10 – the choice of start codon differs from both Glimmer and GeneMark

14 – gene appears truncated based on BLAST analysis

59 – some coding potential is missed

76, 86 – not called by GeneMark, relatively short genes, small gaps

93 – not called in GeneMark

99 – no Starterator data

101 – no coding potential

102 – chose unique start codon

104 – no coding potential

107 – no coding potential

112 – no coding potential