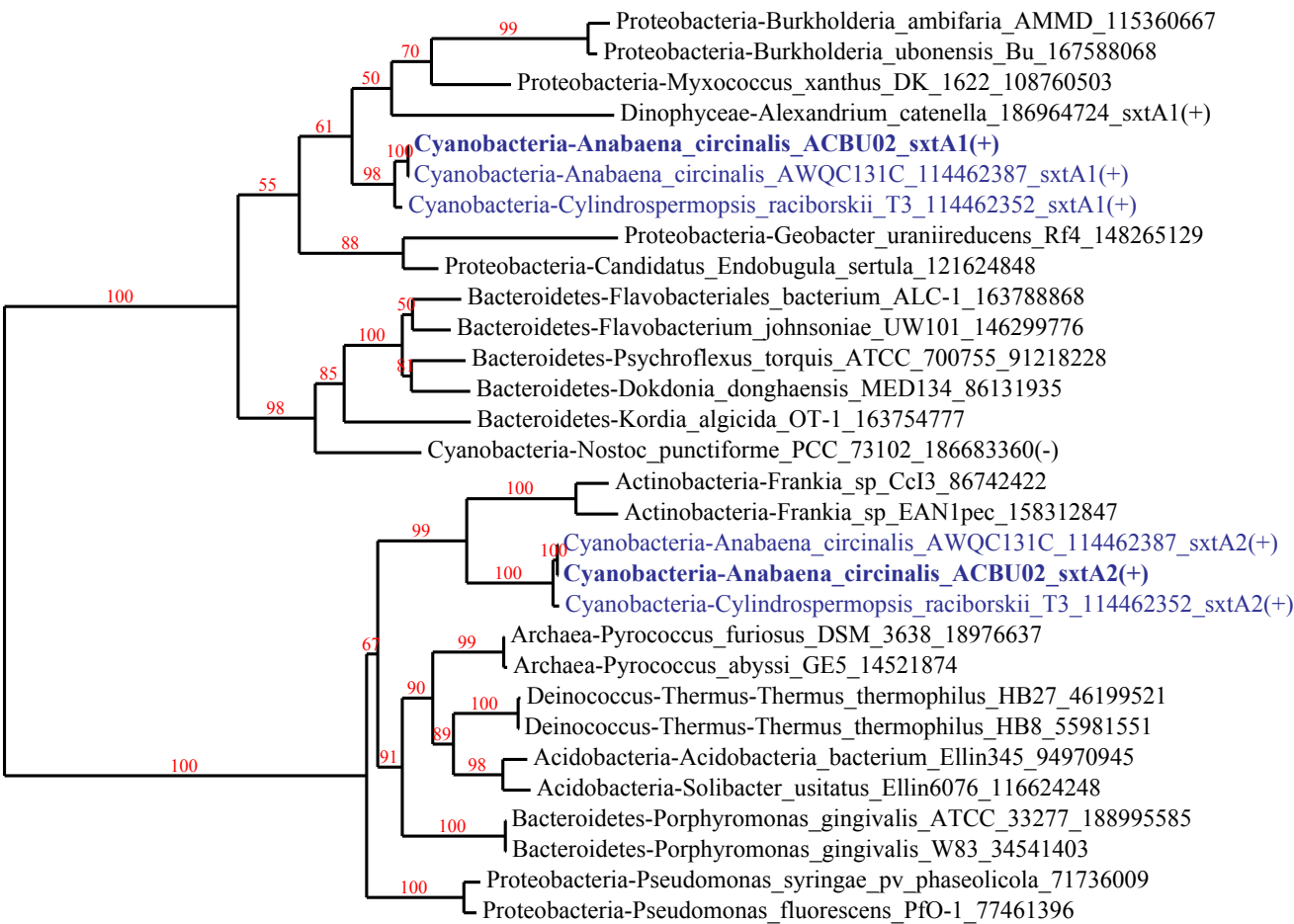
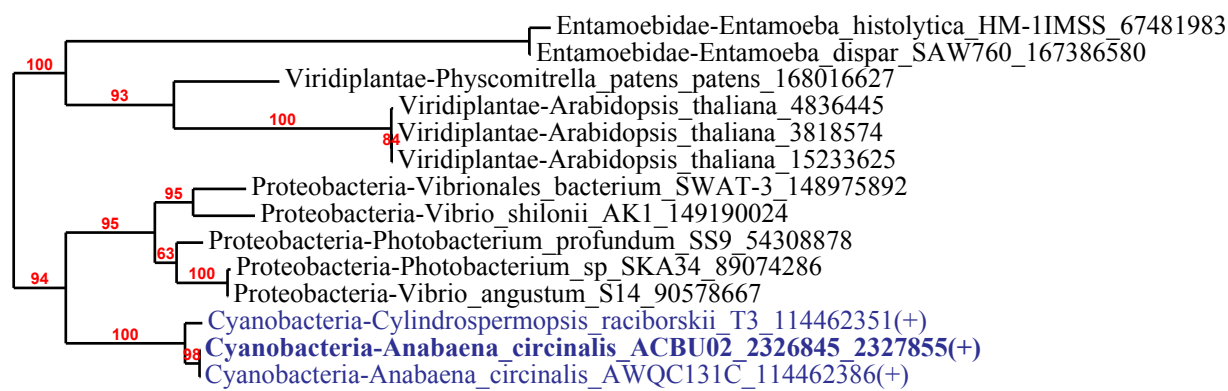


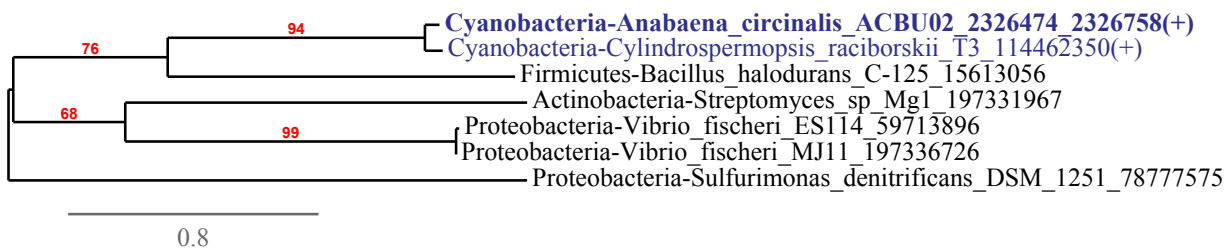
sxtA: polyketide synthase



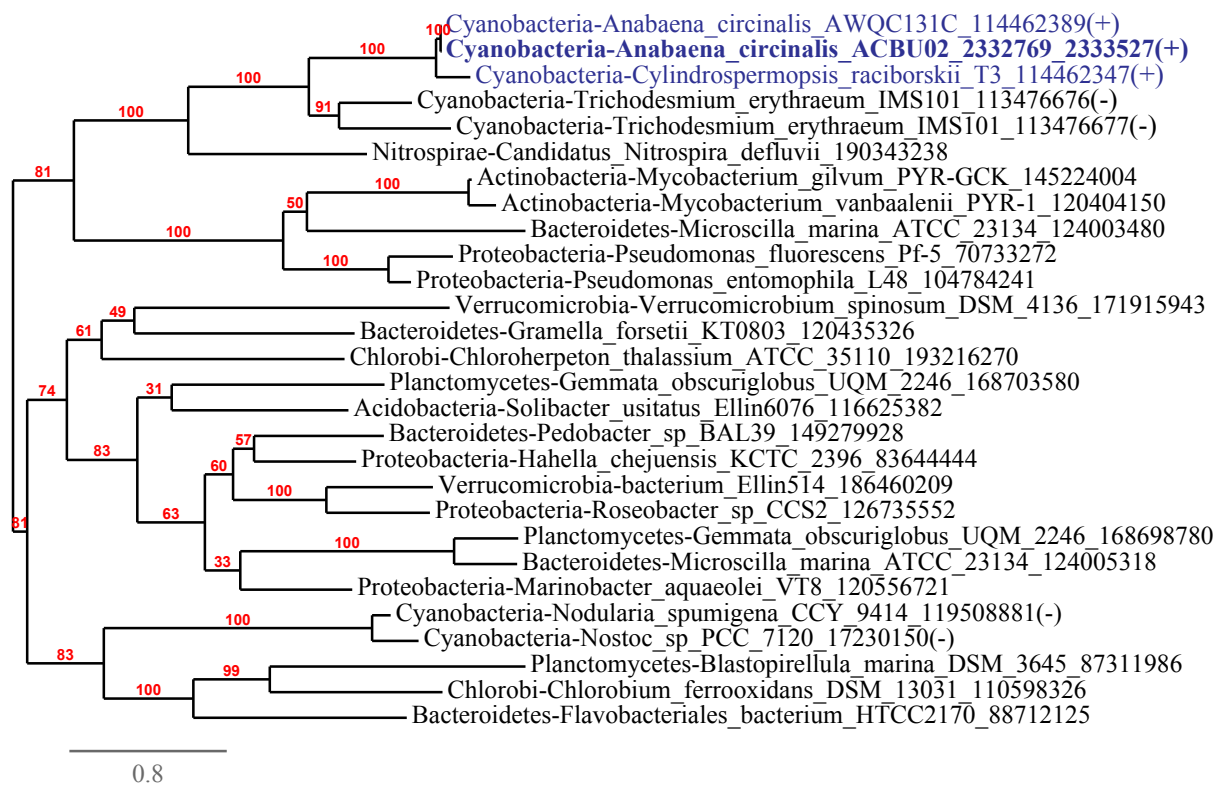
sxtB: Cytidine deaminase



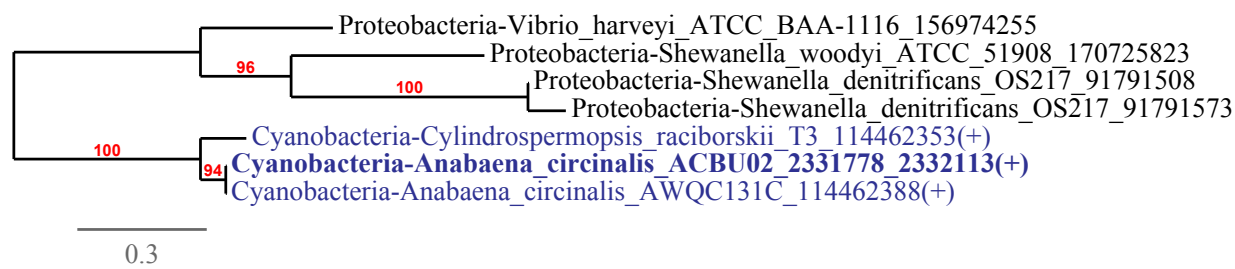
sxtC: Amidohydrolase



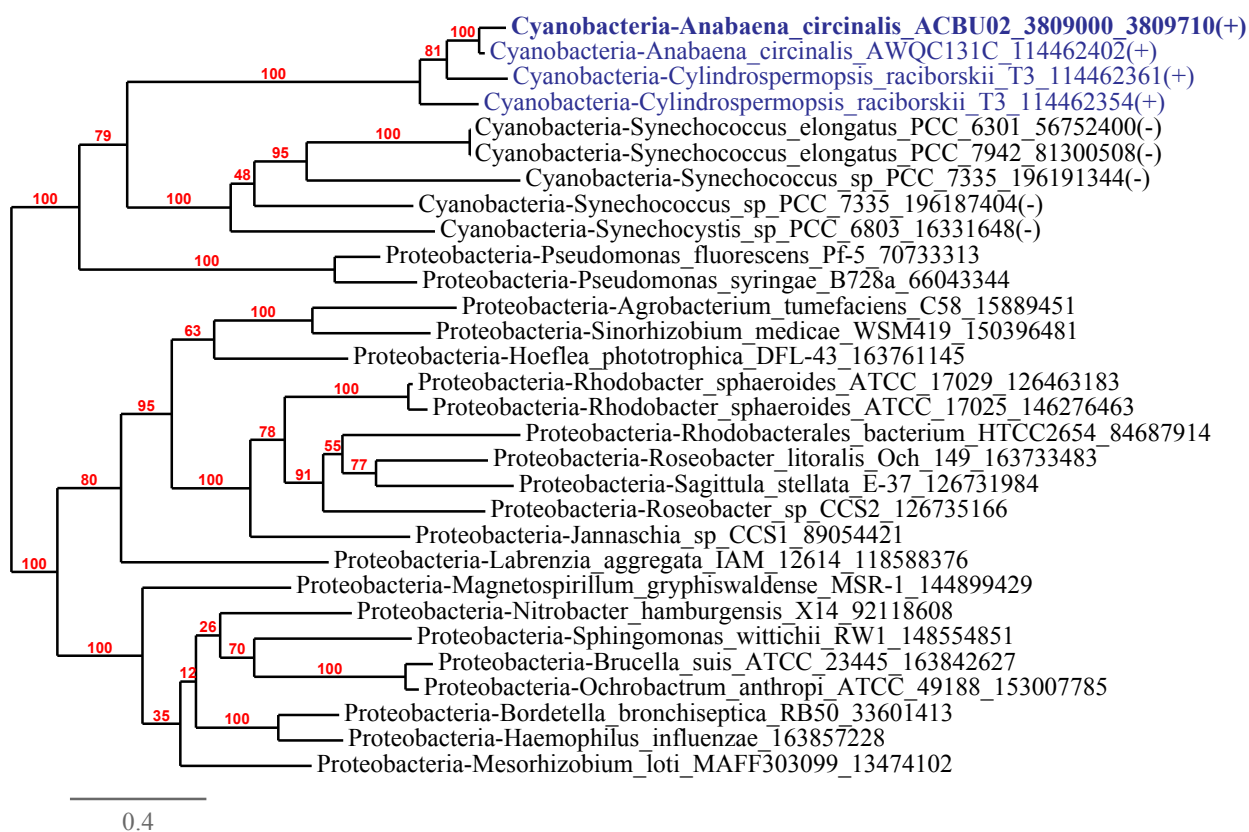
sxtD: sterole desaturase



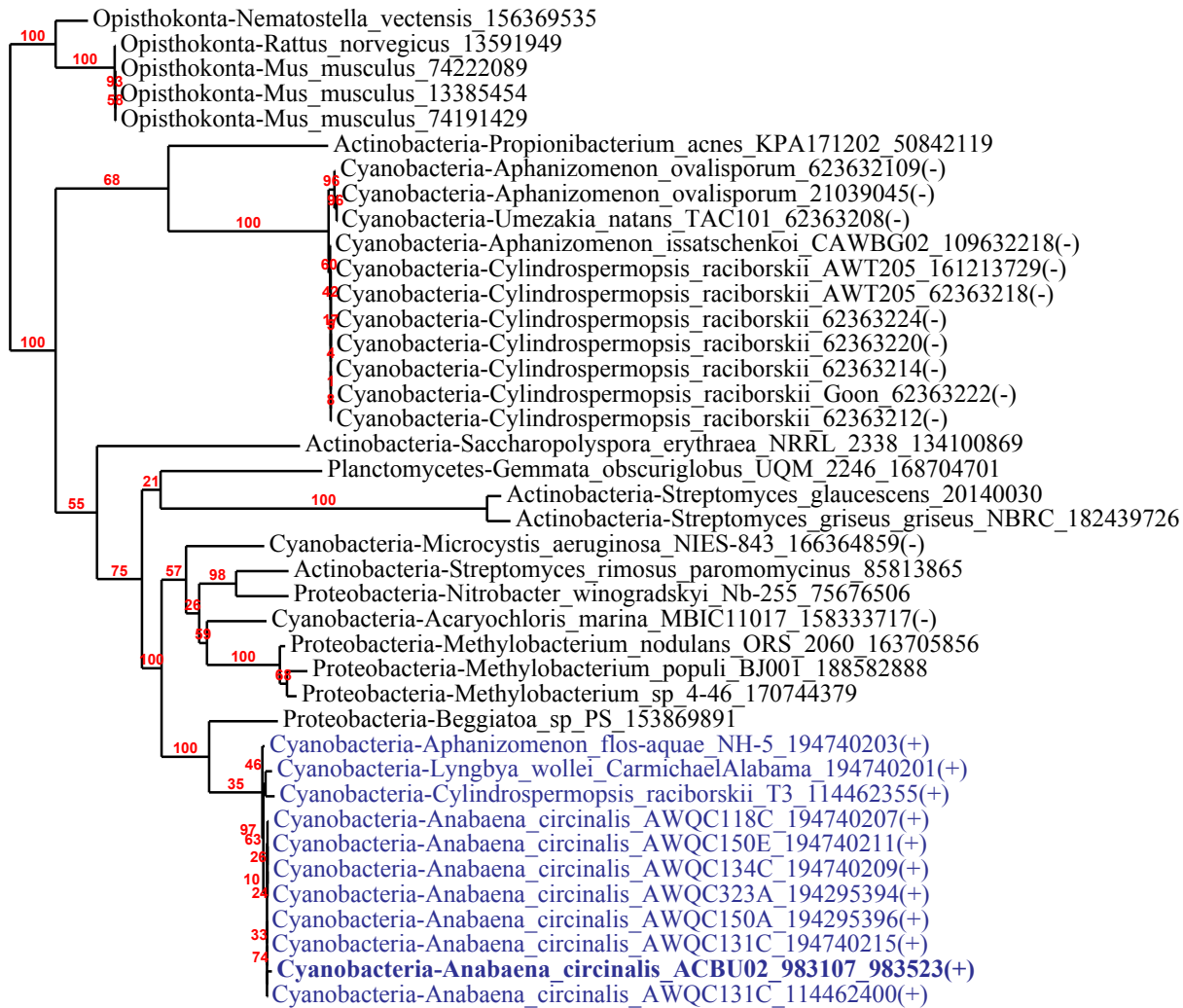
sxtE: Chaperone-like protein



sxtF, sxtM: sodium-driven multidrug and toxic compound extrusion protein

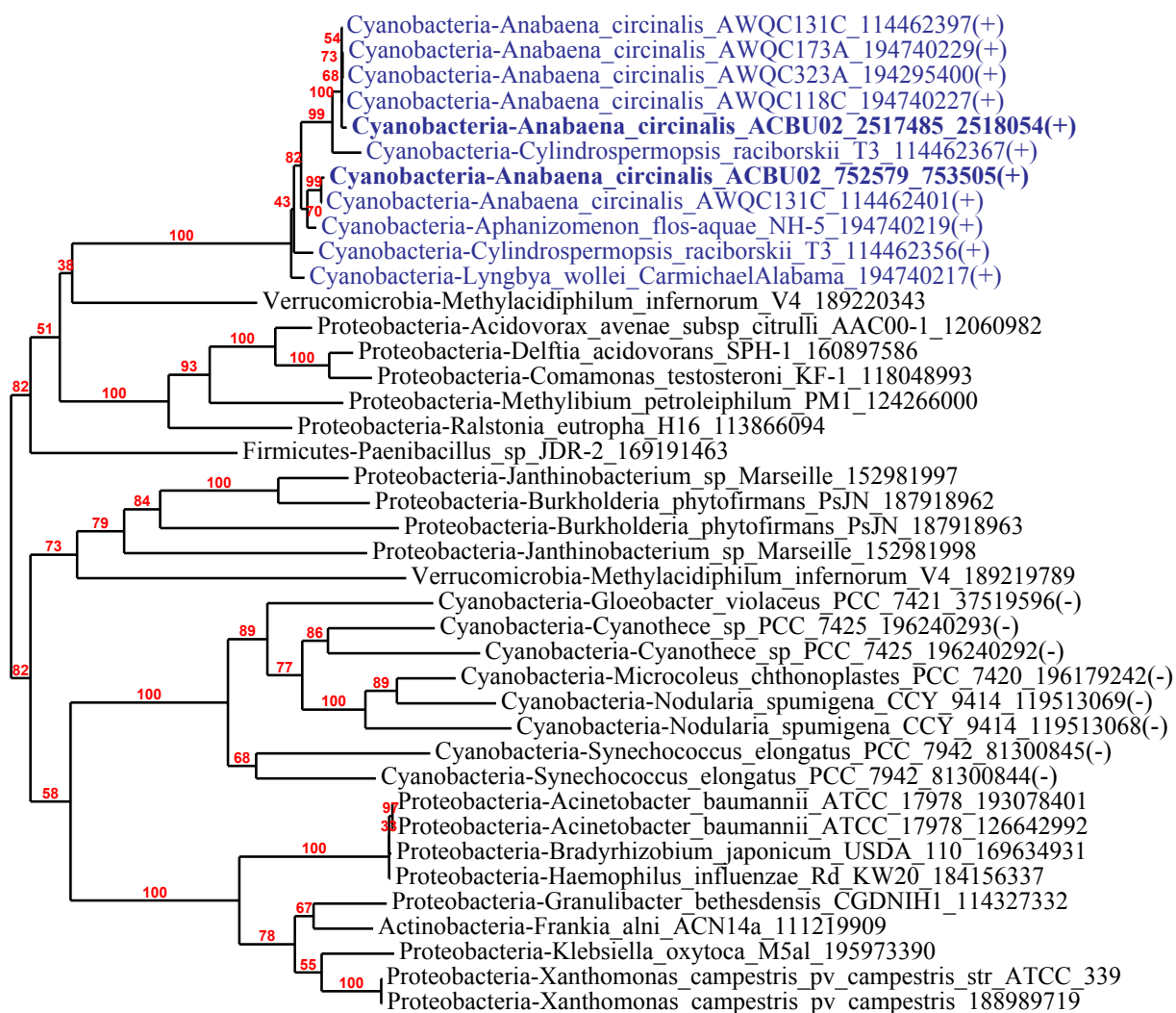


sxtG: Amidinotransferase



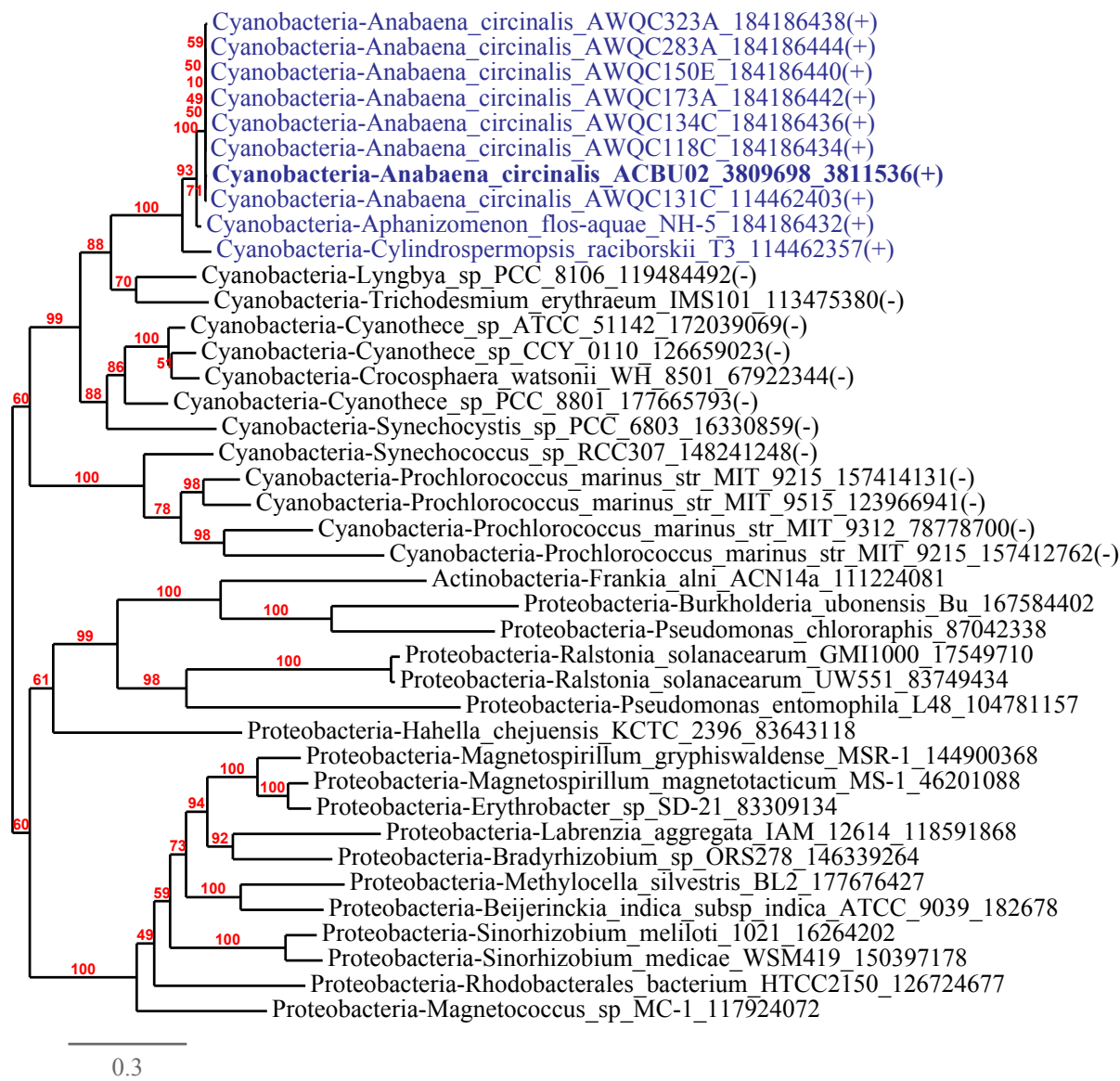
0.9

sxtH, sxtT: Phenylpropionate dioxygenase

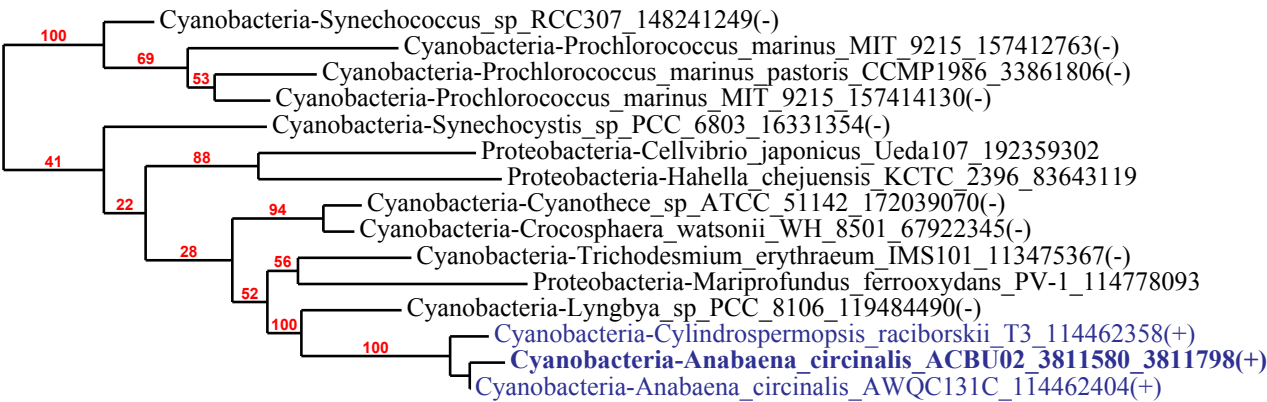


0.7

sxtl: O-carbamoyltransferase

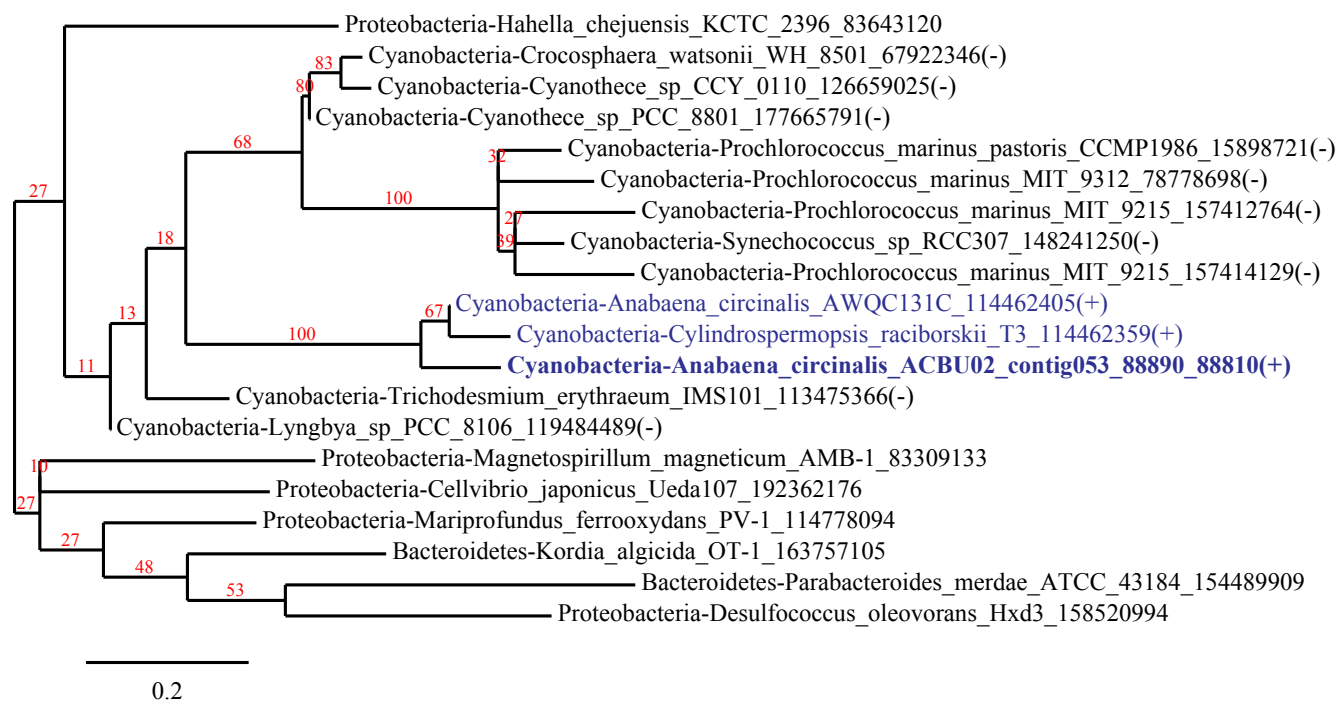


sxtJ: Hypothetical protein

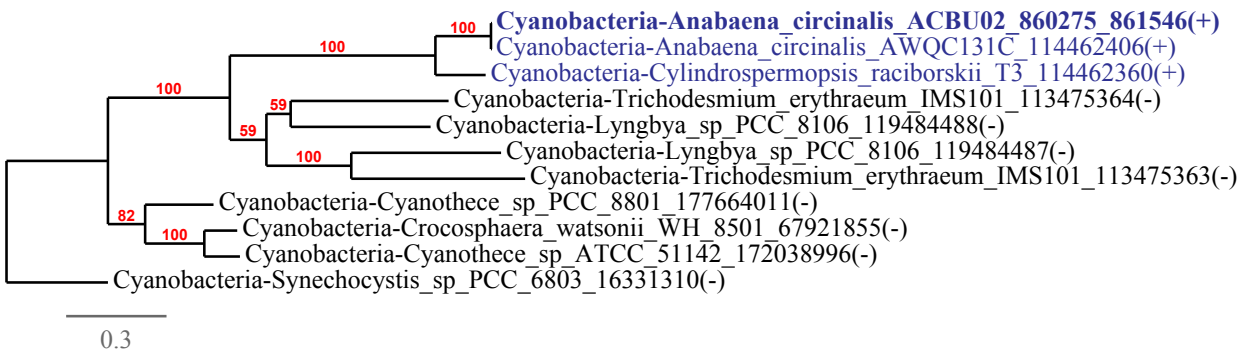


0.4

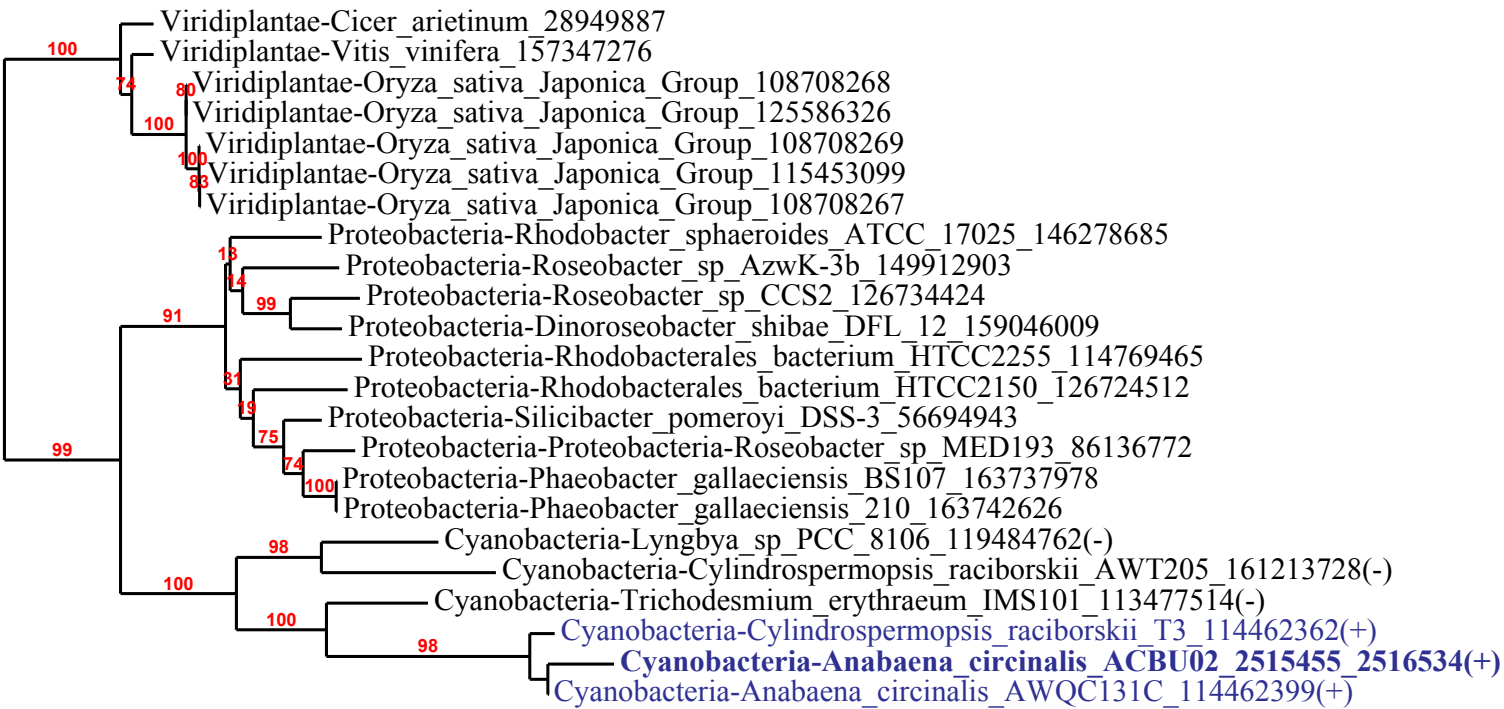
sxtK: Hypothetical protein



sxtL: GDSL-lipase

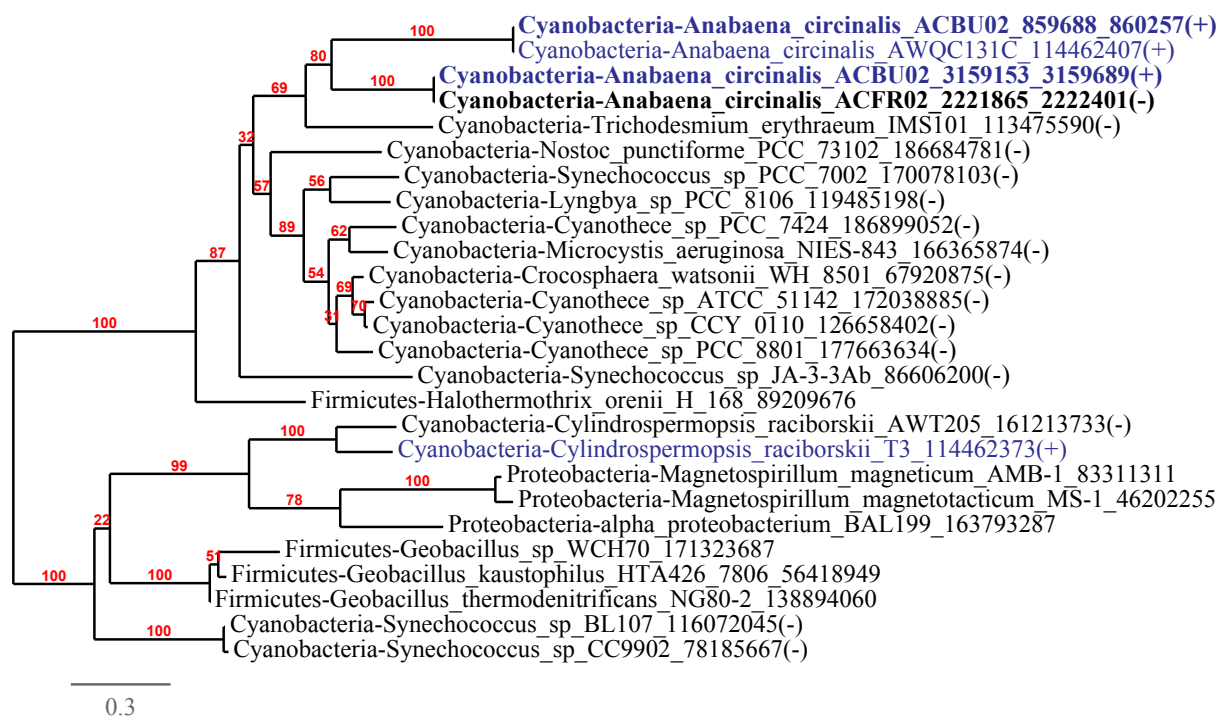


sxtN: Sulfotransferase

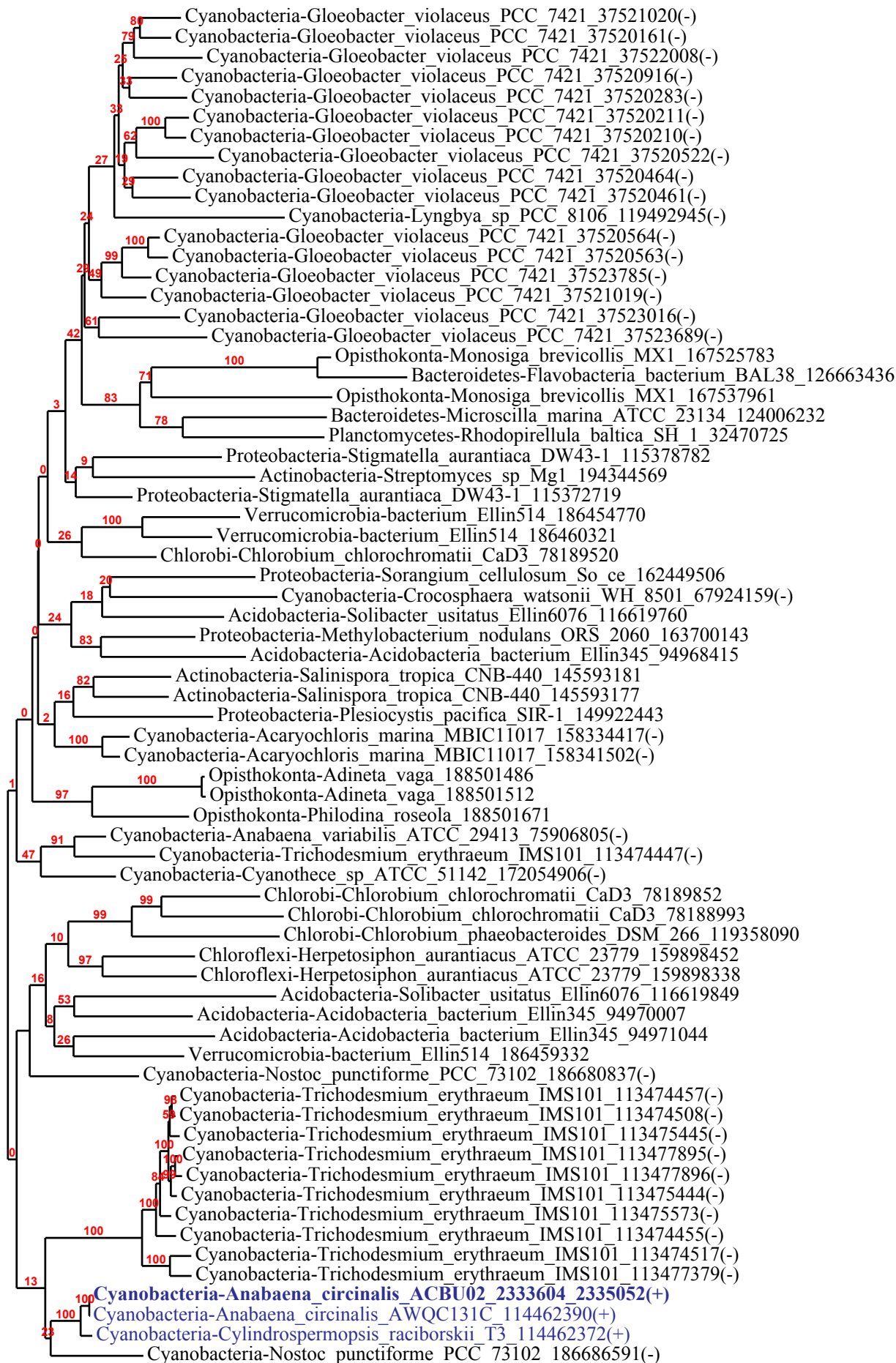


0.4

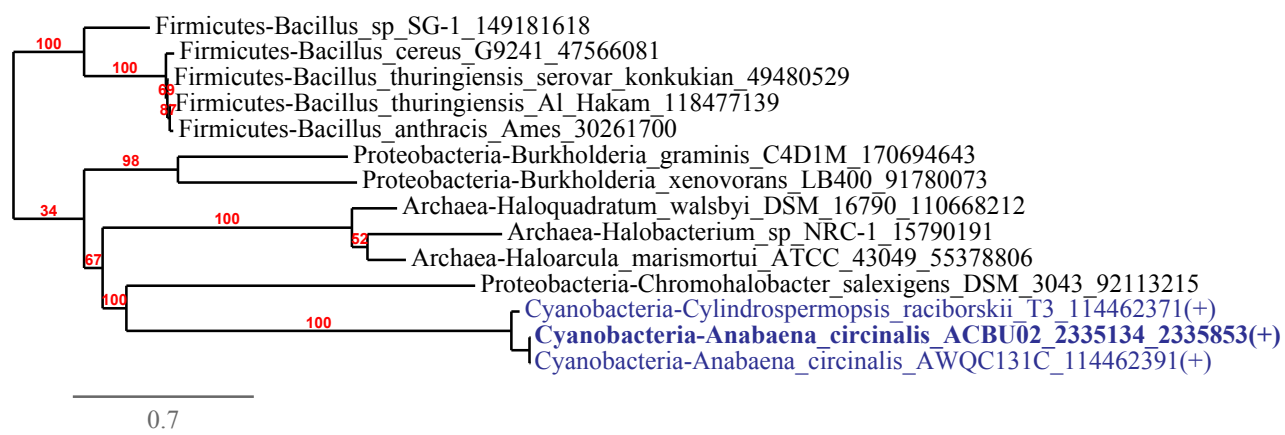
sxtO: Adenylylsulfate kinase



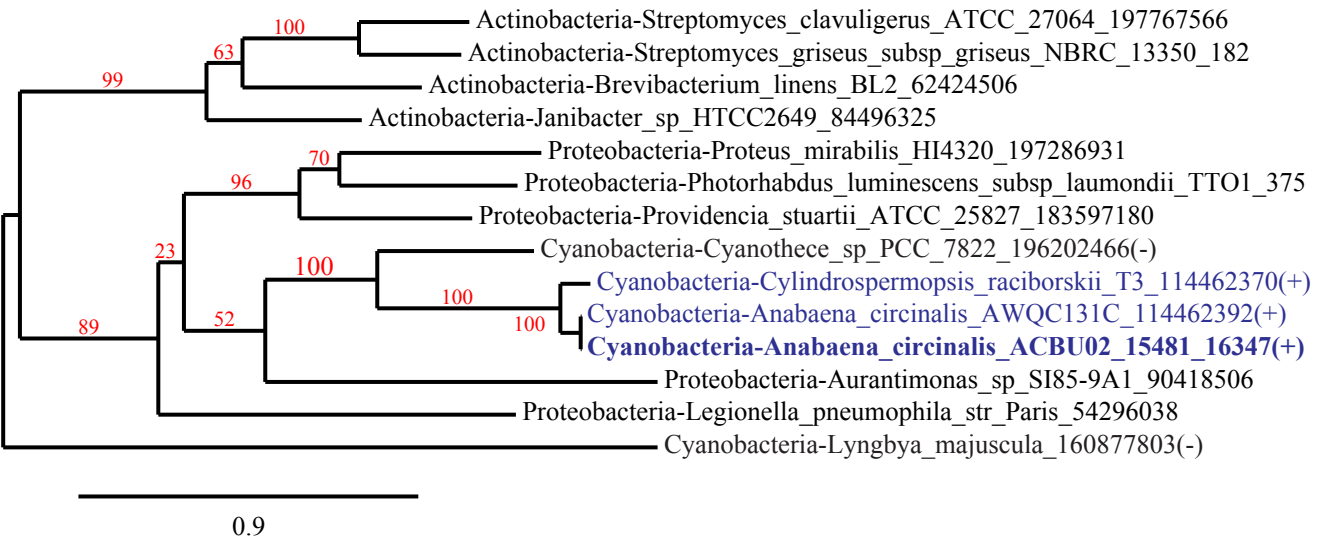
sxtP: Saxitoxin-binding protein



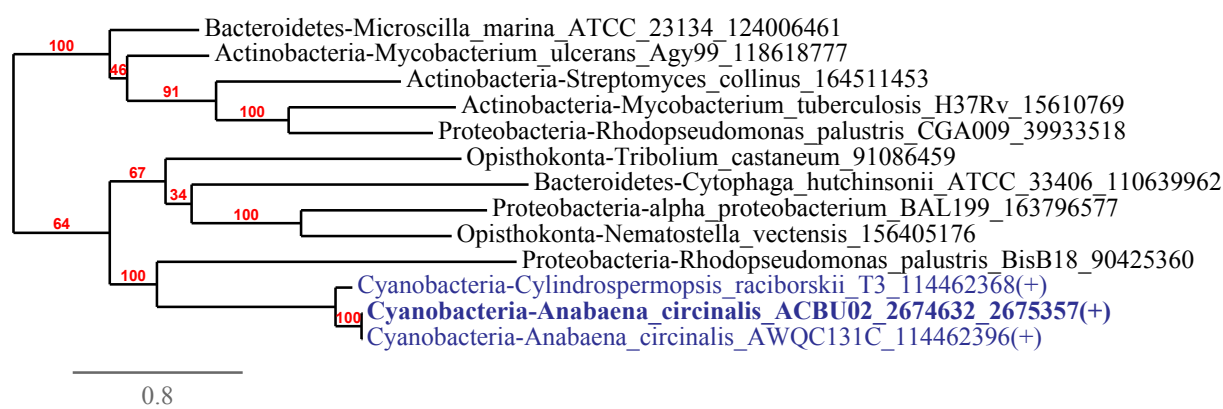
sxtQ: Unknown



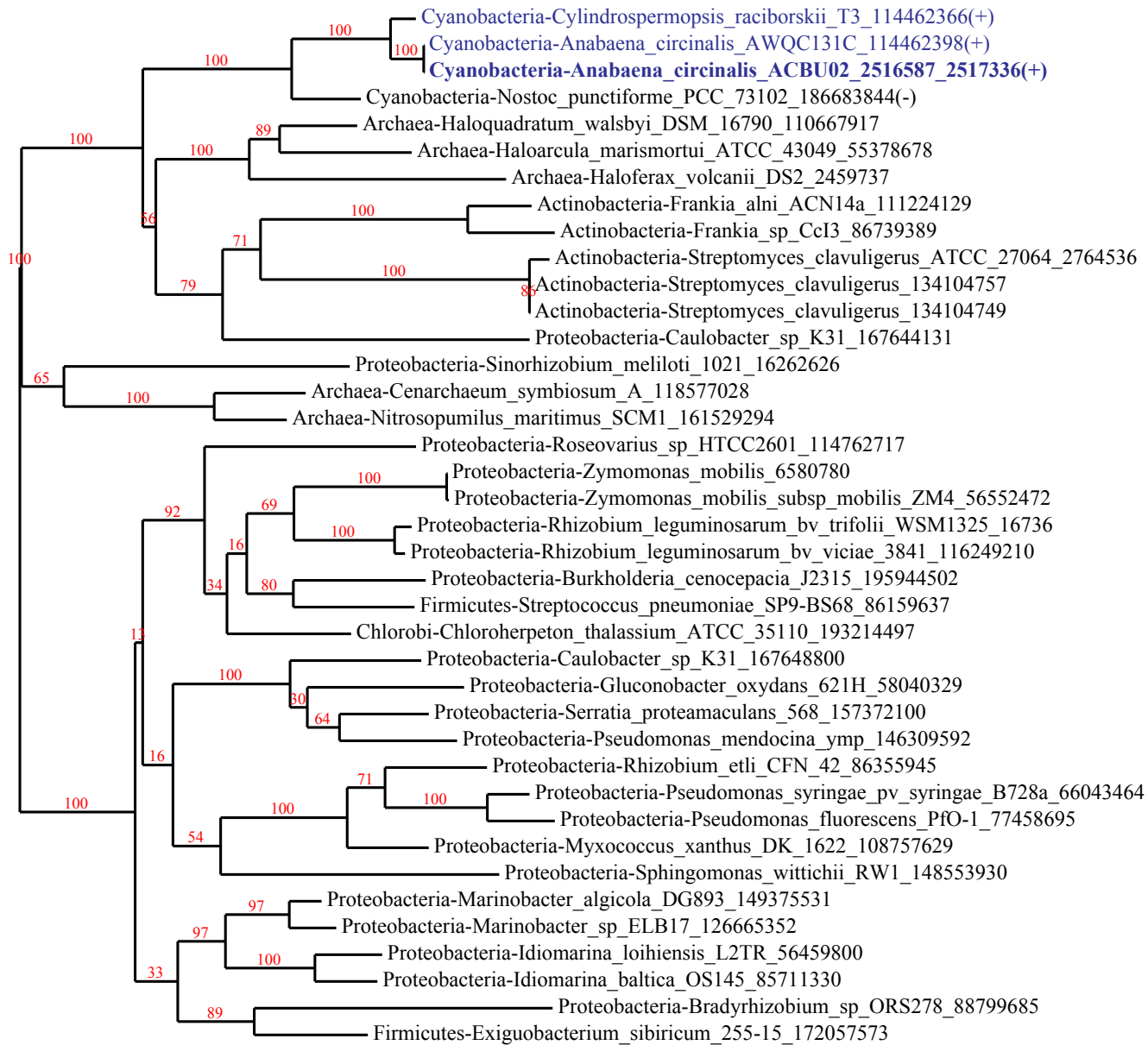
sxtR: Acyl-CoA N-acyltransferase



sxtS: Phytanoyl-CoA dioxygenase

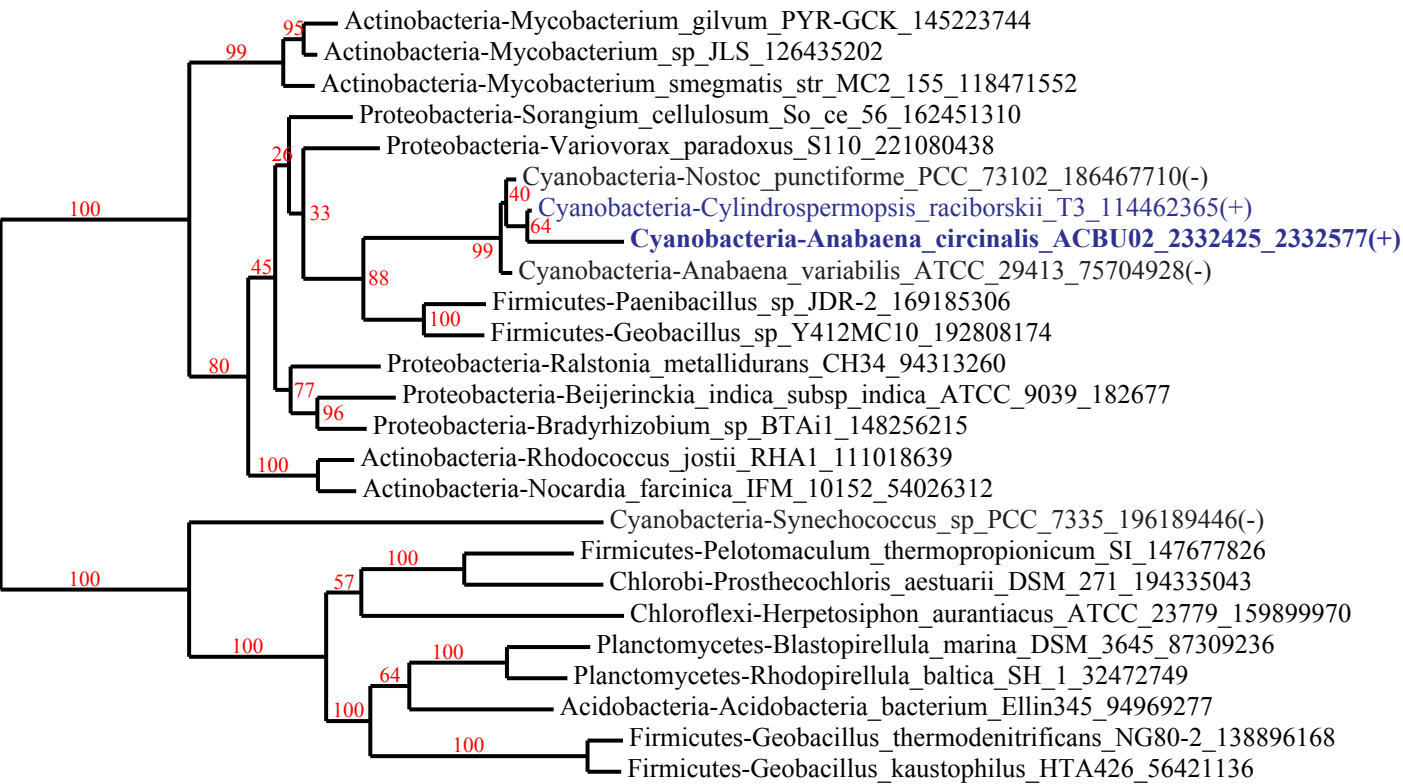


sxtU: Short-chain alcohol dehydrogenase



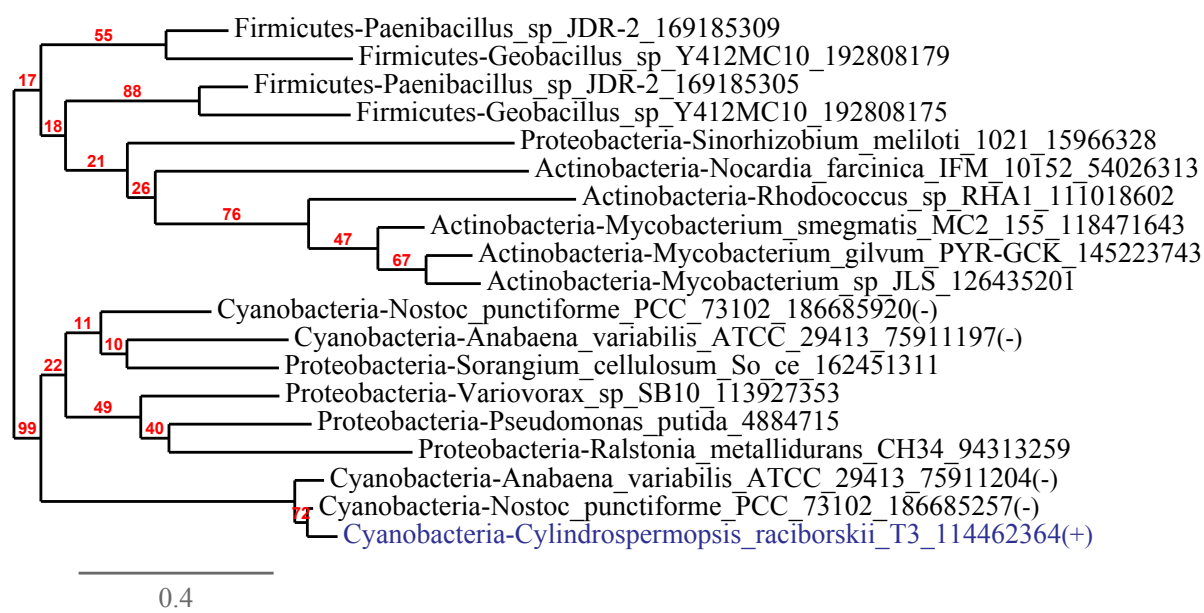
0.3

sxtV: succinate dehydrogenase/fumarate reductase

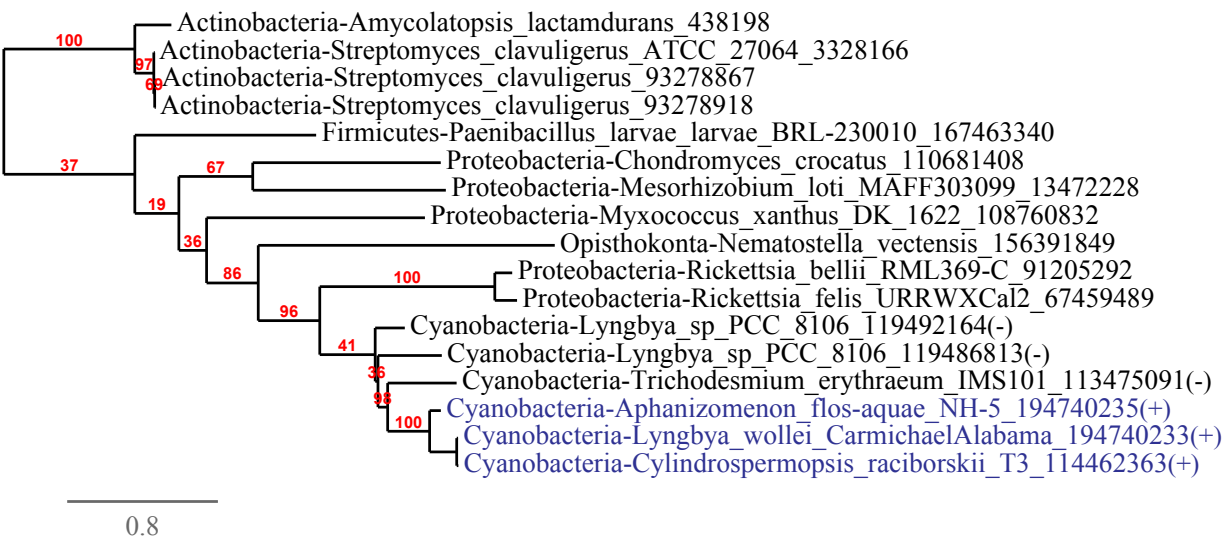


0.9

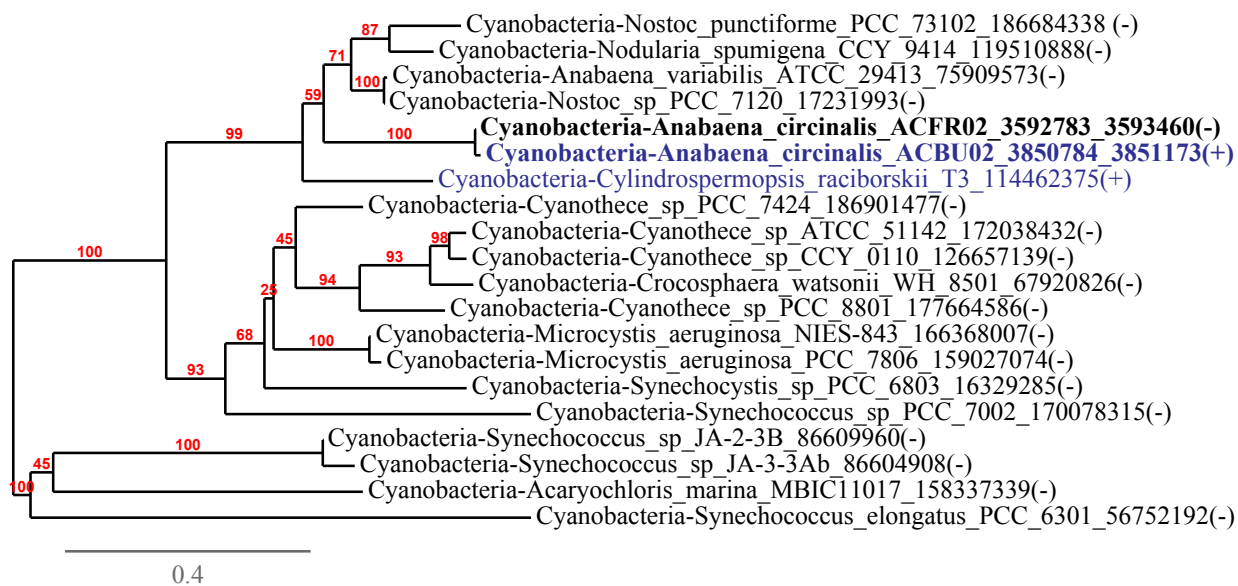
sxtW: ferredoxin



sxtX: Cephalosporin hydroxylase



sxtY: Phosphate uptake regulator



sxtZ: Two-component sensor histidine kinase

