**LOCUS**
AT4G27240 (TAIR GRAMENE Accession)

**ID**
AT4G27240.1

**CHROMOSOME**
4

**LOCATION**
13639969 ~ 13641798

**STRAND**
FORWARD

**SHORT_DESCRIPTION**
zinc finger (C2H2 type) family protein

**DESCRIPTION**
"similar to zinc finger protein-related [Arabidopsis thaliana] (TAIR: AT5G54630.1); similar to Zinc finger, C2H2-type [Medicago truncatula] (GB: ABE93174.1)"

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<th>GO Term</th>
<th>GO Slim Term</th>
<th>GO ID</th>
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**GENE SEQ**
get gene genomic seq

**5'UTR**
5' UTR Sequence

**PROMOTER SEQ**
Paralogue gene identifier
AT5G54630

Orthologue gene identifier
LOC_Os01g57650

**Species**
Oryza sativa

**Identity**
76%
53%

**Cross Promoter Analysis**
Analyze

**Select what you want to assay**
- Transcription Factor Binding Site
- Tandem Repeat
- CpNpG

See additional file 1, figure S4
See additional file 1, figure S5
See additional file 1, figure S6