

# Outline

RNAseq samples

Control genes

Replicates separated

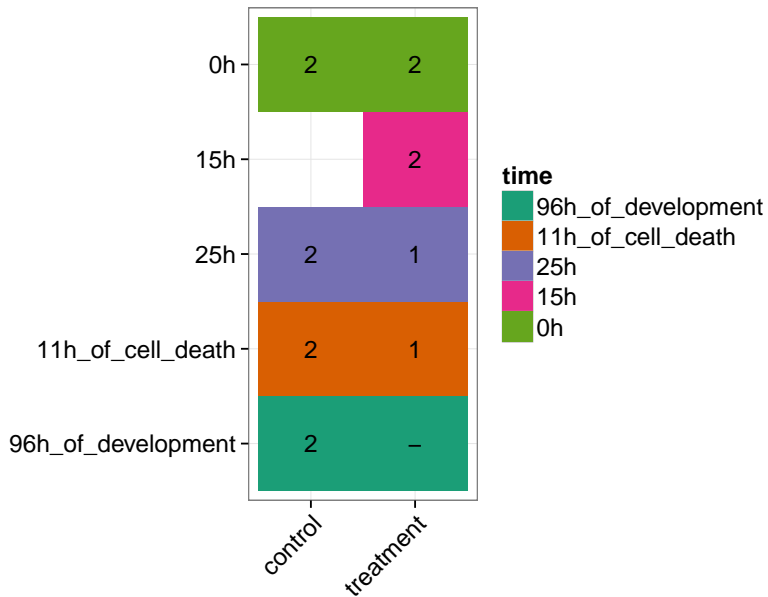
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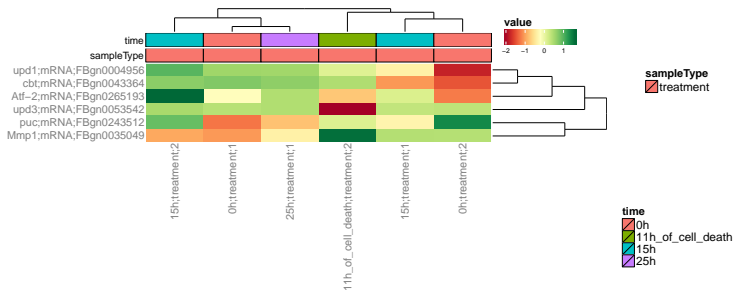
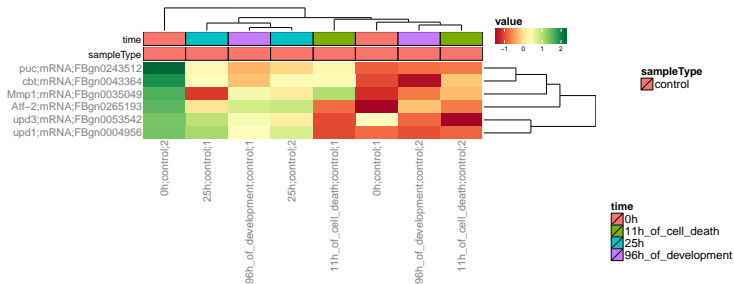
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# FPKM control genes

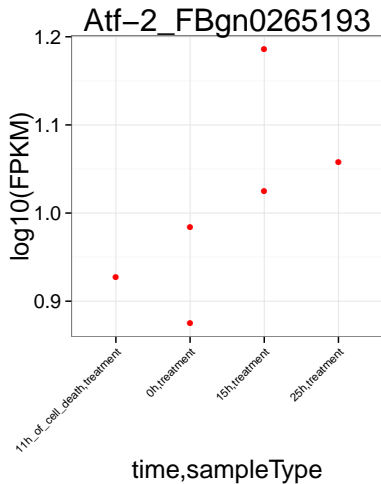
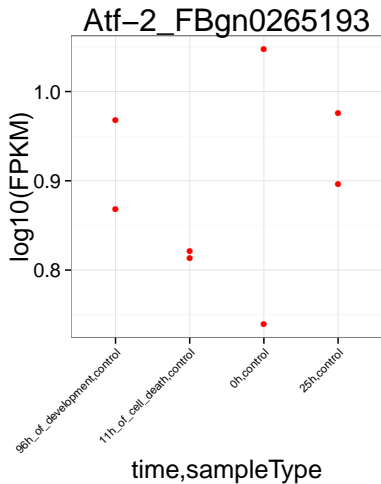
	96h develop., control,1	96h develop., control,2	11h cell death, control,1	11h cell death, control,2	0h, control,1	0h, control,2	25h, control,1	25h, control,2
cbt FBgn0043364	7.05	4.74	8.5	7.17	5.47	14.4	8.53	8.4
Mmp1 FBgn0035049	18.84	14.09	21.6	15.6	12.49	25.04	12.76	17.25
Atf-2 FBgn0265193	9.3	7.38	6.5	6.62	5.49	11.15	7.87	9.47
puc FBgn0243512	5.93	4.81	8.01	4.93	4.55	21.76	8.09	6.79
upd1 FBgn0004956	1.85	1.15	1.13	1.21	1.23	2.74	2.44	2.14
upd3 FBgn0053542	0.18	0.08	0.07	0.05	0.17	0.33	0.25	0.15

	11h of cell death, treatment,2	0h, treatment,1	0h, treatment,2	15h, treatment,1	15h, treatment,2	25h, treatment,1
cbt FBgn0043364	10.15	10.49	8.16	8.54	10.42	10.4
Mmp1 FBgn0035049	23.9	14.88	19.43	19.45	15.15	16.86
Atf-2 FBgn0265193	8.45	9.65	7.5	10.59	15.34	11.41
puc FBgn0243512	7.91	4.87	12.25	6.75	10.35	5.74
upd1 FBgn0004956	3.09	3.78	1.32	2.5	4.54	3.76
upd3 FBgn0053542	0.0	0.31	0.57	0.4	0.34	0.46

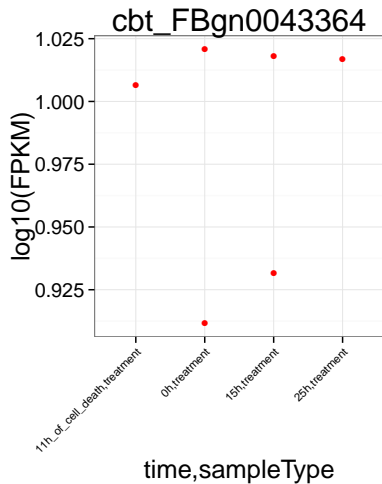
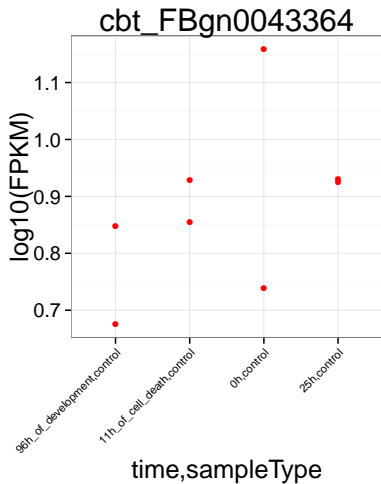
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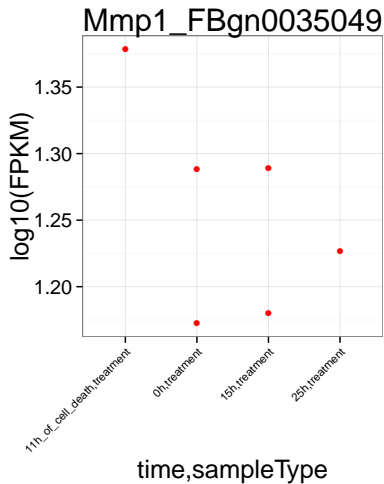
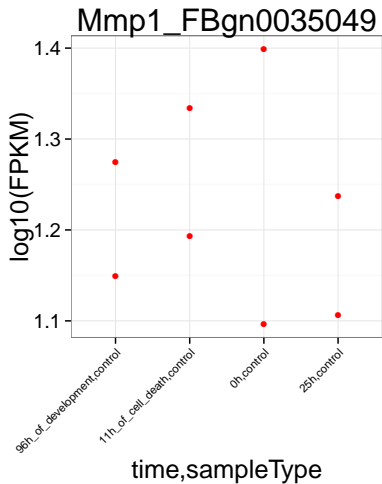


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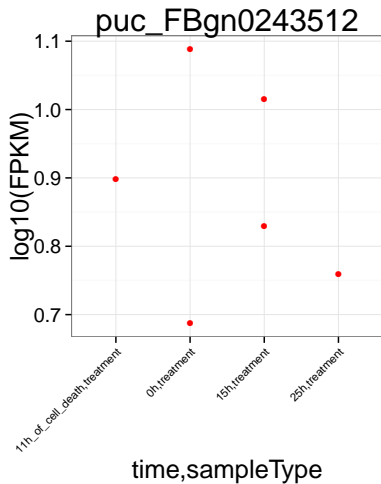
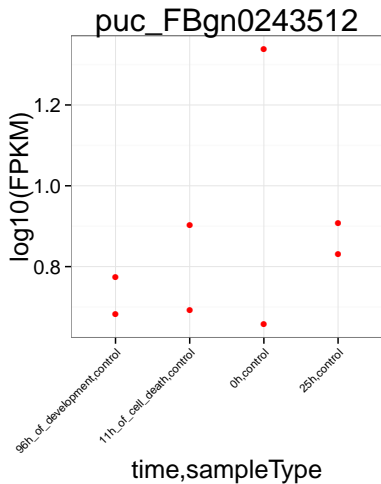




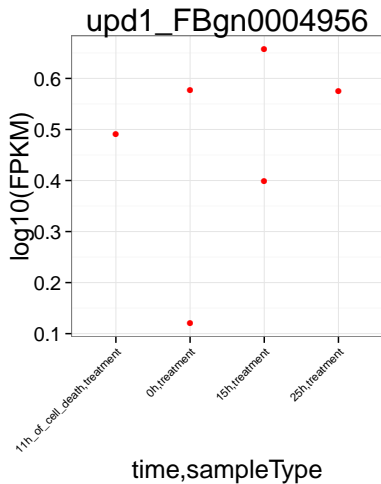
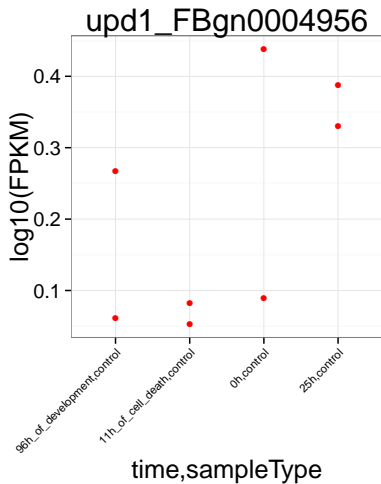
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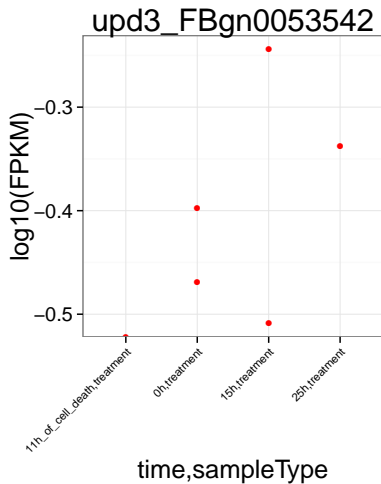
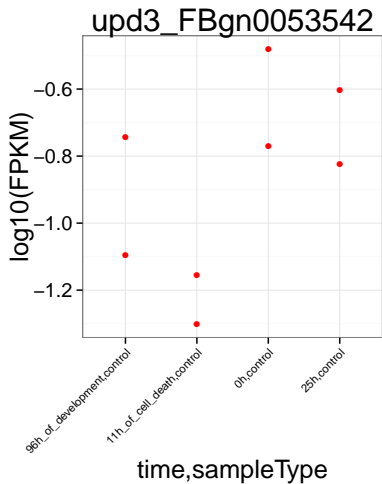
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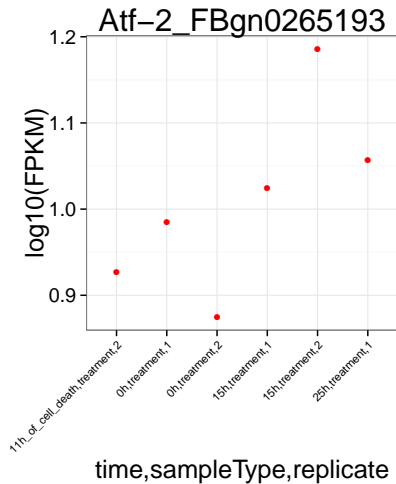
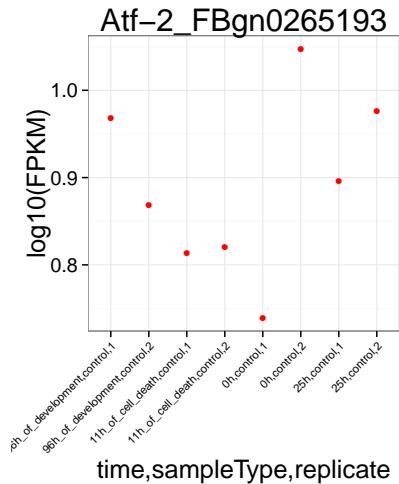
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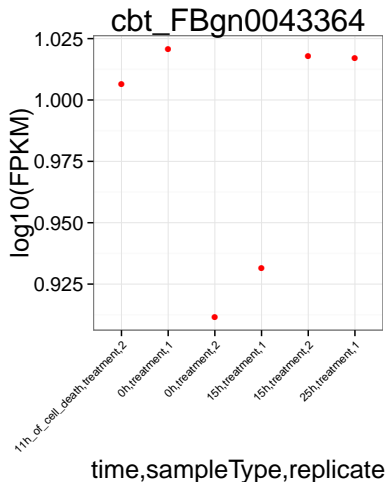
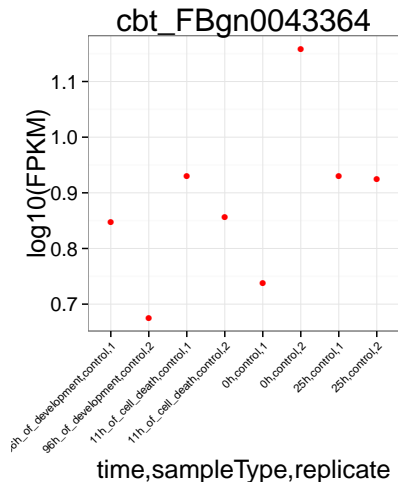
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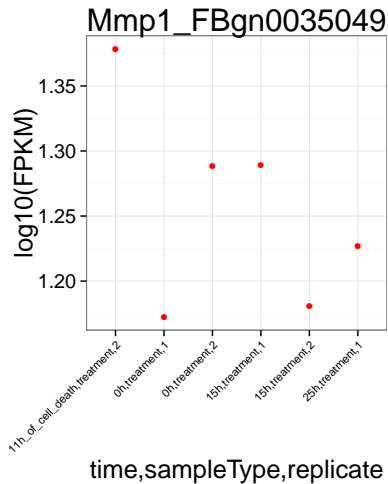
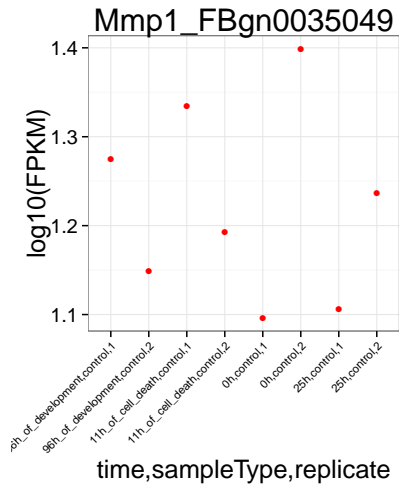
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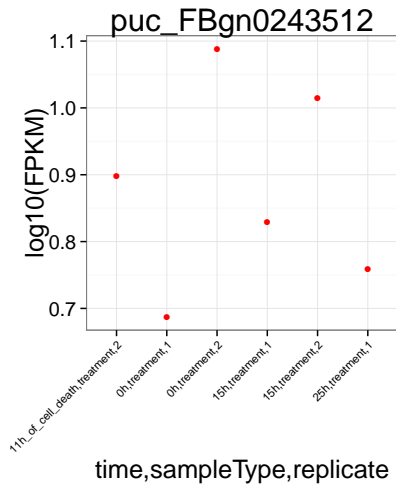
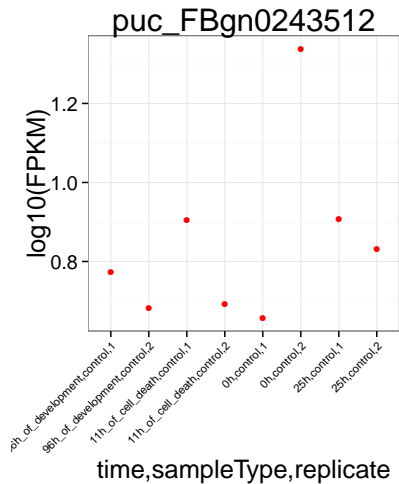


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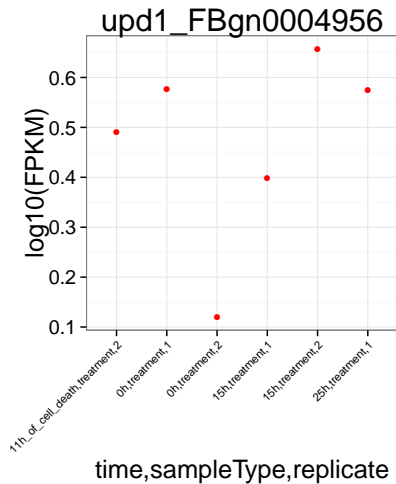
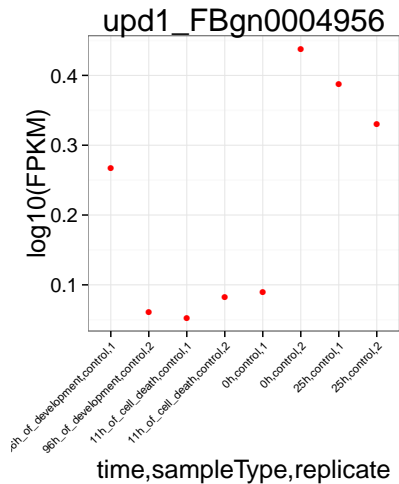




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