



Split Table

Low vs. Others | Middle vs. Others | High vs. Others

TF	Coeff	High 0	High 1	Other 0	Other 1	Avg. High	Avg. Other	Diff	Score
POU1F1	0	112	3	453	0	0.026	0.000	0.026	8.13e-3
IRF7	0	111	4	450	3	0.035	0.007	0.028	0.034
SMAD4	0	115	0	438	15	0.000	0.033	-0.033	0.050
POU2F1	0	115	0	436	17	0.000	0.038	-0.038	0.058
SMAD3	0	115	0	439	14	0.000	0.031	-0.031	0.085
PBX1	0	111	4	448	5	0.035	0.011	0.024	0.087
FOXL1	0	113	2	452	1	0.017	0.002	0.015	0.106
HMGA2	0	113	2	452	1	0.017	0.002	0.015	0.106
RFX1	0	115	0	442	11	0.000	0.024	-0.024	0.132
MEIS1	0	115	0	442	11	0.000	0.024	-0.024	0.132
NR2F1	0	115	0	443	10	0.000	0.022	-0.022	0.135
SP2	0	115	0	443	10	0.000	0.022	-0.022	0.135
POU3F2	0	115	0	443	10	0.000	0.022	-0.022	0.135
SMAD2	0	115	0	443	10	0.000	0.022	-0.022	0.135
SMAD1	0	115	0	443	10	0.000	0.022	-0.022	0.135
NR2F2	0	115	0	441	12	0.000	0.026	-0.026	0.138
SP1	0	115	0	441	12	0.000	0.026	-0.026	0.138
PDX1	0	110	5	445	8	0.043	0.018	0.026	0.152
VSX2	0	113	2	451	2	0.017	0.004	0.013	0.184
IRF9	0	114	1	453	0	-0.004	0.000	-0.004	0.202
HINFP	0	114	1	453	0	0.009	0.000	0.009	0.202
XBP1	0	114	1	453	0	0.009	0.000	0.009	0.202
RUNX1	0	115	0	444	9	0.000	0.020	-0.020	0.216
POU3F1	0	115	0	444	9	0.000	0.020	-0.020	0.216
POU2F2	0	115	0	444	9	0.000	0.020	-0.020	0.216

Total number of genes most likely going through this state is 568

Intercept coefficient is 0; This hour6 state output distribution is Normal( $\mu = 0.122, \sigma = 0.282$ )

GO Split Table | Copy Table | Save Table | Run DECOD

