

	dDDH with <i>Brachybacterium epidermidis</i> strain Q2903 (in %)	C.I. (in %)	G+C content difference (in %)
Brachybacterium muris DSM 15460	31,5	[29,1 - 34]	0,7
Brachybacterium paraconglomeratum KCTC 9916	22,7	[20.4 - 25.2]	1,8
Brachybacterium aquaticum DSM 28796	22,6	[20.3 - 25.1]	1,61
Brachybacterium massiliense MT5	22,3	[20.0 - 24.8]	0,22
Brachybacterium squillarum M-6-3	22,1	[19.9 - 24.6]	2,35
Brachybacterium faecium DSM 4810	22	[19.7 - 24.4]	1,62
Brachybacterium saurashtrense DSM23186	21,7	[19.5 - 24.2]	2,32
Brachybacterium timonense Marseille-P4339	21,6	[19.3 - 24.0]	3,14
Brachybacterium vulturis VM2412	21,5	[19.3 - 24.0]	0,41
Brachybacterium nesterenkovii CIP 104813	21,5	[19.2 - 23.9]	2
Brachybacterium ginsengisoli KCTC 29226	21,4	[19.1 - 23.8]	1,14
Brachybacterium avium VR2415	21,3	[19.1 - 23.8]	0
Brachybacterium endophyticum M1HQ-2	20,6	[18.4 - 23.0]	0,62