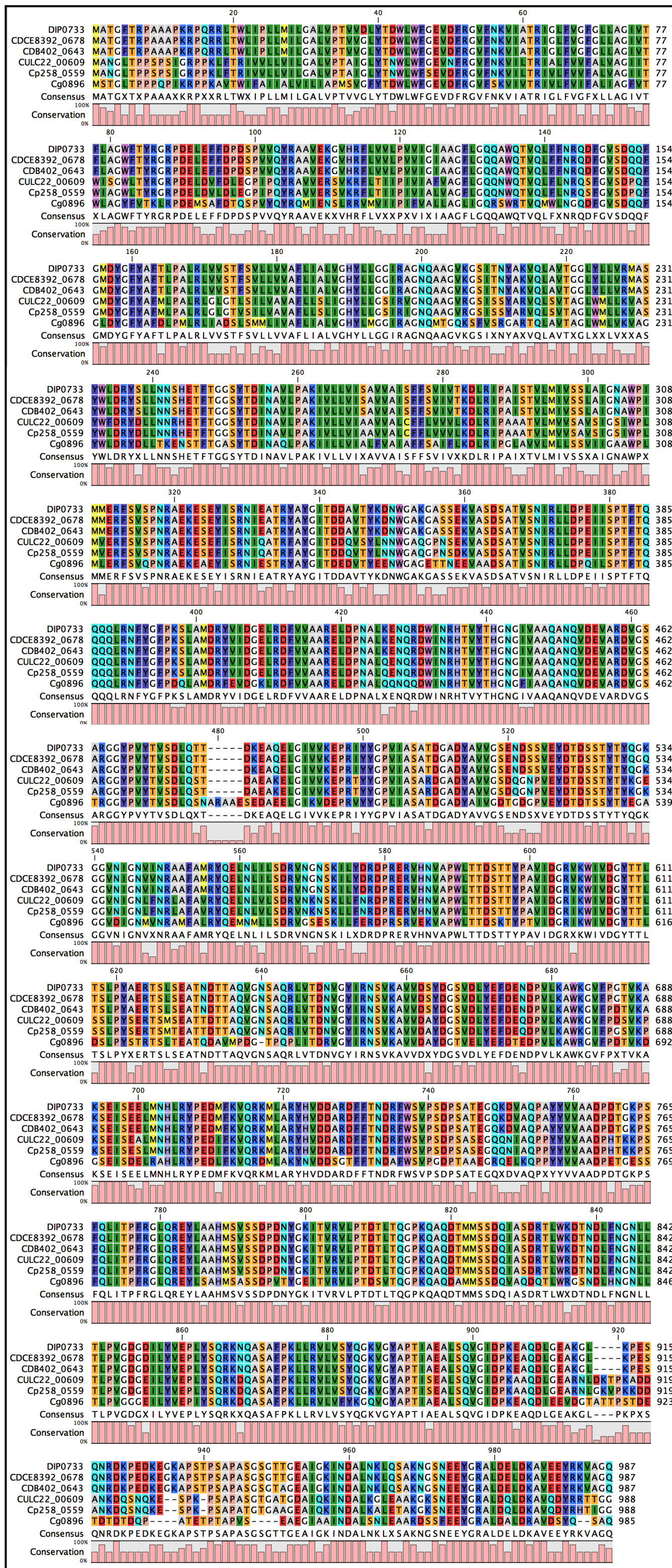


a



b

Strain	Accession	Query cover	Score	Identity	Gaps
<i>C. diphtheriae</i> CDC-E8392 <i>CDC8392_0678</i>	AEX71676.1	100%	2010 bits (5207)	984/987(99%)	0/987(0%)
<i>C. diphtheriae</i> INCA 402 <i>CDB402_0643</i>	AEX45950.1	100%	2006 bits (5198)	984/987(99%)	0/987(0%)
<i>C. ulcerans</i> BR-AD22 <i>CULC22_00609</i>	AEG83325.1	99%	1666 bits (4315)	797/988(81%)	3/988(0%)
<i>C. pseudotuberculosis</i> 258 <i>Cp258_0559</i>	AFK16201.1	99%	1665 bits (4312)	798/990(81%)	7/990(0%)
<i>C. glutamicum</i> ATCC 13032 <i>Cg0896</i>	CAF19491.1	99%	1396 bits (3614)	670/987(68%)	10/987(1%)