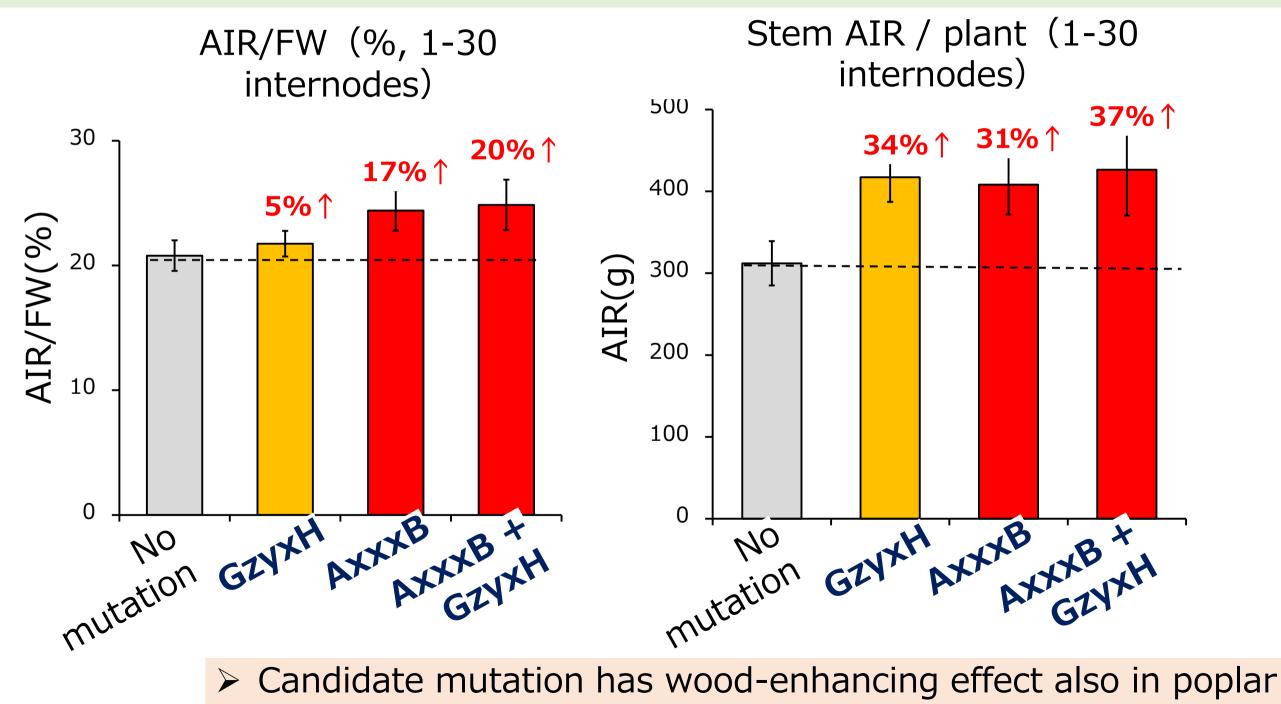
Observation of cross sections of transgenic poplar for NST transcription factor gene with candidate mutations



Pinkish color: Lignin autofluorescence under UV irradiation in pseudo color

Candidate mutation has wood-enhancing effect also in poplar

 Cell wall amount of transgenic poplar for NST transcription factor genes with candidate mutations.

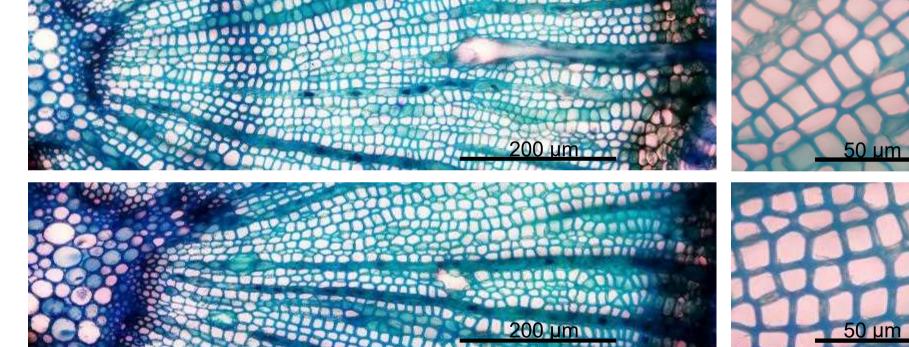


 Observation of cross sections of transgenic larch for NST transcription factor gene with candidate mutations

introduced Hyperactive LkNAC1 introduced

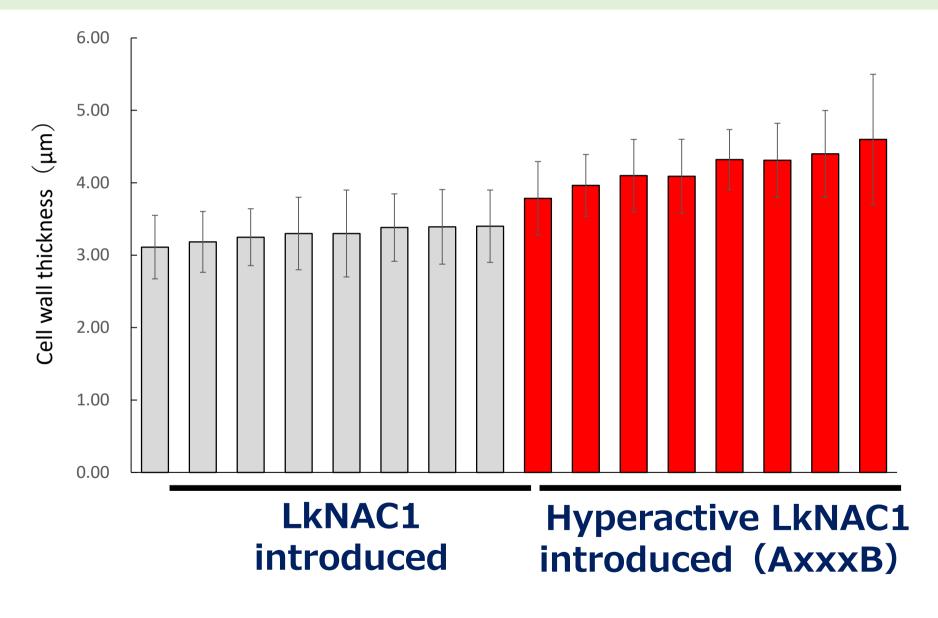
(AxxxB)

LkNAC1



> Candidate mutation has wood-enhancing effect also in larch

 Larch cell wall thickness with NST transcription factor genes with candidate mutations.



Candidate mutation has wood-enhancing effect also in larix

Three major technologies

Gene optimization

Super-distant hybrids

Symbiotic micro-organisms

Development of new crops and biomass plants through distant hybridization

1. Development of efficient cell fusion technology using peptide PEG-lipid conjugates

`N

∽______ 0______ 0⊝

sperm

egg

2. New hybrid Eucalyptus with expanded suitable cultivation areas

3. Creation of new rice and wheat plants and verification of their potential



Development of new crops and biomass plants through distant hybridization

1. Development of efficient cell fusion technology using peptide PEG-lipid conjugates

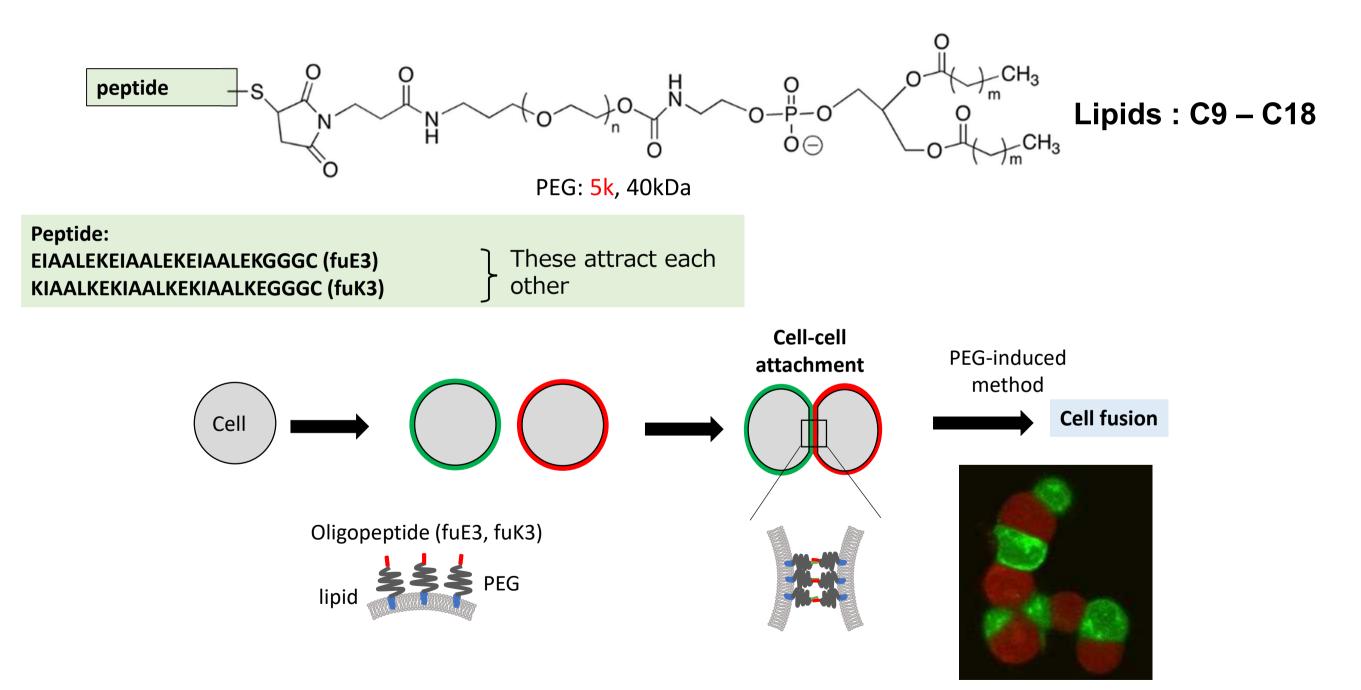
 $\mathcal{H}_{\mathsf{H}} \rightarrow \mathcal{H}_{\mathsf{o}} \rightarrow$

2. New hybrid Eucalyptus with expanded suitable cultivation areas

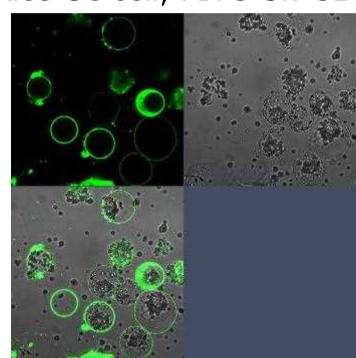
3. Creation of new rice and wheat plants and verification of their potential

peptide +s

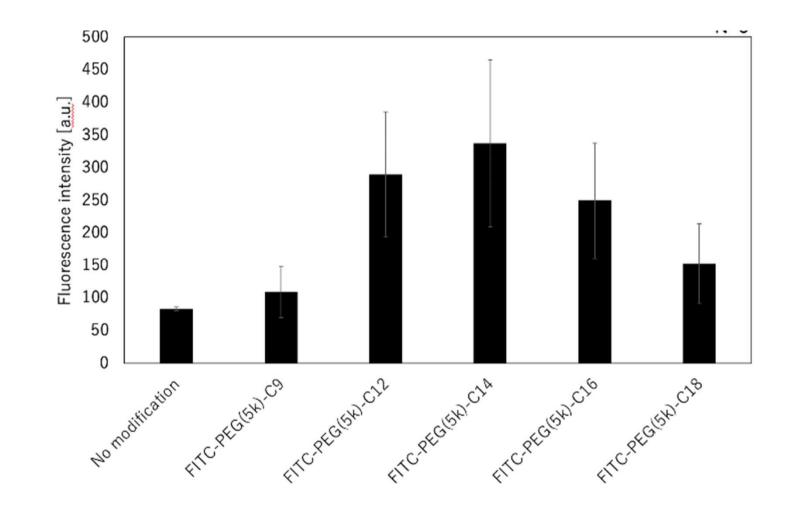
4. Creation of new biomass plants and verification of their potential



The length of carbon chain was optimized in rice Oc cells.

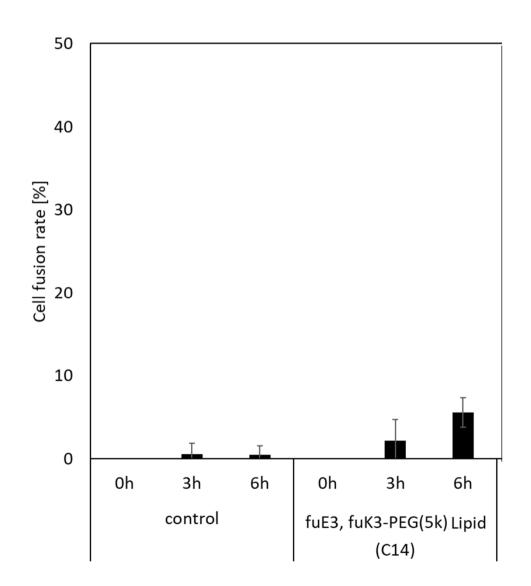




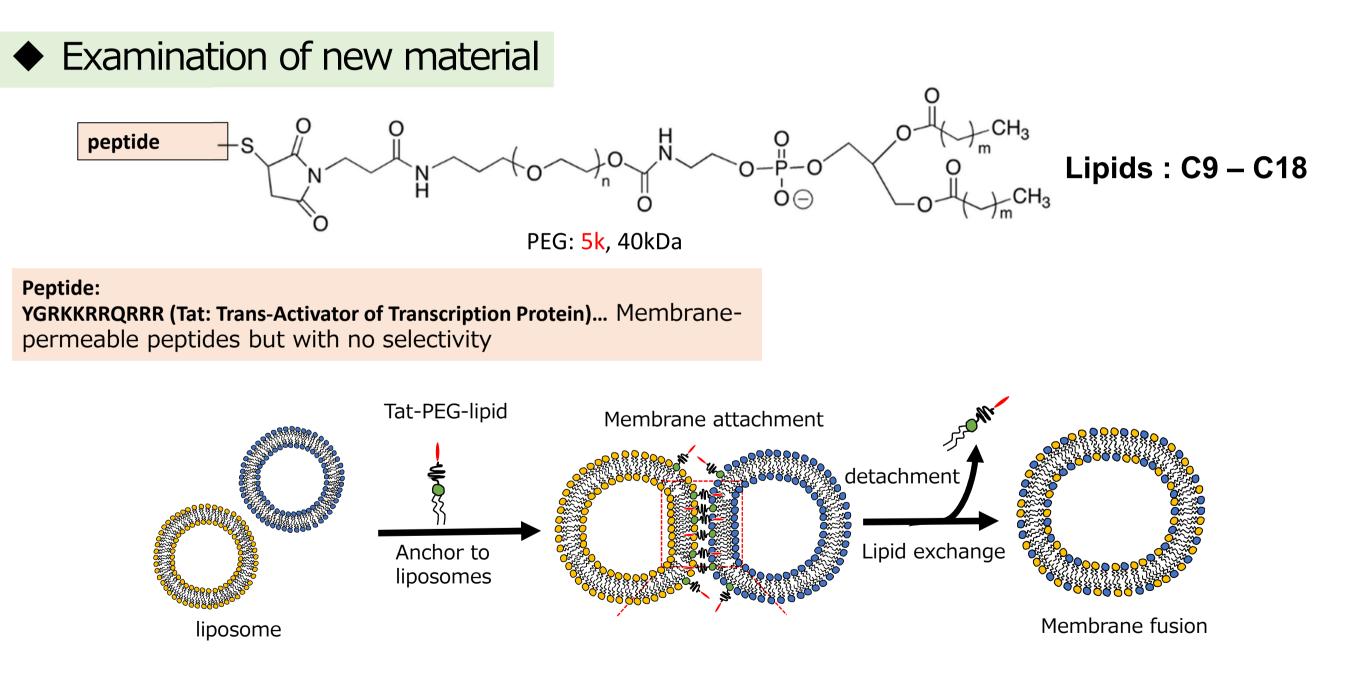


Determine optimal structure for plant cells

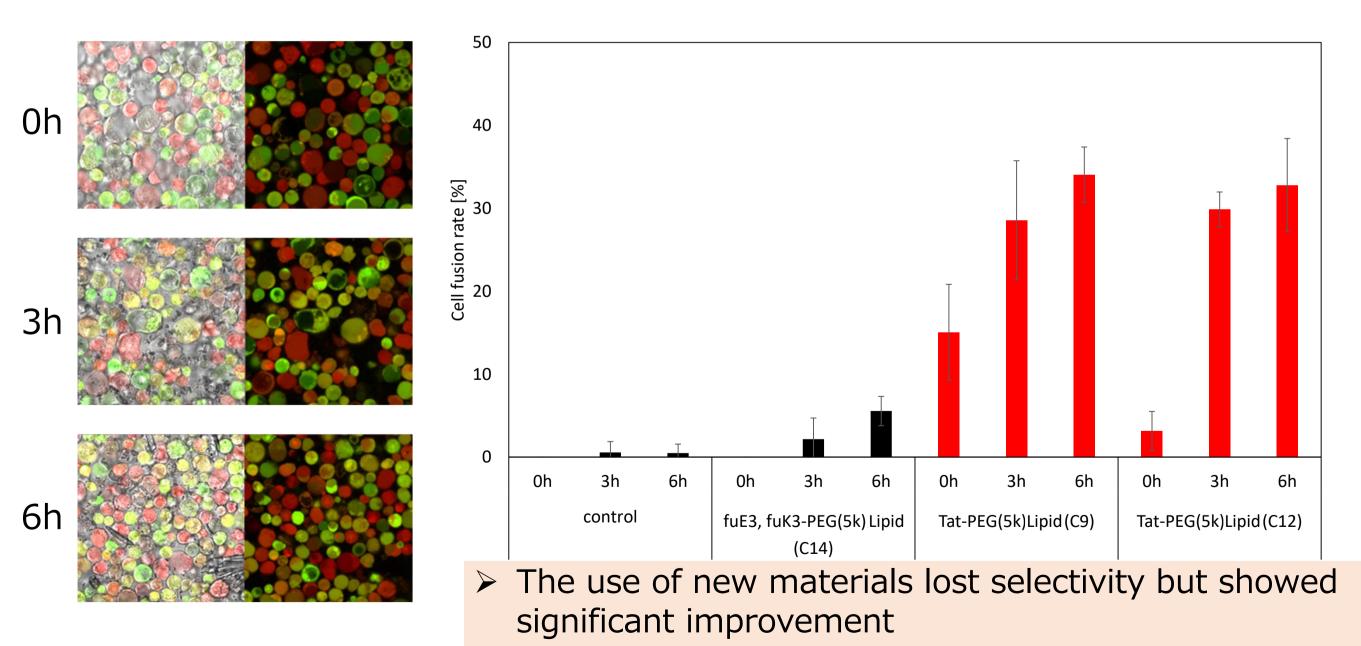
• Examination in rice protoplast



Improvement of cell fusion was observed but the effect was limited
23



Examination of new material in rice protoplast



Development of new crops and biomass plants through distant hybridization

1. Development of efficient cell fusion technology using peptide PEG-lipid conjugates

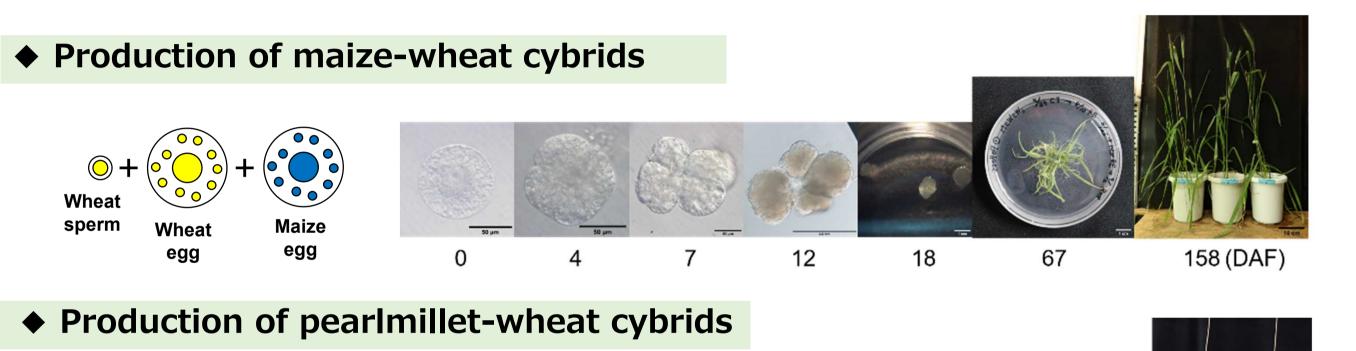
2. New hybrid Eucalyptus with expanded suitable cultivation areas

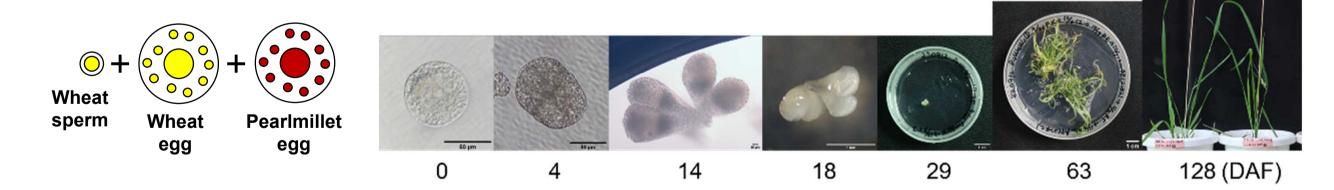
3. Creation of new rice and wheat plants and verification of their potential

4. Creation of new biomass plants and verification of their potential

sperm

egg



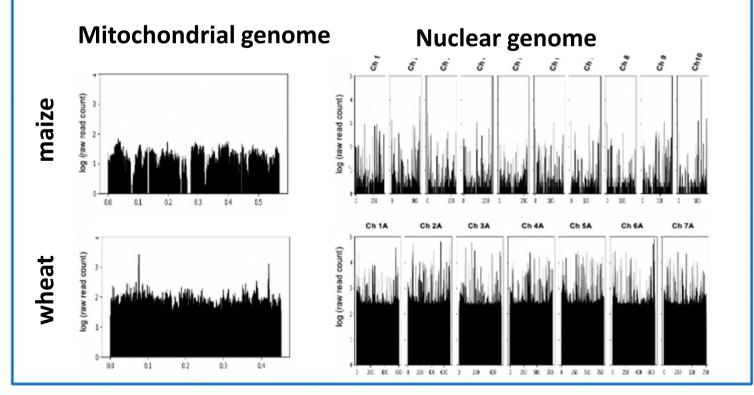


Basic elemental technologies have been established.

• Genome composition of maize-wheat cybrids

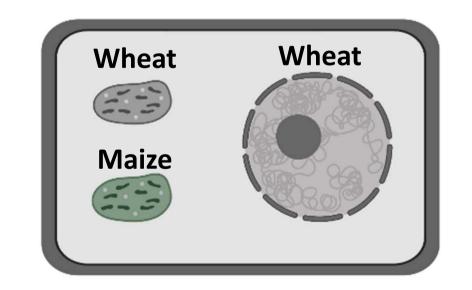
- Nuclear genome: almost wheat
- Mitochondria genome: Maize + wheat

Mapping profiles of sequence reads from maize-wheat cybrid genome Mitochondrial genome





Maize-wheat cybrid



Confirmed the intended cybrid is created.

Calli from rice-setaria hybrid zygotes and their genome composition



The elemental technology to create up to a fusion callus could be established.

Rice nuclear genome

《Future plans》

- Analysis of nuclear and cytoplasmic genome abundance in hybrid plants
 - **Genome analysis**
 - Maize \times Wheat : Done for 6 lines
 - Pearlmillet × Wheat : Underway for 11 lines raw data obtained
 - □ FISH analysis

Examination of agronomic and physiological traits

- □ Maize-wheat seeds, propagated
- □ Photosynthetic potential and mode
- □ Characteristics of cell wall

Rice-C4 Plant callus redifferentiation and determination of genome composition





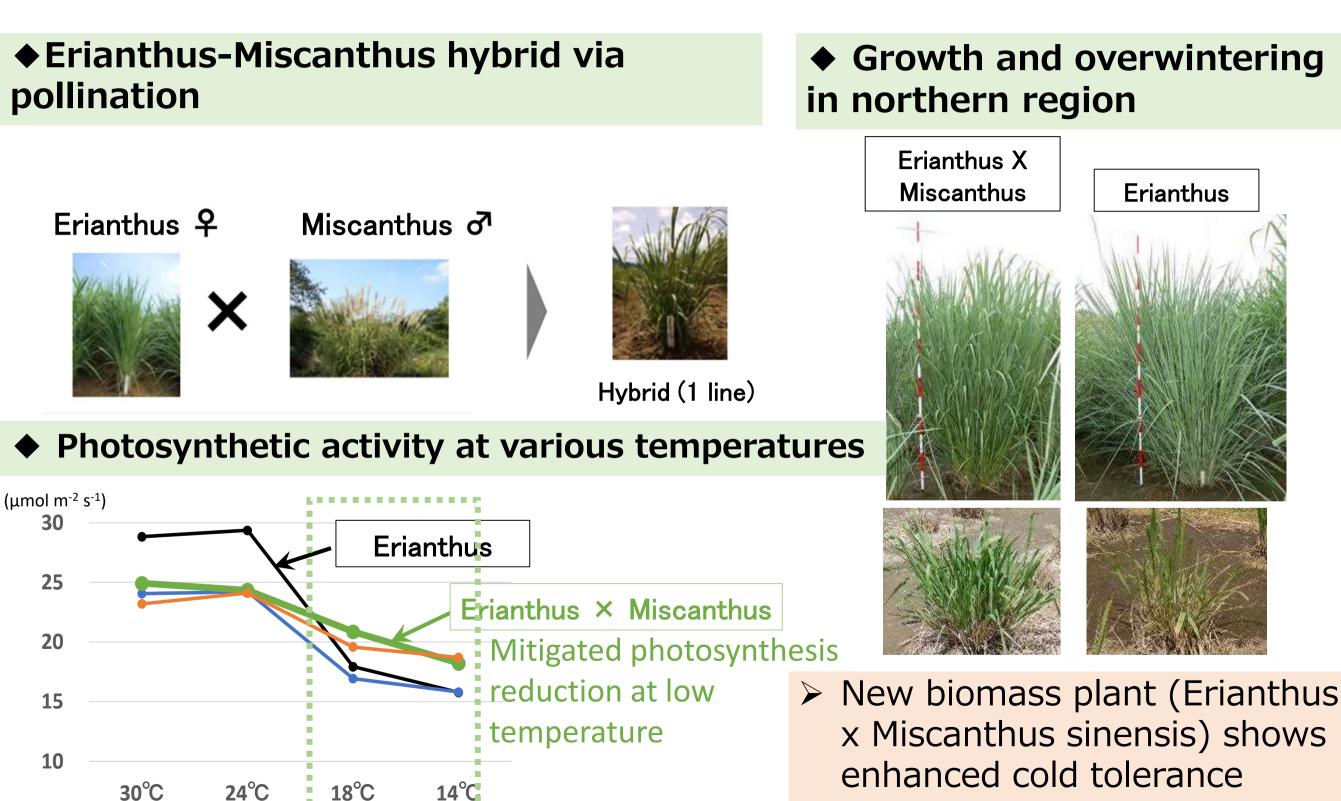
Development of new crops and biomass plants through distant hybridization

1. Development of efficient cell fusion technology using peptide PEG-lipid conjugates

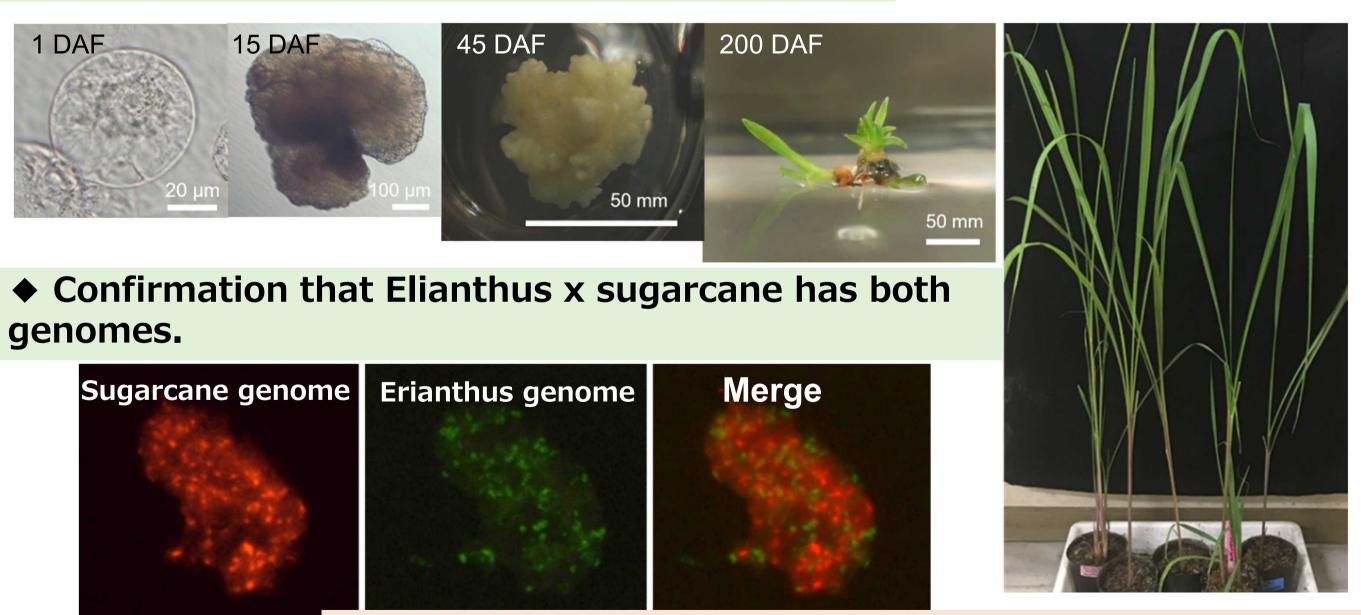
2. New hybrid Eucalyptus with expanded suitable cultivation areas

3. Creation of new rice and wheat plants and verification of their potential

4. Creation of new biomass plants and verification of their potential

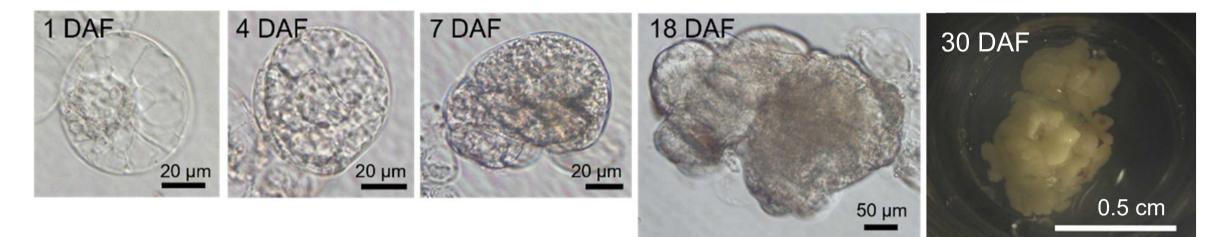


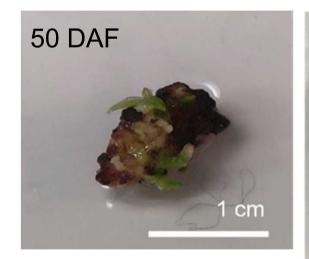
Erianthus × sugarcane by in vitro fertilization

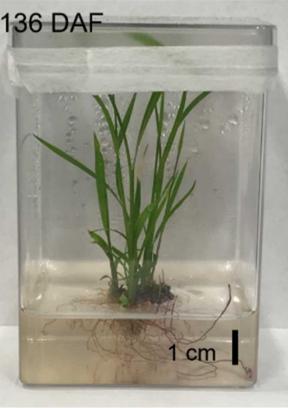


Basic elemental technologies have been established.

Possible Erianthus-Sugarcane/Miscanthu hybrid by in vitro fertilization









Basic elemental technologies have been established.

« Future plans »

Trait evaluation for cold tolerance, biomass production capacity, and photosynthetic potential

Cultivation trials to evaluate cold tolerance, biomass traits, etc. of hybrids of Erianthus x Miscanthus will be conducted in Ishigaki Island, Tottori and Akita.



Propagation and basic growth characterization of Erianthus-Miscanthus hybrid and possible Erianthus-Sugarcane/Miscanthu hybrid

Three major technologies

Gene optimization

Super-distant hybrids

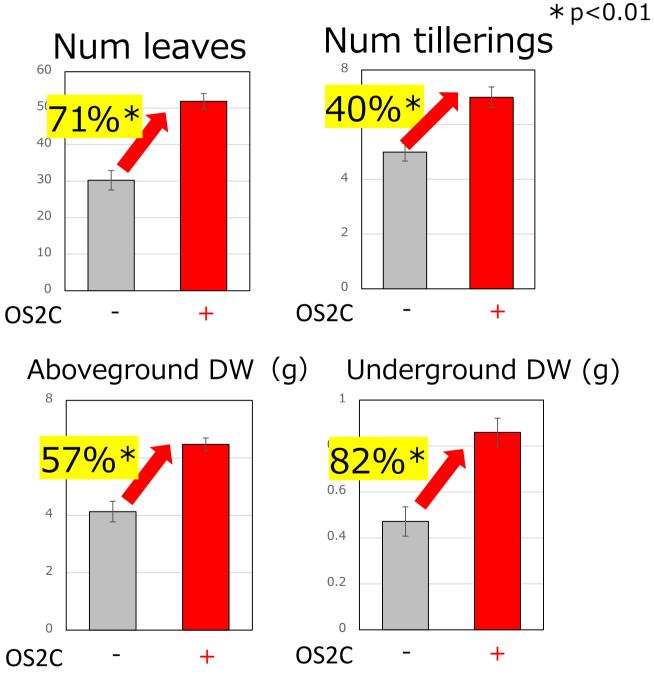
Symbiotic micro-organisms



No inoculation



OS₂C inoculated



Clear growth-promoting effect was confirmed in rice

Growth promotion in rice (harvest stage)

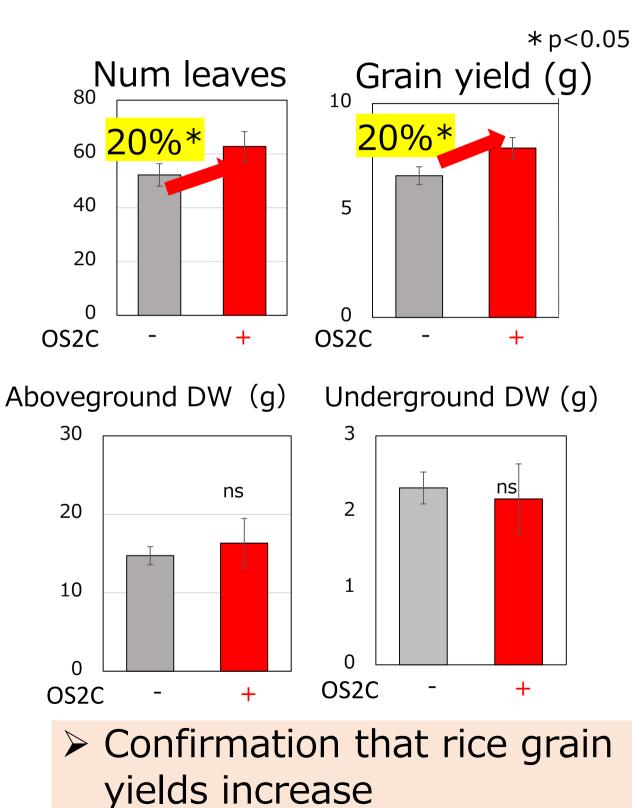
After 5 monts (N=5 [10 individuals])



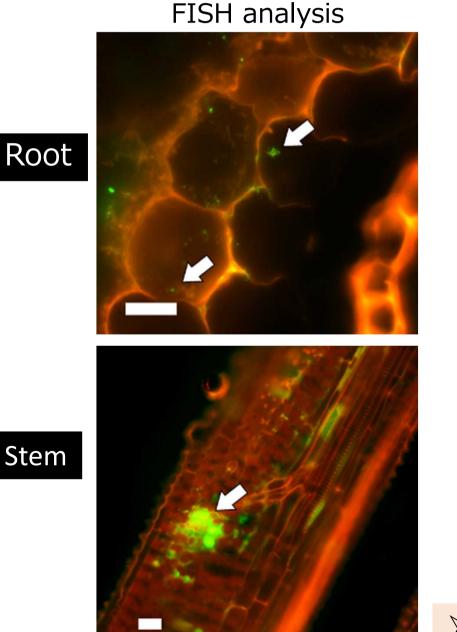
No inoculation



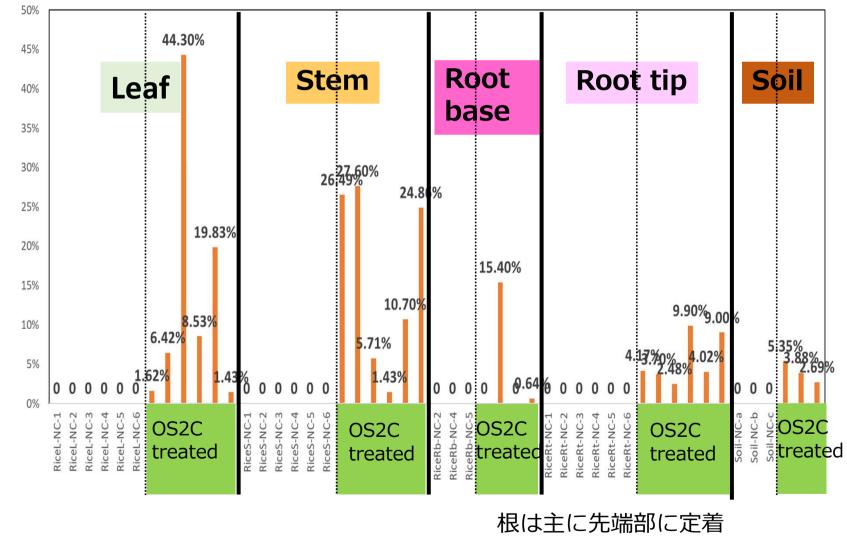
OS2C inoculated



Confirmation of the survival of treated bacteria

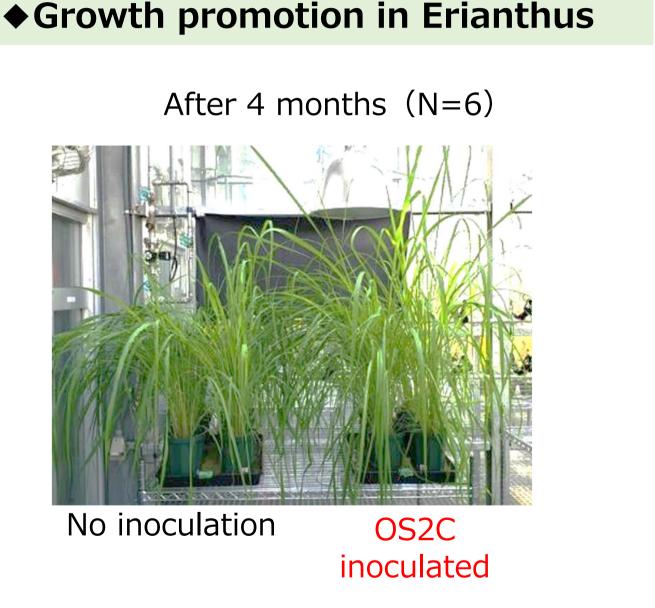


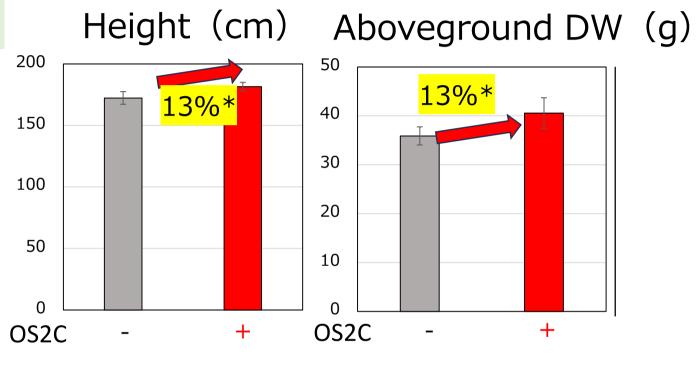
Occupancy of OS2C strain (16SrRNA gene sequence) in each sample



Confirmation of treated bacteria on plants grown for approximately 3 months

*p<0.05



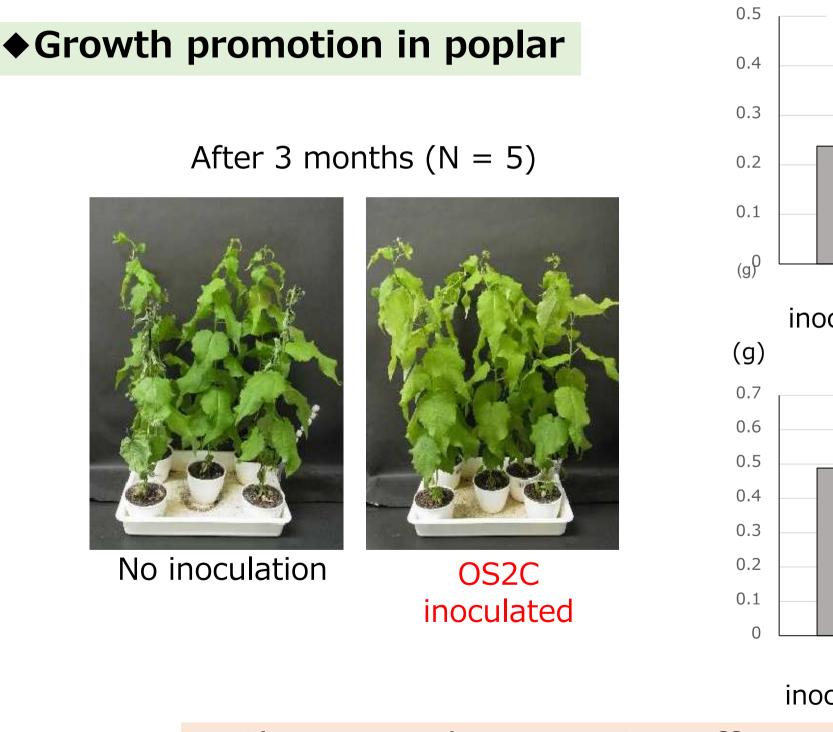


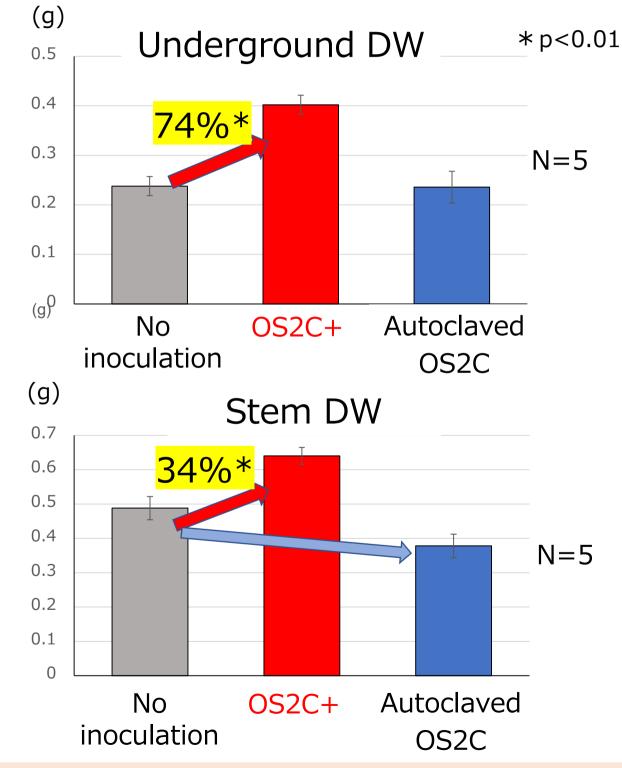
Underground DW (g)

+

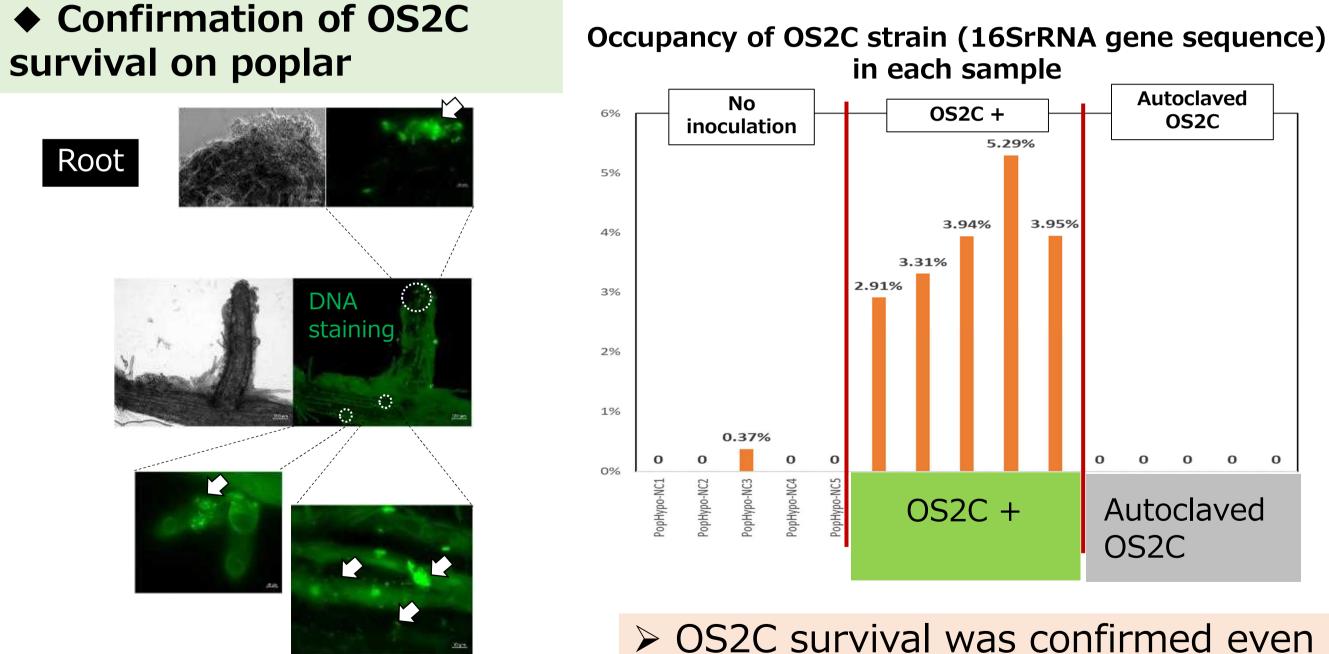
OS₂C

Effect was confirmed even in practical biomass plants



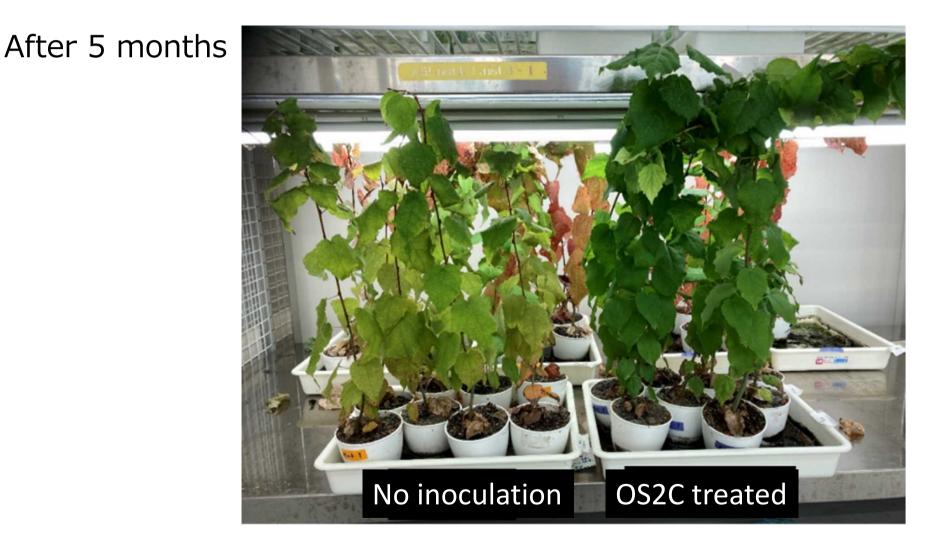


Clear growth-promoting effect was confirmed even in trees



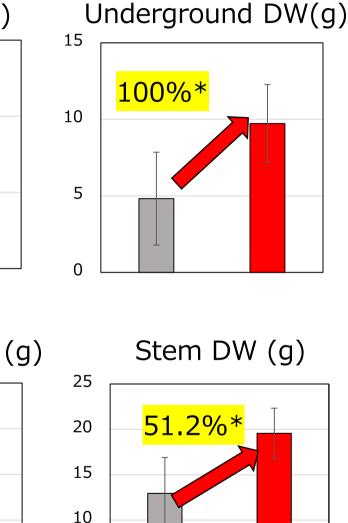
in trees grown for about 3 months

Effect of OS2C to suppress dormancy in poplar



OS2C might suppress dormancy of poplar. Study of this mechanism is underway.

Height (cm) 150 Growth promotion in Eucalyptus 10% 100 After 5 months (N=5)50 0 Aboveground DW (g) 50 <mark>37.1%</mark> 40 30 20 10 0 No inoculation OS2C inoculated



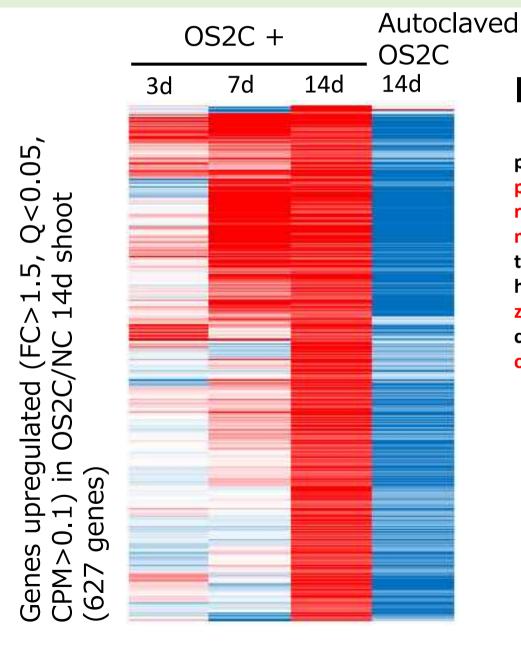
> 実用樹木でも明らかな生育促進効果を確認

5

0

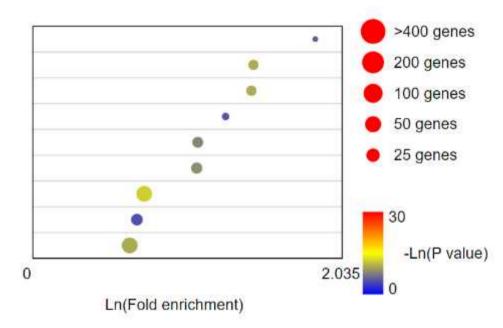
*p<0.05

RNA-seq analysis in rice to elucidate the mechanism of growth promotion



Enrichment analysis for upregulated genes

plant-type cell wall organization peroxidase activity response to oxidative stress metal ion transport transporter activity hydrolase activity, hydrolyzing ... zinc ion binding carbohydrate metabolic process oxidation-reduction process



OS2C may activate reactive oxygen sequestration in plant

Summary

- Approximately 30% wood enhancement in larch introduced with an hyperactive NST transcription factor
- New cell fusion-promoting material showed breakthrough high effectiveness
- Erianthus x Miscanthus hybrid showed improved cold tolerance compared to Erianthus
- Partial elucidation of the mechanism of the growth-promoting effect of OS2C plant symbiotic microorganisms