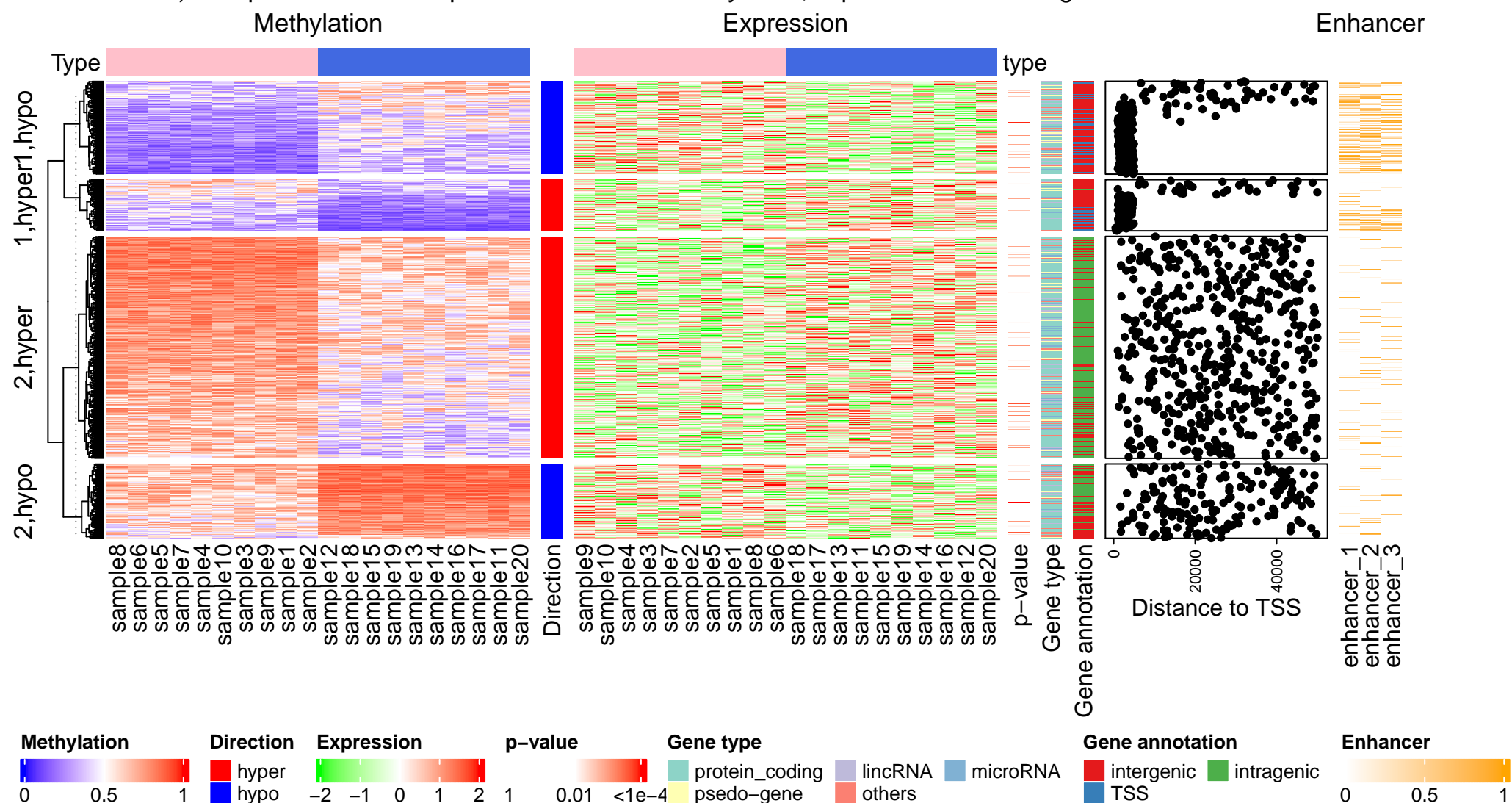


A) Comprehensive correspondence between methylation, expression and other genomic features



B) global summary statistics ...

