Figure 3. BLAST of a putative *wrbA* flavoprotein isolated from C580 *in silico* sequence from *Translate Reverse* tool (http://www. bioinformatics.org/sms2/rev\_trans.html). A: Phylogenetic tree in terms of BLAST with BLASTn alghortim from NCBI; B: Alignment with *in silico* sequence and *wrbA* flavoprotein of *E. coli* O157:H7 EC4115; C: Analysis of a transcrit of *wrbA* flavoprotein *E. coli* O157:H7 EC4115 gene with *Emsembl Bacteria* (http://bacteria.ensembl.org/ index.html); D: Genetic code.

