

Region	True Positives Count	False Positives Count	False Negatives Count	Unassigned %	Notes
v2	19	3	1	1.25	C.beijerinckii --> C. unas
v3	17	3	3	0.84	B. adolescentis, D. radiodurans, C.beijerinckii --> C. unas
v4	15.5	4.5	4.5	0.59	S. aureus --> S. unas(x2), S. epidermidis, E. facalis --> E. unass, L. gasseri, C.beijerinckii --> C. unassigned
v67	17	3	3	1.1	Bifido adolescentis, L. gasseri, N. meningitidis, A.baumannii->L johnsonii?
v8	14.5	4	5.5	0.5	Strep dysgalactiae, P. carotovorum, Enterobacteriaceae. Unas, C. butyricum
v9	2	0	18	0.13	Only positively ID'd E.coli and A.baumannii

Table 2. Results of sequencing the ATCC 20 strain mix. Number of True Positives (TP), False Positives (FP), and False Negatives (FN) identified by variable region