## **Supplemental Data for**

Pathway Profiling in *Mycobacterium tuberculosis*: Elucidation of a Cholesterol-Derived Catabolite and the Enzymes that Catalyze its Metabolism<sup>\*</sup>

## Suzanne T. Thomas,<sup>1,#</sup> Brian C. VanderVen,<sup>2,#,@</sup> David R. Sherman,<sup>3</sup> David G. Russell,<sup>2</sup> and Nicole S. Sampson<sup>1,@</sup>

<sup>1</sup>Department of Chemistry, Stony Brook University, Stony Brook, NY 11794; <sup>2</sup>Department of Microbiology and Immunology, Cornell University, Ithaca, NY 14853; <sup>3</sup>Seattle Biomedical Research Institute, Seattle, WA 98109

Running title: Cholesterol metabolic profile of the M. tb  $\Delta$ igr mutant

<sup>@</sup>To whom correspondence should be addressed: Brian C. VanderVen, Department of Microbiology and Immunology, Cornell University, Ithaca, NY 14853, USA, E-mail: bcv8@cornell.edu; Nicole S. Sampson, Department of Chemistry, Stony Brook University, Stony Brook, NY 11794-3400, USA, E-mail: nicole.sampson@stonybrook.edu

**Table S1.** Cholesterol-derived metabolite parent ions ( $MH^+$  or  $MNa^+$ ) identified using LC/MS/MS are listed. Data were analyzed with XCMS. The exact masses were utilized to determine molecular formulae. Unnatural <sup>13</sup>C isotopomers were used to identify cholesterol-derived compounds. Isotopomers and fragmentation were used to determine likely structures. Compound numbers correspond to the structures shown in Figure S2.

A. WT H37Rv and <i>igr</i> complement ions						
Exact Mass	Molecular Formula	Calculated Mass	Retention Time (min)	Number of <sup>13</sup> C labels	Compound	Integrated Peak Intensity
299.1638	C19H23O3	299.1647	61.9	3	<b>4</b> + O - 2H	1.3E+07
303.1950	$C_{19}H_{27}O_3$	303.1960	66.5	3	<b>5</b> + O	7.2E+06
285.1845	$C_{19}H_{25}O_2$	285.1855	69.9	3	4	3.0E+08
287.2001	$C_{19}H_{27}O_2$	287.2011	71.0	3	5	1.8E+08
B. <i>∆igr</i> ions						
Exact Mass	Molecular	Calculated	Retention	Number of	Compound	Integrated
	Formula	Mass	Time (min)	<sup>13</sup> C labels	•	Peak
						Intensity
355.1746	$C_{18}H_{27}O_7$	355.1757	57.0	3		9.7E+06
297.1691	$C_{16}H_{25}O_5$	297.1702	57.8	3	6	1.9E+08
295.1534	$C_{16}H_{23}O_5$	295.1545	58.0	3	7	1.2E+08
283.1897	$C_{16}H_{27}O_4$	283.1909	59.1	3		1.3E+08
281.1742	$C_{16}H_{25}O_{4}$	281.1753	62.8	3		6.7E+07
355.2109	$C_{19}H_{31}O_6$	355.2121	62.8	3		5.6E+07
398.1989	$C_{13}H_{30}O_7$	298.1992	62.9	3		6.1E+07
709.4133	-		62.9	3		3.0E+06
295.1899	$C_{17}H_{27}O_4$	295.1909	63.4	3		4.1E+08
277.1431	$C_{16}H_{21}O_4$	277.1440	65.8	3		2.8E+07
323.1850	$C_{18}H_{27}O_5$	323.1859	66.2	3		6.9E+07
311.1846	$C_{17}H_{27}O_5$	311.1858	67.4	3	1	2.6E+09
309.1692	$C_{17}H_{25}O_5$	309.1702	68.4	3	8	8.7E+08
279.1950	$C_{17}H_{27}O_3$	279.1960	70.4	3		2.9E+08
337.2006	$C_{19}H_{29}O_5$	337.2015	70.8	3		8.9E+07
744.5028	-		70.9	3		3.7E+06
293.1745	$C_{17}H_{25}O_4$	293.1753	71.2	3		2.3E+07
325.2004	$C_{18}H_{29}O_5$	325.2015	71.5	3	9	6.6E+07
305.2107	$C_{19}H_{29}O_3$	305.2117	71.9	3		1.5E+07
343.2265	$C_{22}H_{31}O_3$	343.2273	72.3	4	10	2.7E+07
345.2420	$C_{22}H_{33}O_3$	345.2430	72.3	4	11	1.6E+07
351.2161	$C_{20}H_{31}O_5$	351.2172	72.7	3		8.6E+06
371.2574	$C_{24}H_{35}O_{3}$	371.2586	73.6	4	12	5.4E+07
429.2991	$C_{27}H_{41}O_4$	429.3005	73.7	5	13	4.9E+06

**Figure S1**. TLC of *M. tuberculosis* extracts cultured with LDL-<sup>14</sup>C-[1,7,15,22,26]-cholesterol. 10,000 cpm were loaded per lane. TLC was resolved with 75:25 (toluene:acetone v/v) and visualized by autoradiography. **A.** *M. tuberculosis* H37Rv cell pellet. **B.** *M. tuberculosis* H37Rv culture medium. **C.** *M. tuberculosis*  $\Delta igr$  cell pellet. **D.** *M. tuberculosis*  $\Delta igr$  culture medium. **E.** *igr* complement cell pellet. **F.** *igr* complement culture medium.



Figure S2. Assigned structures of  $\Delta igr$  metabolite ions identified by LC/MS/MS.

