





Species: Malus domestica

Common name	Malus domestica
Cultivar/subspecies	Not available
PLAZA identifier	mdo
Source	GDDH13 v1.1
NCBI taxonomy	3750
Mitochondrion	Not available
Chloroplast	Not available
Publication	28581499
Color code	



Description

Malus domestica, commonly known as apple, is a fruit of the Rosaceae family. It is very widely cultivated fruit species, with thousands of different cultivars. The genome consists of 17 chromosomes, and it has the one of the highest number of genes yet reported.

Gene Overview

Species	Gene type(s) in	Gene type(s) information				Functional annotation information	
	#Coding genes	#RNA genes	#Pseudo genes	#Transposons	#Genes with GO	#Genes with InterPro	
Malus domestica	45,116	7,625	0	0	34,373	37,067	

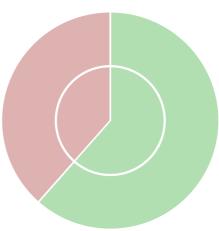
Gene Family Information

GF type	Data type	Multi-species	Species specific, multicopy	Species specific, singlecopy (orphan)
Homologous gene family	Gene Family Counts	7,811	247	2,999
	Gene Counts	41,285	832	2,999
Orthologous gene family	Gene Family Counts	8,373	21	0
	Gene Counts	33,254	56	0

GO Evidence Stats

Evidence code	Description	#Genes	#Entries
IEA	Inferred from Electronic Annotation	27,559	65,723
ISO	Inferred from Sequence Orthology	22,375	105,054





Gene List

Apply filter(s) / Configure data types

Search and download are disabled due to table size. Apply filter to reduce table size.

Gene ID	•	Species	Gene Type	Homologous gf	Orthologous gf
MD00G1000100		Malus domestica	coding	HOM05D000285	ORTHO05D000241
MD00G1000200		Malus domestica	coding	HOM05D006875	ORTH005D008416
MD00G1000300		Malus domestica	coding	HOM05D007271	ORTH005D009217
MD00G1000400		Malus domestica	coding	HOM05D000130	ORTH005D000257
MD00G1000500		Malus domestica	coding	HOM05D000184	ORTH005D000032
MD00G1000600		Malus domestica	coding	HOM05D001237	ORTHO05D009110
MD00G1000700		Malus domestica	coding	HOM05D001193	ORTHO05D001016
MD00G1000800		Malus domestica	coding	HOM05D002287	ORTHO05D002588
MD00G1000900		Malus domestica	coding	HOM05D138137	ORTH005D248012
MD00G1001000		Malus domestica	coding	HOM05D000375	ORTH005D000240







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