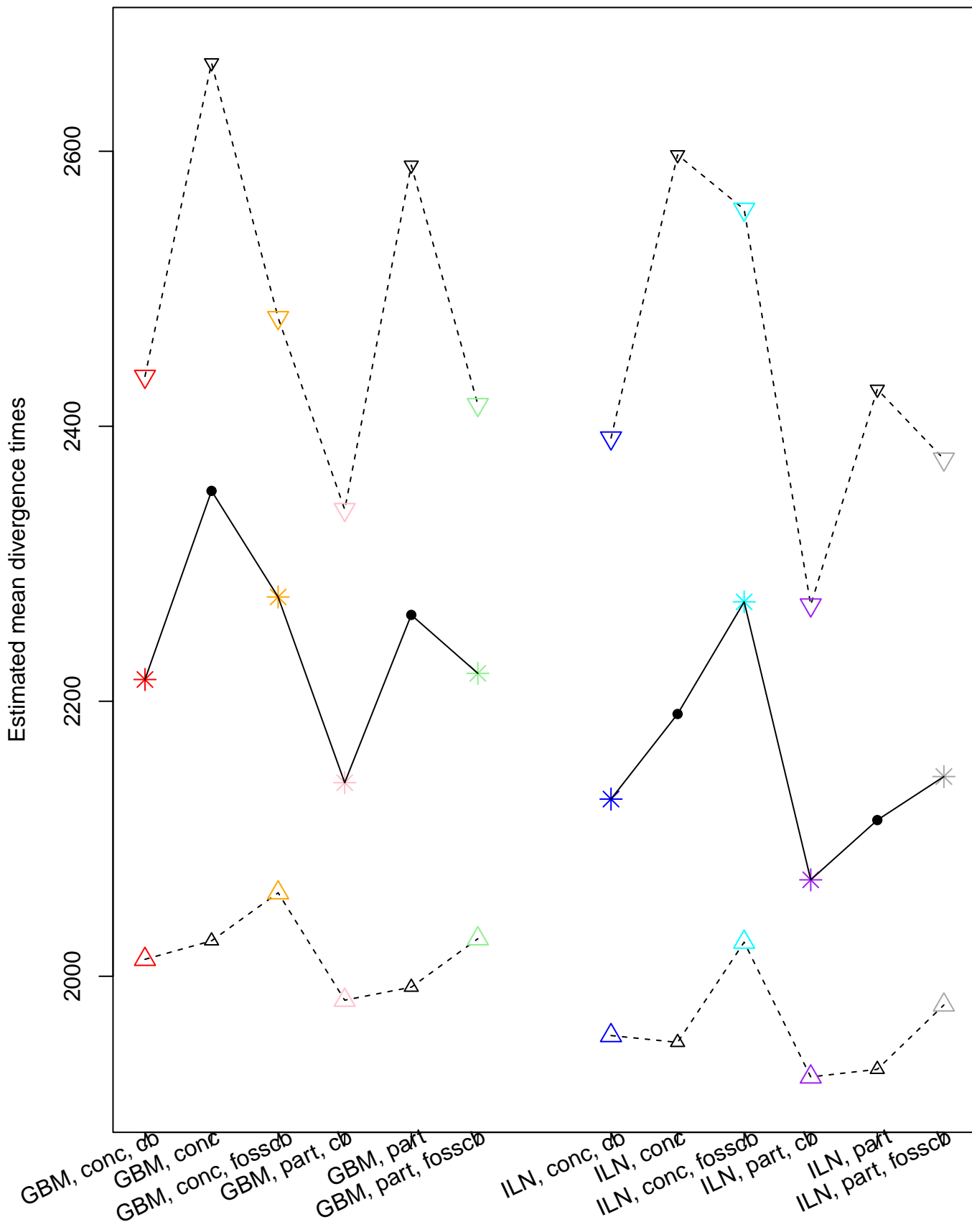


Node t_n309 | CG-OXYPHOTOBACTERIA



Node t_n423 | CG-OXYPHOTOBACTERIA-DUP

