

lg3.1: 122.193 Kb

lg3.2: 6.2 Mb

0 Kb

0.001 Kb

contig-0001436: 6.313 Mb

The diagram illustrates a genomic alignment. A top horizontal line represents a genomic region, with two segments marked by double slashes. Below these marks are labels '0 Kb' and '0.001 Kb'. Above the line, 'lg3.1: 122.193 Kb' is aligned with the first segment, and 'lg3.2: 6.2 Mb' is aligned with the second segment. A bottom horizontal line represents another genomic region, labeled 'contig-0001436: 6.313 Mb'. Two light gray shaded triangles connect the top and bottom lines, indicating a mapping or overlap between the two regions.