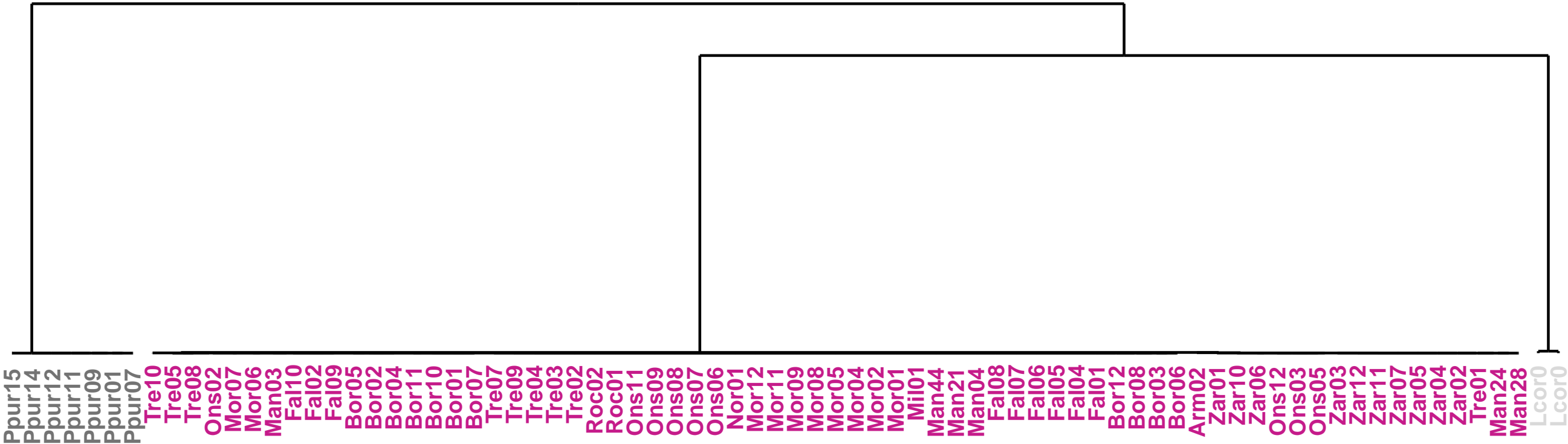


(a)

Maerl species mitogenome CDS alignment

(60.9 % of sites were invariant)



(b)

Maerl species plastome CDS alignment

(73.1 % of sites were invariant)

