Caption:   
Circular representation of the B. licheniformis ATCC 14580 chromosome. Circles are numbered from 1 (outermost) to 7 (innermost). Circles 1 and 2 show the locations of predicted CDSs on the + and - strands, respectively; circle 3, %G+C; circle 4, GC skew ((G-C/G+C)); circle 5, homology with B. subtilis 168; circle 6, homology with B. halodurans; circle 7 shows positions of nine copies of insertion sequence element IS3Bli1 and a putative transposase gene; small green bars inside circle 7 denote the positions of possible prophage elements.

Question: What information is shown in circle 4 of the circular chromosome?   
   
A: locations of predicted CDSs on the + strand   
B: locations of predicted CDSs on the - strand   
C: %G+C content   
D: GC skew

Answer: GC skew ((G-C/G+C))