

Bhattacharya A, Bhowmick P, Ganguli S, Mitra A K. Evolutionary Insights into the Enzymes involved in the Biosynthesis of the Volatile Organic Compounds Isoprene and Pinene in Plants. Plant Science Today. 2023; 10(2): 253–262. <https://doi.org/10.14719/pst.2115>

SUPPLEMENTARY FILES

Evolutionary Insights into the Enzymes involved in the Biosynthesis of the Volatile Organic Compounds Isoprene and Pinene in Plants

SUPPLEMENTARY TABLE 1

List of Plants Emitting VOCs

1. *Acacia leucophloea/Vachellia leucophloea*
2. *Adhatoda vasica*
3. *Albizia lebbeck*
4. *Atrocarpus heterophyllus*
5. *Azadirachta indica*
6. *Bombax ceiba*
7. *Butea monosperma*
8. *Casuarina equisetifolia*
9. *Citrus limon*
10. *Cocos nucifera*
11. *Dalbergia sissoo*
12. *Diospyros melanoxylon*
13. *Eucalyptus globulus*
14. *Ficus benghalensis*
15. *Ficus religiosa*
16. *Hibiscus rosasinensis*
17. *Lantana camara*
18. *Madhuca longifolia*
19. *Mangifera indica*
20. *Morus alba*
21. *Peltophorum pterocarpum*
22. *Psidium guajava*
23. *Pterocarpus marsupium*
24. *Pterospermum acerifolium*
25. *Schleichera oleosa*
26. *Syzygium jambolanum/Syzygium cumini*
27. *Tecoma stans*
28. *Tectona grandis*

Name of Plant	Common Name	Family	Type of Plant	VOC produced	Area of growth in West Bengal	Reference
<i>Acacia leucophloea/ Vachellia leucophloea</i>	White Bark Acacia	Mimosaceae	Deciduous	Moderate emitter – Isoprene, pinene, other VOCs	Murshidabad, Birbhum, Bankura, Purba Bardhaman, Paschim Bardhaman, Purba Medinipur, Paschim Medinipur	(1)
<i>Adhatoda vasica</i>	Malabar Nut/ Adulsa/ Vasaka	Acanthaceae	Evergreen	High emitter - Isoprene, α -pinene, other VOCs	Kakrajhore forest in West Medinipur, West Rarrh region, Lower Bengal	(2, 3)
<i>Albizia lebbeck</i>	Shirish	Mimosaceae	Deciduous	Moderate emitter - Isoprene, α -pinene, other VOCs	Plains of West Bengal	(4)
<i>Atrocarpus heterophyllus</i>	Jackfruit	Moraceae	Evergreen tree	High emitter - Isoprene, monoterpenes	Throughout West Bengal	(5, 6)
<i>Azadirachta indica</i>	Neem	Meliaceae	Deciduous	Moderate emitter - Isoprene, α -pinene, other VOCs	Widespread	(7)
<i>Bombax ceiba</i>	Malabar silk cotton tree/ Shimul	Malvaceae	Deciduous	Moderate emitter - Isoprene, monoterpenes	Throughout plains to 800m	(8)
<i>Butea monosperma</i>	Flame of the forest tree/ Palash	Fabaceae	Deciduous	High emitter - Isoprene	Purulia, West Medinipore	(9 – 11)
<i>Casuarina equisetifolia</i>	Australian pine	Casuarinaceae	Evergreen	High emitter - Isoprene, α -pinene, other VOCs	Throughout West Bengal	(12)
<i>Citrus limon</i>	Lemon	Rutaceae	Deciduous	Moderate emitter - Isoprene, α -pinene, other VOCs	Red lateritic zone of West Bengal	(13)
<i>Cocos nucifera</i>	Coconut	Arecaceae	Evergreen	High emitter - Isoprene	Purba Bardhaman, Howrah, 24 Parganas, Midnapore	(14)

<i>Dalbergia sissoo</i>	Shisham	Papilionaceae	Deciduous	High emitter - Isoprene, α -pinene, other VOCs	South West Bengal	(4)
<i>Diospyros melanoxylon</i>	East Indian ebony/ Kendu	Ebenaceae	Deciduous or evergreen, , depending on habitat	Moderate emitter - Isoprene, monoterpenes	South West Bengal	(4)
<i>Eucalyptus globulus</i>	Gum tree/Stringybark	Myrtaceae	Mostly Evergreen	High emitter - Isoprene, α -pinene, other VOCs	Laterite soils of South and South Western Districts	(4)
<i>Ficus benghalensis</i>	Banyan	Moraceae	Evergreen	Moderate emitter – Isoprene, α -pinene, other VOCs	Throughout West Bengal	(15)
<i>Ficus religiosa</i>	Sacredfig	Moraceae	Deciduous tree	High emitter - Isoprene, α -pinene, other VOCs	From the coast up to 1200m, Bankura	(16)
<i>Hibiscus rosasinensis</i>	Hibiscus	Malvaceae	Evergreen	High emitter - Isoprene, α -pinene, other VOCs	Hooghly, Purba and Paschim Bardhaman, Birbhum	(17)
<i>Lantana camara</i>	Lantana	Verbenaceae	Evergreen shrub	High emitter - Isoprene, α -pinene, other VOCs	Howrah, purba and Paschim Bardhaman, Birbhum	(18)
<i>Madhuca longifolia</i>	Mahua	Sapotaceae	Evergreen to semi-evergreen	High emitter - Isoprene	Purulia, Bankura, West Midnapore and Jhargram.	(19)
<i>Mangifera indica</i>	Mango	Anacardiaceae	Evergreen	High emitter - Isoprene, α -pinene, other VOCs	Murshidabad and Malda districts	(20)
<i>Morus alba</i>	White mulberry	Moraceae	Deciduous	High emitter - Isoprene, α -pinene, other VOCs	Birbhum	(21)
<i>Peltophorum pterocarpum</i>	Yellow flame tree/ Radhachura	Fabaceae	Deciduous	High emitter - Isoprene	Birbhum	(22)
<i>Psidium guajava</i>	Guava	Myrtaceae	Evergreen	High emitter - Isoprene, α -pinene, other VOCs	North and South 24 Parganas, Nadia	(23)
<i>Pterocarpus marsupium</i>	Malabar kino/ Venkai	Fabaceae	Deciduous	High emitter - Isoprene, monoterpenes	Wild and also planted in South Bengal	(24)

<i>Pterospermum macerifolium</i>	Kanak Champa	Sterculiaceae	Deciduous	Moderate emitter - Isoprene, α-pinene, other VOCs	Eastern districts of Bengal	(25)
<i>Schleichera oleosa</i>	Kusum	Sapindaceae	Deciduous	Moderate emitter – Isoprene, α-pinene, other VOCs	Found throughout the hotter parts of south West Bengal, in red lateritic soil.	(4)
<i>Syzygium jambolanum</i> / <i>Syzygium cumini</i>	Malabar plum	Myrtaceae	Evergreen	High emitter - Isoprene, α-pinene, other VOCs	Throughout West Bengal	(26)
<i>Tecoma stans</i>	Yellow bells	Bignoniaceae	Semi-evergreen shrub	Moderate emitter – Isoprene, α-pinene, other VOCs	Purba and Paschim Bardhaman	(27)
<i>Tectona grandis</i>	Teak	Lamiaceae	Deciduous	High emitter - Isoprene	Lower Hill forests in North Bengal, Jhargram	(28)

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SUPPLEMENTARY TABLE 2
Data for enrichment profile – MEP Pathway

Name of Enzyme	Number of families common to gene-specific and protein-specific trees	Common families
Isoprene Synthase	5	Salicaceae Euphorbiaceae Fabaceae Lamiaceae Poaceae
Isopentenyl Diphosphate Isomerase	3	Brassicaceae Lamiaceae Nyssaceae
Hydroxymethylbutenyl diphosphate reductase	0	N/A
Hydroxymethylbutenyl diphosphate synthase	0	N/A
Methylerythritol 2,4-cyclodiphosphate synthase	0	N/A
CDP-ME kinase	2	Lamiaceae Solanaceae
Diphosphocytidylyl methyl erythritol synthase	2	Brassicaceae Poaceae
1-deoxyxylulose-5-phosphate reductoisomerase	0	N/A
1-deoxyxylulose 5-phosphate synthase	1	Solanaceae

Data for enrichment profile – MVA Pathway

Name of Enzyme	Number of families common to gene-specific and protein-specific trees	Common families
Pinene Synthase	2	Lamiaceae Pinaceae
Geranyl diphosphate synthase	13	Apocynaceae Asteraceae Brassicaceae Cannabaceae Fabaceae Lamiaceae Lauraceae Magnoliaceae Pinaceae Plantaginaceae Rosaceae Solanaceae Vitaceae
Mevalonate pyrophosphate decarboxylase/ Diphosphomevalonate decarboxylase	1	Brassicaceae
Phosphomevalonate kinase	1	Brassicaceae
Mevalonate 5 kinase	1	Brassicaceae
3-hydroxy-3-methylglutaryl-CoA reductase/ HMG CoA reductase	4	Solanaceae Poaceae Brassicaceae Malvaceae
3-hydroxy-3-methylglutaryl-CoA synthase/ HMG CoA synthase	0	N/A
Acetoacetyl CoA thiolase	0	N/A

SUPPLEMENTARY 3

- PHYLOGENETIC TREES

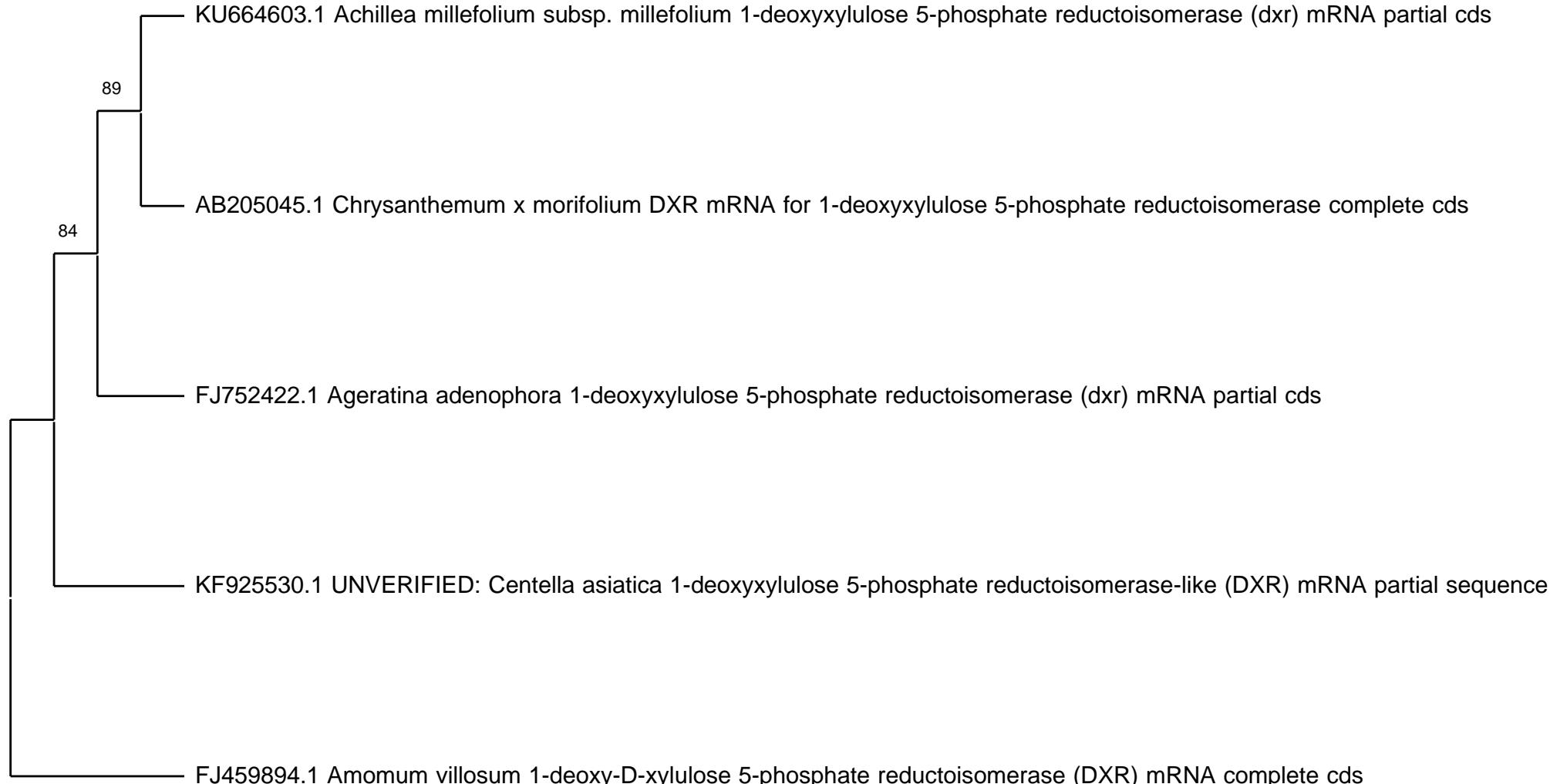
METHYLERYTHRITOL PHOSPHATE (MEP) PATHWAY

GENBANK PHYLOGENETIC TREES

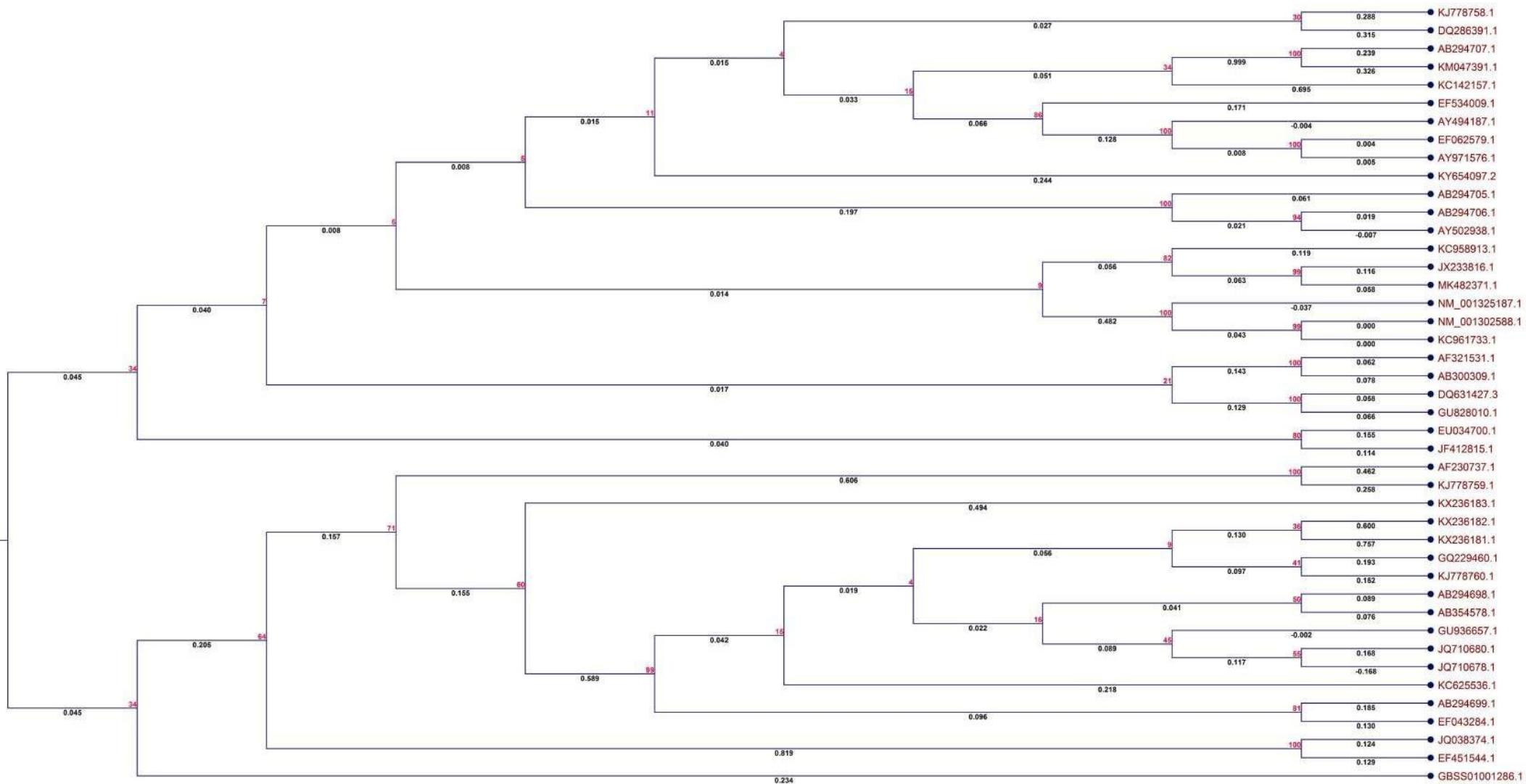
Deoxyxylulose 5-phosphate (DXP) Synthase



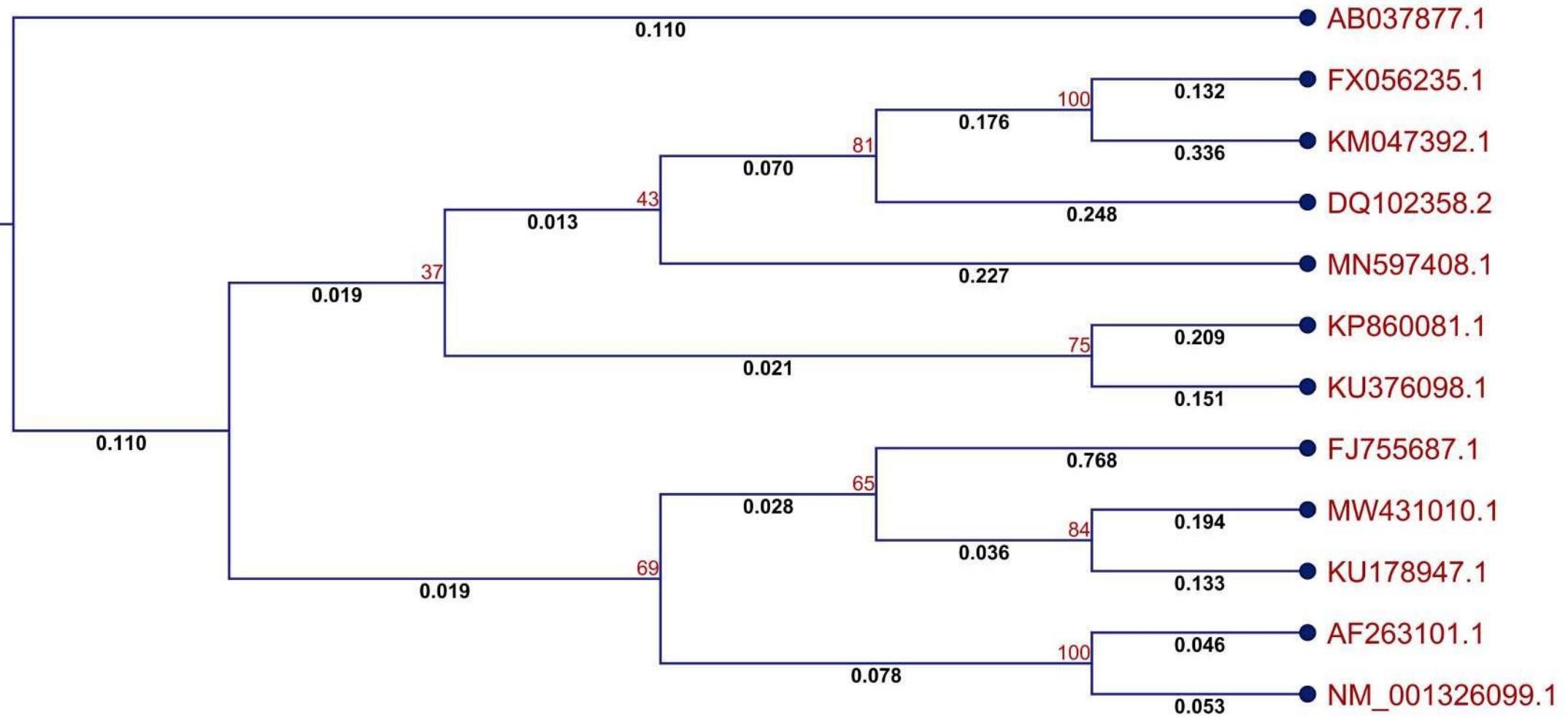
Deoxyxylulose 5-phosphate (DXP) Reductoisomerase



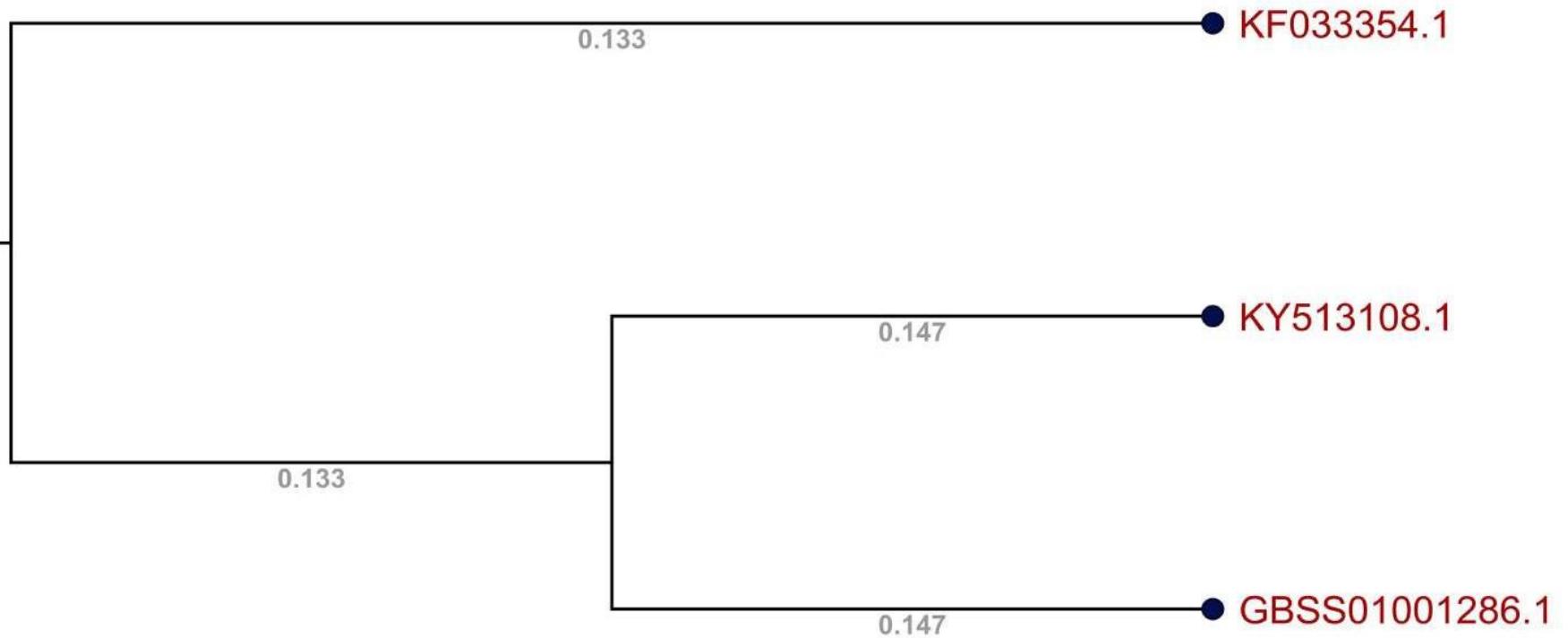
Diphosphocytidyl methyl erythritol (CDP-ME) Synthase



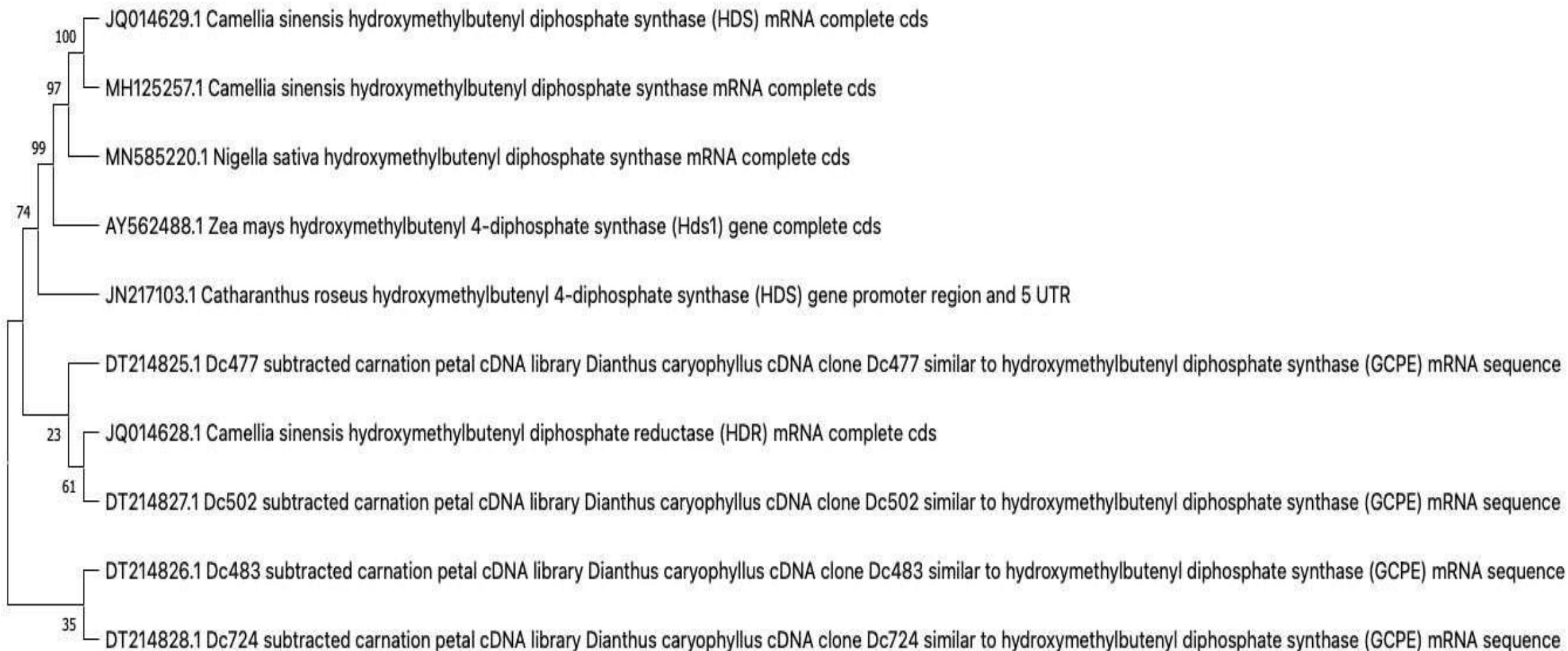
Diphosphocytidyl methyl erythritol (CDP-ME) Kinase



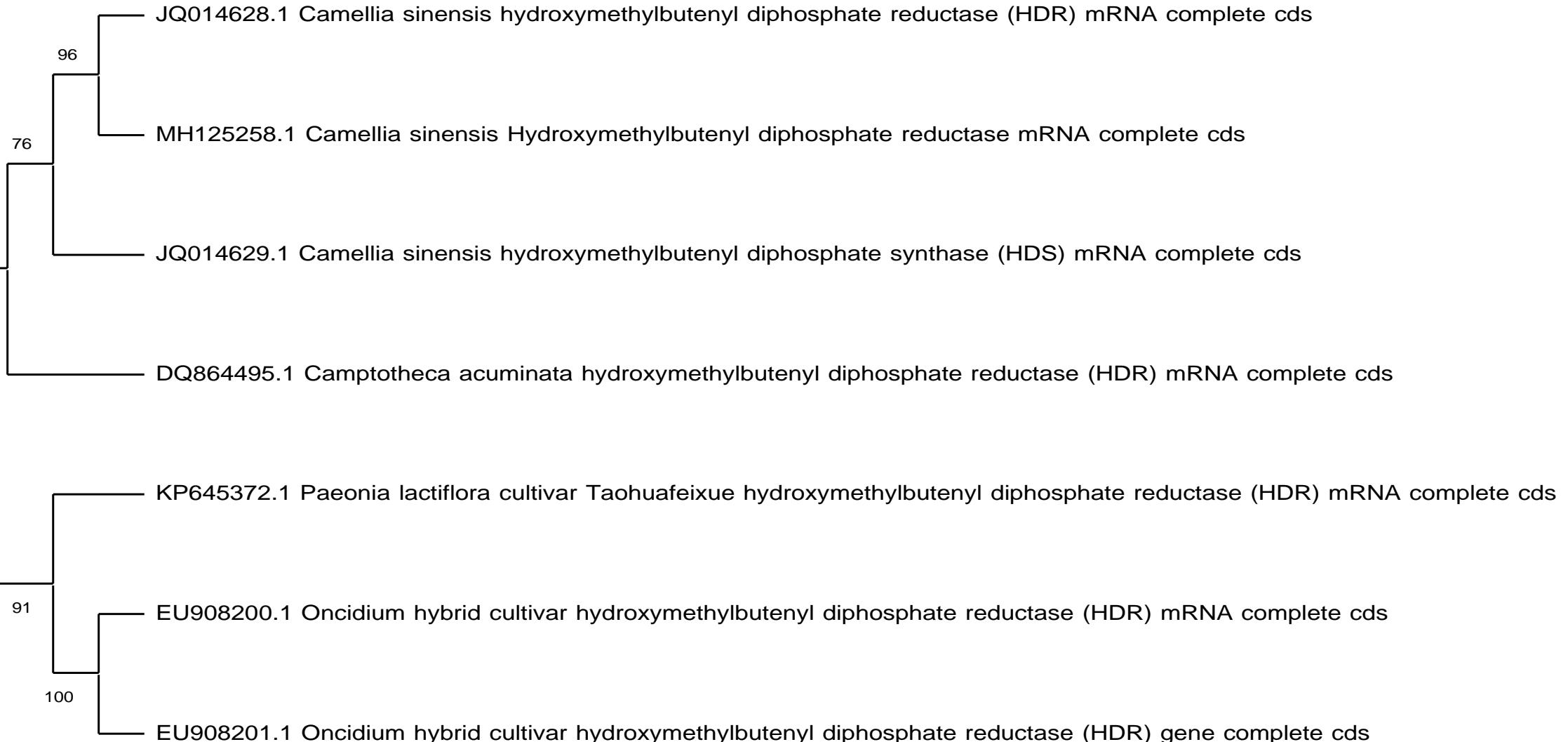
Methylerythritol 2,4-cyclodiphosphate (ME-cPP) Synthase



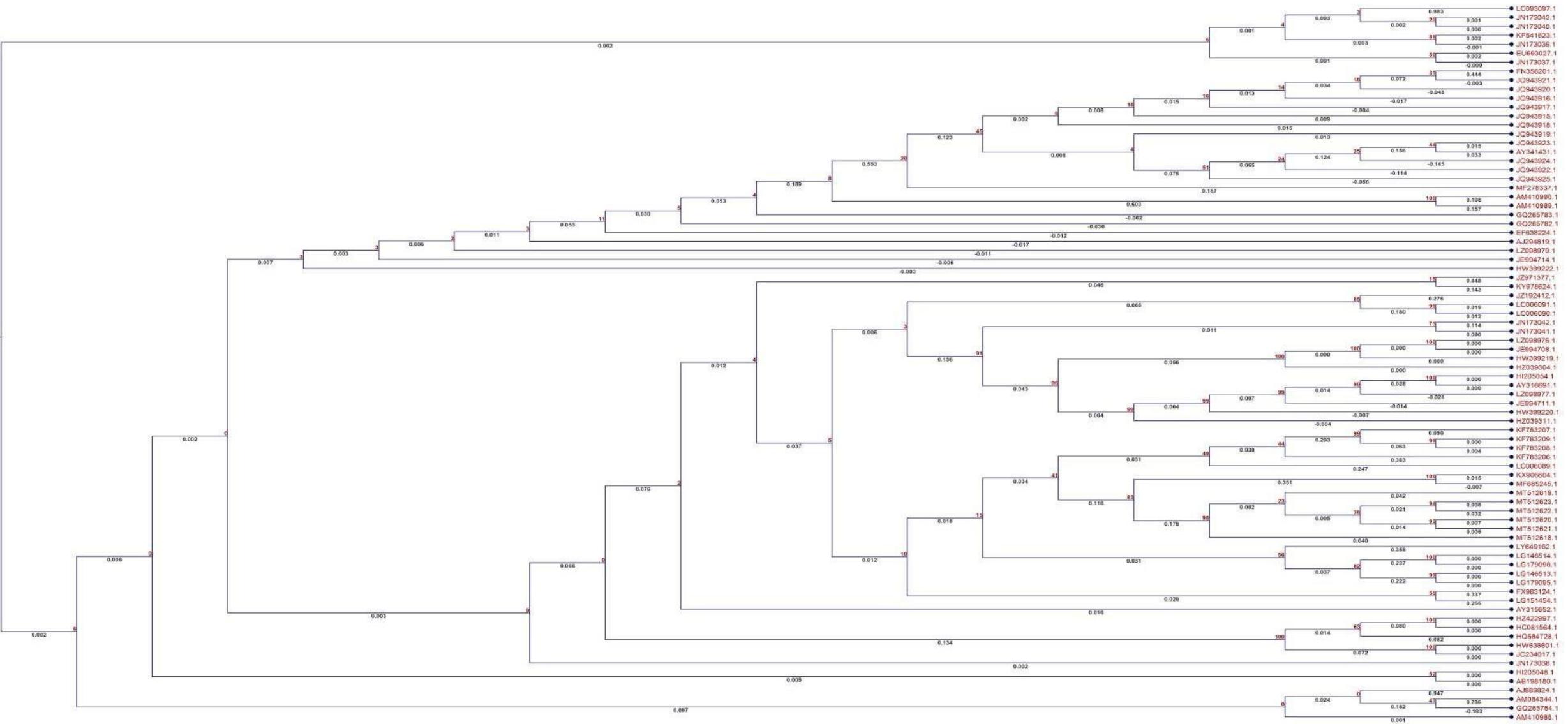
Hydroxymethylbutenyl diphosphate (HMBPP) Synthase



Hydroxymethylbutenyl diphosphate (HMBPP) Reductase

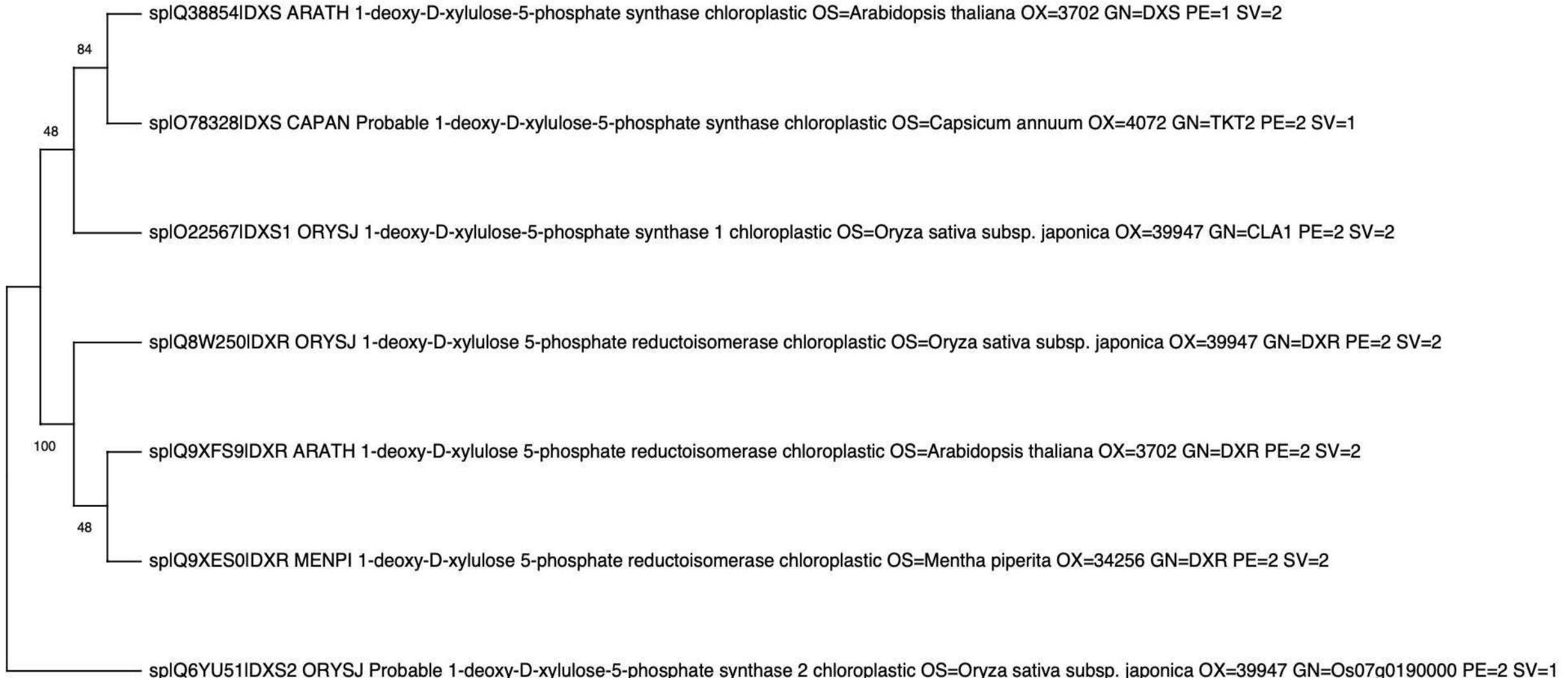


Isoprene Synthase

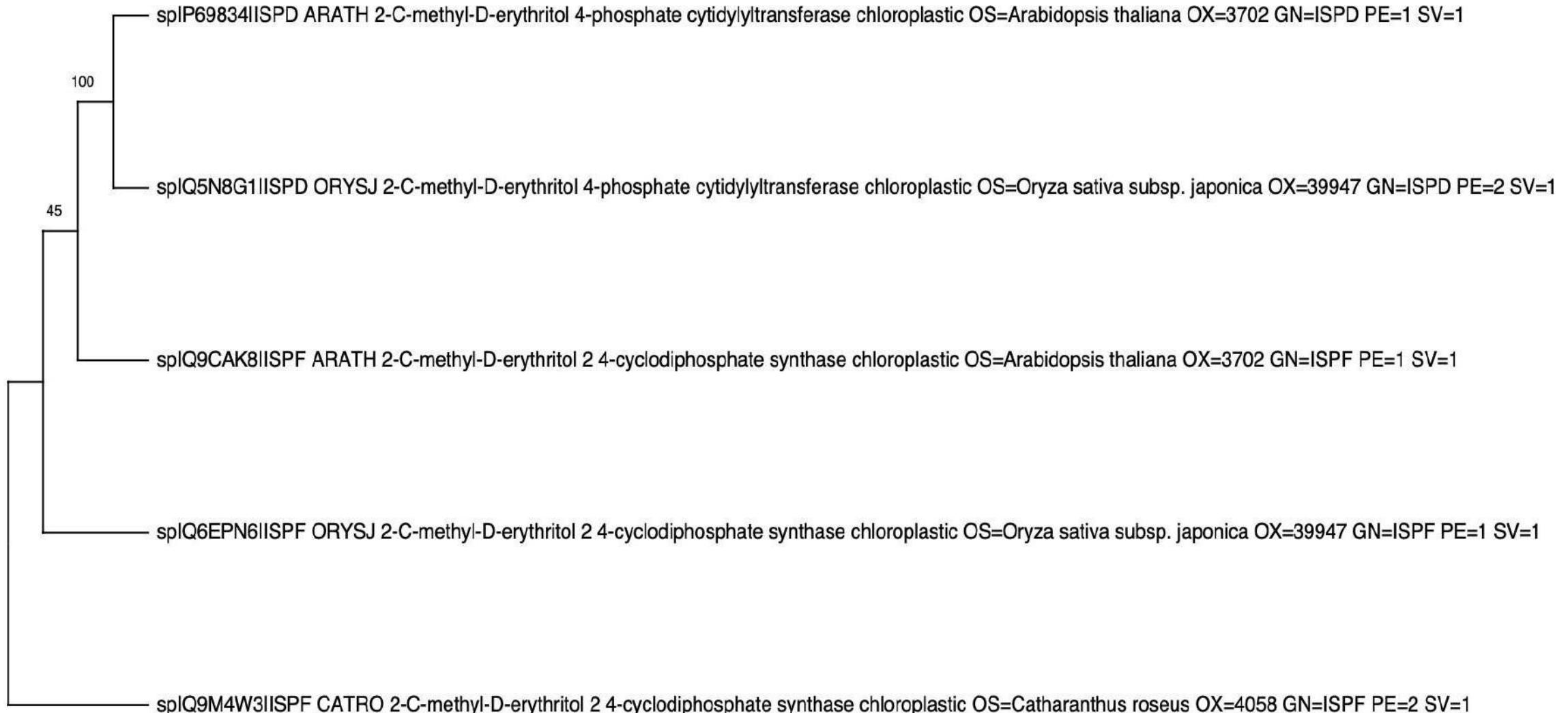


UNIPROT PHYLOGENETIC TREES

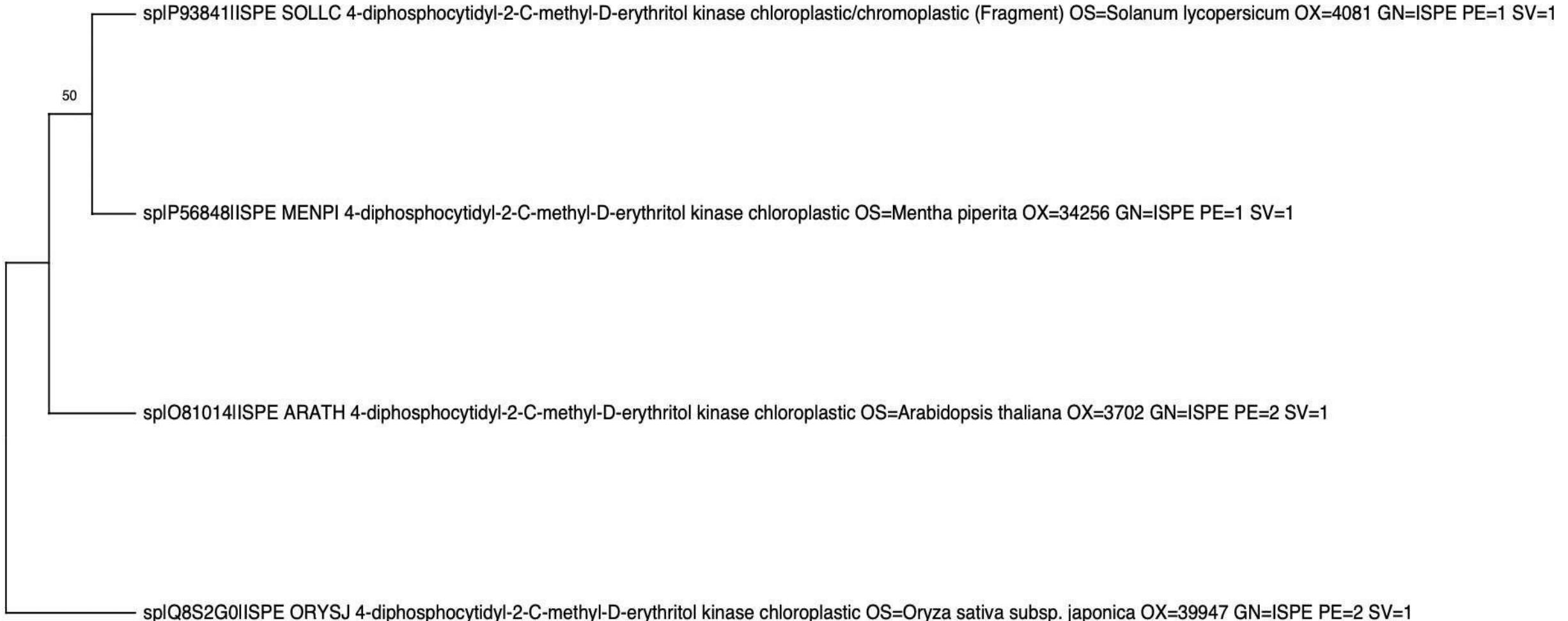
Deoxyxylulose 5-phosphate (DXP) Synthase



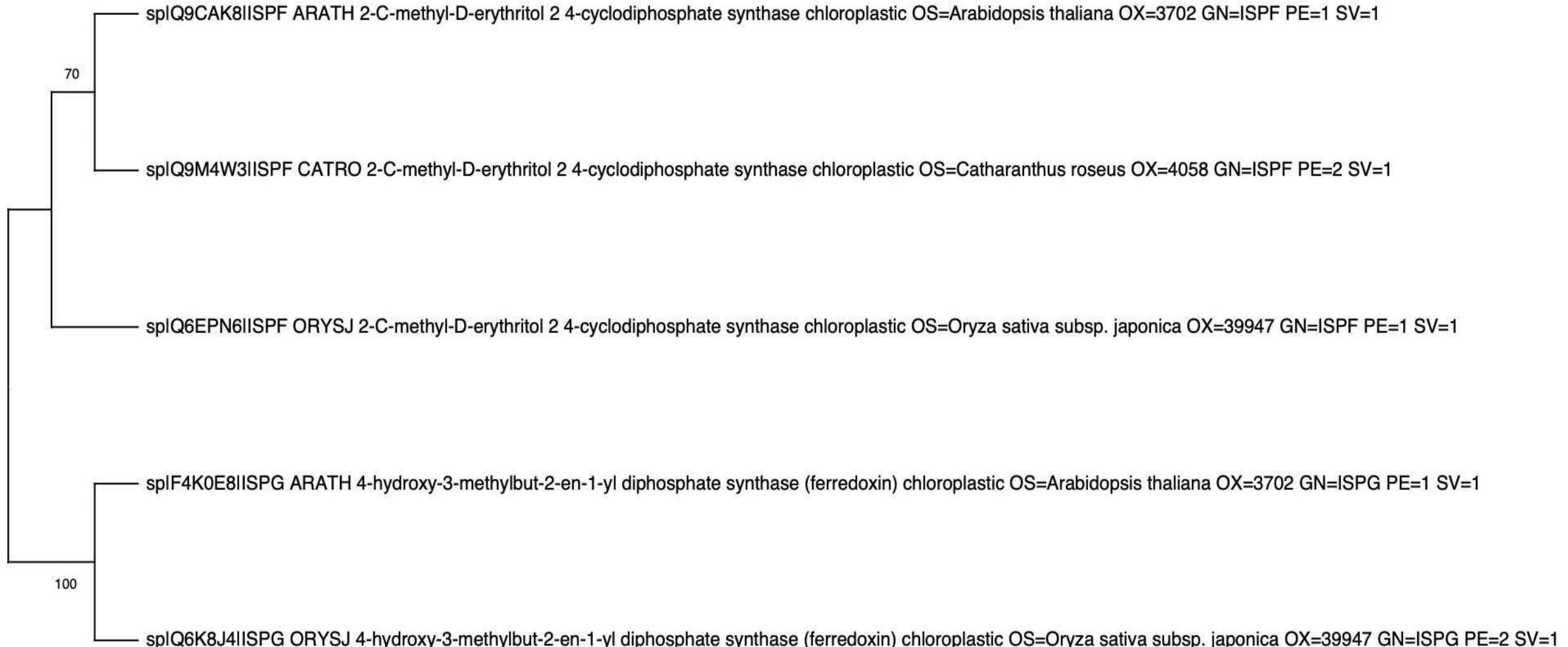
Diphosphocytidylyl methyl erythritol (CDP-ME) Synthase



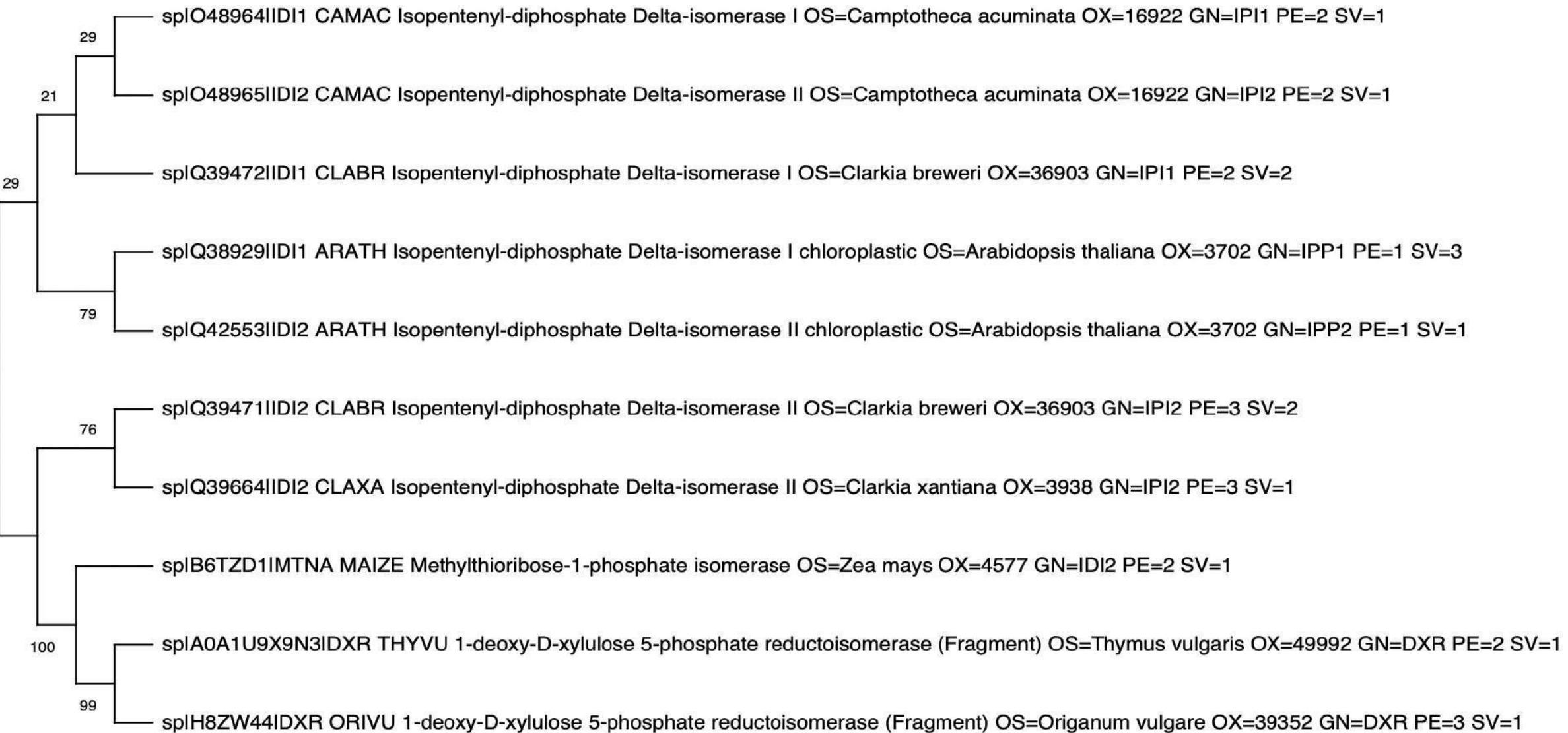
Diphosphocytidyl methyl erythritol (CDP-ME) Kinase



Methylerythritol 2,4-cyclodiphosphate (ME-cPP) Synthase



Isopentenyl diphosphate Isomerase



Isoprene Synthase

sp|P86851|SQE1 PANGI Squalene synthase 5E1 OS=Panax ginseng OX=4054 GN=SQE1 PE=2 SV=1

sp|O29267|DXS1 QRY SJ 1-deoxy-D-xylulose-5-phosphate synthase 1 chloroplastic OS=Oryza sativa subsp. japonica OX=39947 GN=CLAD1 PE=2 SV=2

sp|O29854|DXS ARATH 1-deoxy-D-xylulose-5-phosphate synthase 2 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=DXS PE=1 SV=2

sp|O29851|DXS ARATH 1-deoxy-D-xylulose-5-phosphate synthase 3 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=DXS PE=1 SV=2

sp|P46413|GPPS1 CAPAN Geranylgeranyl pyrophosphate synthase 7 chloroplastic OS=Capsicum annuum OX=4072 GN=GPPS1 PE=1 SV=1

sp|O24089|GPPS1 CAPAN Geranylgeranyl pyrophosphate synthase large subunit 1 chloroplastic OS=Capsicum annuum OX=4072 GN=GPPS1 PE=1 SV=2

sp|O24113|GPPS2 CAPAN Geranylgeranyl pyrophosphate synthase large subunit 2 chloroplastic OS=Capsicum annuum OX=4072 GN=GPPS2 PE=1 SV=2

sp|O24066|GPPS3 CAPAN Geranylgeranyl pyrophosphate synthase 7 chloroplastic OS=Capsicum annuum OX=4072 GN=GPPS3 PE=1 SV=2

sp|O24070|GPPS1 ARATH Geranylgeranyl pyrophosphate synthase 1 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS1 PE=1 SV=1

sp|O24071|GPPS2 ARATH Geranylgeranyl pyrophosphate synthase 2 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS2 PE=1 SV=1

sp|O24072|GPPS3 ARATH Geranylgeranyl pyrophosphate synthase 3 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS3 PE=1 SV=1

sp|O24087|GPPS4 ARATH Geranylgeranyl pyrophosphate synthase 4 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS4 PE=2 SV=2

sp|O24088|GPPS5 ARATH Geranylgeranyl pyrophosphate synthase 5 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS5 PE=1 SV=2

sp|O24089|GPPS6 ARATH Geranylgeranyl pyrophosphate synthase 6 mitochondrial OS=Arabidopsis thaliana OX=3702 GN=GPPS6 PE=1 SV=2

sp|O24090|GPPS7 ARATH Geranylgeranyl pyrophosphate synthase 7 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS7 PE=1 SV=2

sp|O24091|GPPS8 ARATH Geranylgeranyl pyrophosphate synthase 8 chloroplastic OS=Arabidopsis thaliana OX=3702 GN=GPPS8 PE=1 SV=2

sp|P86802|GPPS1 ARATH Heterodimeric geranylgeranyl pyrophosphate synthase large subunit 1 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS1 PE=1 SV=2

sp|Q12133|GPPS2 ARATH Heterodimeric geranylgeranyl pyrophosphate synthase large subunit 2 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS2 PE=1 SV=2

sp|Q12134|GPPS3 ARATH Heterodimeric geranylgeranyl pyrophosphate synthase 7 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS3 PE=1 SV=2

sp|O12135|GPPS4 ARATH Geranylgeranyl pyrophosphate synthase 8 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS4 PE=1 SV=2

sp|O12136|GPPS5 ARATH Geranylgeranyl pyrophosphate synthase 9 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS5 PE=1 SV=2

sp|O12137|GPPS6 ARATH Geranylgeranyl pyrophosphate synthase 10 mitochondrial OS=Arabidopsis thaliana OX=4072 GN=GPPS6 PE=1 SV=2

sp|O12138|GPPS7 ARATH Geranylgeranyl pyrophosphate synthase 11 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS7 PE=1 SV=2

sp|O12139|GPPS8 ARATH Geranylgeranyl pyrophosphate synthase 12 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS8 PE=1 SV=2

sp|O12140|GPPS9 ARATH Geranylgeranyl pyrophosphate synthase 13 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS9 PE=1 SV=2

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sp|O12142|GPPS11 ARATH Geranylgeranyl pyrophosphate synthase 15 chloroplastic OS=Arabidopsis thaliana OX=4072 GN=GPPS11 PE=1 SV=2

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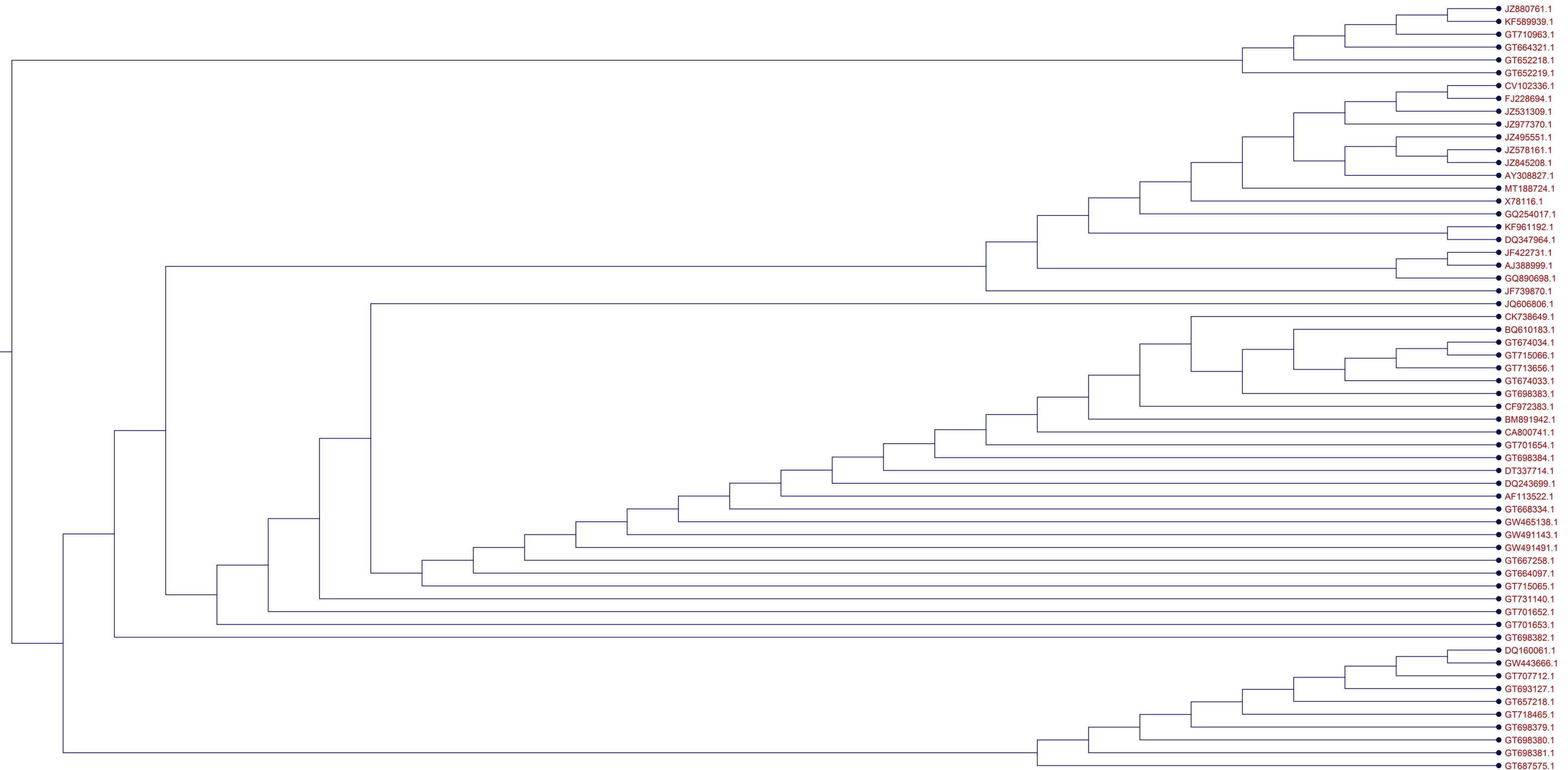
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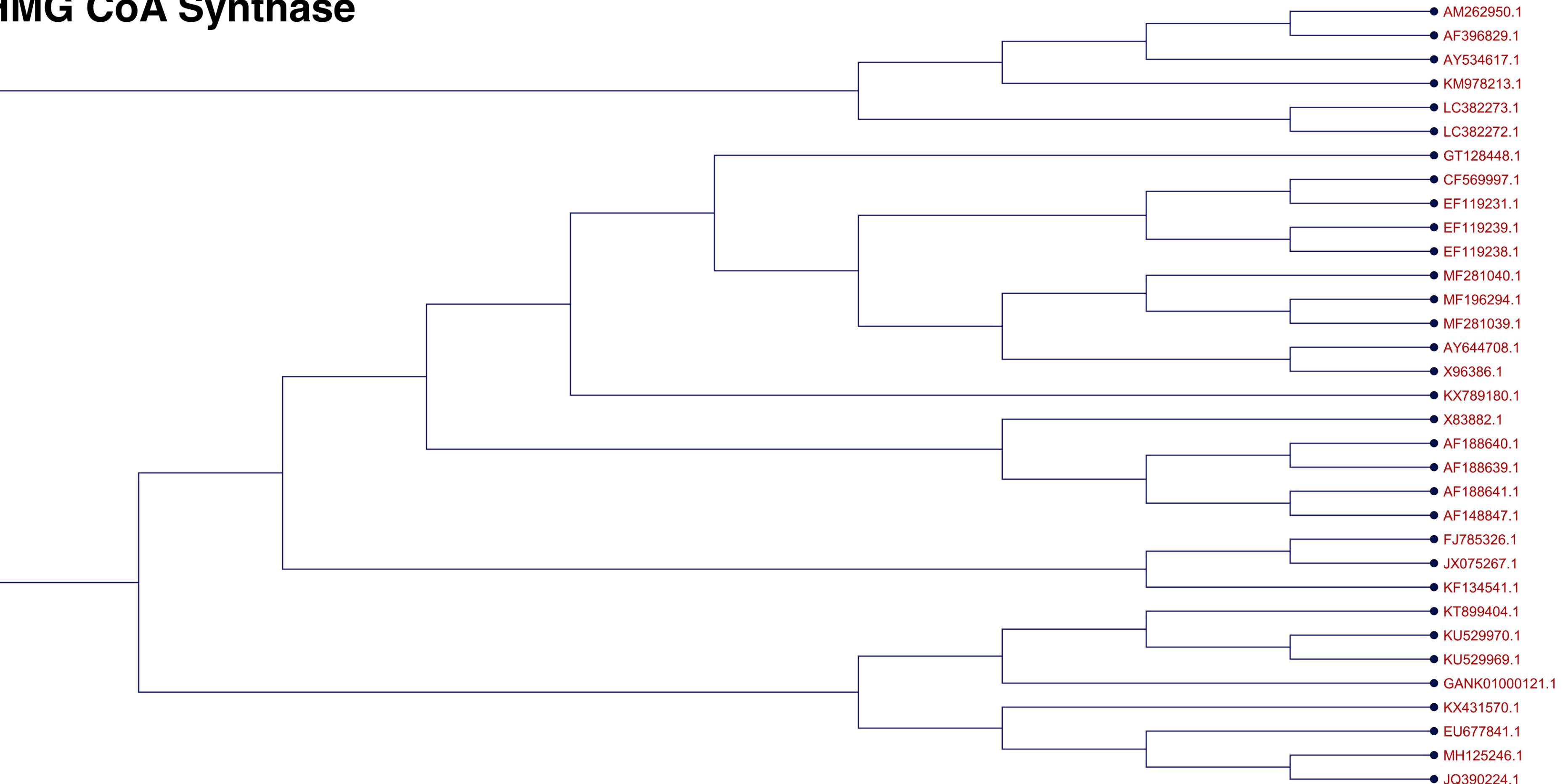
MEVALONIC ACID (MVA) PATHWAY

GENBANK PHYLOGENETIC TREES

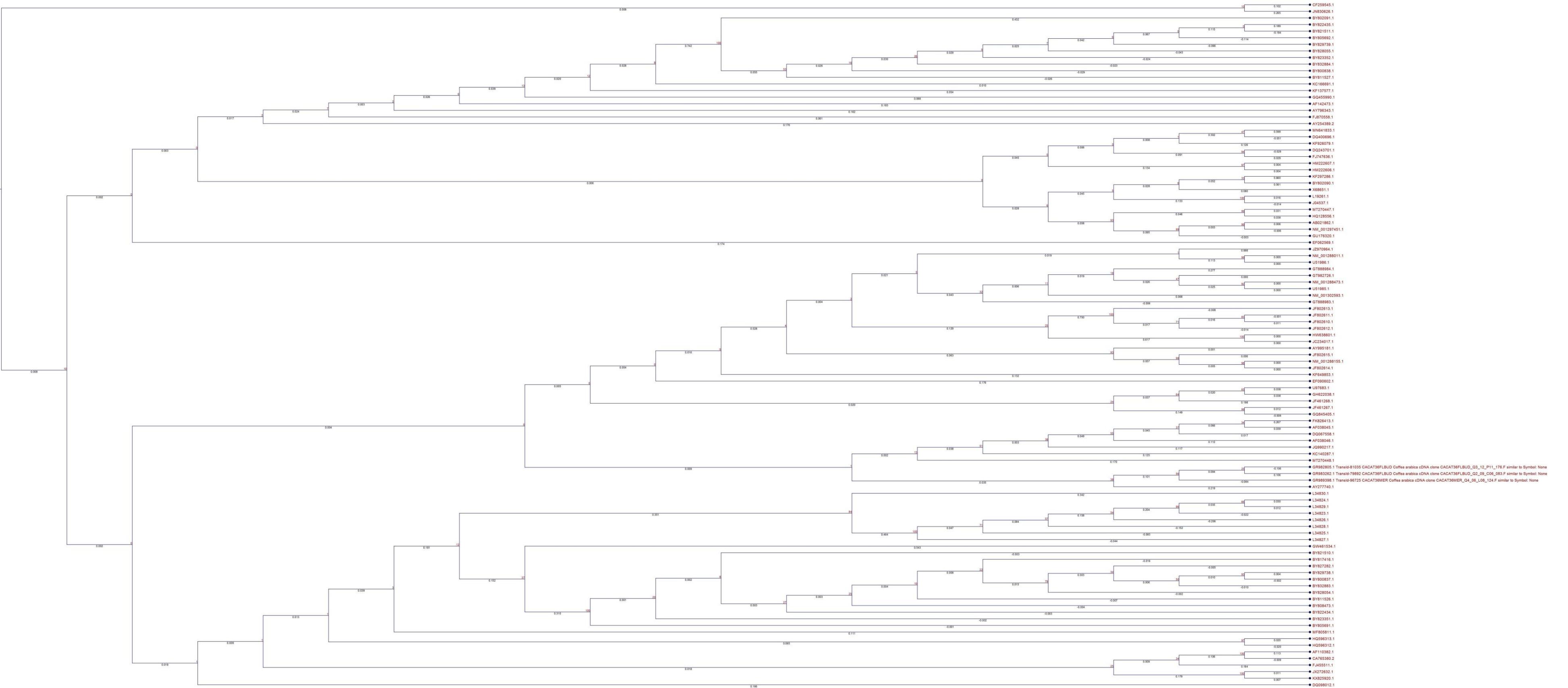
Acetoacetyl CoA thiolase



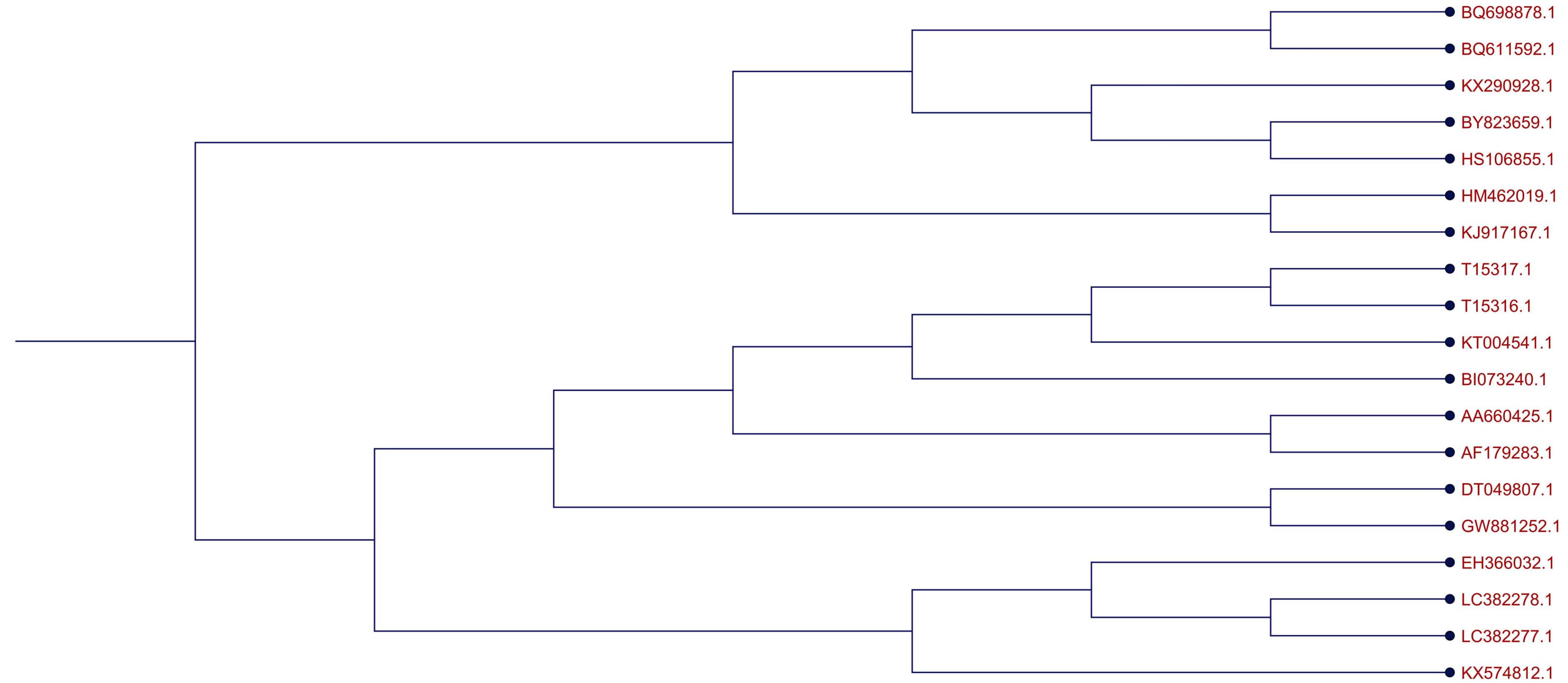
HMG CoA Synthase



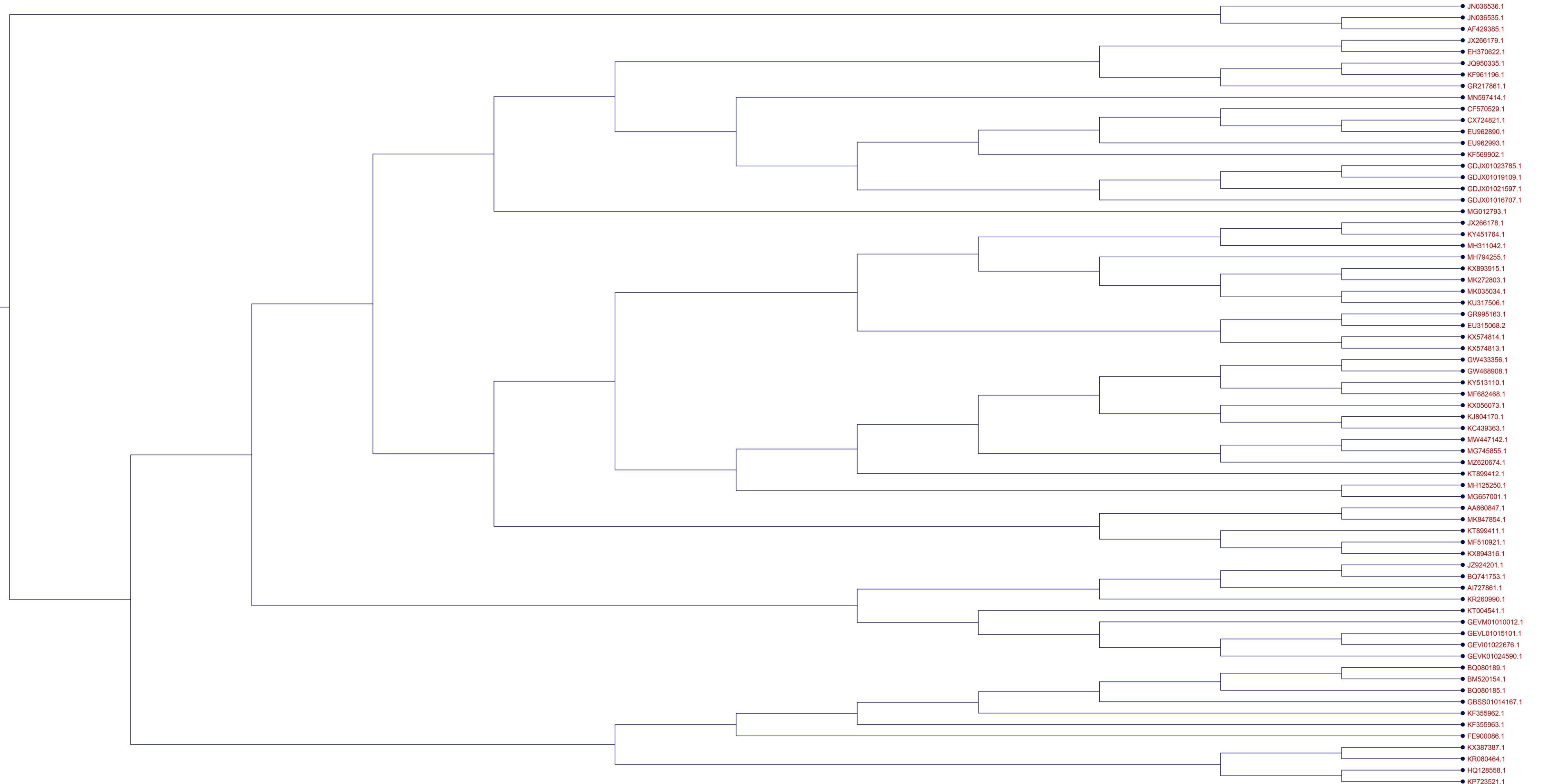
HMG CoA reductase



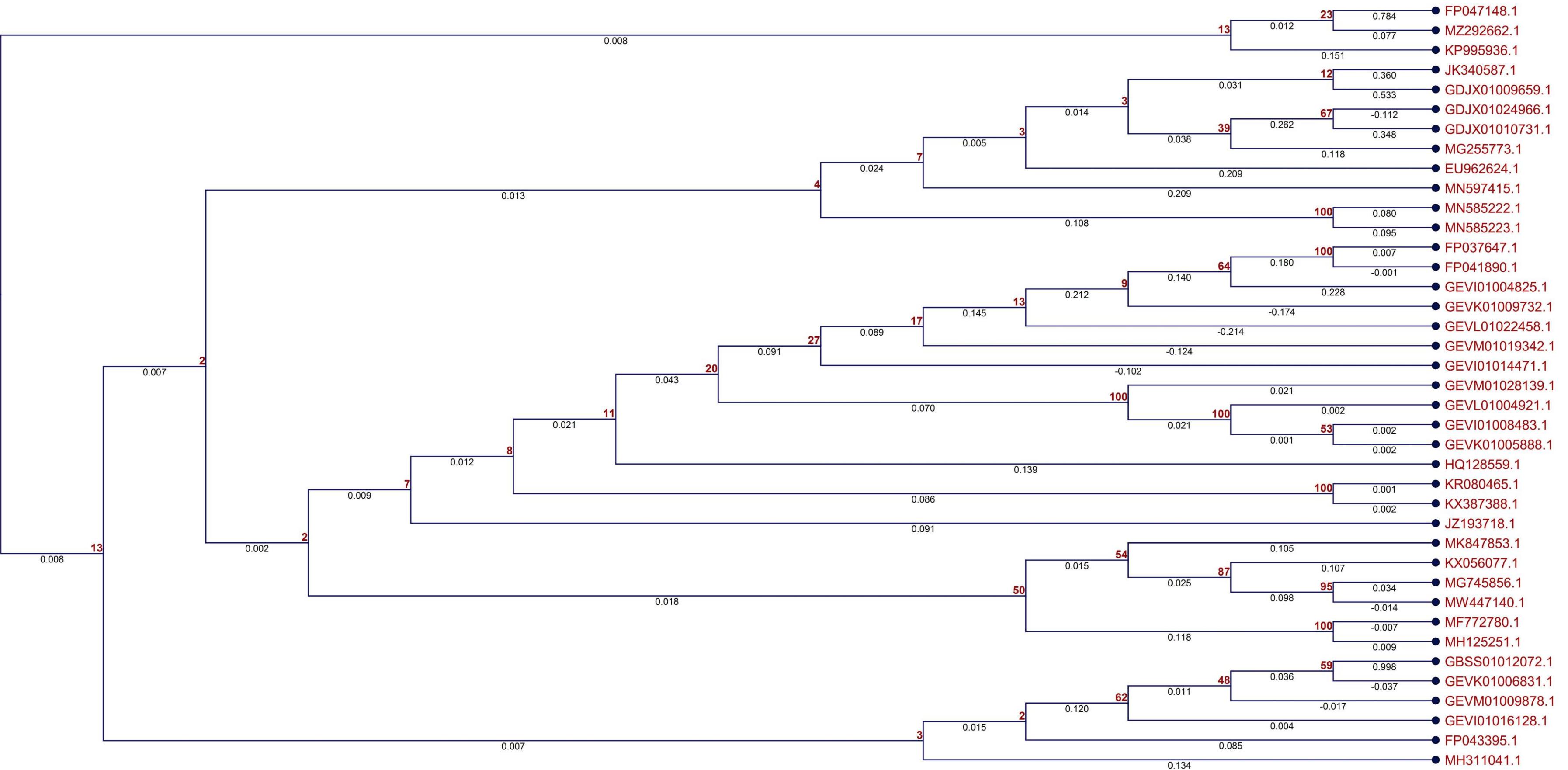
Mevalonate 5 kinase



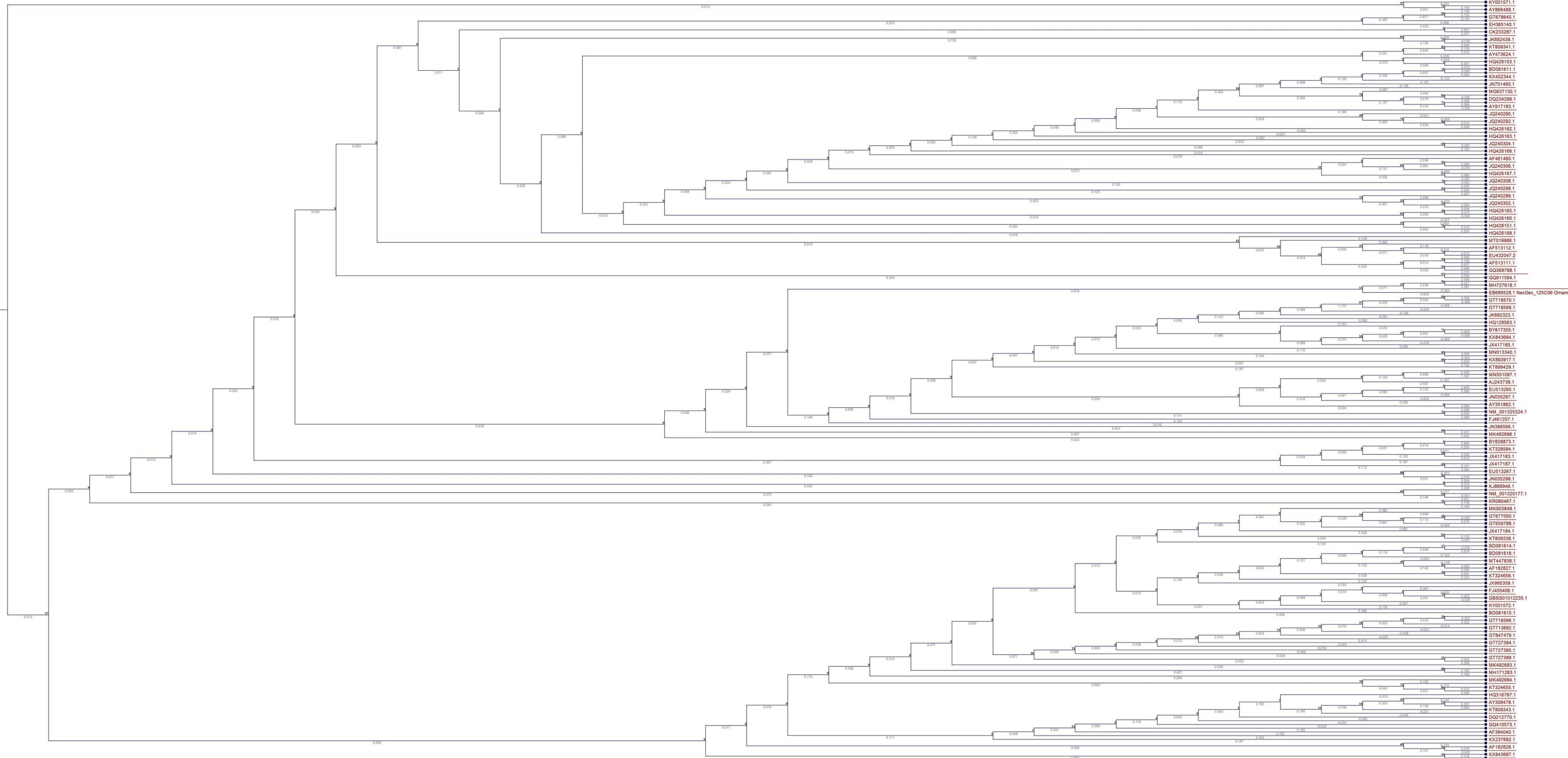
Phosphomevalonate Kinase



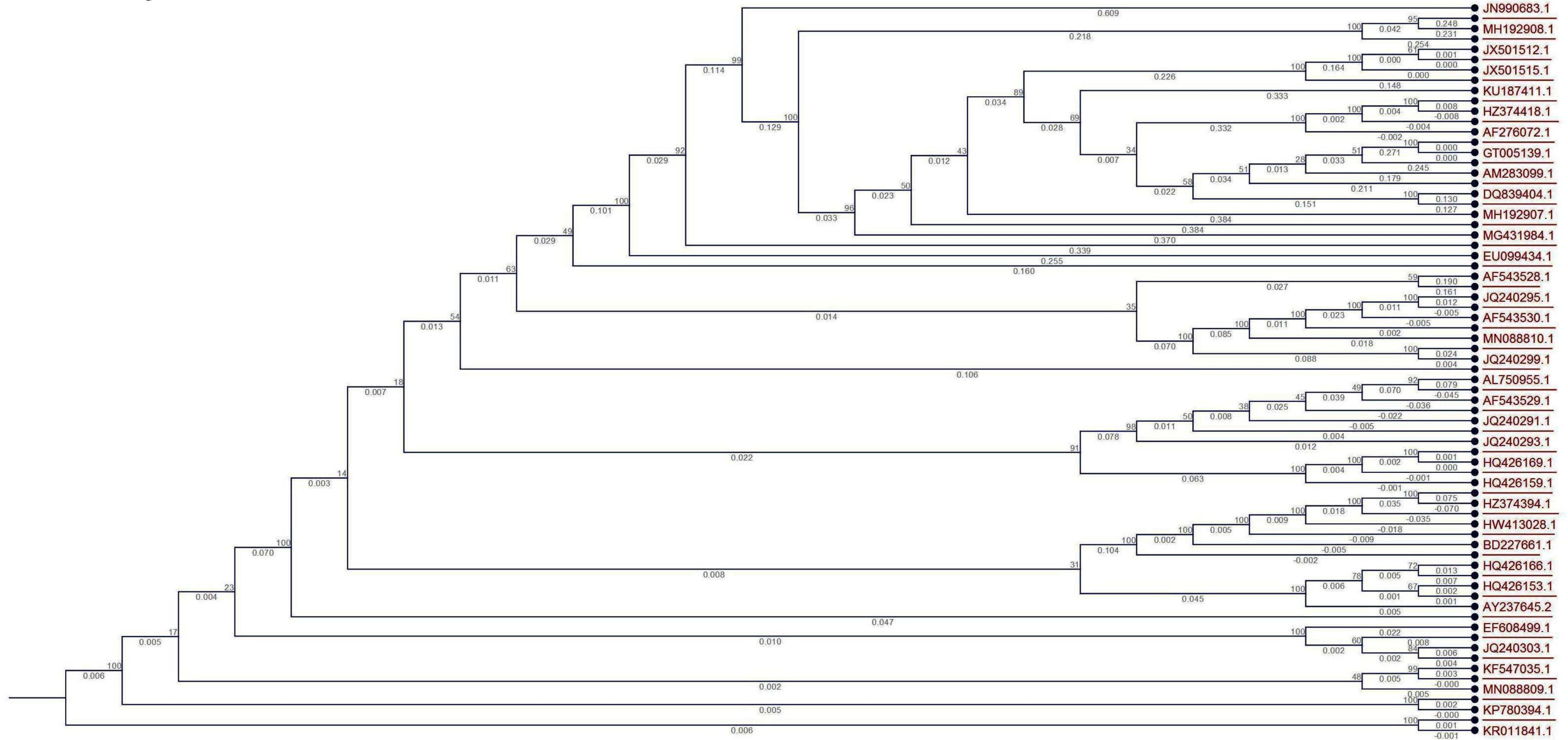
Diphosphomevalonate decarboxylase



Geranyl diphosphate synthase

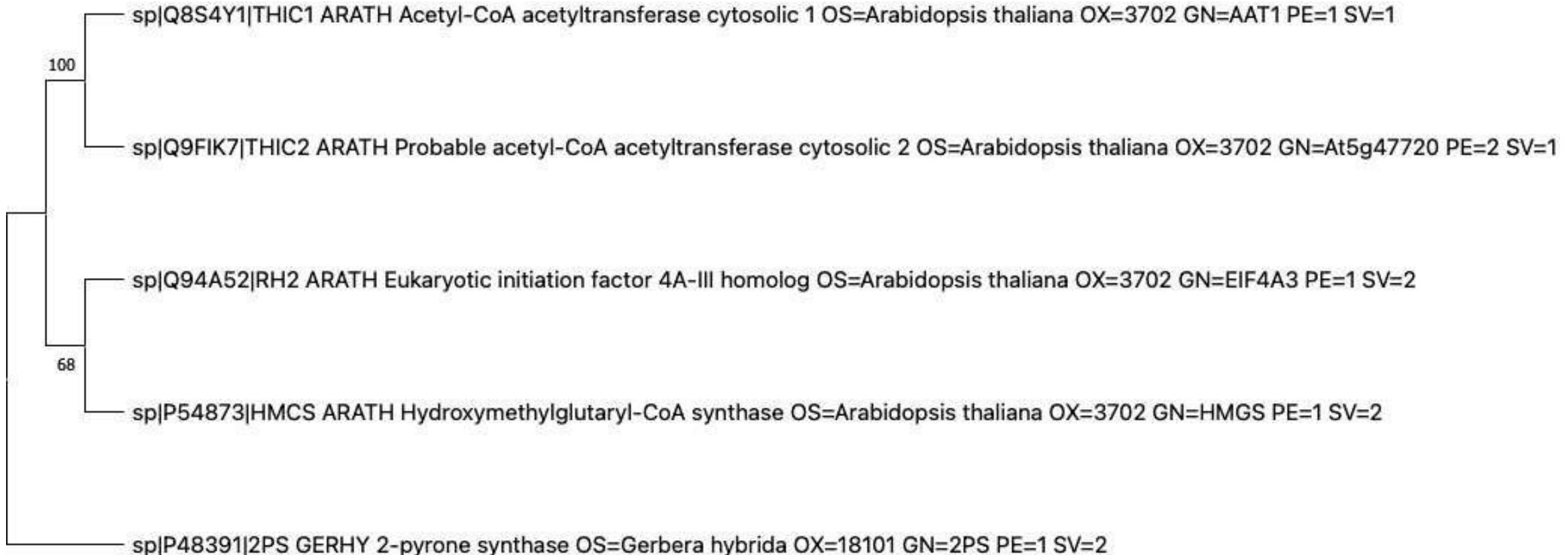


Pinene synthase



UNIPROT PHYLOGENETIC TREES

Acetoacetyl CoA thiolase



HMG CoA synthase

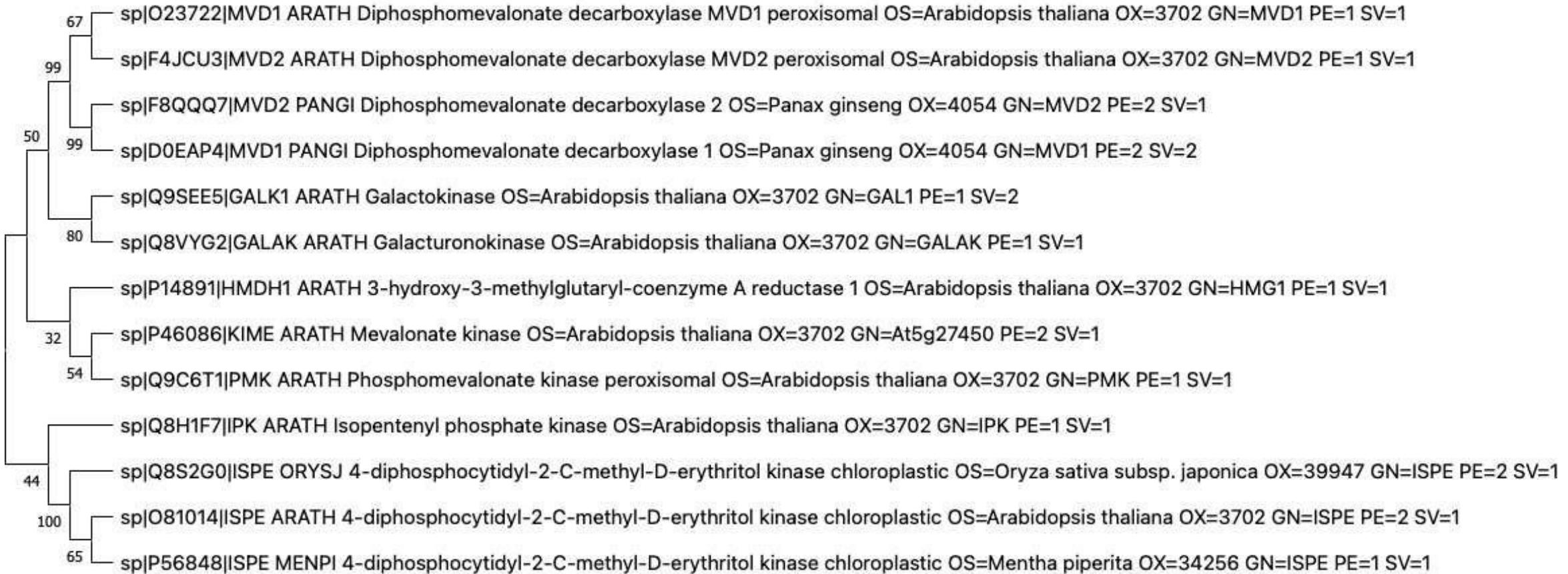
E

sp|P43256|HMDH2_ARATH 0.16454
sp|P14891|HMDH1_ARATH 0.1342
sp|Q03163|HMDH_CATRO 0.15495

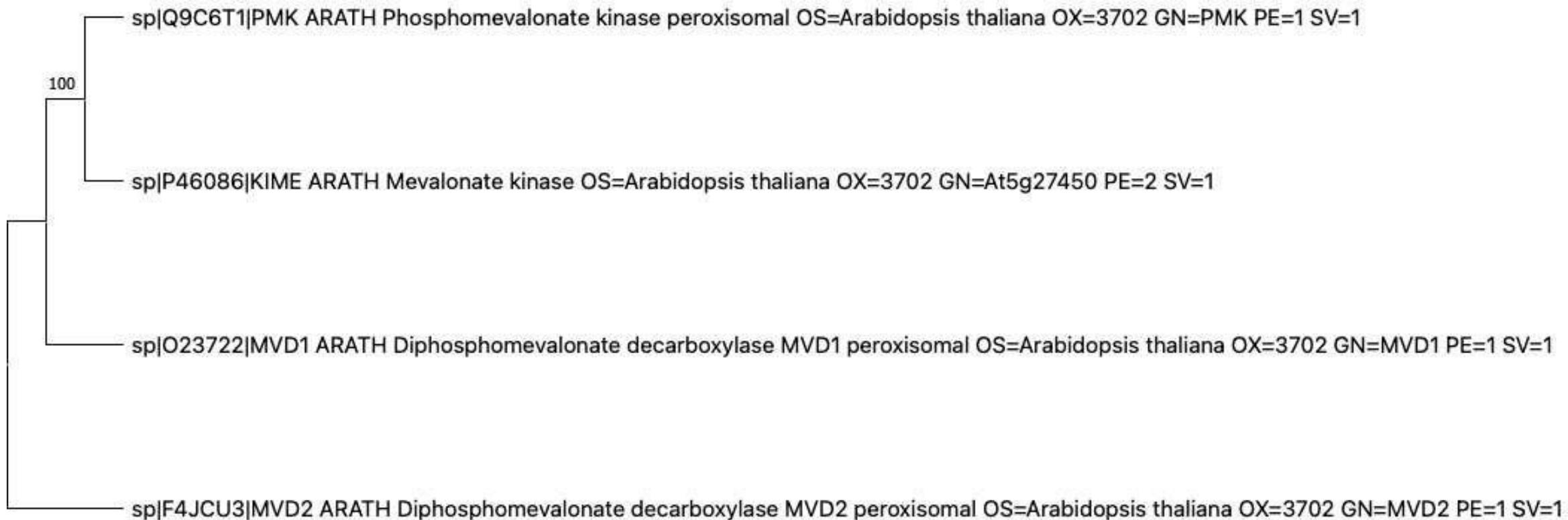
HMG CoA reductase



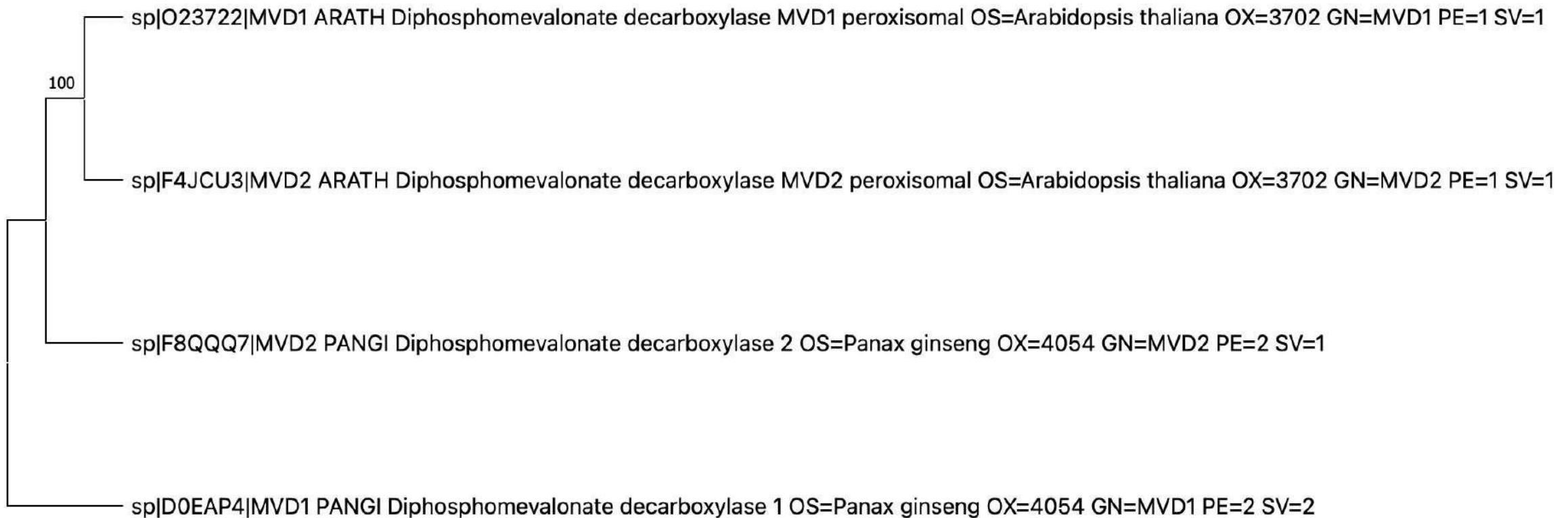
Mevalonate 5 kinase



Phosphomevalonate kinase



Diphosphomevalonate decarboxylase



Pinene synthase

