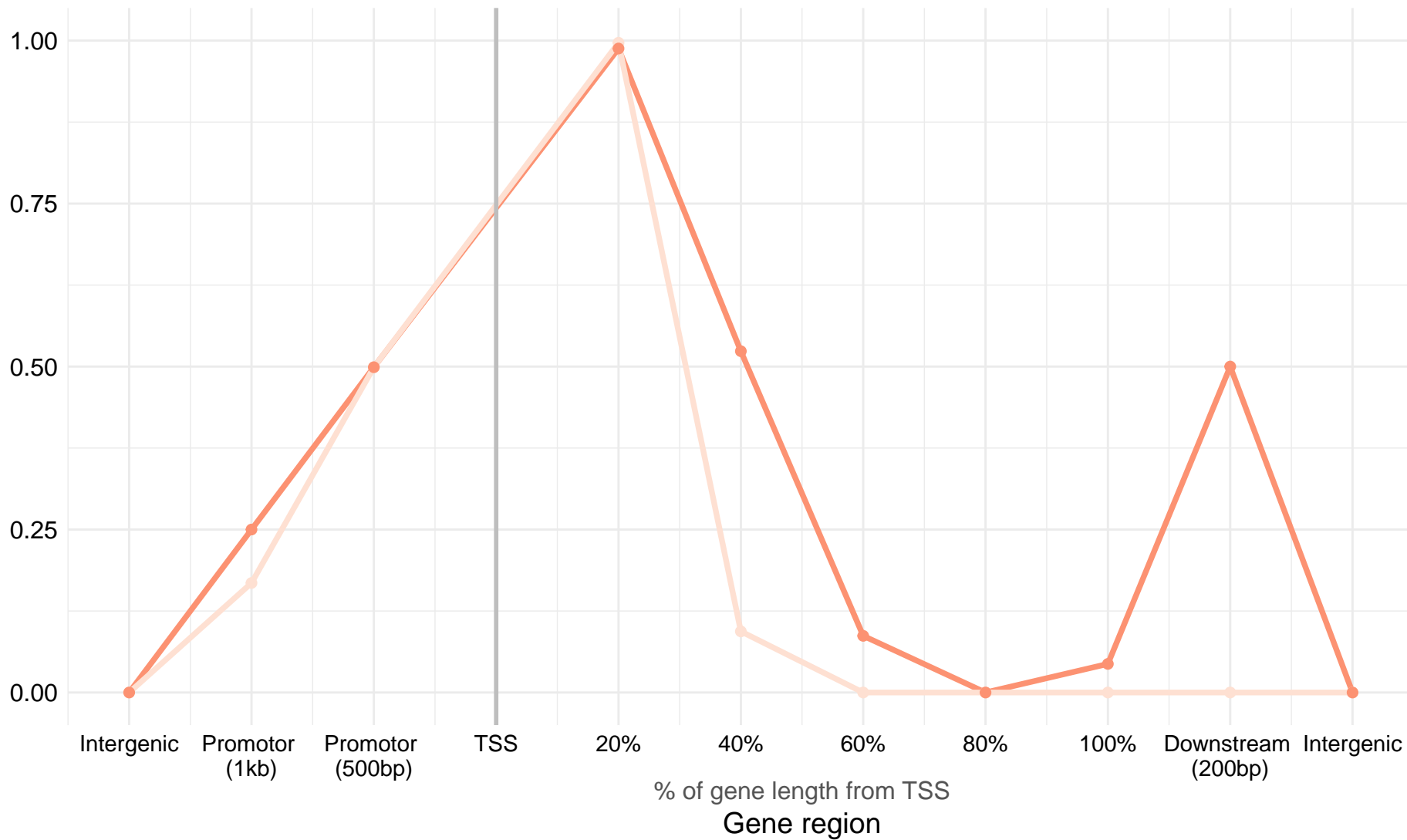


Average proportion of gene region



Expression Level

No Expression (n = 2)

Low Expression (n = 2)