

mag.tooShort  
1.75%

mag.tooShort  
(29.9% complete, 7.5% contamination)  
198 of 7 010 sequences which span 0.0

(29.9% complete, 7.5% contamination)  
(Markers reside on 198 of 7,010 sequences which span 0.06 of 1.93 (3.0%) Mb)

