Hurricane CoverSheet

Please note that thee are three rather large gaps in the annotation.

Gap1: between features 42 and 43 (262 bp)

Gap2: between features 70 and 71 (405)

Gap3: between features 95 and 96 (427bp)

While potential ORFs exist in these regions, I have found no strong evidence supporting the addition of a gene. Large gap3 was in part created by deleting a called feature with with very poor evidence that overlapped substantially with well supported features encoded in the opposite direction.