Vibrio coralliilyticus: Comparison of Anti-Microbial and Antibiotic Resistance with Sister Phylotypes Isolated from Puerto Rico

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Vibrio coralliilyticus ATCC BAA-450, a temperature-dependent coral pathogen first isolated from the Indian Ocean, induces bleaching in *Pocillopora damicornis* at temperatures greater than 24.5°C. Currently, we are comparing the anti-microbial and antibiotic resistance and susceptibility profiles of this V. coralliilyticus strain and possible sister phylotypes isolated from the southern coast of Puerto Rico. We hypothesize that V. corallilyticus ATCC BAA-450 and the sister phylotypes will display similar antimicrobial and antibiotic resistance. We have several lines of evidence to support this hypothesis. First, five bacterial isolates, obtained from the surface mucopolysaccharide layer of visually-diseased Pseudopterogorgia americana, showed 97-99% 16S rDNA homology to V. corallilyticus. Genomic profiling through repetitive element PCR (REP-PCR) demonstrated that one isolate had a similar profile to ATCC BAA-450, while the other four were different (but similar to one another). Second, we examined whether bacteria isolated from healthy and diseased *P. americana* colonies were able to inhibit *V.* corallilyticus grown at 24 and 27°C. We observed that twelve (12/140) isolates inhibited V. coralliilyticus at 24°C, while only five strains inhibited growth at 27°C. Third, four of the homologous strains were screened against a subset of the coral isolates that exhibited bioactivity in anti-microbial tests. Three of the homologous strains responded similarly to the ATCC strain and showed a high level of resistance to the antimicrobial compounds produced by the coral isolates. Lastly, V. corallilyticus ATCC BAA-450 and the three sister phylotypes were screened against 26 known antibiotics; they all exhibited similar resistance and susceptibility profiles (resistant to 10-14 antibiotics). To summarize, both V. corallilyticus ATCC BAA-450 and three of the Caribbean phylotypes showed similar levels of resistance in our anti-microbial (temperature-dependent) and antibiotic assays. This study will further contribute to our understanding of the pathogenicity of V. coralliilyticus and the similarity among sister phylotypes in a broader ecological context.

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