HighStump is a very typical cluster B1 phage. I don’t think you’ll find anything unusual about it. Large gaps (>100 bp) that we couldn’t find good support for including a gene were 9-10, 10-11, 20-21 (divergent), 26-27 (divergent), 54-53, 61-60, 67-66, 67-68 (divergent), 68-69, 76-77, and 99-98. Starts that were particularly tough calls were 43 and 53.