MikeLieIn

Areas of Uncertainty

66 - Added to fill gap but covers a small blip of coding potential and has BLAST hits.

130/131 - Annotated as fusion protein based on other phage annotations. Not sure if the possible function is still tail assembly chaperone

189/190 - Very large gap but no coding potential

208 - 38bp overlap which is larger than what is normally called but it had alignments and the next start would miss some coding potential