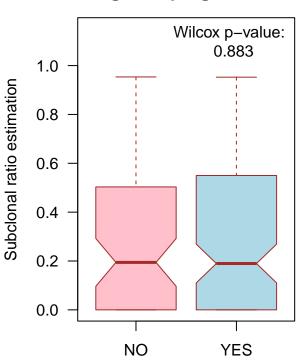
(A) Subclonality estimation against progression



(B) Purity estimation against progression

