

strain

	1	2	3	4	5	6	7	8	9	10	11	12
A	blank	MD1855_1	3.19_1	RW_1	MD1855_1	3.19_1	RW_1	WTlac_1	lacUV5_1	MD1855_1	blank	blank
B	WTlac_1	lacUV5_1	MD1855_2	3.19_2	MD1855_2	3.19_2	RW_2	WTlac_2	lacUV5_2	3.19_1	RW_1	WTlac_2
C	RW_2	WTlac_2	lacUV5_2	MD1855_1	MD1855_1	3.19_1	RW_1	WTlac_1	lacUV5_1	lacUV5_1	MD1855_2	3.19_2
D	3.19_1	RW_1	WTlac_1	lacUV5_1	MD1855_2	3.19_2	RW_2	WTlac_2	lacUV5_2	RW_2	WTlac_2	lacUV5_2
E	MD1855_2	3.19_2	RW_2	WTlac_2	MD1855_1	3.19_1	RW_1	WTlac_1	lacUV5_1	MD1855_1	3.19_1	RW_1
F	lacUV5_2	MD1855_1	3.19_1	RW_1	MD1855_2	3.19_2	RW_2	WTlac_2	lacUV5_2	WTlac_1	lacUV5_1	MD1855_2
G	WTlac_1	lacUV5_1	MD1855_2	3.19_2	MD1855_1	3.19_1	RW_1	WTlac_1	lacUV5_1	3.19_2	RW_2	WTlac_2
H	blank	RW_2	WTlac_2	lacUV5_2	MD1855_2	3.19_2	RW_2	WTlac_2	lacUV5_2	lacUV5_2	blank	blank

Figure 1 is a heatmap illustrating the relative abundance of 12 bacterial taxa across 8 different media types (A-H). The taxa are listed on the y-axis, and the media types are listed on the x-axis. The color scale represents relative abundance, ranging from 0.0 (white) to 1.0 (black). The taxa are: *MR*, *MR*, *MR*, *MR*, *MR*, *MR*, *MR*, *MR*, *MR*, *MR*, *MR*, and *MR*. The media types are: 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, and 12.

[illegible]