

Forward



Reverse complement



p-value:	1e-1278
Number of Target Sequences with motif	4083
Percentage of Target Sequences with motif	21.53%

**Supplemental Figure 3: *de novo* ZNF217 motif**

Identification of *de novo* motif for ZNF217 binding sites in MCF7 cells. HOMER *de novo* motif analysis using the central most 100 base pairs of all 18,965 ChIP-seq binding regions reveal the indicated motif (shown in both forward and reverse complement).