



Figure S2. Comparison of *N. seriolae* UTF1 genome and two other *N. seriolae* draft genomes, ZJ0503 (gi: 696559939) and N-2927 (gi: 780022598). The BLASTN-based ring image was generated by BLAST Ring Image Generator (BRIG) version 0.95 [32]. The innermost two rings show GC content (black) and GC skew (purple). The second and third innermost rings represent a BLASTN comparison with draft genome of ZJ0503 (blue) and N-2927 (red). Bars indicate the position of mobile-element related genes in the *N. seriolae* UTF1 genome such as transposases (black), endonuclease DDE (blue) and integrase (green).