GP126 (61193-61259) tRNA was called only by tRNAScan-SE with an Infernal score just barely above the cutoff (17.1). Called as pseudo tRNA because anticodon couldn’t be elucidated.

Gp159 (74793-74629) was added in to fill a gap and had some coding potential as shown in GeneMark-S output. Small Blast match to Pegleg gp142. Consider for possible deletion?

The following gaps were investigated, but no significant evidence of uncalled genes was found:

**Idx** **Length** **Description**

1 1773 Idx\_1\_78351\_to\_80123

2 562 Idx\_2\_38606\_to\_39167

3 289 Idx\_3\_58759\_to\_59047

4 286 Idx\_4\_58246\_to\_58531

5 281 Idx\_5\_59198\_to\_59478

6 259 Idx\_6\_525\_to\_783

7 257 Idx\_7\_56574\_to\_56830

8 244 Idx\_8\_57731\_to\_57974

9 224 Idx\_9\_1\_to\_224

10 203 Idx\_10\_64827\_to\_65029

11 199 Idx\_11\_49853\_to\_50051

12 194 Idx\_12\_48525\_to\_48718

13 188 Idx\_13\_73829\_to\_74016

14 170 Idx\_14\_57370\_to\_57539

15 166 Idx\_15\_41457\_to\_41622

16 139 Idx\_16\_59812\_to\_59950

17 132 Idx\_17\_70551\_to\_70682

18 127 Idx\_18\_1900\_to\_2026

19 114 Idx\_19\_73232\_to\_73345

20 110 Idx\_20\_59626\_to\_59735

21 108 Idx\_21\_60447\_to\_60554