

**S3 Table. T6SS locus genes in *A. baumannii* strains under study.**

	DSM30011		ATCC17978			Ab242/244/825	
Protein Description	Locus	Length (AA)	Locus	Accession	Length (AA)	Locus	Length (AA)
Conserved hypothetical (Signal peptide)	11700*	229	A1S_1292	YP_001084322	230	Ab244_01	230
TssB	11695	167	A1S_1293	YP_001084323	49	<i>tssB</i>	167
			A1S_1294	YP_001084324	79		
TssC	11690	493	A1S_1295	YP_001084325	443	<i>tssC</i>	493
TssD/Hcp	11685	167	A1S_1296	YP_001084326	167	<i>hcp</i>	167
TssE	11680	158	A1S_1297	YP_001084327	158	<i>tssE</i>	158
TssF	11675	603	A1S_1298	YP_001084328	185	<i>tssF</i>	603
			A1S_1299	YP_001084329	231		
TssG	11670	332	A1S_1300	YP_001084330	298	<i>tssG</i>	332
Putative membrane protein	11665*	470	A1S_1301	YP_001084331	421	Ab244_02	470
TssM	11660	1274	A1S_1302	YP_001084332	1041	<i>tssM</i>	1274
			A1S_1303	YP_001084333	188		
TagF	11655	319	A1S_1304	YP_001084334	275	<i>tagF</i>	319
TagN	11650	255	A1S_1305	YP_001084335	209	<i>tagN</i>	255
PAAR domain containing-protein	11645	87	A1S_1306	YP_001084336	87	Ab244_03	87
TssH/ClpV domain	11640	892	A1S_1307	YP_001084337	896	<i>tssH</i>	896
TssA	11635	364	A1S_1308	YP_001084338	273	<i>tssA</i>	364
TssK	11630	454	A1S_1309	YP_001084339	388	<i>tssK</i>	454
TssL	11625	268	A1S_1310	YP_001084340	268	<i>tssL</i>	268
Hypothetical protein	11620*	200	A1S_1311	YP_001084341	85	Ab244_04	200
Putative D-Ala D-Ala Carboxypeptidase	11615*	317	A1S_1312	YP_001084342	127	Ab244_05	317

\* genes detected only in species of the genus *Acinetobacter*