



Figure S2: Characterization of LRR protein fragments from *Flavobacterium psychrophilum*. **(A)** Computational approach for high-resolution mapping of ORF fragments. **(B)** Histogram of fragment ends ($n = 206$) as a function of position in DNA (green: non-coding; magenta: leucine-rich repeat motif region). Multiple repeat units were concatenated into a single unit for visualization purposes.