## **Supplementary Information**

In silico identification of enhancers on the basis of a combination of transcription factor binding motif occurrences

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Supplementary Figures

Supplementary Figures S1

Supplementary Figures S2

Supplementary Tables

Supplementary Table S1

Supplementary Dataset

Supplementary Dataset S1

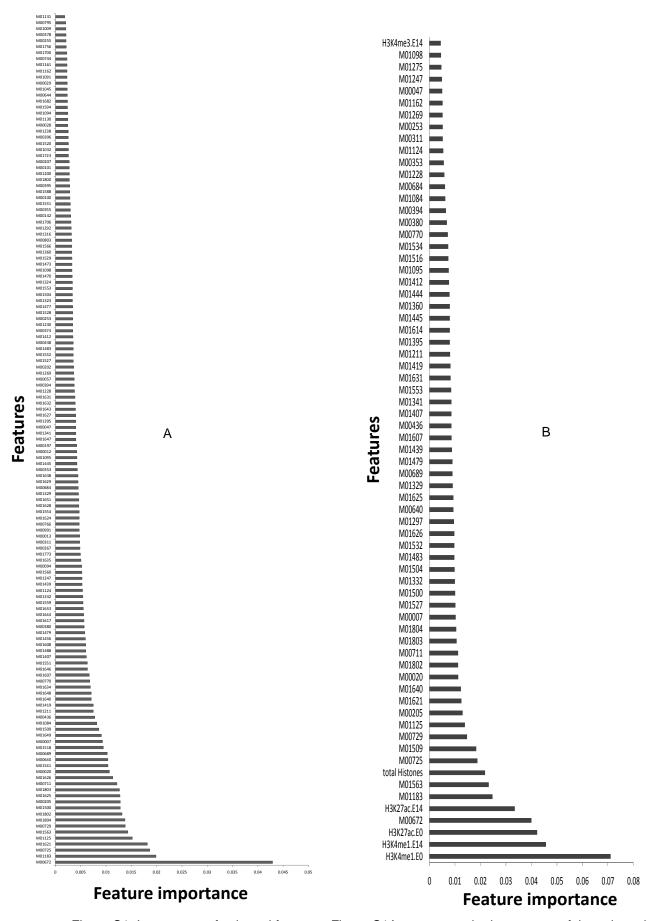


Figure S1. Importance of selected features. Figure S1A represents the importance of the selected features in the model incorporating TF binding motif occurrence features in Table 3. Figure S1B represents the importance of the selected features in the model incorporating the feature group TF binding motif occurrence and the chromatin state in Table 3. The prefix M represents Position Weight Matrix (PWM).

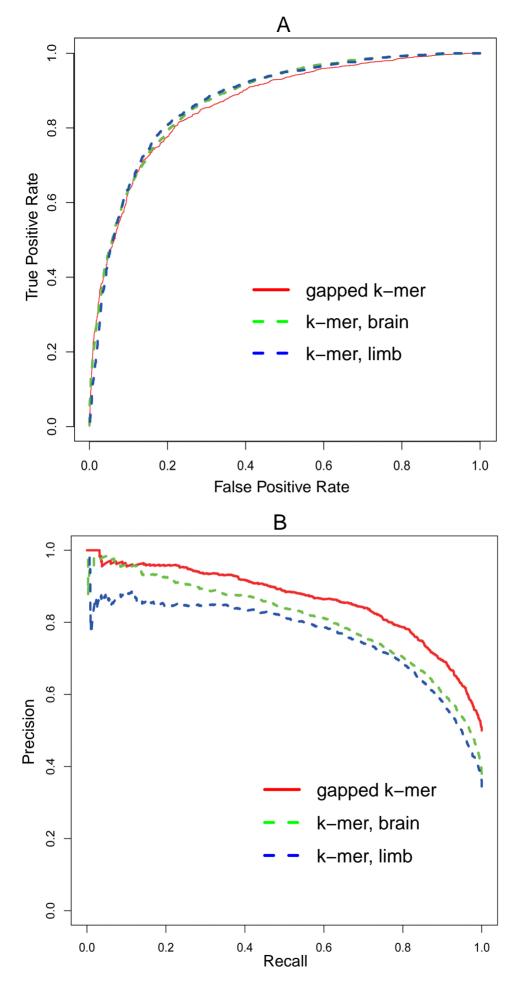


Figure S2. The receiver operator characteristic (ROC) and precision-recall curves for different datasets. Figure S2A represents the ROC curves for different datasets. Figure S2B shows the precision-recall curves for different datasets. The value of AUC is 0.8742 for the mouse embryo brain dataset (green dash line), 0.8738 for the mouse limb dataset (blue dash line) and 0.8643 for the binding sites of P300 protein (red line).