

Supplemental Table 2: Summary of novel transcript analysis

Summary of reads, junctions and genes	Single-end 1		Paired-end 1		Paired-end 2	
	E18	P7	E18	P7	E18	P7
Reads	2,302	2,689	621	746	794	1,084
Junctions	1,800	2,177	151	189	504	631
Genes*	1,367	1,596	143	172	448	536
Total genes	2,396		295		811	

*Numbers in this row showed how many genes the detected novel junctions belonged to.