



Figure S11. Mean coverage (relative to the genome average) of DNase I hypersensitive zones (A-C) and GC content (D-F) as a function of the distance to the closest U-domain border in K562 (A,D), GM06990 (B,E) and BG02 (C,F), for four U-domain size categories : $L < 0.8\text{Mb}$, $0.8\text{Mb} < L < 1.2\text{Mb}$, $1.2\text{Mb} < L < 1.8\text{Mb}$ and $1.8\text{Mb} < L < 3\text{Mb}$ from light to dark curves.