

GO terms	Description
GO:0008356	asymmetric cell division
GO:0043433	negative regulation of DNA-binding transcription factor activity
GO:2000037	regulation of stomatal complex patterning
GO:0010182	sugar mediated signaling pathway
GO:0009553	embryo sac development
GO:0043565	sequence-specific DNA binding
GO:0019900	kinase binding
GO:0003700	DNA-binding transcription factor activity
GO:0090057	root radial pattern formation
GO:0009742	brassinosteroid mediated signaling pathway
GO:0009733	response to auxin
GO:0006351	DNA-templated transcription

GO terms	Description
GO:0019760	glucosinolate metabolic process
GO:0009759	indole glucosinolate biosynthetic process
GO:0009862	systemic acquired resistance, salicylic acid mediated signaling pathway
GO:0009737	response to abscisic acid
GO:0046982	protein heterodimerization activity
GO:0000976	transcription cis-regulatory region binding
GO:1901141	regulation of lignin biosynthetic process
GO:0009970	cellular response to sulfate starvation
GO:0009094	L-phenylalanine biosynthetic process
GO:0005730	nucleolus