Table S1. Characteristics of G-diketone and other aromatics analyzed by chromatography

Short Name	Structure	IUPAC Name	Molecular Formula	Molecular Weight (g/mol)	Retention time in HPLC (min)	Retention time in GC (min)*
G- diketone	OH OH	1-(4-hydroxy-3- methoxyphenyl) propane-1,2- dione	C ₁₀ H ₁₀ O ₄	194.180	4.20	14.0
GP-1	ОН	2-hydroxy-1-(4- hydroxy-3- methoxyphenyl) propan-1-one	C10H12O4	196.200	2.86	15.8
GP-2	HO OH	1-hydroxy-1-(4- hydroxy-3- methoxyphenyl) propan-2-one	C10H12O4	196.200	Unknown	14.8

threo-GD	НОд ОН	threo-1-(4- hydroxy-3- methoxyphenyl) propane-1,2-diol	C ₁₀ H ₁₄ O ₄	198.220	2.08	15.4
erythro- GD	НО ОН ОН ОН	erythro-1-(4- hydroxy-3- methoxyphenyl) propane-1,2-diol	C10H14O4	198.220	Unknown	15.2
Vanillic acid	OH OH OH	4-hydroxy-3- methoxybenzoic acid	C8H8O4	168.150	2.54	14.4
Vanillin	O H OH	4-hydroxy-3- methoxybenzald ehyde	C ₈ H ₈ O ₃	152.150	2.84	12.0

* Retention time in GC is for TMS derivatized compound.



Figure S1. HPLC-UV analysis of extracellular compounds found in the media of cultures grown on glucose plus G-diketone at timepoints 0 hours and 75.5 hours and comparison to standards.



Figure S2. Mass spectra of extracellular compounds identified in the media of cultures grown on glucose plus G-diketone at timepoint 75.5 hours analyzed via GC-MS and comparison to GP-1 and threo-GD standards. We have also included the published spectrum for GP-2 from Mitchell (2014) (1) for reference to our proposed identification of GP-2.

Table S2. Quantification of extracellular vanillic acid and vanillin in cultures grown with the

indicated aromatic substrates

Substrates*	Vanillic acid (µM)**	Vanillin (µM)**
G-diketone plus glucose	21 ± 1	None detected
GP-1 plus glucose	28 ± 4	3 ± 0
Glucose only	None detected	None detected

* Substrates were normalized to having 0.5 gCOD/L of the aromatic compound plus 0.5 g COD/L of glucose.

** Reported concentrations are average and standard deviations of six separate cultures after overnight incubation.

Table S3. Quantification of vanillin and vanillic acid in G-diketone and GP-1 preparations

Contaminant/Substrate ratio (HPLC peak area)	G-diketone (custom synthesized)	GP-1 (Key Organics)
Vanillin	1:77	None detected
Vanillic acid	1:1765	None detected

Dataset S1. RPKM data, differential expression testing, and gene location and annotation information from RNA-Seq analysis of cultures grown in the presence of glucose and G-type aromatics.



Figure S3. Number of genes that significantly differ in transcript abundance during growth in the presence of glucose alone compared to glucose plus each indicated aromatic substrate. Genes with a q-value < 0.01 are considered significant. Panel A displays counts of genes with increased transcript abundance compared to the glucose control, while Panel B displays counts of genes with decreased transcript abundance.



Figure S4. Genomic neighborhoods of *N. aromaticivorans* genes associated with β -O-4 linked aromatic dimer degradation. Shown are position and genes linked to transcripts with increased abundance when cells were grown in the presence of G-diketone and glucose compared to glucose alone.



Figure S5. Reaction velocity vs. substrate concentration used to calculate K_m and k_{cat} values for LigL, LigN, and LigD on GGE, G-diketone, and GD.



Figure S6. Growth of individual 12444\[]LigLNDO deletion strains on glucose and glucose plus G-diketone compared to that of the 12444\[]SacB parent strain.

Table S4. Multiple-reaction monitoring (MRM) of compounds quantified using HPLC-MS in this study.

Compound	MW (g/mol)	Parent (-) m/z	Transition 1	Transition 2	Transition 3
G-diketone	194.19	193.1	193.1 -> 136.1	193.1 -> 107.1	193.1 -> 122.1
			CE22	CE30	CE25
GP-1	196.2	195.2	195.1 -> 180.1	195.2 -> 136.0	195.2 -> 108.0
			CE15	CE22	CE25
Vanillic acid	168.15	167	167.0 -> 152.1	167.0 -> 107.9	167.0 -> 123.0
			CE19	CE19	CE14

References:

Myers, KS, Vera, JM, Lemmer, KC, Linz, AM, Landick, R, Noguera, DR, Donohue, TJ. 2020. Genome-wide identification of transcription start sites in two Alphaproteobacteria, *Rhodobacter sphaeroides* 2.4. 1 and *Novosphingobium aromaticivorans* DSM 12444. Microbiology Resource Announcements 9(36) e00880-20.