

## **Colby College** Digital Commons @ Colby

Undergraduate Research Symposium (UGRS)

Student Research

2007

## Characterization of AFN1, a Gene Associated with Cereal Grain Germination

Tenzin Tsewang Colby College

Russell Johnson Colby College

Follow this and additional works at: https://digitalcommons.colby.edu/ugrs



Part of the Cell Biology Commons, and the Molecular Genetics Commons

#### **Recommended Citation**

Tsewang, Tenzin and Johnson, Russell, "Characterization of AFN1, a Gene Associated with Cereal Grain Germination" (2007). Undergraduate Research Symposium (UGRS). 43. https://digitalcommons.colby.edu/ugrs/43

This Article is brought to you for free and open access by the Student Research at Digital Commons @ Colby. It has been accepted for inclusion in Undergraduate Research Symposium (UGRS) by an authorized administrator of Digital Commons @ Colby.



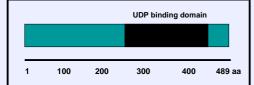
## Characterization of AFN1, a gene associated with cereal grain germination Tsewang, TD, Gilg, IC, and Johnson, RR



Colby College, Waterville, ME 04901

#### **Abstract**

The AFN1 gene is transiently expressed in germinating oat grains. As AFN1 is not expressed in dormant oat grains during imbibition, we hypothesize that AFN1 may be involved in stimulating the germination process. Sequence analysis of an AFN1 cDNA clone indicates that the AFN1 polypeptide is similar to a previously identified abscisic acid (ABA) glucosyl transferase. This suggests that AFN1 may be acting to glucosylate ABA, thereby inactivating it. As the hormone ABA is known to inhibit germination, ABA glucosylation/inactivation could lead to germination in grains expressing AFN1. To test this hypothesis, we have constructed an expression plasmid that encodes an MBP::AFN1 (maltose binding protein) fusion protein. E. coli cells carrying the expression plasmid were found to produce the MBP::AFN1 fusion protein as a substantial fraction of total protein. We are currently in the process of purifying the MBP::AFN1 fusion protein by affinity chromatography, so that it can be assayed for ABA glucosyl transferase activity. We also wish to test the effect of AFN1 gene expression during grain imbibition on the germination behavior of the grains. To this end, we have constructed plasmids for the overexpression and RNAi-based suppression of AFN1 in transgenic plants. These plasmids have been introduced into oat cells by particle bombardment and we are in the process of regenerating transgenic plants for study.



The AFN1 cDNA encodes a 489 amino acid polypeptide.

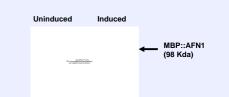
The AFN1 polypeptide contains UDP binding domain which is also present in (ABA) glucosyl transferases. This suggests that AFN1 may act to glucosylate ABA, thereby inactivating it.

# DORMANT NONDORMANT 0 3 6 12 24 36 48 0 3 6 12 24 36 48 hours .1.7kb AFN1 germination completed AFN1 mRNA is produced in germinating grains.

# **EcoRI** AFN1 pMALc2x/AFN1 7.3 kb The pMALc2x/AFN1 expression plasmid has been

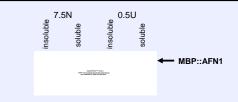
constructed.

#### Results



Production of AFN1 fusion protein in E. coli cells. BL21(DE3)pLvsS E. coli cells were transformed with pMALc2x/AFN1. Cell cultures were grown to an OD<sub>600</sub> of 0.5 and expression was induced by addition of IPTG. The culture was allowed to incubate for 4 hours at 37°. Crude protein was obtained from the E. coli cells by sonication and analyzed on

MBP::AFN1 fusion protein has been produced in E. coli cells containing pMALc2x/AFN1 plasmid.



Production of AFN1 fusion protein in E. coli cells. BL21(DE3)pLysS E. coli cells were transformed with pMALc2x/AFN1. Cell cultures were grown to an OD<sub>600</sub> of 0.5 and expression was induced by addition of IPTG. The culture was allowed to incubate for 4 hours at 37°. Crude protein was obtained from the E. coli cells by sonication in 7.5N and 0.5U buffer. Soluble and insoluble fractions were suspended in 1x SDS gel loading buffer, heated at 95-100°C for 10 minutes and analyzed on polyacrylamide gels.

A substantial amount of MBP::AFN1 fusion protein has been found in the 0.5U buffer soluble fraction.

### Acknowledgments

This work was supported in part by NSF Grant IOB-0443676, by NIH Grant P20-RR-016463 from the INBRE program of the National Center for Research Resources, and by a Colby College Natural Sciences Division Research Grant.