

**Information on EC 6.2.1.12 - 4-coumarate-CoA ligase**

PROTEIN VARIANTS	ORGANISM	UNIPROT	COMMENTARY	LITERATURE
C403A	<a href="#">Arabidopsis thaliana</a>	-	activity reduced to 45% of wild-type level, no effect on Km values for ATP and caffeate	<a href="#">651262</a>
E401Q	<a href="#">Arabidopsis thaliana</a>	-	activity reduced to 21% of wild-type level, no effect on Km values for ATP and caffeate	<a href="#">651262</a>
K211S	<a href="#">Arabidopsis thaliana</a>	-	activity reduced to 2.7% of wild-type level	<a href="#">651262</a>
K320A	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	new substrate specificity, accepts ferulate as substrate	<a href="#">652146</a>
K320L	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	new substrate specificity, accepts ferulate as substrate	<a href="#">652146</a>
K445T	<a href="#">Arabidopsis thaliana</a>	-	activity reduced to 0.2% of wild-type level	<a href="#">651262</a>
K457S	<a href="#">Arabidopsis thaliana</a>	-	activity reduced to 4% of wild-type level	<a href="#">651262</a>
K540N	<a href="#">Arabidopsis thaliana</a>	-	no activity	<a href="#">651262</a>
L356A	<a href="#">Arabidopsis thaliana</a>	-	only little effect on activity	<a href="#">653597</a>
M293A	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	new substrate specificity, accepts ferulate as substrate	<a href="#">652146</a>
M293A/V294L	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	substitution of valine decreases Km for ferulate 2-3fold compared to single methionine substitution mutant	<a href="#">652146</a>
M293F	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	no reaction with ferulate, increases Km for caffeate	<a href="#">652146</a>
M293P	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	new substrate specificity, accepts ferulate as substrate	<a href="#">652146</a>
M293P/K320L	<div> <div></div> <div>2 entries</div> </div>			
M293P/V294M	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	substitution of valine decreases Km for ferulate 2-3fold compared to single methionine substitution mutant	<a href="#">652146</a>
M293W	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	no reaction with ferulate	<a href="#">652146</a>
M293Y	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	no reaction with ferulate, increases Km for caffeate	<a href="#">652146</a>

N256A/M293P/K320L	<a href="#">Arabidopsis thaliana</a>	-	effectively uses ferulate and cinnamate as substrates, 10fold reduced activity with caffeate	<a href="#">653597</a>
R449Q	<a href="#">Arabidopsis thaliana</a>	-	activity reduced to 3% of wild-type level, Km values significantly increased	<a href="#">651262</a>
V294L	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	slightly reduced activity with caffeate, no activity with ferulate	<a href="#">652146</a>
V294M	<a href="#">Arabidopsis thaliana</a>	<a href="#">Q9S725</a>	slightly reduced activity with caffeate, no activity with ferulate	<a href="#">652146</a>
V355A	<a href="#">Arabidopsis thaliana</a>	-	strongly reduced activity	<a href="#">653597</a>
V355A/L356A	<a href="#">Arabidopsis thaliana</a>	-	strongly reduced activity	<a href="#">653597</a>
E340A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
G333A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
G337A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
G342A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
I346A	<a href="#">Glycine max</a>	-	4CL1-mutant, very low activity with all substrates	<a href="#">652422</a>
L344A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
M338A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
M348A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
P285A	<a href="#">Glycine max</a>	-	4CL1-mutant, very low activity with all substrates	<a href="#">652422</a>
P286A	<a href="#">Glycine max</a>	-	4CL1-mutant, minor changes in activity	<a href="#">652422</a>
P343A	<a href="#">Glycine max</a>	-	4CL1-mutant, very low activity with all substrates	<a href="#">652422</a>
Q334A	<a href="#">Glycine max</a>	-	4CL1-mutant, very low activity with all substrates	<a href="#">652422</a>
T331A	<a href="#">Glycine max</a>	-	4CL1-mutant, minor changes in activity	<a href="#">652422</a>
T339A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
V284A	<a href="#">Glycine max</a>	-	4CL1-mutant, minor changes in activity	<a href="#">652422</a>
Y336A	<a href="#">Glycine max</a>	-	4CL1-mutant, activity almost completely abolished	<a href="#">652422</a>
H237A	<a href="#">Nicotiana tabacum</a>	<a href="#">O24146</a>	the mutant shows reduced activity compared to the wild type enzyme	<a href="#">746513</a>

K197A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
K441A	Nicotiana tabacum	O24146	inactive	746513
K443A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
K526A	Nicotiana tabacum	O24146	inactive	746513
M344A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
R435A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
T193A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
T336A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
Y239A	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
Y239F	Nicotiana tabacum	O24146	the mutant shows reduced activity compared to the wild type enzyme	746513
A251S	Plagiochasma appendiculatum	A0A0D4C3L0	the mutant enzyme shows a lower binding affinity and catalytic efficiency than the wild type enzyme	746000
A251S/M247Y	Plagiochasma appendiculatum	A0A0D4C3L0	the mutant enzyme shows a lower binding affinity and catalytic efficiency than the wild type enzyme	746000
M247Y	Plagiochasma appendiculatum	A0A0D4C3L0	the catalytic efficiency of, 4-coumaric, caffeic and dihydro-4-coumaric acids are higher in the mutant than in the wild type enzyme	746000
V388del	Populus tomentosa	H9AZ21, H9AZ22, H9AZ23, H9AZ24, Q941M3	deletion of V338 residues in isoform 4CL1 results in activity towards sinapate	746098
Y236F	Populus tomentosa	H9AZ21, H9AZ22, H9AZ23, H9AZ24, Q941M3	the mutant of isoform 4CL2 exhibits high substrate catalytic efficiency for 4-coumarate but very low substrate affinities and specificities for caffeate and ferulate	746098
F239A	Solanum lycopersicum	-	site-directed mutagenesis, the mutant strain produces more naringenin chalcone compared to the wild-type	728481
F239N	Solanum lycopersicum	-	site-directed mutagenesis, the mutant strain produces more	728481

			naringenin chalcone compared to the wild-type	
F239R	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces more naringenin chalcone compared to the wild-type	<a href="#">728481</a>
F239S	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant shows increased activity in vivo, and in vitro with coumaric acid, but reduced activity with ferulic acid compared to the wild-type enzyme. The mutant strain produces more naringenin chalcone compared to the wild-type	<a href="#">728481</a>
F269G	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces less naringenin chalcone compared to the wild-type	<a href="#">728481</a>
F269I	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces less naringenin chalcone compared to the wild-type	<a href="#">728481</a>
F269L	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces more naringenin chalcone compared to the wild-type, loss of feedback inhibition by naringenin	<a href="#">728481</a>
F269L/K415T	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant shows increased activity in vivo, and in vitro with coumaric acid, but reduced activity with ferulic acid compared to the wild-type enzyme	<a href="#">728481</a>
F269V	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces less naringenin chalcone compared to the wild-type	<a href="#">728481</a>
Q274D	<a href="#">+ 2 entries</a>			
Q274E	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces less naringenin chalcone compared to the wild-type	<a href="#">728481</a>
Q274G	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces more naringenin chalcone compared to the wild-type	<a href="#">728481</a>
Q274H	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant shows increased activity in vivo, and in vitro with coumaric acid, but reduced activity with ferulic acid compared to the wild-type enzyme, loss of feedback	<a href="#">728481</a>

			inhibition by naringenin, the mutant strain produces more naringenin chalcone compared to the wild-type	
Q274S	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces less naringenin chalcone compared to the wild-type	<a href="#">728481</a>
V186A	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces more naringenin chalcone compared to the wild-type	<a href="#">728481</a>
V186G	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant shows highly increased activity in vivo, and in vitro with coumaric acid, but reduced activity with ferulic acid compared to the wild-type enzyme. The mutant strain produces highly increased naringenin chalcone content compared to the wild-type	<a href="#">728481</a>
V186I	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces highly increased naringenin chalcone content compared to the wild-type	<a href="#">728481</a>
V186I/V187L	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant shows increased activity in vivo, and in vitro with coumaric acid compared to the wild-type enzyme, but no activity with ferulic acid	<a href="#">728481</a>
V186L	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, the mutant strain produces more naringenin chalcone compared to the wild-type	<a href="#">728481</a>
V235M/A325G	<a href="#">Solanum lycopersicum</a>	-	site-directed mutagenesis, highly unstable inactive mutant	<a href="#">728481</a>
<a href="#">additional information</a>	<a href="#">+ 11 entries</a>			