	L O		٥	0	B2	B1	<	
6.3	ш О	ш •	۰	•				β-alanine biosynthesis I
	• •	•	•	•	ŏ	•		1,2-dichlorobenzene degradation
	•	•		•	Ŏ	•		1,2,4-trichlorobenzene degradation
	•	•		•		•		1,2,4,5-tetrachlorobenzene degradation
	• •	0		•			•	1,3-dichlorobenzene degradation
	•	•	•	•	•	•	•	2-deoxy-D-ribose degradation I
	0 0	•		•		•		2-hydroxypenta-2,4-dienoate degradation
₿.3	9	•		•				2-O-α-mannosyl-O-glycerate degradation
	9 9	•		-			-	3-chlorotoluene degradation I
	• •	•	•	•	•	-		3-hydroxybutyryl-CoA dehydrogenase 3-phenylpropanoate and 3-(3-hydroxyphenyl)propanoate degradation to 2-hydroxypentadienoate
	-	0	8	•				3-риенуургоран оаге ано 3-(3-нуогохуриенуу дооран оаге оед адааси то 2-нуогохуреналияноаге 4-hydroxyphenylacetate degradation
			•	•				4-methylphenyl adenosylcobamide biosynthesis from adenosylcobinamide-GDP
	•	0		•	•			5'-deoxyadenosine degradation I
	0 0	0	•	•	•	•	•	5'-deoxyadenosine degradation II
b .4		•	•	•	•	•		acetate formation from acetyl-CoA (succinate)
۲.۲	0 0			0	•		0	acetoacetate degradation (to acetyl CoA)
	•			•				acrylate degradation II
	•	•	0	0	•	•		aerobactin biosynthesis
	•	0		•			•	anthranilate degradation IV (aerobic)
	• •	•	•	•				arginine deiminase test
	•	•		•		•		autoinducer Al-2 degradation
	• •	0		•			0	chlorobenzene degradation
1 5	0 0	•		•		-		cinnamate and 3-hydroxycinnamate degradation to 2-hydroxypentadienoate
	9 0	-	•	-	•	-		CMP-N-acetylneuraminate biosynthesis II (bacteria)
1	• •	•	-	•		-		conversion of succinate to propanoate curcumin degradation
		•		-				curcumin degradation D-allose degradation
1	0 0	•	=	0	•		-	D-gulosides conversion to D-glucosides
	•	•	=		ě			D-mannose degradation I
	• •	•		-	ă		F	D-threonate degradation
15	-	•	•	•	•	•	-	dTDP-β-L-rhamnose biosynthesis
113	•	•		•				Ferredoxin:NAD+ oxidoreductase
			•	•	•	•		formaldehyde assimilation III (dihydroxyacetone cycle)
	•	•	•	•		•		fructoselysine and psicoselysine degradation
		0		•	•	•		glycerol degradation II
	•	•		•		•		glycine betaine biosynthesis III (plants)
	•	•		•	•			heme degradation IV
1	•	•	•	•				heme degradation V
∥6	-			•	•	•	•	indigo biosynthesis
-11	0 0	•		0	•	L.		kojibiose degradation
-[[• •	•	•	-				L-arginine degradation XIII (reductive Stickland reaction)
	0 0	•	•	-		-		L-asparagine biosynthesis III (IRNA-dependent) L-carnitine degradation III
1			•	-	•			L-histidine degradation II
			•	-	•	÷	0	L-histidine degradation VI
1			•	•	•	•		L-idonate degradation
1.3	9 -			•	•	•		L-isoleucine biosynthesis III
		0	•		•	•		L-methionine salvage from L-homocysteine
	•	•	•	0	•	•	0	L-threonate degradation
	•	•	0	•				L-threonine degradation III (to methylglyoxal)
		•		•				lathyrine biosynthesis
	•	•		•	•	•		lipid A-core biosynthesis (E. coli K-12)
	•	•		•				lipoprotein posttranslational modification
, ,	• •	•		0	•	•	•	maltose degradation
4.2	• •	•	•	•	•	•		N-acelylneuraminate and N-acelylmannosamine degradation II
	• •	•		-		-		NAD salvage pathway I (PNC VI cycle) nitrobenzene degradation II
-	• •	•	•	•				nitrobenzene degradation II phenol degradation I (aerobic)
		-	-	•				phenol degradation i (aetrobic) phenyl adenosy(cobamide biosynthesis from adenosy(cobinamide-GDP
1			H			F		phenyl adendsylcodaniae dosynthesis indiri adendsylcodaniae-corr phenylacetate degradation I (aerobic)
	• •		0	-				phenylethylamine degradation I
				-	ŏ	6		polymyxin A biosynthesis
7.	• •	•	•	•	Ŏ			putrescine degradation II
	•	•		•	Ŏ			pyruvate fermentation to (S)-lactate
	•			•	•	•		S-methyl-5-thioadenosine degradation I
		0	0		•	•		S-methyl-L-methionine cycle
	•	•	•	•		•		salicylate biosynthesis I
	•			•	•	•		salmochelin biosynthesis
	•	•		•		•		salmochelin degradation
lb.	• •	•	•	•	•	•	0	sucrose degradation I (sucrose phosphotransferase)
٣.		•	•	•		•		sucrose degradation III (sucrose invertase)
	• •	•		•				sucrose degradation IV (sucrose phosphorylase)
1	• •	•	•	•	•	•	•	sulfoquinovose degradation I
1	•	•			•	Ļ	-	trehalose degradation III
		-	H	-	•	-	-	trehalose degradation IV trehalose degradation V
	•	•		-		÷		trenaiose degradation V vancomycin resistance I
				. 67	-			
	• -	-	-					vancomyon resistance i vanillin and vanillate degradation l