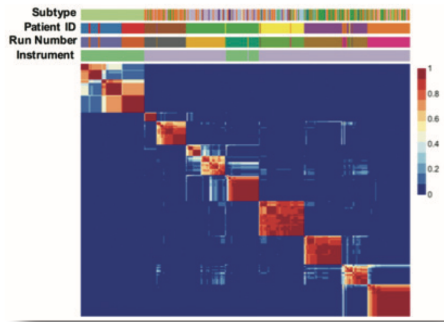
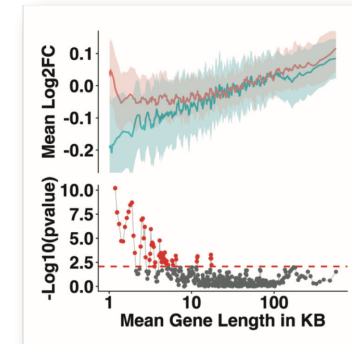


Decoding big genomic datasets: Biases, hypotheses & regulation

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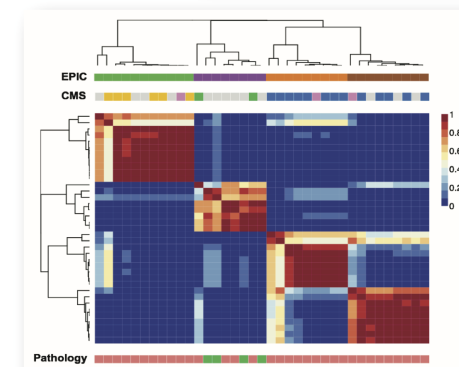
Biases

Batch Effects

Hypothesis

Integration

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