

Contig	Completeness	Score	Total Proteins	Region Position	Most Common Phage	GC (%)
NODE_1_length_437545_cov_15 4.117295	incomplete	10	9	386566-391945	PHAGE_Bacill_G_NC_023719(2)	49.20%
NODE_3_length_307816_cov_12 7.929648	incomplete	30	18	47496-65362	PHAGE_Staphy_SPbeta_like_NC_029119(3)	46.89%
NODE_4_length_235316_cov_12 5.443460	incomplete	20	8	201421-209815	PHAGE_Prochl_P_SSM2_NC_006883(3)	49.51
NODE_3_length_132037_cov_18 6.274235	intact	150	47	90384-129768 info_outline	PHAGE_Lactob_LF1_NC_019486(16)	44.95%
NODE_8_length_136053_cov_13 2.574234	questionable	70	16	66279-80812	PHAGE_Entero_IME_EFm5_NC_028826(2)	44.8
<i>L. paracasei</i> T0902						
NODE_1_length_437545_cov_14 8.551080	incomplete	10	9	386566-391945	PHAGE_Bacill_G_NC_023719(2)	49.2
NODE_3_length_307816_cov_11 8.004826	incomplete	20	10	242422-251804	PHAGE_Staphy_SPbeta_like_NC_029119(3)	46.37
NODE_4_length_235316_cov_11 7.084246	incomplete	20	8	201421-209815	PHAGE_Prochl_P_SSM2_NC_006883(3)	49.51

Contig	Completeness	Score	Total Proteins	Region Position	Most Common Phage	GC (%)
NODE_8_length_136053_cov_11 9.565740	incomplete	60	15	56818-68446	PHAGE_Entero_IME_EFm5_NC_028826(2)	45.5
<i>L. paracasei</i> T1301						
NODE_1_length_437545_cov_14 4.968924	incomplete	10	9	386566-391945	PHAGE_Bacill_G_NC_023719(2)	49.2
NODE_3_length_307816_cov_12 9.130097	incomplete	20	10	242422-251804	PHAGE_Staphy_SPbeta_like_NC_029119(3))	46.37
NODE_4_length_235316_cov_12 5.059548	incomplete	20	8	25502-33896	PHAGE_Prochl_P_SSM2_NC_006883(3)	49.51
NODE_8_length_136053_cov_12 9.613623	questionable	70	16	66279-80812	PHAGE_Entero_IME_EFm5_NC_028826(2)	44.87
<i>L. paracasei</i> T1304						
NODE_1_length_437546_cov_14 5.587285	incomplete	10	9	386566-391945	PHAGE_Bacill_G_NC_023719(2)	49.2
NODE_2_length_307625_cov_12 9.244541	incomplete	20	10	242455-251804	PHAGE_Staphy_SPbeta_like_NC_029119(3))	46.37

Contig	Completeness	Score	Total Proteins	Region Position	Most Common Phage	GC (%)
NODE_3_length_235315_cov_12 6.925271	incomplete	20	8	25502-33896	PHAGE_Prochl_P_SSM2_NC_006883(3)	49.51
NODE_9_length_136053_cov_13 4.368528	questionable	70	16	66279-80812	PHAGE_Entero_IME_EFm5_NC_028826(2)	44.87
<i>L. paracasei</i> T1901						
NODE_1_length_437548_cov_14 8.094406	incomplete	10	9	386568-391947	PHAGE_Bacill_G_NC_023719(2)	49.2
NODE_2_length_307816_cov_11 4.422400	incomplete	30	18	47496-65362	PHAGE_Staphy_SPbeta_like_NC_029119(3))	46.89
NODE_3_length_235315_cov_11 8.819834	incomplete	20	8	201420-209814	PHAGE_Prochl_P_SSM2_NC_006883(3)	49.51
NODE_9_length_136053_cov_13 4.123323	incomplete	60	15	56818-68446 info_outline	PHAGE_Entero_IME_EFm5_NC_028826(2)	45.5