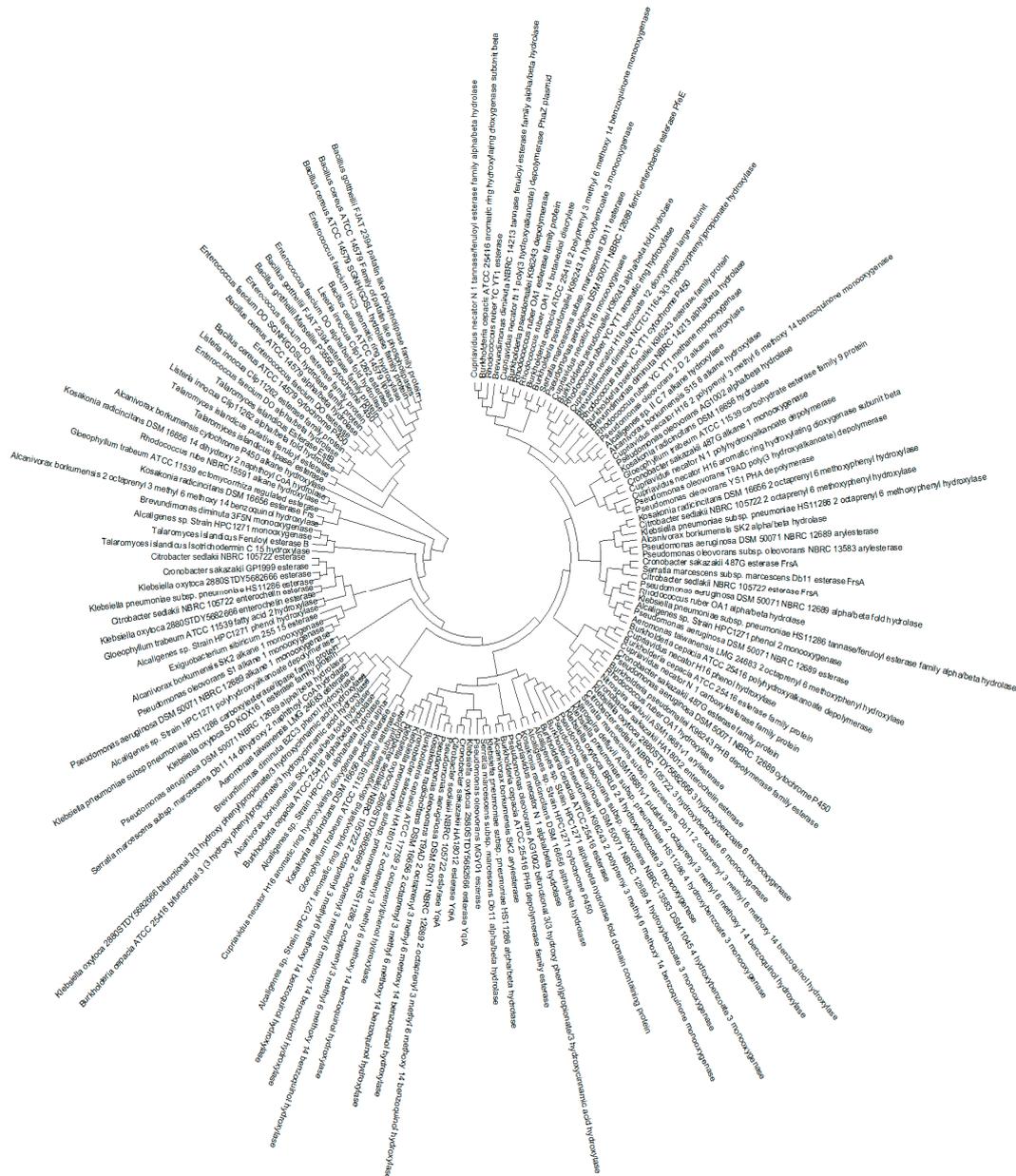


# Potential for and Distribution of Enzymatic Biodegradation of Polystyrene by Environmental Microorganisms

Liyuan Hou and Erica L.-W. Majumder



**Figure S1.** Phylogenetic tree of DNA sequences of all subclass enzymes from target microorganisms listed in Table 2 including cytochrome P450, monooxygenase, aromatic ring hydroxylase, esterase, and alpha/beta hydrolase.

**Table S1.** Potential PS degrading microorganisms and representative enzymes listed in the literature.

Kingdom	Species	Representative Potential Enzymes
Bacteria	<i>Bacillus cereus</i> ATCC 14579	Alpha/beta hydrolase, SGNH/GDSL hydrolase family protein, cytochrome P450
	<i>Cytobacillus gottheilii</i> ASM163634v1	Esterase family protein, cytochrome P450
	<i>Enterococcus faecium</i> DO	Alpha/beta hydrolase, SGNH/GDSL hydrolase family protein, aromatic ring hydroxylase, esterase family protein
	<i>Listeria innocua</i> Clip11262	Esterase family protein, alpha/beta fold hydrolase
	<i>Exiguobacterium sibiricum</i> 255-15	Esterase
	<i>Cupriavidus necator</i> N-1	Alpha/beta hydrolase, carboxylesterase family protein, polyhydroxyalkanoate depolymerase, phenol hydroxylase, monooxygenase, aromatic ring-hydroxylating dioxygenase subunit alpha
	<i>Burkholderia cepacia</i> ATCC 25416	Esterase family protein, alpha/beta fold hydrolase, polyhydroxyalkanoate depolymerase, PHB depolymerase family esterase, 2-polyprenyl-3-methyl-6-methoxy-1,4-benzoquinone monooxygenase
	<i>Burkholderia pseudomallei</i> K96243	PHB depolymerase family esterase, depolymerase, alpha/beta hydrolase, 2-polyprenyl-3-methyl-6-methoxy-1,4-benzoquinone monooxygenase
	<i>Klebsiella pneumoniae</i> subsp. <i>pneumoniae</i> HS11286	Alpha/beta hydrolase, carboxylesterase/lipase family protein, 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase, 4-hydroxybenzoate 3-monooxygenase, 2-octaprenyl-6-methoxyphenyl hydroxylase
	<i>Kosakonia radicincitans</i> DSM 16656	Hydrolase, alpha/beta hydrolase, 2-octaprenyl-6-methoxyphenyl hydroxylase, 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase
	<i>Cronobacter sakazakii</i> ASM98282v1	Esterase, esterase family protein, 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase, alkane 1-monooxygenase
	<i>Klebsiella oxytoca</i> ASM102219v1	Esterase, esterase family protein, 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase, bifunctional 3-(3-hydroxyphenyl)propionate/3-hydroxycinnamic acid hydroxylase, 4-hydroxybenzoate 3-monooxygenase
	<i>Citrobacter sedlakii</i> NBRC 105722	Esterase, 2-octaprenyl-6-methoxyphenyl hydroxylase, 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase, 3-hydroxybenzoate 6-monooxygenase
	<i>Alcaligenes</i> sp. Strain HPC1271	polyhydroxyalkanoate depolymerase, alpha/beta hydrolase, aromatic-ring-hydroxylating dioxygenase subunit beta, monooxygenase, phenol 2-monooxygenase, phenol hydroxylase, cytochrome P450
	<i>Aeromonas taiwanensis</i> LMG 24683	Esterase, 2-octaprenyl-6-methoxyphenyl hydroxylase
	<i>Brevundimonas diminuta</i> 48290_B02	Alpha/beta hydrolase, monooxygenase, phenol hydroxylase
	<i>Serratia marcescens</i> subsp. <i>marcescens</i> Db11	Esterase, alpha/beta hydrolase, 2-octaprenyl-3-methyl-6-methoxy-1,4-benzoquinol hydroxylase
	<i>Pseudomonas aeruginosa</i> PAO1	Esterase, alpha/beta hydrolase, 4-hydroxybenzoate 3-monooxygenase, cytochrome P450, alkane 1-monooxygenase
	<i>Rhodococcus ruber</i> ASM274172v1	Esterase, alpha/beta hydrolase, aromatic ring hydroxylase, cytochrome P450, alkane hydroxylase
	Eukaryota	<i>Gloeophyllum trabeum</i>
<i>Talaromyces islandicus</i>		Feruloyl esterase B, putative feruloyl esterase A

**Table S2.** DNA sequences of higher potential subclass enzymes from target microorganisms including cytochrome P450, monooxygenase, and aromatic ring hydroxylase.

Enzymes from Species	Accession NO.	
<i>Bacillus cereus</i> ATCC 14579_cytochrome P450	WP_000062123.1	<p>ATGTCAATGAAAAACAAAGTTGGGTAAAGCATAGAGGATGGCATTAAATTTAG            CTTACAGCTCAGTTTAAAGAAGATGCGTATGAAATTTATAAAGAATCGCGAAA            AAAACAACCTATATTTTGTAAACCAAGTTGAAATCGGTAAAGAATGGCTC            ATTACTAGATATGAAGATGCTCTGCCACTTTTAAAAGATAATCGTTTAAAAAA            AGATTGGACAAATGTGTTTTCTCAAGATATAAAGAACATGTATCTTTCCGTTG            ATAATAGTGACCCTTAACAACACATATGCTAAATTCAGATCCACCTAACCA            CAGTCGTTTACGATCTTTAGTTCAAAAAGCTTTTACACCGAAGATGATTGCAC            AATTAGACGGAAGAATTCAGAGAATAGCAGATGATTTGATAAGTGATATAGA            GCGAAAAGGTACATTAATCTTGTGGATGATTATTCATTTCCATTACCAATTA            TTGTAATAAGCGAGATGCTCGGTATTCCAAAAGAAGATCAAGCGAAATTTAG            GATTTGGTCTCATGCTGTTATTGCATCACCGAAACACCTGAAGAAATAAAA            GAGACTGAAAAACAATCTGAATTTATTACATATCTTCAATATTTAGTTGA            TATTAAGAAGAAAGAGCCAAAAGAAGACTTGGTGAGTGCTTTAATACTTGCA            GAGAGTGAAGGGCATAAACTTAGCGCTCGGGAACCTATATTCAATGATAATGC            TATTAATTGTCGCAGGACATGAGACGACAGTGAATTTAATTACAAATACGGT            ATTAGCACTTCTTGAATAATCCAAATCAATTACAGTTATTAAGAATAATCCAA            AACTAATTGATTCCGGCTATTGAGGAAGGATTGCGTTATTATTCTCCAGTTGAG            GTTACAACCTGCAAGATGGGCAGCGGAACCTTTTCAAATTCACCATCAAACAA            TACAGAAAGGAGATATGGTTATTATTGCATTGGCTTCAGCGAACCGTGATGA            AACAGTATTTGAAAATCCAGAAATATTTGATATTACACGGGAGAACAACCGT            CACATTGCCTTTGGTCATGGTAGTCATTTCTGCTTAGGAGCTCCACTTGCGAGG            TTAGAAGCAAAGATTGCTATTACTACTTTGTTAATCGAATGCCTGAACTACA            AATAAAAGGGAATCGTGAAGAAATTAATGGCAAGGTAACCTATTTAATGCGT            TCTTTAGAGGAATTACCTTTAACTTTCTAG            ATGACTGAAGTAAAACACTTTCCAAAAGAAGAGGGTCTCGATCATAGCCTTG            ATTTATTAAGAAGGCTATTTGTTTATTACAAATCGCAGTACTGCATTTCAA            TCAGATATTTTTGAAACACGCCTGCTGGGAGAGCGGGTTATTTGTCTGCGAGG            CAAGGAAGCTGCATCAATCTTTACGATACGCAGAAATTCAAACGAGAAGGT            GCAGCCCCAAATCGAATTCAGAAAACCTGTTAGGGAAAAACGGGGTACAA            TCACTGGATGGTGAAGCACATCAGCATCGGAAGGCAATGTTTATGTCGTTAAT            GACACCTGCTTCTCCAAAGAATGCGTGCTTTATAAAAAAAGAGTGGGAC            ATTGCGGCCAAGAAGTTTTCAAGAAAAGAATTATCCTTTACGAAGAAG            CAAAAGGGTGCTTTGTAAAGCAGCTTGTGACTGGGCTGGTGTTCCTCTTAAA            GAGAACAATGTTGAGGAAACGGCCAATTTACTTGGATTACTATTTGAAACAC            CCGCAGCACTCGGCCCTAAGCATTGGCAAGGCAGACACGCACGGACAAAAT            TGGAAAAATGGCTTAAGGAATTAGTCATAGAAGTTCGAAACGGCAAATTCCT            CCTCCTGAAGATAAATCCTTATTCATTATTTCAAATCACCGCAATTTGGATG            GTGAGCTGCTTGTGACAGACATCGCCGCTGTTGAACTATTAATATTTGCGT            CCTATTGTCGCGGTAGCCGTGTATATTTGTTTTACGGCGCTTGCTGTATACCAG            CATCCAAAAGAAGCAGGCAAACCTGCGCGGGTATGATGAAAATCTTCTTCAGA            ACTTTGTACAAGAGGTCAGGAGATTTTATCCATTTTTCCGTTTGCACCAGCA            AGAGTAAAAGCGGACTTCACATGGAATGGATATCTTTTGAGGAAAATACAT            TAACACTCCTCGATTTATATGGAACAAACCACCACCCTAAATTATGGGATAA</p>
<i>Bacillus gottheilii Marseille- P3555_cytochrome P450</i>	WP_080848536.1	<p>ATGACTGAAGTAAAACACTTTCCAAAAGAAGAGGGTCTCGATCATAGCCTTG            ATTTATTAAGAAGGCTATTTGTTTATTACAAATCGCAGTACTGCATTTCAA            TCAGATATTTTTGAAACACGCCTGCTGGGAGAGCGGGTTATTTGTCTGCGAGG            CAAGGAAGCTGCATCAATCTTTACGATACGCAGAAATTCAAACGAGAAGGT            GCAGCCCCAAATCGAATTCAGAAAACCTGTTAGGGAAAAACGGGGTACAA            TCACTGGATGGTGAAGCACATCAGCATCGGAAGGCAATGTTTATGTCGTTAAT            GACACCTGCTTCTCCAAAGAATGCGTGCTTTATAAAAAAAGAGTGGGAC            ATTGCGGCCAAGAAGTTTTCAAGAAAAGAATTATCCTTTACGAAGAAG            CAAAAGGGTGCTTTGTAAAGCAGCTTGTGACTGGGCTGGTGTTCCTCTTAAA            GAGAACAATGTTGAGGAAACGGCCAATTTACTTGGATTACTATTTGAAACAC            CCGCAGCACTCGGCCCTAAGCATTGGCAAGGCAGACACGCACGGACAAAAT            TGGAAAAATGGCTTAAGGAATTAGTCATAGAAGTTCGAAACGGCAAATTCCT            CCTCCTGAAGATAAATCCTTATTCATTATTTCAAATCACCGCAATTTGGATG            GTGAGCTGCTTGTGACAGACATCGCCGCTGTTGAACTATTAATATTTGCGT            CCTATTGTCGCGGTAGCCGTGTATATTTGTTTTACGGCGCTTGCTGTATACCAG            CATCCAAAAGAAGCAGGCAAACCTGCGCGGGTATGATGAAAATCTTCTTCAGA            ACTTTGTACAAGAGGTCAGGAGATTTTATCCATTTTTCCGTTTGCACCAGCA            AGAGTAAAAGCGGACTTCACATGGAATGGATATCTTTTGAGGAAAATACAT            TAACACTCCTCGATTTATATGGAACAAACCACCACCCTAAATTATGGGATAA</p>

<p><i>Enterococcus faecium</i> IHC3_aromati c ring hydroxylase</p>	<p>RCT70246.1</p>	<p>CCCTGAGCTCTTTCAGCCGGATCGTTTTTCAAACCTGGAAGGACAGTCCATTCA GTTTTATTCTCAAGGCGGAGGTGACCATGATTTTGGTCATCGCTGTGCAGGC GAATGGGTGACGATTGAAATTATGAAGGAAACGCTGAATTTTCTTGTGAATA AAATTAGCTTCGCCATTCCAGATCAGGACTTAAGCTATAGCTTTAATGATATT CCAGCTCTTCTCACAGTAAAATCATTATGAAAGAAATTCATTTAAAATAA ATGATAAATAAAAATCGTGTTAAGGAGTGTCTGACGATGAGCGAAGCAAAC CAAGAACGGTCTGGACAAGAAATAGAATAATCAAAGAACGTATTCTTGCTG CCTTGAAAATGGTGATTGATCCTGAATTAGGAATCGACATCGTCAATTTAGGC TTGATTTATGATATAGAATTCAATCCTGAAAATGGAGAAAACAGTCATTAAGA TGACTCTAACAACATATGGGCTGTCCATTAGCAGATATTTAACAGAATCTATC CATGACGCATTAAGAAGTGCCAGAAGTAACGAAAGAAGAAGTAAAATA GTCTGGTAG</p>
<p><i>Cupriavidus necator</i> H16_phenol hydroxylase</p>	<p>WP_010810421.1</p>	<p>ATGAACACCCCGTCGGTTCCTGTTTGAAGCCACGCCGCGCTATGTGCGGGT CGAGGGCCGACCCCGAAGGGTTTGTGCAGTTCGCCTTCAGCGTGGCCGAC CCGGAGCTCAACGTCGAACTGATCATGCCGGAACCGATGTTTGAAGCCTTCT GCTGCGTCAACCGGTGCGCTTCCTGCCCCGCTGGAAACCGCGCCGACGCC GCAAGCCGACGACTGA</p>
<p><i>Cupriavidus necator</i> H16_monoox ygenase</p>	<p>WP_010810419</p>	<p>ATGTCAGCCACCGCTGCCAACGTCTATATCGCCCTGCAGAACAACGACGACA CCCGACCCATCATCGACGCCATACCGAGGCCAACCCGCATGCGGTGGTGTG GCAGTTTCCCGCCATGGTCAAGATCGACGCCCGCCACCTGACCATCGTG CGCGAACTGGTGGCCGGCAAGCTCGGCCGCGACTGGGACCTGCAGGAGATC CACCTGAACCTGATCTCGCTGTGCGGAAACATCGACGAAGACGAAGACGCCT TCACGCTGCGCTGGAACGCCTGA</p>
<p><i>Cupriavidus necator</i> H16_2- polyprenyl-3- methyl-6- methoxy-1,4- benzoquinone monooxygenase</p>	<p>WP_010814779.1</p>	<p>ATGGACACCCTGATCAAAGAATTCGACGTGGCATTGCGCGCCATTGCCGGCG CCACGCGCACCGCGCGCGCAACCCGGCTGACCGGCTGGCGCCGGACACCG AACAGATGAGCGCCGATGAACGCCGCCACGTGGCCGGGCTGATGCGCATCA ACCATGTCGGCGAGGTCTGCGCGCAGGCCCTGTACCAGGCCCAGAAGCTCAC CGCGCGGACCGCGCGGTGCGCGCGCAGATGGATGCCGCCGCGCGCAAGA GGAAGACCACCTGGCCTGGTGCGCCGAGCGCCTGCGCGAGCTGGGCTCGCGC CCGAGCCTGCTCAACCCGCTGTGGTACGCCGGCGCCTTCGCCATCGGCTGGAT GGCCGGCCGCGCCGGCGACCGCGTCAGCCTGGGCTTCGTGCGCGAGACCGAG CGCCAGGTCGAGCACCACTGGGCGGGCACCTGGACCGCTGCCCGAGGCC GACGGCCGCTCGCGCGCCATCCTCGAGCAGATGCGCGACGACGAGATCCGCC ACGGCAACGCCGCGCGCGATGCCGGCGGGATCCCGTTACCTGCCCGGTGCG GGCGCTGATGCGCGCGCCTCGCGCGTATGACCACCGCCGCCTACCGGATC TGA</p>
<p><i>Cupriavidus necator</i> H16_benzoate 1,2- dioxygenase large subunit</p>	<p>WP_013956903.1</p>	<p>ATGATCCCCATCTATCCGGAGGGCAAGGCCCGTTTCGCCCGGCGCAAAGC GCCTCGACCTGGATACTTCTTGGTGAAGACAAGGCAAGCGGCGACTACCG GCTGCACCGCGCGGCCTTCACTGACGAAGAGCTGTTTGAACCTCGAGATGAAG CACATCTTCGAGGGCAACTGGATCTACCTGGCGCACGAAAGCCAGATCCCGA ACAACAACGACTACTACACCACCCATATCGGCCGCCAGCCGGTGGTGATCGC GCGCAATCGCCAGGGCGAACTGAACGCCTTCATCAACCGGTGCAGCCACCGC GGCGCCATGCTGTGCCGCCACAAGCGCGGCAACAAAGCCACCTACACCTGCC CGTTCCACGGCTGGACCTTCAACAACAGCGGCAAGCTGCTGAAGGTGAAGGA TCCCGAGAACGCCGGCTACCCTGACTGCTTCAACAAGGAAGGCTCGCACGAC CTGAAGAAGGTGGCGCGCTTCGAGAATAACCGCGGCTTCCTGTTGCGCAGCC TGAACGCTGACGTGCCGCCGCTGAAGGATTTCTGGGCGAAGCGGCGCGCGT CATCGACATGATCGTCGACCAGTCGGCCGACGGGCTCGAGGTGCTGCGCGGC</p>

*Cupriavidus  
necator*  
H16\_aromatic  
ring-  
hydroxylating  
dioxygenase  
subunit alpha

WP\_010811430.1

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*Cupriavidus  
necator*  
H16\_aromatic  
-ring-  
hydroxylating  
dioxygenase  
subunit beta

WP\_041688019.1

*Burkholderia cepacia* ATCC 25416\_2-polyprenyl-3-methyl-6-methoxy-1,4-benzoquinone monooxygenase

WP\_027788457.1

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*Burkholderia cepacia* ATCC 25416\_aromatic-ring-hydroxylating dioxygenase subunit beta

WP\_027792273.1

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*Burkholderia cepacia* ATCC 25416\_bifunctional 3-(3-hydroxyphenyl)propionate/3-hydroxycinnamic acid hydroxylase

WP\_027787836.1

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TTCCCGAAGCTCGGCCACGGCATCGTGCACGCGCAAGCCGACGGCACGCTGC

*Burkholderia  
cepacia* ATCC  
17759\_2-  
octaprenylph  
enol  
hydroxylase

POM19962.1

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CTCGTGTCCGCCGCGCTCGGCTGA

*Burkholderia  
pseudomallei*  
K96243\_2-  
polyprenyl-3-  
methyl-6-  
methoxy-1,4-  
benzoquinone  
monooxygenase

WP\_004527787.1

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*Burkholderia  
pseudomallei*  
K96243\_4-  
hydroxybenz  
oate 3-  
monooxygena  
se

WP\_004556451.1

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*Klebsiella  
pneumoniae*  
subsp.  
*pneumoniae*  
HS11286\_2-  
octaprenyl-3-  
methyl-6-  
methoxy-1,4-  
benzoquinol  
hydroxylase

WP\_002894733.1

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*Klebsiella  
pneumoniae*  
subsp.  
*pneumoniae*  
HS11286\_4-

WP\_004151384.1

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hydroxybenz  
oate 3-  
monooxygena  
se

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*Klebsiella*  
*pneumoniae*  
subsp.  
*pneumoniae*  
HS11286\_2- WP\_002916482.1  
octaprenyl-6-  
methoxyphen  
yl  
hydroxylase

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*Kosakonia*  
*radicincitans*  
DSM 16656\_2- WP\_007370239.1  
octaprenyl-6-  
methoxyphen  
yl  
hydroxylase

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*Kosakonia  
radicincitans*  
DSM 16656\_2-  
octaprenyl-3-  
methyl-6-  
methoxy-1,4-  
benzoquinol  
hydroxylase

WP\_007373869.1

*Cronobacter  
sakazakii*  
HA18012\_2-  
octaprenyl-3-  
methyl-6-  
methoxy-1,4-  
benzoquinol  
hydroxylase

WP\_166492126.1

<p><i>Cronobacter sakazakii</i> 487G_alkane 1- monooxygenase</p>	<p>WP_105944427.1</p>	<p>CAGGCGCGCCTGGTGATCGGTGCGGACGGTGCCAGCTCGCAGGTACGTAAT GGCAGGTATCGGCGTTAACGCCTGGCAGTATCAGCAGTCATGCCTGTTGATT AGCGTGGAATGCGAGCAGCCGCCGGCGACAGCACCTGGCAGCACTTTACCC CGAACGGGCCGCATGCGTTTTTACC GTTGTGTTGATAACCGTGCGTTCGTC TGGTATGACCGCCCGGCACGTGTTTCGTCAGCTTCAGGCGATGACCATGACAC AGCTTGGGCGAGAGATTGCCGCCGCAATTCCTGCCCGGCTTGGCCGCGTGAC GCCGTCGCCTGCGGCGCGTTTTCCGCTGGTGCCTCGCCATGCGCTGCGTTATG TGCAGCCGGGCGTTCGCGCTAATTGGCGACGCCGCGCATAACCATCATCCGT GGCGGGCAGGGGGTGAATCTCGGTTATCGCGATGTGGACGCGCTGCTGAAG GTGGTGATTGATGCCCGCAATGCGGCGGAAGACTGGGCTTCCGGCGCGGTGC TTAAGCGCTATCAGCGTTCGTCGATGCCGGATAACCTGCTGATGCAAAGCGG TATGGATCTTTTCCATGCCGGTTTTACTGCGAAACTGAAACCGCTGCGCGTGC TGCGCAATCTTGGGCTTATCGCCGCCAGCGCTCAGGCGTGCTGAAACGCCA GGCGCTGCGGTATGCGCTCGGTTTATGA ATGCGCTGCCCGCAGGCGTTCGCGCCGCGCATTATGGCCTCGCGCACCCGCTTT TGTGACGGATAACAGTGACGACGCCCGCCGCTTCGCGCAAACCGGGCTGGCG GCTCAGGCCGCGCAGTTCCGCGCGCAGGGCCACCGGCTTACGGGCGAGACG CTGGAAGATTATATCGCCCGGTTTCGATGTGCATCTCGGCACCCCGCGGGAAA CGCTCGCCACGCTGTCGCGCGATAACGGCGCTCGCACGCGCCACCGATCTGTC GTTCCAGGTGCATTCCATCGATCCGCCGACCCGTTTATCTTGCCTCCATCTC ACTGCTGGCGCAGCAGGTGGCCCCCGCGCTTGGCTGGAAACAGGCCGACCCG GGCGCTGATGCCTCCACCCTTAACCCGATCAAGGAAACCTTATGA ATGACAAATCAACCAACGGAATGCCATTGTGCGCGGAGGAATGGTCCGC GGCGACTGGCGCTGGGGCTGGCACAGCACGGATTTGCGGTAACGGTGATCG AGCACGCAGAACCTGCACCGTTTGTGCTGACAGCCAGCCTGACGTGCGGAT CTCGGCGATTAGTGGGCTTCGGTATCATTGCTTAAAGGTTTAGGCATCTGGG ACGCGGTACAGGCTATGCGTTGCCATCCTTACCGCAGACTGGAAACGTGGGA GTGGAAACGGCGCATGTGGTGTGTTGACGCCGCTGAACTTAAAGTACCGTTG CTTGGCTACATGGTGGAAAACACTGTCCTGCAACAGGCGCTGTGGCAGGCGC TGGAAGCGCATCCGAAAGTAAACGTTACGTGTGCCAGGCTCGCTGATTGCATT ACATCGAGATAATGATCTTCAGGAGCTGGAGCTGAAAGGCGGTGAAGTGATC CAGGCCAAGCTGGTGATTGGCGCTGACGGTGCAAATTCGCAGGTGCGGCAA TGGCGGGAATTGGCGTTCATGCCTGGCAGTATGCGCAGTCGTGTATGTTGATT AGCGTACAGTGCAGAAATGATCCCGGCGACAGCACCTGGCAGCAGTTTACTC CGGACGGGCGCGTGCCTTTTTGCCGCTGTTGATAACTGGGCGTTCGCTGGTG TGGTATGACTCTCCGGCGCGTATTCGCCAGTTGCAGAAATATGAATATGGCGCA GCTCCAGGCGGAAATCGCGAAGCATTTCGCCGTCGCGTCTGGGTTACGTGACA CCGCTTGCCGCTGGTGCCTTTCCGCTGACACGTGCCATGCGTTGACGATGATG GCAACCAGGGCTTACGCTGGTGGGCGATGCCGCGCACACCATTATCCGTTG GCGGGCAGGGGTTAACCTCGGTTATCGCGATGTTGATGCGCTGATTGACGT TCTGGTGAATGCCCCGAGCTACGGTGAAGCGTGGGCCAGTTATCCTGTCTCA AGCGTTACCAGATGCGGCGTATGGCGGATAACTTCATTATGCAAAGCGGTAT GGATTGTTTTATGCCGGATTCAGCAATAATCTGCCACCCTGCGTTTTATGCG TAATCTCGGATTAATGGCGGCGGAGCGTGCTGGCGTGTGAAACGTCAGGCC CTGAAATATGCGTTAGGGTTGTAG ATGGCAATACAACACCCTGACATCCAGCCTGCTGTTAACCATAGCGTTCAGG TGGCGATCGTGGTGCCGGTCCGGTCCGGCTGATGATGGCGAACTATCTCGG CCAGATGGGCATTGACGTGCTGGTGGTGGAGAACTCGATAAGTTGATCGAC</p>
<p><i>Klebsiella oxytoca</i> 2880STDY568 2666_2- octaprenyl-3- methyl-6- methoxy-1,4- benzoquinol hydroxylase</p>	<p>WP_022645356.1</p>	<p>AGCGTACAGTGCAGAAATGATCCCGGCGACAGCACCTGGCAGCAGTTTACTC CGGACGGGCGCGTGCCTTTTTGCCGCTGTTGATAACTGGGCGTTCGCTGGTG TGGTATGACTCTCCGGCGCGTATTCGCCAGTTGCAGAAATATGAATATGGCGCA GCTCCAGGCGGAAATCGCGAAGCATTTCGCCGTCGCGTCTGGGTTACGTGACA CCGCTTGCCGCTGGTGCCTTTCCGCTGACACGTGCCATGCGTTGACGATGATG GCAACCAGGGCTTACGCTGGTGGGCGATGCCGCGCACACCATTATCCGTTG GCGGGCAGGGGTTAACCTCGGTTATCGCGATGTTGATGCGCTGATTGACGT TCTGGTGAATGCCCCGAGCTACGGTGAAGCGTGGGCCAGTTATCCTGTCTCA AGCGTTACCAGATGCGGCGTATGGCGGATAACTTCATTATGCAAAGCGGTAT GGATTGTTTTATGCCGGATTCAGCAATAATCTGCCACCCTGCGTTTTATGCG TAATCTCGGATTAATGGCGGCGGAGCGTGCTGGCGTGTGAAACGTCAGGCC CTGAAATATGCGTTAGGGTTGTAG ATGGCAATACAACACCCTGACATCCAGCCTGCTGTTAACCATAGCGTTCAGG TGGCGATCGTGGTGCCGGTCCGGTCCGGCTGATGATGGCGAACTATCTCGG CCAGATGGGCATTGACGTGCTGGTGGTGGAGAACTCGATAAGTTGATCGAC</p>
<p><i>Klebsiella oxytoca</i> 2880STDY568</p>	<p>WP_001577817.1</p>	<p>ATGGCAATACAACACCCTGACATCCAGCCTGCTGTTAACCATAGCGTTCAGG TGGCGATCGTGGTGCCGGTCCGGTCCGGCTGATGATGGCGAACTATCTCGG CCAGATGGGCATTGACGTGCTGGTGGTGGAGAACTCGATAAGTTGATCGAC</p>

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onal 3-(3-  
hydroxy-  
phenyl)propi  
onate/3-  
hydroxycinna  
mic acid  
hydroxylase

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*Klebsiella*  
*oxytoca*  
2880STDY568  
2666\_3-  
hydroxybenz  
oate 6-  
monooxygena  
se

WP\_032728552.1

*Klebsiella*  
*oxytoca* BRL6-  
2\_4-  
hydroxybenz  
oate 3-  
monooxygena  
se

WP\_024273698.1

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CGGCTGCGATGGCTTTCACGGCGTGTCCCGTCAAAGTATTCTCCTGGGGTGC  
TGAAAGAGTACGAAAGCGTCTGGCCGTTTGGCTGGCTGGGGCTGCTGGCCGA  
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GTTCTTTGTAGTCAGCGCTCACTAACTCGAAGCCGTTATTACCTCCAGGTCCC  
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*Citrobacter*  
*sedlakii* NBRC  
105722\_2-  
octaprenyl-6-  
methoxyphen  
yl  
hydroxylase

WP\_042292169.1

*Citrobacter  
sedlakii* NBRC  
105722\_2-  
octaprenyl-3-  
methyl-6-  
methoxy-1,4-  
benzoquinol  
hydroxylase

WP\_042285453.1

AACGTCGGGTTAATGGCTATGGAGCTGTTTCCACCCGCCCGTGACGCTCTGGT  
GAAGCGCACGCTTGGCTGGGTTGCGCGATAA  
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TGCGTAATCTGCTGTGGAAAGGCAAATCGCAGGCGGAATTTTATCGCGGCAT  
GGAGTGGTTATACGGCTGGAAAGAGGATAACTGTCTGCTGCCGCGCTAG

*Citrobacter  
sedlakii* NBRC  
105722\_3-  
hydroxybenz  
oate 6-  
monooxygena  
se

WP\_042284472.1

<p><i>Alcaligenes sp.</i> Strain HPC1271 _aromatic- ring- hydroxylating dioxygenase subunit beta</p>	<p>WP_009463815.1</p>	<p>ATGAGCGAAATCAGCCGTCAGGCCCTTATCGATTTTGTGTATGCGGAAGCCC GCCTGCTGGATGAGCAGCGCTACGAGCAATGGCTCGACCTGTTACCCGAGGA TGGCTACTACTGGATGCCGCTGGTGCACGATCAGCAGGACGCCCGTTGCAC GCGTCCCTGATGCACGAGGACAAGCTCTTGCTGCGTATCCGTGTGGAACGTCT GGCAGGCCGCCGTACCTTTTCCAGCAGCCCAAGAGCCGCAGCCATCACCTG CTCCAGCAACCCACTGTGGAAAGCATGGACGAGGAAAAAGGCCAATACACC GTGCGTTGCGCCTTCCATTACACGGAAACGCGTGGCGACAACCAAGATATCT ATGTGGGTTGGAATACCTACACCCTGGTGCGCCAGGACGACGCCTTGAAGAT CCGTCTGAAACGGGTGGATCTGCTCAATTGCGACGCGCCTTTCGGCAATATCC AGTTGTTTATGTGA ATGACATCCAAGGTTTATATCGGCCTGCAAGACAACGACATGTCGCGCTACA TCGTCGAGGCCATCGAGGAAGACAACCCGACGCGACGGTGATTTATCAGCC AGCCATGATCCGTATTGAGTGTGTGGGTCAGTTGACCGTCAAGCGCGAGACC GTGGAAGAAAAGTATGGGCAGCAATGGGACATGCAGGAGCTGCATCTGAAT CTGATCACCTTGGGTGGCAACGTGGATGAAGATGAGGACCGCTTGTTTTGCA CTGGAACAGCTAA</p>
<p><i>Alcaligenes sp.</i> Strain HPC1271_mo nooxygenase</p>	<p>WP_003800437.1</p>	<p>ATGTCAGACACGAGCTTGCCCACTTTGACAAAGTACGTACGCGTACGCAGTC CGGCCAACGCCCGTTTTGTGGAATTTGATTTTGCCATCGGCCAGCCGGACCTG TTTGTGGAAGTGGTCATGCCCCCGCGCGCTTTGAGCAGTTTTGTCTGCAAAA CAATGTCCAGCCCATGAGCGATGAGCAAATGCGGGTCAATGACGAGAACGA AGAGAAATGGCGTTTTGGTTATGACACTTTGGTTCGGCAACAGTTCGTCAGGCTG AAGCCCAGTAG ATGGCAGTTACTGCAATTACCCCGGACTACCAAGGCTATGCGCGCGACCAGC AGTCGCACTATGGCGACAGCATGTTGCTGTATATAGGCTGGGACGAGCACCT TTTGTCTGCTCGGCCAAGGCCTTTCTGGTTCAGCCCGAGATGACCGTGTCCC AGATGATCCAGCAACTGTTGCCCTGCCGTTTTTGGCCAGCACCTGACTTCGAG CAGATCGAATGGAGCAAGGTGCAATGGAATTTGAACGGCGAGGCGGTTGAA CTGAATCCTGACGACACCTTGGGTTCCAAGGGCTTCGATCACAATCCTTGT GCGTTTCAAGACCCCTGGCCTGAATGGCTACAAGGGTACGGGCGTATAA ATGTTTGGCGATTTTGTCTTTCGCTCACCTATGCAGCTTATGCCCCGTTCTCGT CAGGGGCCGCTTTGGTTTGGGATGCGGATTTCTGCGGTGGGGCGTGGTTAGT TCATAGGCACAGCGATGTACAAGCCGCCCTGCGCAATCCCGATTTAAGTGTG CGGCGTGTGGCGGGTGGGTGGCCAGGCTGGCGCTGACCCCTCCCCTCGTCT GCAAGCCTTCAAGGGCTTGATGGCCAGGACGCTGGTCTTTCTGGACCGCCCC CGTCACACACGCGTCAGACGAGTAGTGAACGCGGCTTTTAGTGCGAAGGCTT TGGAACCCTGCACAGCACGATTGAAAAGCGCGTTACGCATATCCACCTGGA GTTAAAAGACAAGCTGCGGCACGGTTCAGTGGATCTGGTTGATGAGGTCGT CGCCCACTCCCCGCTTGGTCATGATGGACGTGCTCGGCTTGGAGCAACTGCC</p>
<p><i>Alcaligenes sp.</i> Strain HPC1271_phe nol 2- monooxygense</p>	<p>WP_009454452.1</p>	<p>ATGTCAGACACGAGCTTGCCCACTTTGACAAAGTACGTACGCGTACGCAGTC CGGCCAACGCCCGTTTTGTGGAATTTGATTTTGCCATCGGCCAGCCGGACCTG TTTGTGGAAGTGGTCATGCCCCCGCGCGCTTTGAGCAGTTTTGTCTGCAAAA CAATGTCCAGCCCATGAGCGATGAGCAAATGCGGGTCAATGACGAGAACGA AGAGAAATGGCGTTTTGGTTATGACACTTTGGTTCGGCAACAGTTCGTCAGGCTG AAGCCCAGTAG ATGGCAGTTACTGCAATTACCCCGGACTACCAAGGCTATGCGCGCGACCAGC AGTCGCACTATGGCGACAGCATGTTGCTGTATATAGGCTGGGACGAGCACCT TTTGTCTGCTCGGCCAAGGCCTTTCTGGTTCAGCCCGAGATGACCGTGTCCC AGATGATCCAGCAACTGTTGCCCTGCCGTTTTTGGCCAGCACCTGACTTCGAG CAGATCGAATGGAGCAAGGTGCAATGGAATTTGAACGGCGAGGCGGTTGAA CTGAATCCTGACGACACCTTGGGTTCCAAGGGCTTCGATCACAATCCTTGT GCGTTTCAAGACCCCTGGCCTGAATGGCTACAAGGGTACGGGCGTATAA ATGTTTGGCGATTTTGTCTTTCGCTCACCTATGCAGCTTATGCCCCGTTCTCGT CAGGGGCCGCTTTGGTTTGGGATGCGGATTTCTGCGGTGGGGCGTGGTTAGT TCATAGGCACAGCGATGTACAAGCCGCCCTGCGCAATCCCGATTTAAGTGTG CGGCGTGTGGCGGGTGGGTGGCCAGGCTGGCGCTGACCCCTCCCCTCGTCT GCAAGCCTTCAAGGGCTTGATGGCCAGGACGCTGGTCTTTCTGGACCGCCCC CGTCACACACGCGTCAGACGAGTAGTGAACGCGGCTTTTAGTGCGAAGGCTT TGGAACCCTGCACAGCACGATTGAAAAGCGCGTTACGCATATCCACCTGGA GTTAAAAGACAAGCTGCGGCACGGTTCAGTGGATCTGGTTGATGAGGTCGT CGCCCACTCCCCGCTTGGTCATGATGGACGTGCTCGGCTTGGAGCAACTGCC</p>
<p><i>Alcaligenes sp.</i> Strain HPC1271_phe nol hydroxylase</p>	<p>WP_009454448.1</p>	<p>ATGTCAGACACGAGCTTGCCCACTTTGACAAAGTACGTACGCGTACGCAGTC CGGCCAACGCCCGTTTTGTGGAATTTGATTTTGCCATCGGCCAGCCGGACCTG TTTGTGGAAGTGGTCATGCCCCCGCGCGCTTTGAGCAGTTTTGTCTGCAAAA CAATGTCCAGCCCATGAGCGATGAGCAAATGCGGGTCAATGACGAGAACGA AGAGAAATGGCGTTTTGGTTATGACACTTTGGTTCGGCAACAGTTCGTCAGGCTG AAGCCCAGTAG ATGGCAGTTACTGCAATTACCCCGGACTACCAAGGCTATGCGCGCGACCAGC AGTCGCACTATGGCGACAGCATGTTGCTGTATATAGGCTGGGACGAGCACCT TTTGTCTGCTCGGCCAAGGCCTTTCTGGTTCAGCCCGAGATGACCGTGTCCC AGATGATCCAGCAACTGTTGCCCTGCCGTTTTTGGCCAGCACCTGACTTCGAG CAGATCGAATGGAGCAAGGTGCAATGGAATTTGAACGGCGAGGCGGTTGAA CTGAATCCTGACGACACCTTGGGTTCCAAGGGCTTCGATCACAATCCTTGT GCGTTTCAAGACCCCTGGCCTGAATGGCTACAAGGGTACGGGCGTATAA ATGTTTGGCGATTTTGTCTTTCGCTCACCTATGCAGCTTATGCCCCGTTCTCGT CAGGGGCCGCTTTGGTTTGGGATGCGGATTTCTGCGGTGGGGCGTGGTTAGT TCATAGGCACAGCGATGTACAAGCCGCCCTGCGCAATCCCGATTTAAGTGTG CGGCGTGTGGCGGGTGGGTGGCCAGGCTGGCGCTGACCCCTCCCCTCGTCT GCAAGCCTTCAAGGGCTTGATGGCCAGGACGCTGGTCTTTCTGGACCGCCCC CGTCACACACGCGTCAGACGAGTAGTGAACGCGGCTTTTAGTGCGAAGGCTT TGGAACCCTGCACAGCACGATTGAAAAGCGCGTTACGCATATCCACCTGGA GTTAAAAGACAAGCTGCGGCACGGTTCAGTGGATCTGGTTGATGAGGTCGT CGCCCACTCCCCGCTTGGTCATGATGGACGTGCTCGGCTTGGAGCAACTGCC</p>
<p><i>Alcaligenes sp.</i> Strain HPC1271_cyt ochrome P450</p>	<p>WP_009457883.1</p>	<p>TTTGTCTGTTTTTTCAGGCCTGGTCAGAGCAACTGGCAGGTTTTATCGGACAAG CCACGCCGGATATGCCTTTGCTGAGAAGCACCCAGGACGCCTTGCTGGATAT GGCTGACTTTCTAGGCAACAGTGCCAATCTGCAAGAAGGCGGGTTGGCCTGG CGGCTTTTGCATGATGAGCAGCTTTCCCTTCAAGGCAGGCGCGAGCGCTTGGC ACAGTCCTGTATGTTGCTCTTTGCAAGGCTATGAAACCTCCCAGCATCTGCTTA GCTCCCTGTTTCAAACACTGCTGGCCGACTCCACAACGCTGCAGGACTTGCTA TCGAATCCGCAGAAAATTCTGCCGCGATCAAAGAGGTGCTGCGTCACGACA GCCCGATTTCAGTACACAGCCAGACGTTTGTGATGCGCGACCAGCACTGGCATGG TCAGCATTGCGCAAGGGGCAGTTGCTGCTTTTGTCTTTGGGCGCTGCCAACCC GCGATCCGGCGGTGTTTGCCAATCCCGACGATCTGGACTTCAGCCGCAACAA</p>

<p><i>Alcaligenes sp.</i> II-C-7_alkane hydroxylase</p>	<p>ACJ22769.1</p>	<p>TCAGGCCGAGCTATCCATGGGACACGGAATTCATCATTGCCTGGGAGCCGGT  TTGGTGCAGCTGGAAGCCAAGCTCGTACTTGAGCAATTCCTGCCTTTGCTCCC  TCGCCTGCAAGGGACGCAAGGTGAGCGTATTCGACTCCCCGCCTATCGGGGC  TGGAAACCGCTTGCCGGTTCAATACCGGGCGGCACAGTCATGA  AATACCGGCCATGAGCTGGGGCACAAGACCGACCACCAGAGAAATGGATG  GCCAAGCTATGTCTCGCCCCGGTGTTCATATGGGCACTTCTATGTGGAACATAA  TCGTGGGCATCACGTGAGGGTTCCACGCCGAGGATCCGGCATCCTCCCGG  TTCGGCGAGACGTTCTGGGAGTTCCTGCCGCGACCGTGATCGGCAGCCTTAA  ATCCGCCTGGTCGCTGGAGAAACAGCGGCTGGAGCGGCAGGGGCTTTCGGTA  TGGAGCTGGCATAACGACAATCTGCAGGCCTGGGCATTGTCCGGTGGTGTTC  GGGTGCCCTGATACTGTGGTTGGGCTGGGCGGTGGTGCCGTTCTGTGATTC  AGAGCCTGTTCCGTTTTAGTTGCTGGAGGTGGTCAACTATATCGAGCA  ATGTCTCAATCTGATCTGGCTCACGTTGACGTGACGATCGTCGGCGGCGGTAT  GAGCGGTGCCGTGCTGGCCCTCTCCCTCGCCGCCCTCCGGGGATCGGACGGT  GCCCCCTGCAGATCCTGCTGCTGGAGGCGAGCGCCCCGAGCTCAATGCTC  ATCCCGGCTTCGACGCCCGGCCATCGCCCTCTCGGCGGGCACCTGCGAGGC  GCTGACCCGCCACGGCATCTGGTCCCGTTTACCCCCATTGCACCCCCATCA  CCGACATTCACGTCTCGGATCGGGGCCACTGCGGCCAGGCCAGGCTGTCCGC  GGCGGAATATGGCTTGCCGGCGCTCGGCCAGGTGATCGAACTCTGTCCGCC  GGGATCGAGTTGCAGCGGGCCCTTGCCGCCACCCCGCAGATACGGGTGTGCT  GCCCCGGCCAGCTCGCACAGATTGAGCCCGACGAAGAAGGGGTGACCCTGA  CCCTCGCCACGAACGAGCGCTATCGGACCCAGCTGCTGGTGGCGGGCGGATGG  GGGCAACTCCTTCGTGCGCCAGCACTTCAAGATGCCGGTCACTGCCACGAC  TATGGCCAGAGCGCCATCATCGCCACGGTGAAGACCGCCCTGGACCCCGCCG  GGCGGGCCTTCGAGCGCTTACCGAAGGGGGGCGCTCGCCCTGCTGCCGAT  GCAGGAGGGGCTCTCCTCCCTGGTCTGGAGCCTGGCGGGGACGAGGCACAG  GCCCTGATGGCGCTGGACGACGCGGCGTTTTCTCGCCCCCTGCAGCAAGCCT  TTGGCTGGCGCCTCGGGCGTTTTCGAACGCACCGGCGTGCGCCACCTCTATCCG  CTGGTGCTGACGGTGGCCGATTATCCCCTGGCCCAGCGCACCGTGCTGGTGG  GCAACGCCGCCCATCTGCTGCACCCCATCGCCGTCAGGGCTTCAACCTCGG  CATGCGCGACCTTGACCTGCTCACCCGCGCCGTTGGCCAGGCGCTCCTGGCG  GGGAGGACATCGGCAGCTTCGAGGTGCTGAACGGTACTGGCAATCGCGCC  GGGAGGATCAAGCCCAGACCGTCTGGCTCACCTCCTCCCTGGCCCAGCTCTTC  TCCAACGCCACGCTCCCCTGGTGGCGGGGCGAACCTGGCGCTCTCTGTGAT  GGGGCGACTGCCCTGGCTCAAGGCGCCGCTGGCTTCCCGCACCTGGGGTTC  GTCACCGACGTGTGCCGGCCATGA  ATGATCAAGCCGTTTCCATCAGCATTGAGAACACCCAGGACGGTTTTCGCCAT  CATCGAAGCCATCCTGGCCGACAACCCCGAGGCCAAGAGCAACCCCTTCCCG  GCCATGACCAAGATCGACTGCCCCGGTCCGGCTGGAAATCCCGCGCCGAGAGC  GTTTCCGAACGCCTGGGCCGCGACTGGGATCCTCAGGAAATCCACCTCAGCG  TCATCTCCCTGTCCGTTCCGGTGGACGAGGACGACGGCTCTCTGATCATCTTC  TGGAGATAG</p>
<p><i>Aeromonas taiwanensis</i> LMG 24683_2- octaprenyl-6- methoxyphen- yl hydroxylase</p>	<p>WP_043759840.1</p>	<p>ATGAACATCGACATCCAGGCCGAGGCGATCACGCCGCGGCGTCATACCTTCT  CGCACATCGCGCGGCGCTTCGGCGTCGACAAACCCGCTCGCGCTATGAAGA  GGCGACCTATGACGTGACGCCGGTGGTCAACTTCCACTATCGGCCGACCTAT  ACGCCGAGTTCGAACTGTACGACCCGCGCCGCACCCGCATCGTGATGAACG  ACTGGTACGTCTCCGCGATCCGCGCCAGTTCTATTACGCCTCCTACAACATC  AGCCGCGCGGCCATGCAGCAGTCGTTCCGACGACGCCGTGAACCTCGTCGAGA</p>
<p><i>Brevundimona s diminuta</i> 3F5N_monoo- xygenase</p>	<p>WP_087139175.1</p>	<p>ATGAACATCGACATCCAGGCCGAGGCGATCACGCCGCGGCGTCATACCTTCT  CGCACATCGCGCGGCGCTTCGGCGTCGACAAACCCGCTCGCGCTATGAAGA  GGCGACCTATGACGTGACGCCGGTGGTCAACTTCCACTATCGGCCGACCTAT  ACGCCGAGTTCGAACTGTACGACCCGCGCCGCACCCGCATCGTGATGAACG  ACTGGTACGTCTCCGCGATCCGCGCCAGTTCTATTACGCCTCCTACAACATC  AGCCGCGCGGCCATGCAGCAGTCGTTCCGACGACGCCGTGAACCTCGTCGAGA</p>
<p><i>Brevundimona s diminuta</i> BZC3_phenol hydroxylase</p>	<p>WP_088410947.1</p>	<p>ATGAACATCGACATCCAGGCCGAGGCGATCACGCCGCGGCGTCATACCTTCT  CGCACATCGCGCGGCGCTTCGGCGTCGACAAACCCGCTCGCGCTATGAAGA  GGCGACCTATGACGTGACGCCGGTGGTCAACTTCCACTATCGGCCGACCTAT  ACGCCGAGTTCGAACTGTACGACCCGCGCCGCACCCGCATCGTGATGAACG  ACTGGTACGTCTCCGCGATCCGCGCCAGTTCTATTACGCCTCCTACAACATC  AGCCGCGCGGCCATGCAGCAGTCGTTCCGACGACGCCGTGAACCTCGTCGAGA</p>

*Brevundimona  
s diminuta*  
NCTC11164\_3  
-(3-  
hydroxyphen  
yl)propionate  
hydroxylase

SPU44975.1

AACGCGACCTGATCGCCGATGTGCCCCGGAATGGCGCGACCTGTTGCCCGC  
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CTGGCCCCGCTCTCGCCGATTGGCGCCAGCCGTTACGGCGCGGTTCGGAATTT  
CTAA

*Serratia  
marcescens  
subsp.  
marcescens*  
Db11\_2-  
octaprenyl-3-  
methyl-6-

WP\_025301854.1

ATGAAGACATCTCAAATCGGTATGACGCGGTGGTGGTGGGTTGGCGGCATGG  
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CCTGGGAGTGGGCGTCTGTCGTGCGTGGCGTTTCGACGCGGTGTCCCTGGGGCTG  
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methoxy-1,4-  
benzoquinol  
hydroxylase

AACAGTTCGCGCAGTGCGCCAACCTGACGCTGCTGTGCCCCGCCAGGCTGCA  
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CAACCCGCTGGCCGGGCAGGGCGTCAATCTGGGGTATCGCGATGTGGATGCC  
TTGCTGAACGTGCTGAGCGAGGGCGGGATCAGGGAGAGGACTGGAGCAGC  
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GCAGAGCGGCATGGATCTGTTTTATAACCGCCTTCAGCAATAACCTGGCGCCG  
CTGAACGTCGCGCGCAATCTGGCGCTGATGGCCGCGCAGCGGGCGGGCAAG  
CTGAAAGAACATGCGTTGAAGTACGCGTTGGGATTGTAA

*Pseudomonas  
aeruginosa*  
DSM 50071 =  
NBRC  
12689\_4-  
hydroxybenz  
oate 3-  
monooxygenase

AKO84441.1

ATGAAGACTCAAGTCGCCATCATCGGCGCCGGTCCGTCCGGCCTCCTGCTCG  
GCCAGTTGCTGCACAAGGCCGGCATCGACAACGTGATCCTCGAACGCCAGAC  
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GTCGACCTGCTGCGCGAGGCCGGCGTCGACCGGCGCATGGCGCGCGACGGG  
CTGGTCCACGAAGGCGTGGAGATCGCCTTCGCCGGGCAGCGCCGGCGCATCG  
ACCTGAAGCGCCTGAGCGGCGGCAAGACGGTGTGGTCTACGGCCAGACCG  
AGGTCACCCGCGACCTCATGGAGCCCCGGAAGCCTGCGGCGCCACTACCGT  
CTACCAGGCCCGGAGGTGCGCCTGCACGACCTGCAAGGTGAGCGCCCCTAC  
GTGACCTTCGAACGCGACGGCGAACGGCTGCGCCTGGATTGCGACTACATCG  
CCGGCTGCGATGGCTTCCACGGCATCTCGCGGCAATCGATCCCGGCGGAACG  
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CGCCCTGTGCAGCCAGCGTTCGGCGACCCGACCGCTACTACGTACAGGTG  
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CTGCTGCTGAAGGCCTACCGCGAAGGGCGCGGCGAACTGCTGGAACGCTACT  
CGGCAATCTGCCTGCGGCGGATCTGGAAGGCCGAACGCTTCTCTGGTGGAT  
GACTTCGGTGCTGCATCGCTTCCCCGACACCGACGCGTTACGCCAGCGCATCC  
AGCAGACCGAACTGGAGTACTACCTGGGCTCCGAGGCGGGCCTGGCGACCAT  
AGCCGAGAACTATGTGCGCCTGCCCTACGAGGAAATCGAGTAG

*Pseudomonas  
aeruginosa*  
DSM 50071 =  
NBRC  
12689\_cytochr  
ome P450

WP\_003091710.1

GTGCCTGATCGCAAACCTGAGACTGGGCGAGGAACTGATCTCGCCACTGCACG  
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CTAACCATCCGGGCGTCCGCCGCGACGCCCGGAGGCCGCCGAACTCTACG  
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ATGGCCGACTTCGCGATCCCGCTGACCATCGCGGTGATCTTCGAGCTGCTGGG

*Pseudomonas  
aeruginosa*  
DSM 50071 =  
NBRC  
12689\_2-  
octaprenyl-3-  
methyl-6-  
methoxy-1,4-  
benzoquinol  
hydroxylase

WP\_003120388.1

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ATCGCCCAATGA

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GCTCGACGTGGCGCCGTTCAAGCCGGAGGCGCCCTACGAGCCGCGGGTCAGC  
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GGACGAGGATGGCTTCCGCGCCGCCCTCGGCGAAGCCTTCAACAGCGCCTG  
GGCGCCATCCTGCATGCCGACCGGCGCCTGTGCATCCCGCTGCGCCAGCGGC  
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CACCATCCACCCGCTGGCCGGACAGGGCGTGAACCTGGGCTTCCCTCGATGCG  
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CCGACGAACGCGTGCTGAGTCGCTTCGAGCGGCGGCGGATGCCGACAACCT  
GGGCATGATGACGGCGATGGAAGGCTTCGAGCGCCTGTTCCAGGCCGACCGG  
TTGCCGCTGCGCTGGCTGCGCAACGCCGGCCTGCGCCTGGTTCGACGGCCATC  
ACGAGGCGAAGGCGCTGTTCTCGTCCGCCAGGCGCTTGGCCTCAGCGGCGACCT  
GCCGTCGCTGGCACGAGTCTGA

*Pseudomonas  
aeruginosa*  
DSM 50071 =  
NBRC  
12689\_alkane  
1-  
monooxygena  
se

WP\_003083349.1

ATGTTTGCCTCGCTTTCCTCTGCCTGGATGCTGCGTCTGAAAAAGTACGGCTA  
CTGGATCTGGCTGATCGCGGTGCTCGGCATCCCGCTCAGCTACTGGTGGTTCGC  
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*Rhodococcus*  
*ruber*  
OA1\_hydroxy  
lase

RQM33257.1

GATCCACAAGGACCCGCAACTGGAACAGAACGCCGGCGGCTGCTGCTGGC  
AGCGGTGTGCTATGCCGGCTTCAAGGTGCAACACGTGCGCGGCCACCATGTA  
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CCGAGACCCTGGCCTGGCAGCTCGCCGCCGTCGTCCACGGCAACGCCGACCG

*Rhodococcus*  
*ruber* YC-  
YT1\_aromatic WP\_119699570.1  
ring  
hydroxylase

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CCTGA

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*Rhodococcus*  
*ruber* YC-  
YT1\_cytochro  
me P450

WP\_102031076.1

*Rhodococcus*  
*ruber* YC-  
YT1\_methane

WP\_010596327.1

TTGAGTAGGCAGAGCCTGACCAAGGCCATGCAAAGATCACCGAGCTGACG  
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CCCCATGGAGGAGGAGAAGGACAACCGGGTCTACGGCGCGATGGACGGCGC

monooxygenase

GATCCGCGGCAACATGTTCCGCCAGGTCCAGCAGCGCTGGCTGGAGTGGCAG  
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GCCGATGGCCATCGACGCGGTGCCCAACCCGAGATCCACAACGGTCTCGCG  
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ACAACCACTGCGTGGTTCGACGCGCGATCGGCACGTTTCATCGAATACGGCAC  
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C

*Rhodococcus*  
*ruber*  
NBRC15591\_a BAV60970.1  
lkane  
hydroxylase