The image depicts a bar chart titled "snoID_1022". The x-axis represents the genomic position on chr14, ranging from 31878200 to 31878600. The y-axis represents the read count, ranging from 0 to 4. The chart shows two peaks: one at around 31878300 with a high read count, and another at 31878500 with a moderate read count. The figure illustrates the distribution of read counts across the specified genomic position range.