

Course > Section 2: Biocondu... > IRanges and GRang... > Assessment: Introd...

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Assessment: Introduction to GenomicRanges

We will continue using the ERBS example introduced in the video.

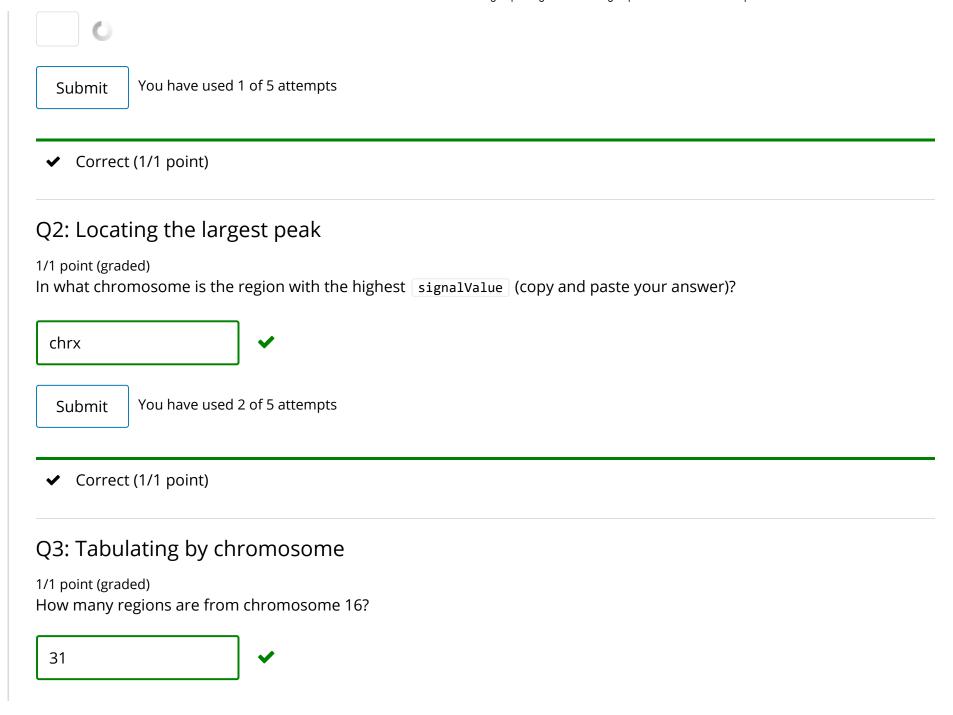
Q1: Statistics on peak scores

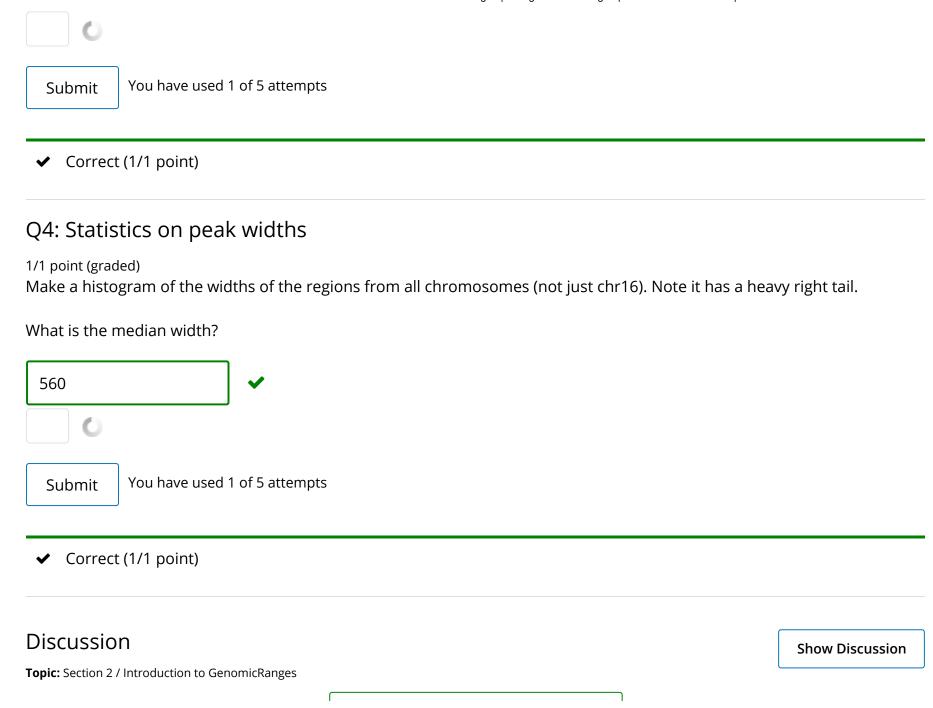
1/1 point (graded)

In the video we used the values() method to extract meta-data on the regions. An alternative, and actually preferred approach going forward, is mcols().

What is the median of the signalValue column for the HepG2 data?

7.024





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