

A

Gene Feature eFP Corepression Treatment SRA Project gPCR Primer BLAST

eFP for BnaA01g270900

gFP Map gFP Chart **eFP Table**

Sample	Origin	Tissue	Stage	Condition	FPKM
Germminating_seed_Seeds.germination_Greenhouse_12_HAG	seed	germinating seed	germination	greenhouse	25.400
Cotyledons_Seeds.germination_Greenhouse_70_HAG	leaf	cotyledon	germination	greenhouse	28.101
Germminating_seed_Seeds.germination_Greenhouse_38_HAG	seed	germinating seed	germination	greenhouse	28.523
Hypocotyls_Seeds.germination_Greenhouse_34_HAG	leaf	hypocotyls	germination	greenhouse	28.525
Radicle_Seeds.germination_Greenhouse_34_HAG	root	radicle	germination	greenhouse	58.988
Cotyledons_Seeds.germination_Greenhouse_48_HAG	leaf	cotyledon	germination	greenhouse	55.888
Hypocotyls_Seeds.germination_Greenhouse_48_HAG	leaf	hypocotyls	germination	greenhouse	42.430
Radicle_Seeds.germination_Greenhouse_48_HAG	root	radicle	germination	greenhouse	21.900
Cotyledons_Seeds.germination_Greenhouse_72_HAG	leaf	cotyledon	germination	greenhouse	48.018
Hypocotyls_Seeds.germination_Greenhouse_72_HAG	leaf	hypocotyls	germination	greenhouse	41.071

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B

BnaA01g270900 Expressions Under Treatments

Biotic Abiotic Chemical

Table Chart

SRA Accession	Sample Accession	FPKM	TPM	Reads Count	Sample Name	Cultivar	Organ	Tissue	Stage	Condition	Treatment Type	Treatment Group	Treatment Reagent
PRJNA274853	SRR1793957	22.668	26.977	3	susceptibility	J602	Stem	stem	flowering	field	biotic	test	Sclerotinia sclerotiorum
PRJNA274853	SRR1793958	36.100	52.791	4	susceptibility	J602	Stem	stem	flowering	field	biotic	control	N
PRJNA274853	SRR1793959	28.113	50.849	15	susceptibility	J602	Stem	stem	flowering	field	biotic	control	N
PRJNA274853	SRR1793960	27.119	48.684	18	susceptibility	J602	Stem	stem	flowering	field	biotic	control	N
PRJNA274853	SRR1793963	26.599	54.013	12	resistance	J664	Stem	stem	flowering	field	biotic	test	Sclerotinia sclerotiorum
PRJNA230825	SRR1151470	18.913	2.014	6	vestar-Lmo-7dca	Westar	Leaf	cotyledon	seedling	NA	biotic	test	Erpophthora maculans

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C

Browse Genes of Brassica napus L.

Gene ID	Gene	Start	End	Strand	Length	Gene Note
BnaA21g000500	chr401	5435	2985	-	2791	protein SCA-like
BnaA21g114400	chr401	5621758	5621209	+	510	guanine oxidoreductase PG2-like (HbPm X2)
BnaA21g115960	chr401	5857856	5856795	-	3882	2nd finger NHR23 domain-containing protein 15-like
BnaA21g117400	chr401	5859447	5856592	-	2756	proteinaceous RNase P 2-like isoform X1
BnaA21g144900	chr1021_random	8486	3489	-	2622	protophylla-chloroplast-180
BnaA22g543401	chr402_random	7428	3409	+	3350	R1-like effector response 5-like
BnaA22g119400	chr1021_random	1607021	1602289	-	5233	systemic-rich repeat secretory protein 12-like
BnaA22g171700	chr1021_random	1617218	1615041	-	2178	systemic-rich repeat secretory protein 13-like
BnaA23g075000	chr1021_random	1954302	1979712	-	5619	RNA polymerase II-associated protein 2-like isoform X1
BnaA24g070100	chr1021_random	9971	4245	+	1929	L-type lectin domain-containing receptor kinase V 5-like

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