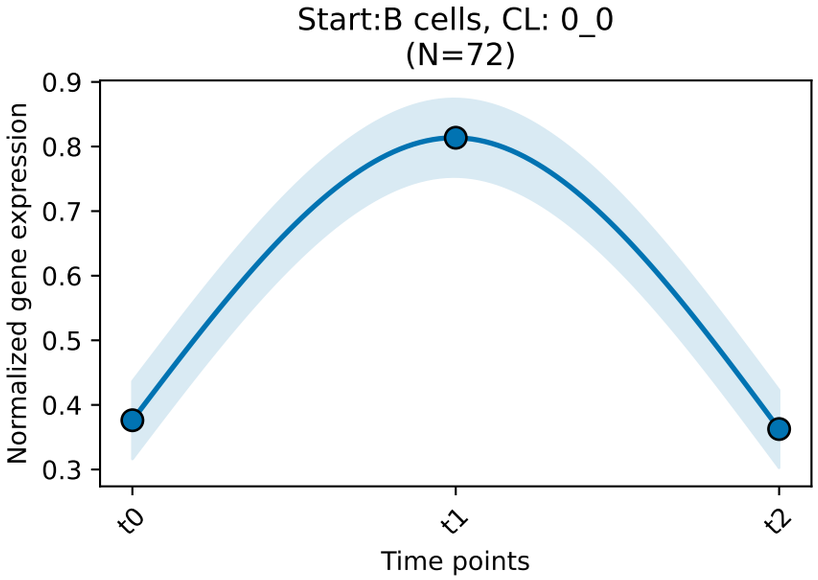
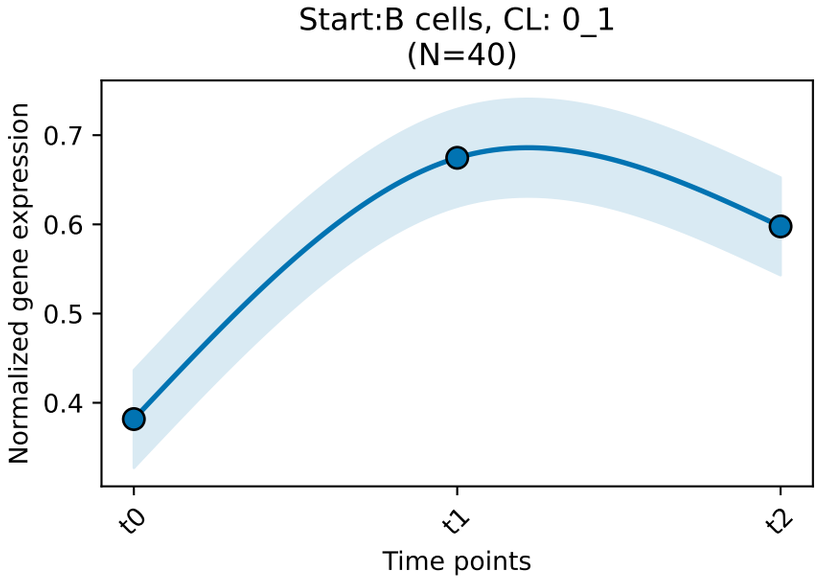
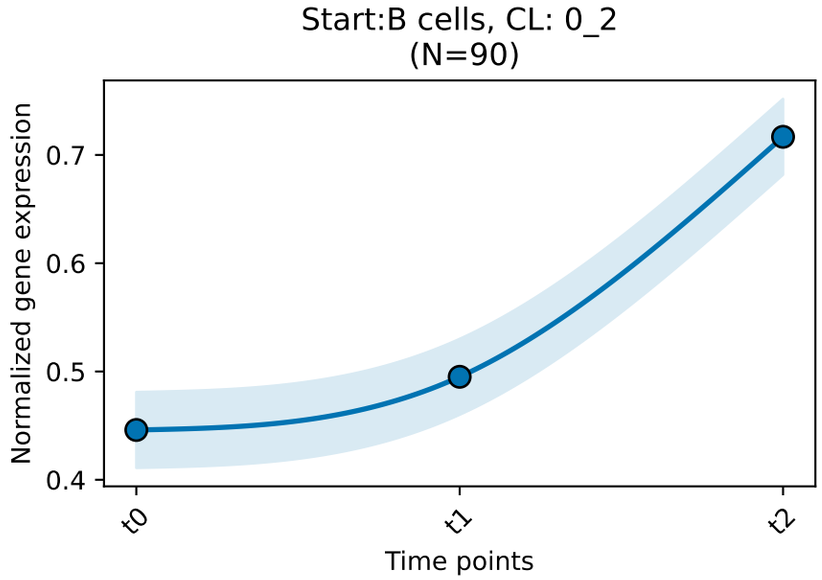
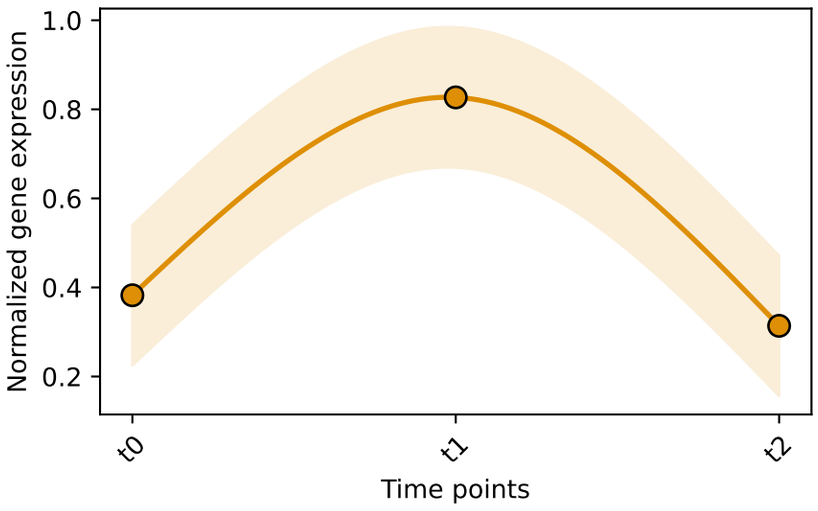
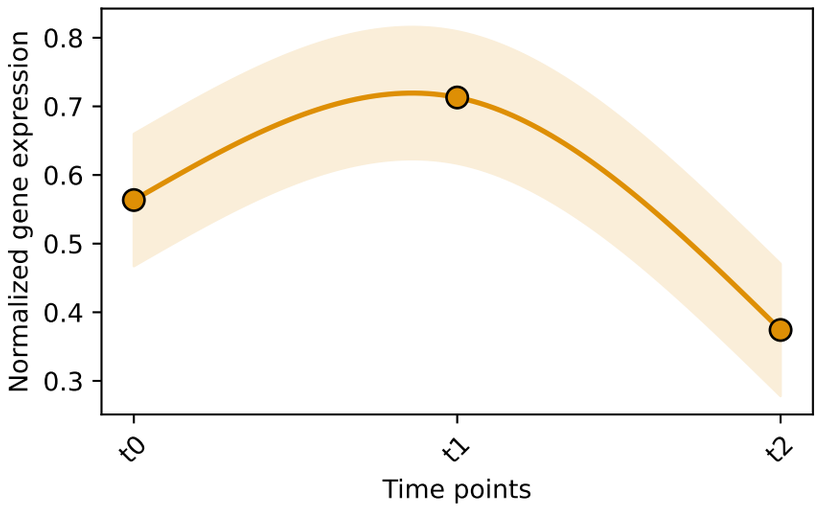
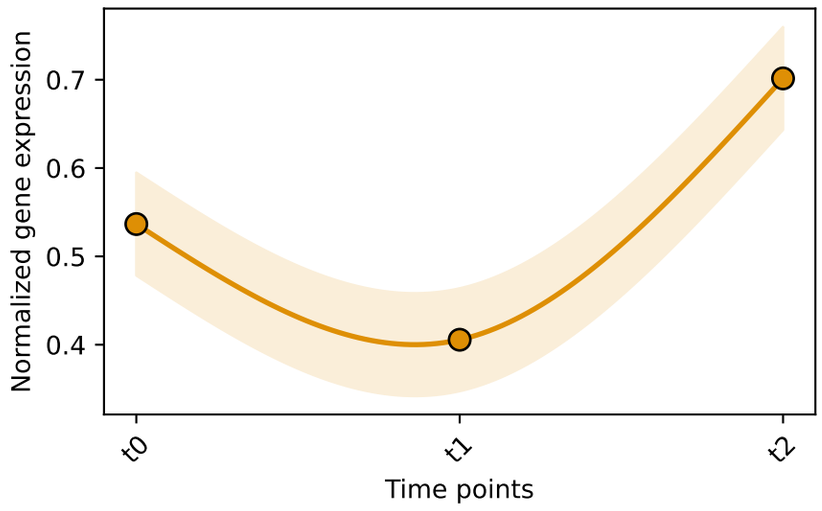

Start:B cells, CL: 0_0
(N=72)
Normalized gene expression
0.9
0.8
0.7
0.6
0.5
0.4
0.3
t0
t1
t2
Time points

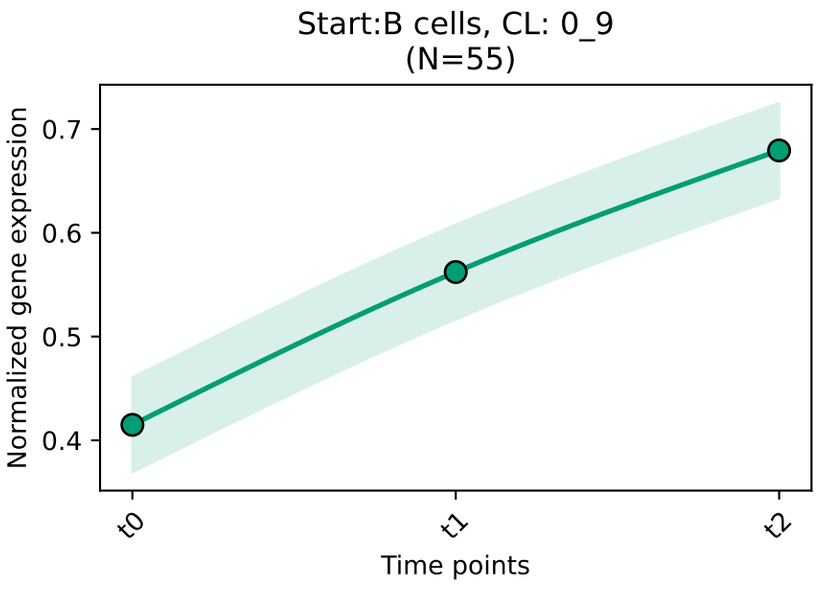

Start:B cells, CL: 0_1
(N=40)
Normalized gene expression
0.7
0.6
0.5
0.4
t0
t1
t2
Time points

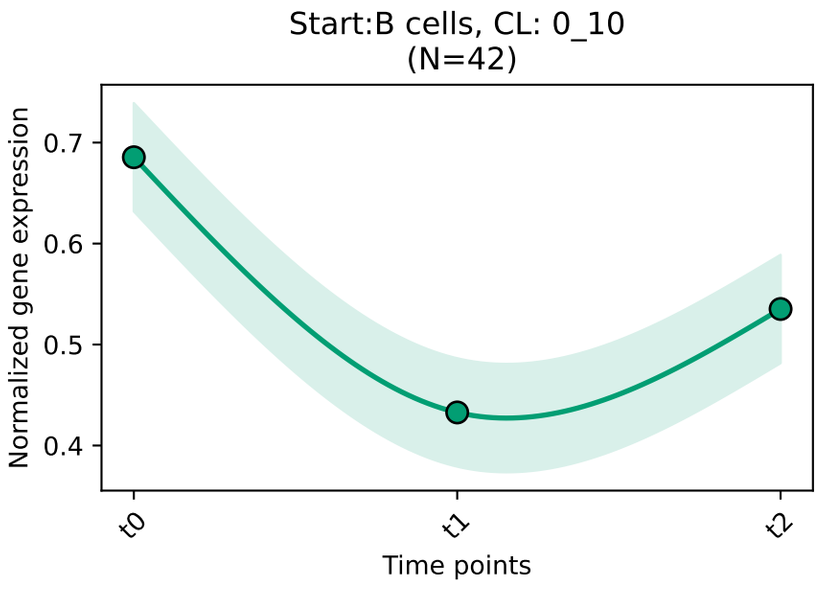

Start:B cells, CL: 0_2
(N=90)
Normalized gene expression
0.7
0.6
0.5
0.4
t0
t1
t2
Time points

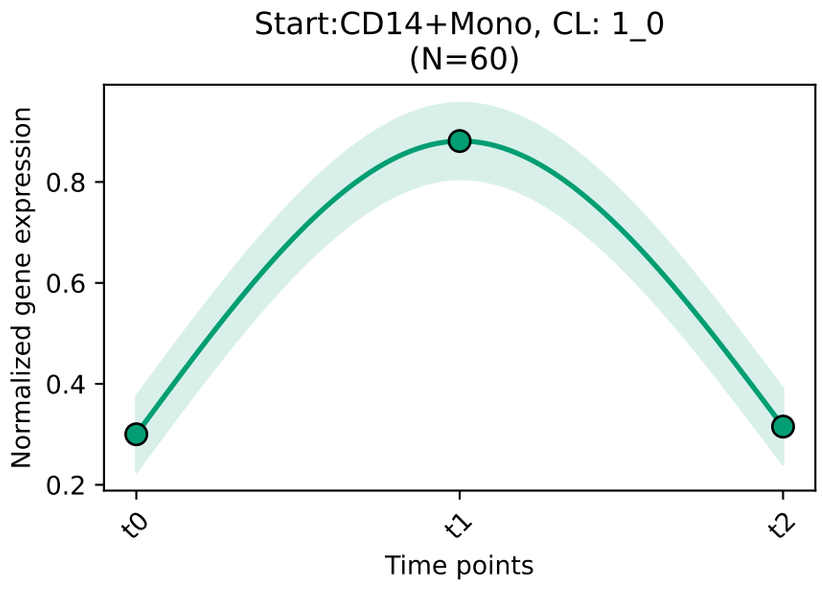

Start:B cells, CL: 0_3
(N=14)
Normalized gene expression
1.0
0.8
0.6
0.4
0.2
t0
t1
t2
Time points


Start:B cells, CL: 0_6
(N=15)
Normalized gene expression
0.8
0.7
0.6
0.5
0.4
0.3
t0
t1
t2
Time points

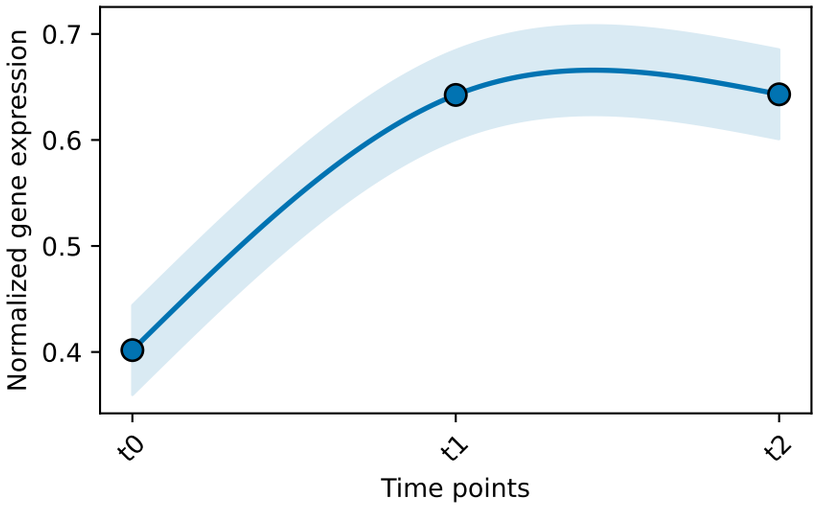

Start:B cells, CL: 0_7
(N=40)
Normalized gene expression
0.7
0.6
0.5
0.4
t0
t1
t2
Time points


Start:B cells, CL: 0_9
(N=55)
Normalized gene expression
0.7
0.6
0.5
0.4
t0
t1
t2
Time points

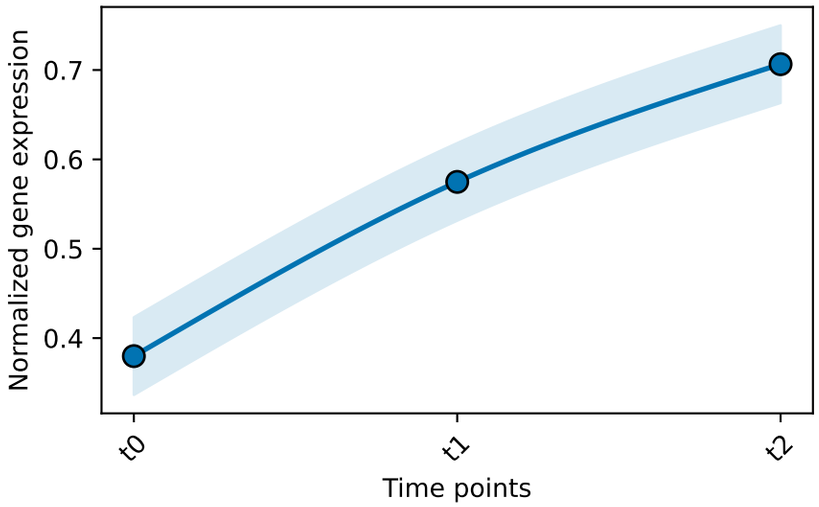

Start:B cells, CL: 0_10
(N=42)
Normalized gene expression
0.7
0.6
0.5
0.4
t0
t1
t2
Time points


Start:CD14+Mono, CL: 1_0
(N=60)
Normalized gene expression
0.8
0.6
0.4
0.2
t0
t1
t2
Time points

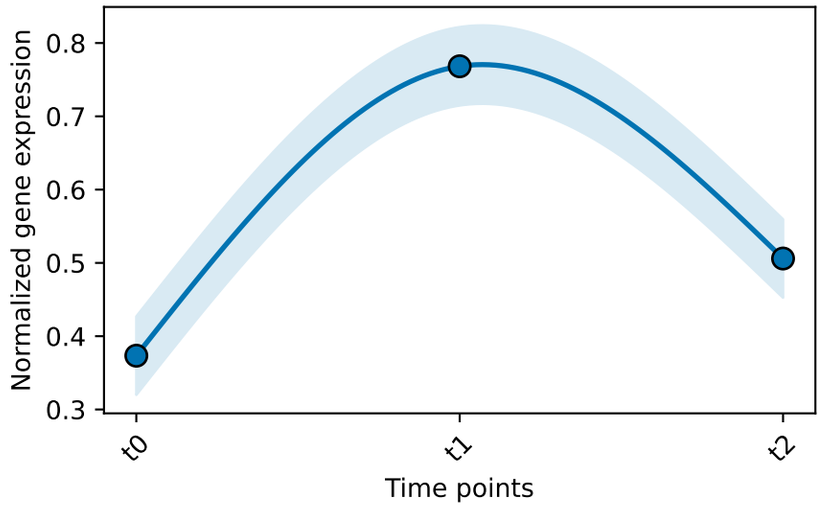
Start:CD14+Mono, CL: 1_1
(N=39)



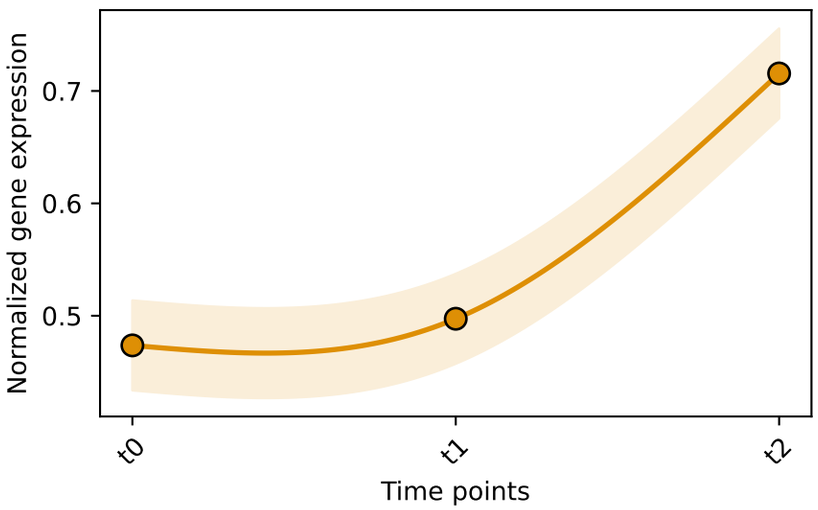
Start:CD14+Mono, CL: 1_4
(N=57)



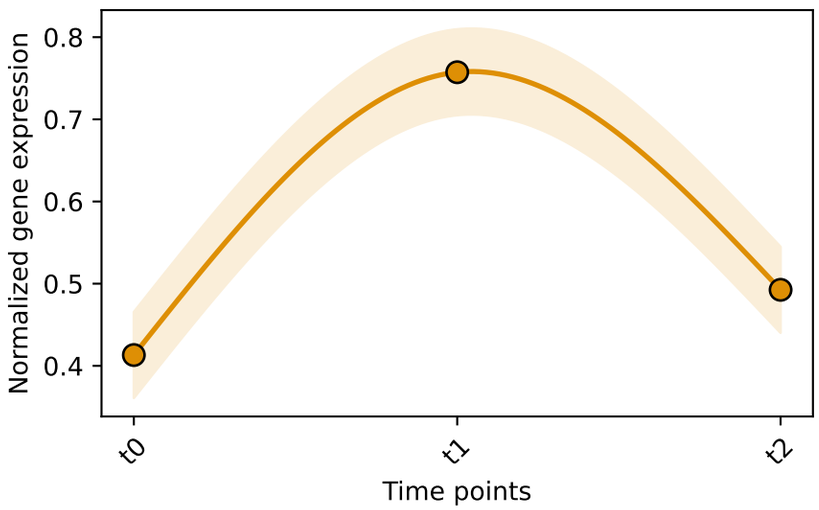
Start:CD14+Mono, CL: 1_6
(N=45)



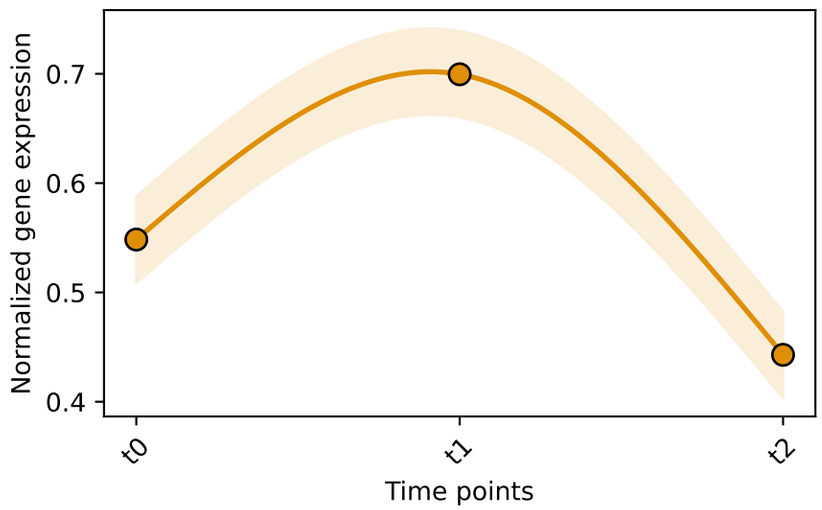
Start:CD14+Mono, CL: 1_7
(N=44)



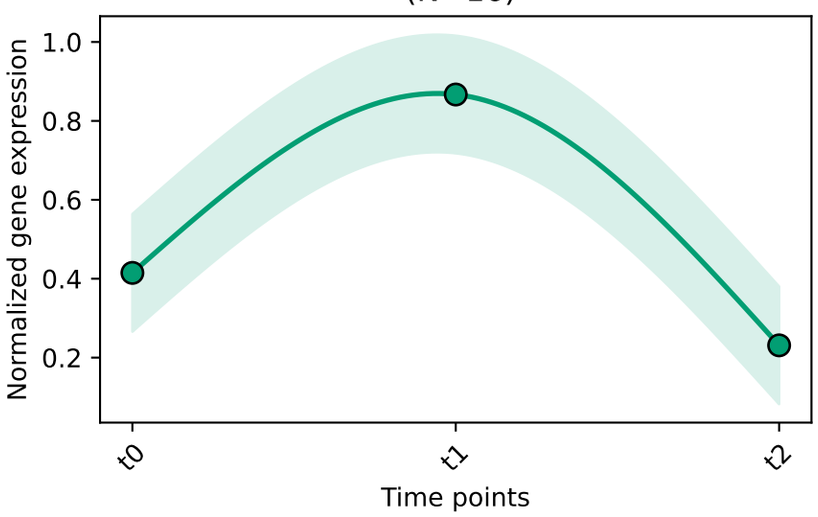
Start:CD14+Mono, CL: 1_8
(N=39)



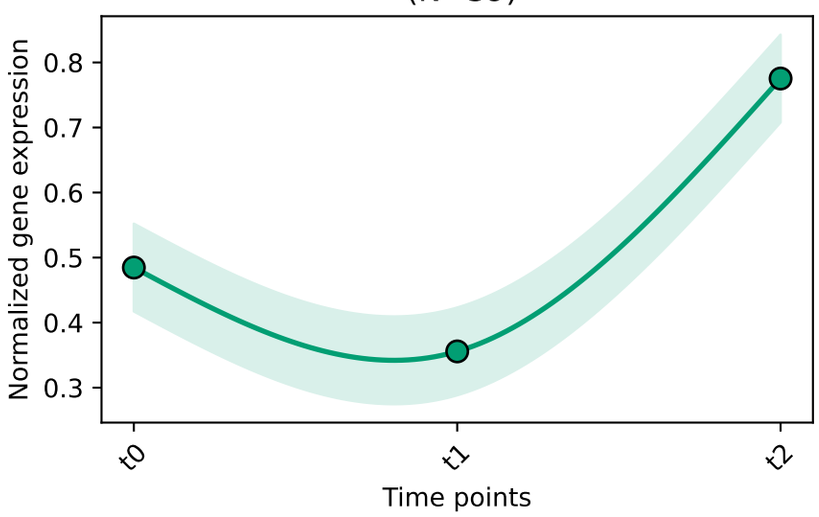
Start:CD14+Mono, CL: 1_11
(N=41)



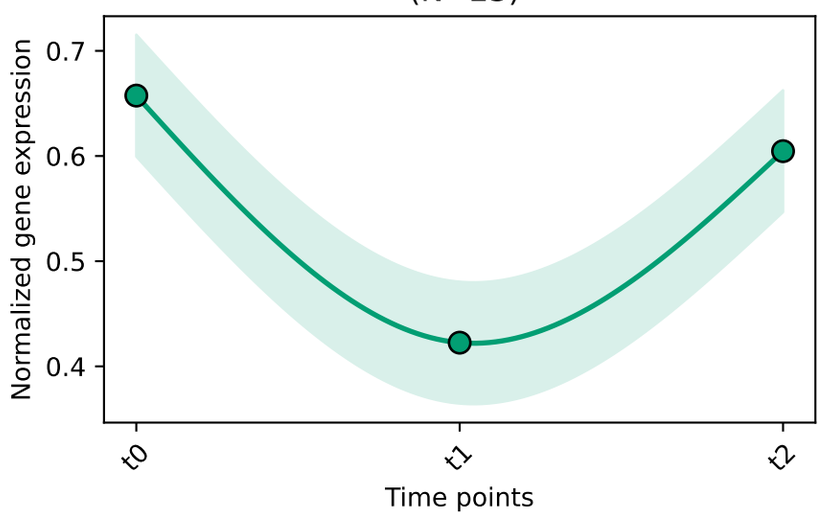
Start:CD14+Mono, CL: 1_13
(N=16)

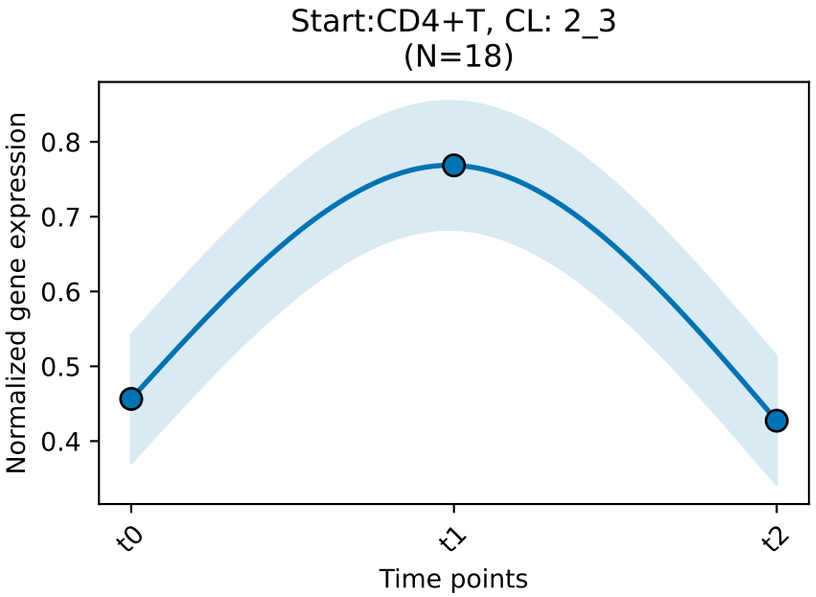


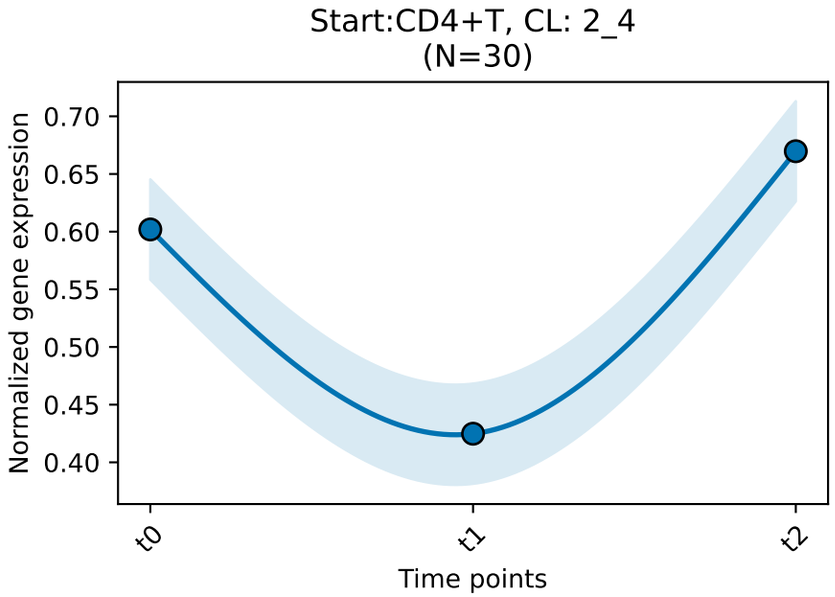
Start:CD4+T, CL: 2_0
(N=39)

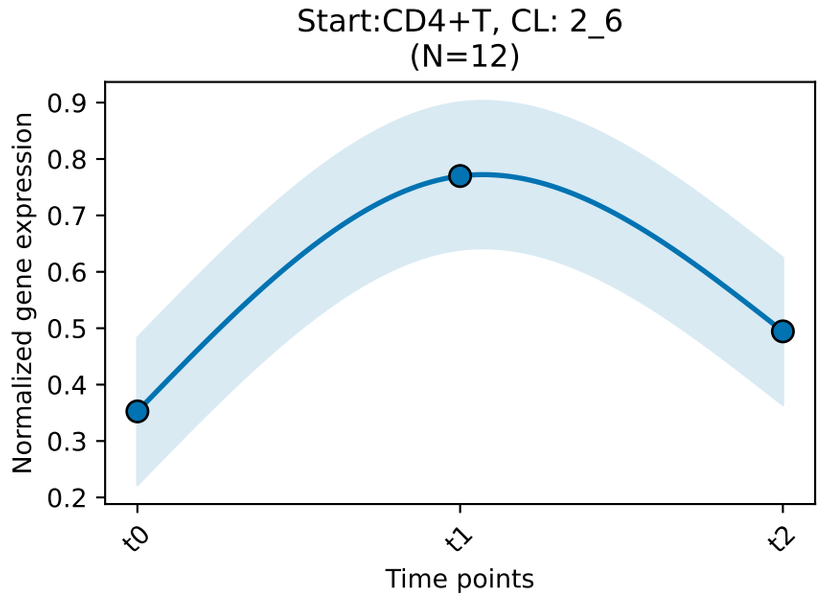


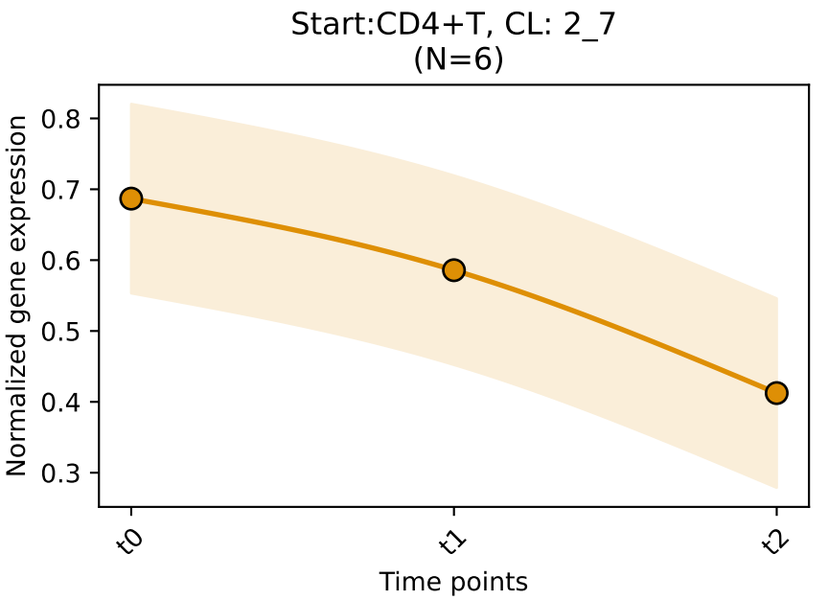
Start:CD4+T, CL: 2_1
(N=23)

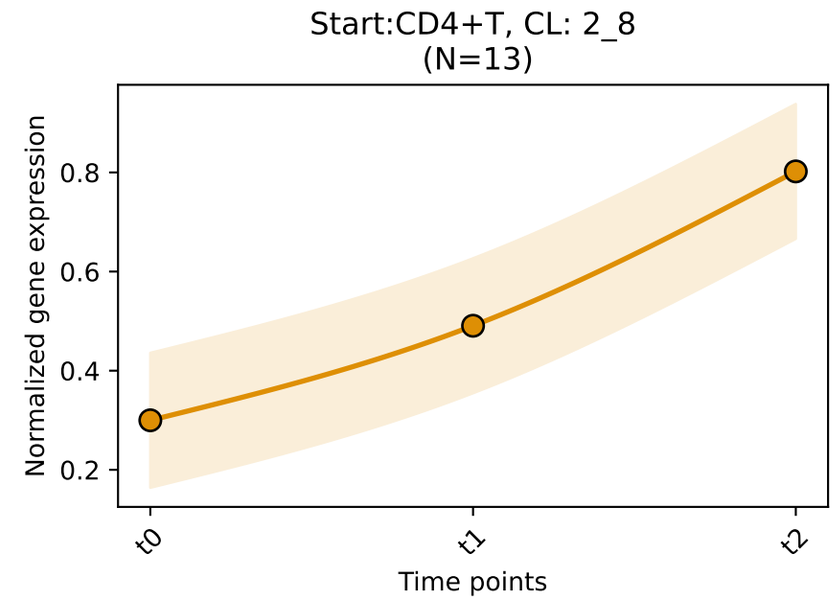


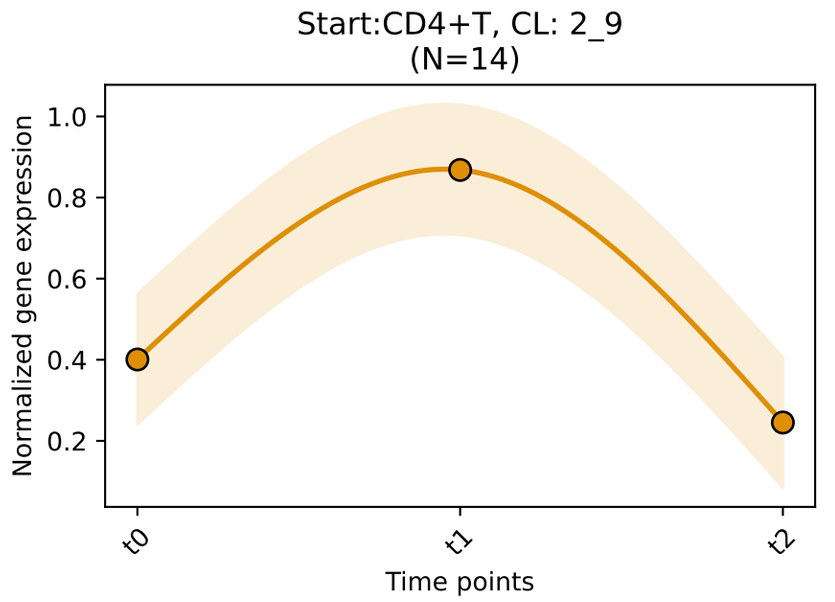

Start:CD4+T, CL: 2_3
(N=18)
Normalized gene expression
0.8
0.7
0.6
0.5
0.4
t0
t1
t2
Time points

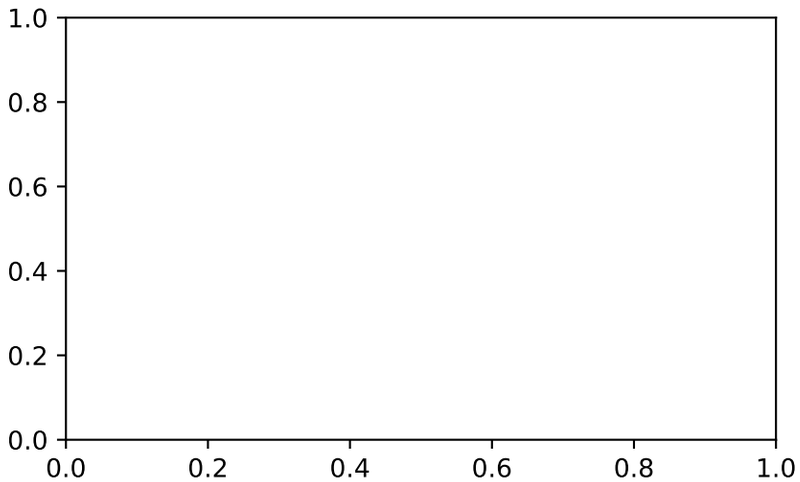

Start:CD4+T, CL: 2_4
(N=30)
Normalized gene expression
0.70
0.65
0.60
0.55
0.50
0.45
0.40
t0
t1
t2
Time points

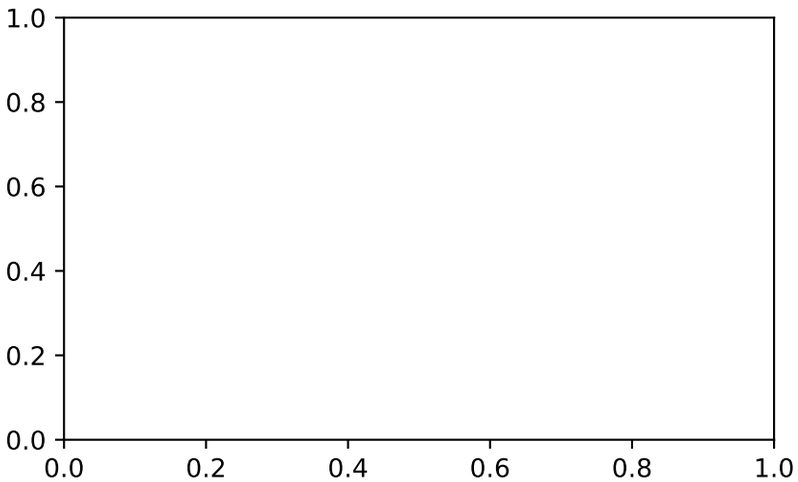

Start:CD4+T, CL: 2_6
(N=12)
Normalized gene expression
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
t0
t1
t2
Time points

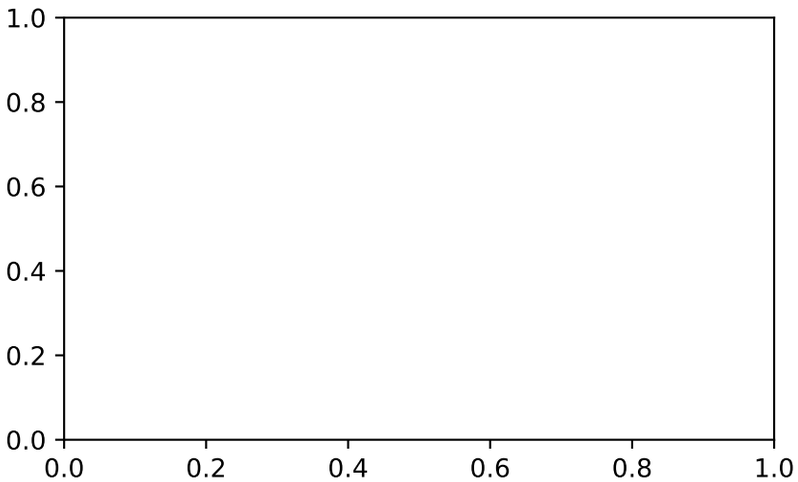

Start:CD4+T, CL: 2_7
(N=6)
gene expression
0.8
0.7
0.6
0.5
0.4
0.3
t0
t1
t2
Time points


Start:CD4+T, CL: 2_8
(N=13)
gene expression
0.8
0.6
0.4
0.2
t0
t1
t2
Time points


Start:CD4+T, CL: 2_9
(N=14)
gene expression
1.0
0.8
0.6
0.4
0.2
t0
t1
t2
Time points


1.0
0.8
0.6
0.4
0.2
0.0
0.0
0.2
0.4
0.6
0.8
1.0


1.0
0.8
0.6
0.4
0.2
0.0
0.0
0.2
0.4
0.6
0.8
1.0


1.0
0.8
0.6
0.4
0.2
0.0
0.0
0.2
0.4
0.6
0.8
1.0