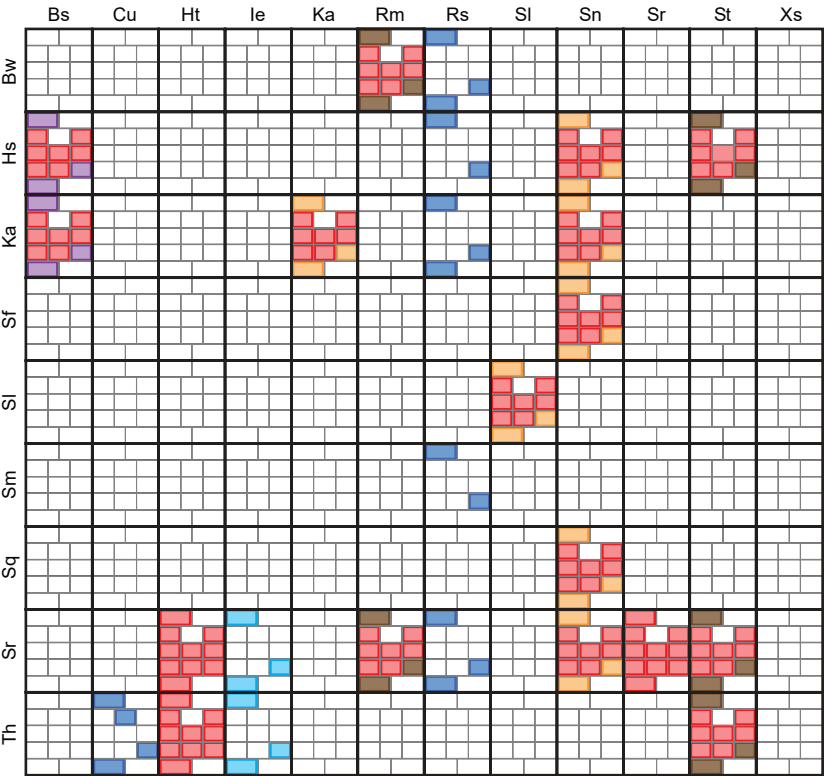


Expression Modeler

a

Genes

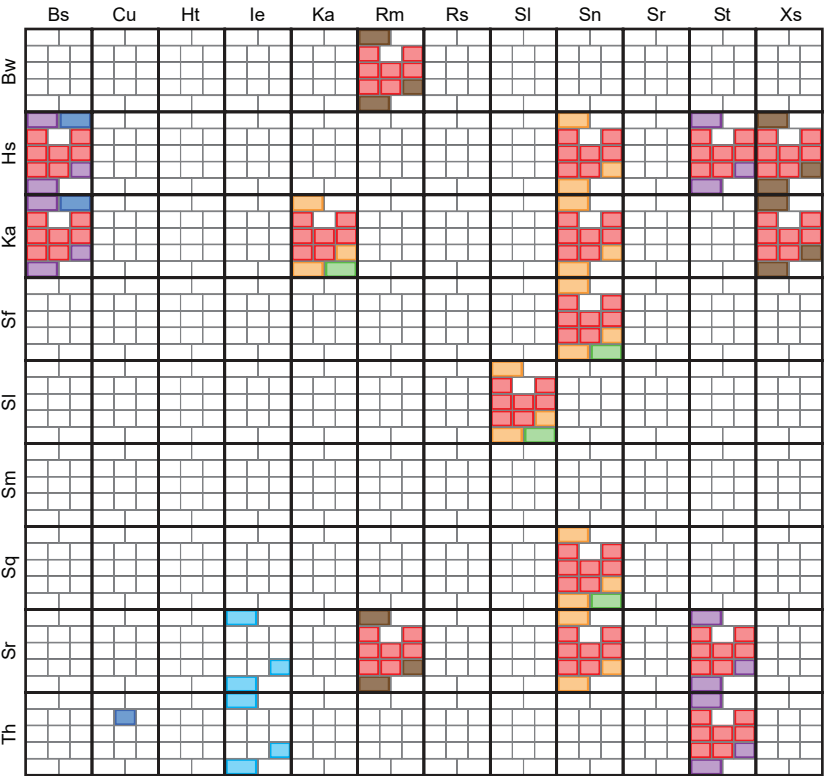
Read Aligner



b

Transcripts

Read Aligner



- counts only
- FPKM only
- TPM only
- counts + FPKM
- counts + TPM
- TPM+ FPKM
- counts+TPM+FPKM

Bl		Bs	
By	Cd	De	
Eb	Er	Lo	
Lv	Nb	No	
Sa		Su	