21/05/31 21/06/04

Prevention training

Meeting with IVA; she proposed that I search in the databases instead of the papers, to look for SNPs as well and some general advice.

Read (Pierson et al., 2015)

Read the TMM paper

Understand how to implement TMM using edgeR.

Clean the data (GTEx) and select the most relevant attributes, save the cleaned version in a separate Rdata file.

Determine the number of reads per tissue.

Apply TMM