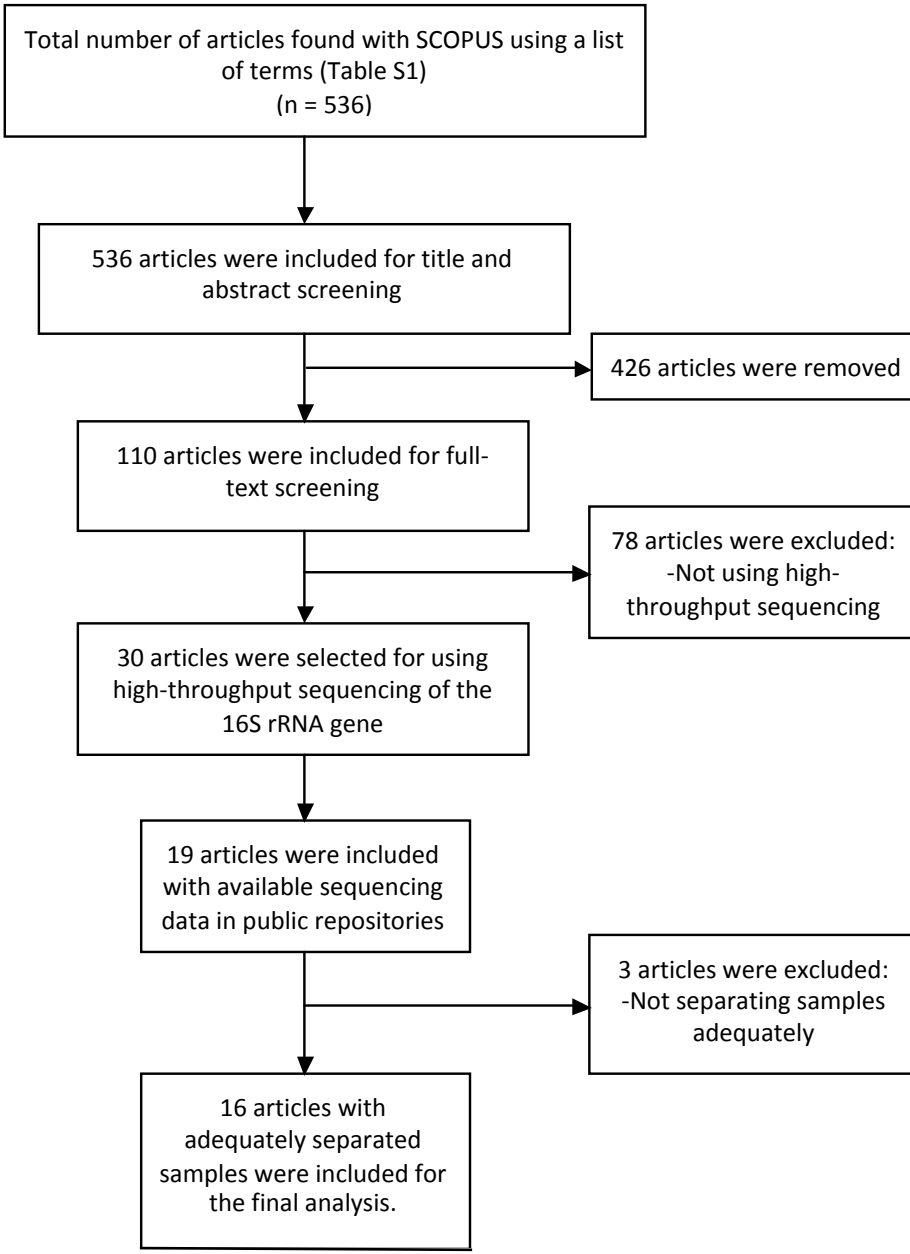
Total number of articles found with SCOPUS using a list of terms (Table S1)
(n = 536)
536 articles were included for title and abstract screening
426 articles were removed
110 articles were included for full-text screening
78 articles were excluded:
-Not using high-throughput sequencing
30 articles were selected for using high-throughput sequencing of the 16S rRNA gene
19 articles were included with available sequencing data in public repositories
3 articles were excluded:
-Not separating samples adequately
16 articles with adequately separated samples were included for the final analysis.