

## GEPHE SUMMARY

**Gephebase Gene**  
CF-2.1 and Cf-2.2

**Entry Status**  
Published

**GepheID**  
GP00000180

**Main curator**  
Courtier

## PHENOTYPIC CHANGE

**Trait Category**  
Physiology

**Trait**  
Pathogen resistance (leaf mold fungus ; root parasitic nematode)

**Trait State in Taxon A**  
Lycopersicon pimpinellifolium - resistant ; Lycopersicon esculentum MoneyMaker - resistance re-acquired from L. pimpinellifolium

**Trait State in Taxon B**  
Lycopersicon esculentum sensitive strains

**Ancestral State**  
Taxon A

**Taxonomic Status**  
Domesticated

	Taxon A	Taxon B
<b>Latin Name</b>	<i>Solanum pimpinellifolium</i>	<i>Solanum lycopersicum</i>
<b>Common Name</b>	-	tomato
<b>Synonyms</b>	Lycopersicon pimpinellifolium; Solanum pimpinellifolium var. racemigerum; currant tomato; Lycopersicon pimpinellifolium (L.) Mill.; Solanum pimpinellifolium L.	Lycopersicon esculentum var. esculentum; Solanum esculentum; Solanum lycopersicum var. humboldtii; tomato; Lycopersicon esculentum Mill.; Solanum esculentum Dunal; Solanum lycopersicum L.; Lycopersicon lycopersicum; Lycopersicum esculentum; Solanum lycopersicon
<b>Rank</b>	species	species
<b>Lineage</b>	cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; asterids; lamiids; Solanales; Solanaceae; Solanoideae; Solaneae; Solanum; Lycopersicon	cellular organisms; Eukaryota; Viridiplantae; Streptophyta; Streptophytina; Embryophyta; Tracheophyta; Euphyllophyta; Spermatophyta; Magnoliophyta; Mesangiospermae; eudicotyledons; Gunneridae; Pentapetalae; asterids; lamiids; Solanales; Solanaceae; Solanoideae; Solaneae; Solanum; Lycopersicon
<b>Parent</b>	Lycopersicon () - (Rank: subgenus)	Lycopersicon () - (Rank: subgenus)
<b>NCBI Taxonomy ID</b>	4084	4081
<b>is Taxon A an Intraspecies?</b>	No	No

## GENOTYPIC CHANGE

**Generic Gene Name**  
-

**Synonyms**  
-

**String**  
-

**Sequence Similarities**  
-

**GO - Molecular Function**  
-

**GO - Biological Process**  
-

**GO - Cellular Component**  
GO:0016021 : integral component of membrane

**Presumptive Null**

**UniProtKB Solanum pimpinellifolium**  
Q41398

**GeneBankID or UniProtKB**  
AAC15779

Yes

#### Molecular Type

Gene Loss

#### Aberration Type

Deletion

#### Deletion Size

-

#### Molecular Details of the Mutation

loss of the two genes Cf-2.1 and Cf-2.2 (see Dixon et al. 1998) in cultivated tomato - resistance re-acquired from related species

#### Experimental Evidence

Linkage Mapping

#### Main Reference

The tomato Cf-2 disease resistance locus comprises two functional genes encoding leucine-rich repeat proteins. (1996)

#### Authors

Dixon MS; Jones DA; Keddie JS; Thomas CM; Harrison K; Jones JD

#### Abstract

In plants, resistance to pathogens is frequently determined by dominant resistance genes, whose products are proposed to recognize pathogen-encoded avirulence gene (Avr) products. The tomato resistance locus Cf-2 was isolated by positional cloning and found to contain two almost identical genes, each conferring resistance to isolates of tomato leaf mould (*C. fulvum*) expressing the corresponding Avr2 gene. The two Cf-2 genes encode protein products that differ from each other by only three amino acids and contain 38 leucine-rich repeat (LRR) motifs. Of the LRRs, 20 show extremely conserved alternating repeats. The C-terminus of Cf-2 carries regions of pronounced homology to the protein encoded by the unlinked Cf-9 gene. We suggest that this conserved region interacts with other proteins involved in activating plant defense mechanisms.

#### Additional References

The tomato Cf-5 disease resistance gene and six homologs show pronounced allelic variation in leucine-rich repeat copy number. (1998)

Dual disease resistance mediated by the immune receptor Cf-2 in tomato requires a common virulence target of a fungus and a nematode. (2012)

## RELATED GEPHE

#### Related Genes

1 (Cf-4/9)

#### Related Haplotypes

No matches found.

## COMMENTS

Diverse family of paralogous genes ; @Introgression