

2 fold change Differential expressed genes in Mycelium vs Sporangium

SEQ_ID	DESCRIPTION	Mycelium	Sporangium	Fold change
PITG_12551	Elicitin precursor (INF1)	8.59442	14.21405	49.16750717
PITG_12556	Elicitin like INF6	8.70743	13.94198	37.6492691
PITG_16999	Conserved hypothetical protein	8.43902	13.65272	37.10907364
PITG_06202	Conserved hypothetical protein	8.89462	13.44095	23.36590958
PITG_11060	Conserved hypothetical protein	9.39722	13.51552	17.36728096
PITG_03742	Conserved hypothetical protein	8.58947	12.6397	16.56691742
PITG_10765	Beta Glucan synthesis associated protein	9.5479	13.48678	15.33635044
PITG_02501	Conserved hypothetical protein	9.54883	13.48195	15.27517128
PITG_13660	Conserved hypothetical protein	8.98762	12.91893	15.25612545
PITG_00840		8.70857	12.61527	14.99801826
PITG_03637	Cellulose binding elicitor lectin	10.00708	13.9069	14.92663097
PITG_12562	Elicitin like protein	9.3965	13.27382	14.69564342
PITG_22317		9.38495	13.25422	14.61387348
PITG_00127		8.6794	12.51402	14.26706505
PITG_02139		8.99817	12.77543	13.71104527
PITG_14950		8.93755	12.67513	13.33904362
PITG_14721		9.30312	13.01768	13.12792206
PITG_08659		9.36217	13.01568	12.58398247
PITG_09370		8.60375	12.25507	12.56480789
PITG_20344		8.22698	11.8772	12.55523109
PITG_15731	ATP Binding cassette (ABC Superfamily)	8.94282	12.58678	12.50095749
PITG_00384		9.06653	12.69478	12.36551189
PITG_16731		10.02318	13.61598	12.0653677
PITG_23156		8.29853	11.88438	12.0073843
PITG_02954		8.59923	12.1364	11.6089592
PITG_16715		8.8556	12.3903	11.58912754
PITG_07482		8.7879	12.29713	11.38634872
PITG_00833		8.53502	12.03052	11.27847385
PITG_00761		8.60703	12.09303	11.20445061
PITG_03639		9.4509	12.90753	10.97868443
PITG_05616		9.47085	12.92202	10.93716335
PITG_06474		10.85657	14.30362	10.90599918
PITG_03721		9.37887	12.81243	10.80454731
PITG_12533		8.85242	12.26722	10.66491127
PITG_14046		7.84205	11.23522	10.50618267
PITG_11853		9.07208	12.46113	10.47624683
PITG_06860		10.66112	14.04915	10.46886635
PITG_21410		9.34402	12.72472	10.4157877
PITG_09065		8.56708	11.93053	10.29198933
PITG_06032		9.1032	12.46512	10.28105736
PITG_07347		9.22832	12.56877	10.12921143
PITG_10750		9.28905	12.61378	10.01946354
PITG_16275		8.43388	11.75842	10.01807404