

No	tRNA	vB_EalM-132			vB_BcoS-136		
		Position	bases	%GC	Position	bases	%GC
1	tRNA-Asn(gtt)	33473,33545	73	49.3	54444,54518	75	49.3
2	tRNA-Asp(gtc)				54522,54596	75	49.3
3	tRNA-Ala(tgc)				54841,54912	72	45.8
4	tRNA-Gln(ttg)				54925,55001	77	46.8
5	tRNA-Trp(cca)				55006,55079	74	45.9
6	tRNA-His(gtg)				55246,55316	71	49.3
7	tRNA-Glu(ttc)				55653,55727	75	48.0
8	tRNA-Ile(gat)				55729,55803	75	37.3
9	tRNA-Tyr(gta)				55807,55891	85	49.4
10	tRNA-Leu(tag)				56065,56139	75	52.0
11	tRNA-Pro(tgg)				56144,56221	78	50.0
12	tRNA-Thr(tgt)				56223,56297	75	42.7
13	tRNA-Arg(acg)				56315,56388	74	47.3
14	tRNA-Ser(tga)				56488,56581	94	48.9
15	tRNA-Ser(gct)				56588,56679	92	58.7
16	tRNA-Arg(tct)	33244,33319	76	46.1	56681,56754	74	50.0
17	tRNA-Gly(tcc)				56758,56828	71	53.5

able showing tRNAs in phages vB_EalM-132 (2) and vB_BcoS-136 (17), their respective positions in the genome, number of bases and %GC content