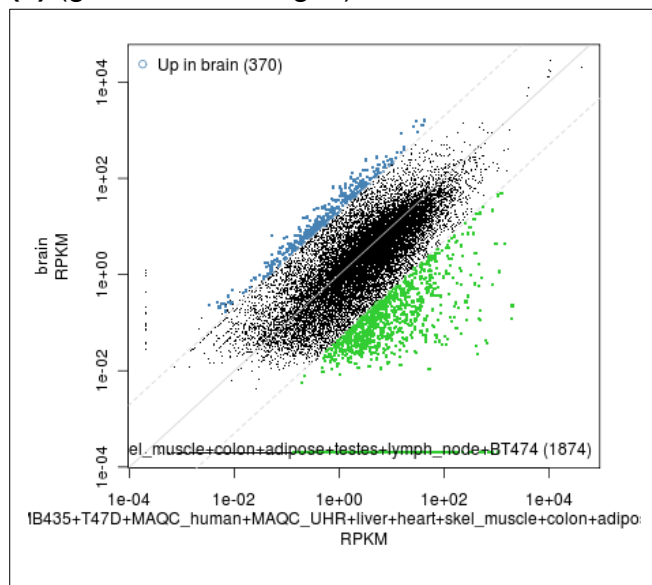
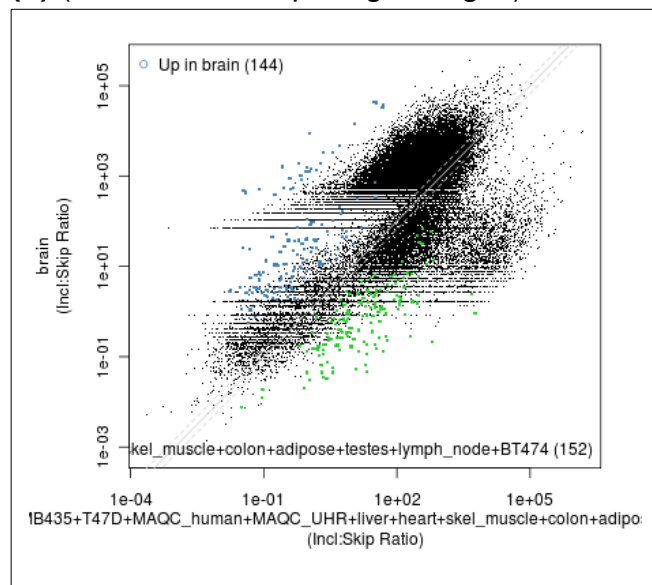


(a) (gene-level changes)**(b)** (cassette exon splicing changes)**(c)****Human_Tissues/brain Up skipped_exons**

Platform=RNASeq P=1e-4 FC=1.5

Showing 144/144 Up skipped_exons

Download: Convert hg18 skipped_exons to Make Background controlling for

chr	strand	reg.exon	gene	ID	eup1	jup1	js
<input type="text"/>			<input type="text"/>		<input type="button" value="All"/> <input type="text"/>	<input type="button" value="All"/> <input type="text"/>	<input type="button" value="All"/> <input type="text"/>
chr9	1	123102155,123102225	GSN	170434	0	9	1617
chr9	1	36199264,36199317	CLTA			12	489
chr6	1	33518208,33518249	SYNGAP1			12	184
chr9	1	123102164,123102225	GSN			2	1617
chr9	1	123102201,123102225	GSN			0	1617
chr17	1	37914116,37914133	ATP6V0A3			2	226
chr11	1	118031780,118031812	PHLDB1			2	238
chr15	-1	70282417 70282583	PKM2	190189	18511	270	544

Open Readmap Browser

Human_Tissues

- CLTA
- [chr9:36199264-36199317](#)