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Hidden Markov Models Predict Epigenetic Chromatin Domains

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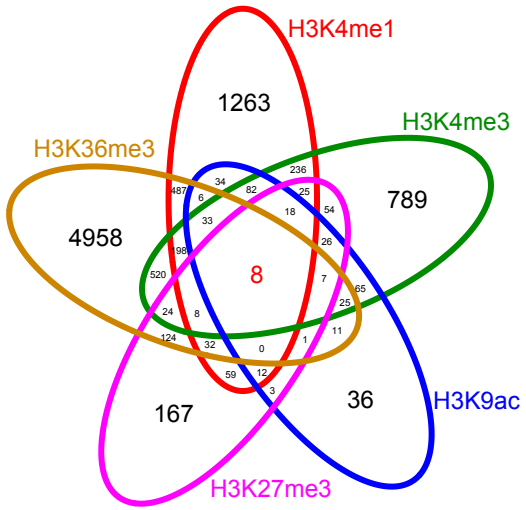
Accessibility

Supplemental Figure 4

(A)



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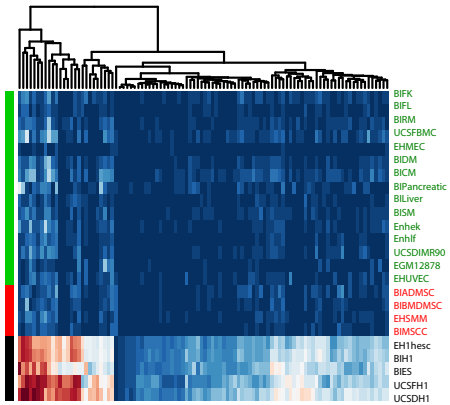

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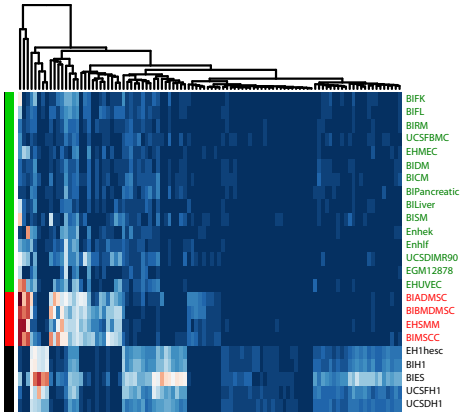

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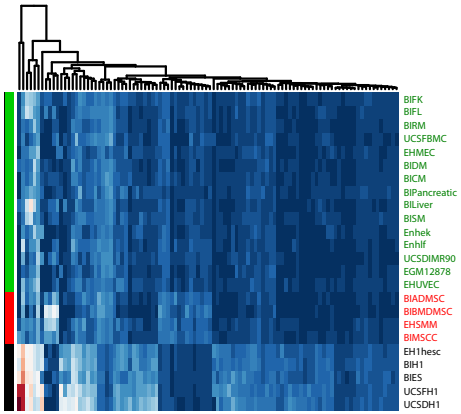
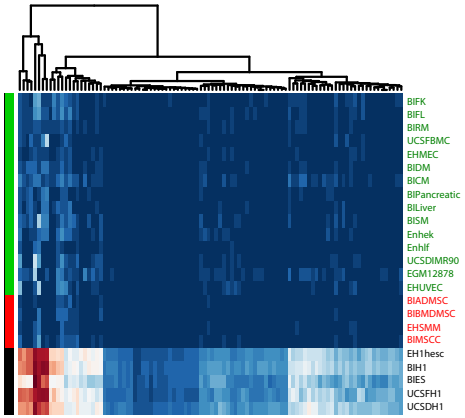
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