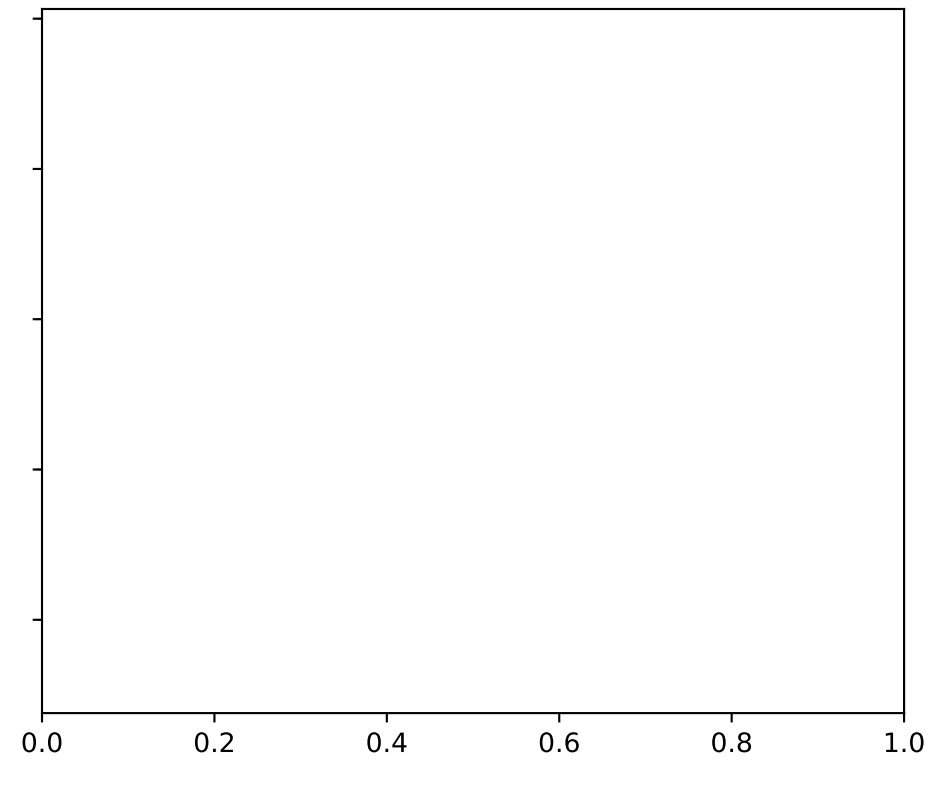
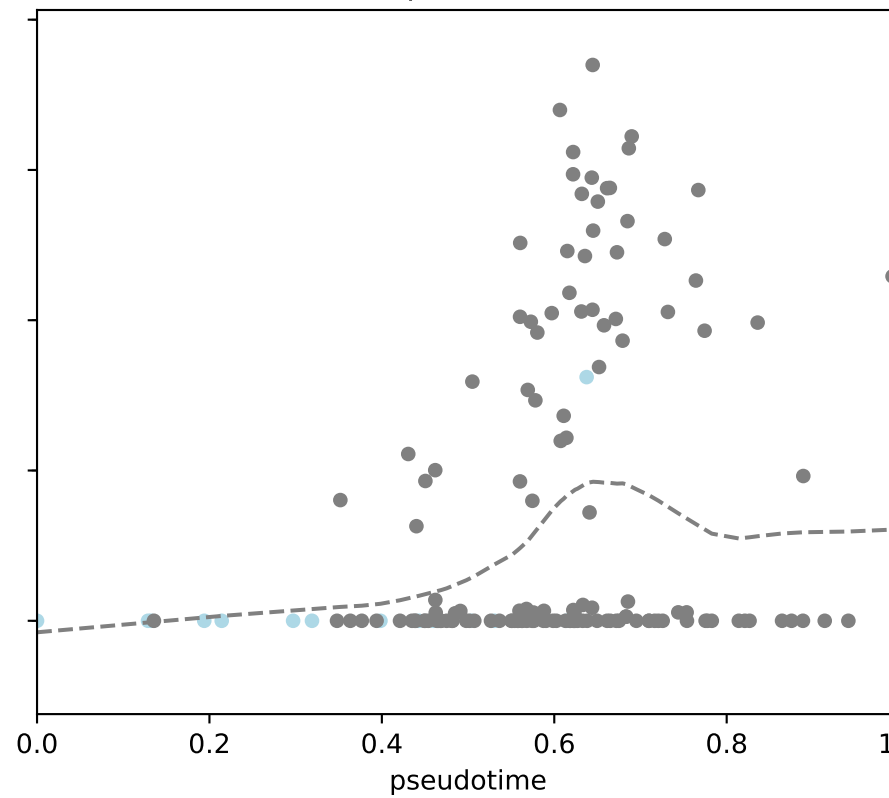
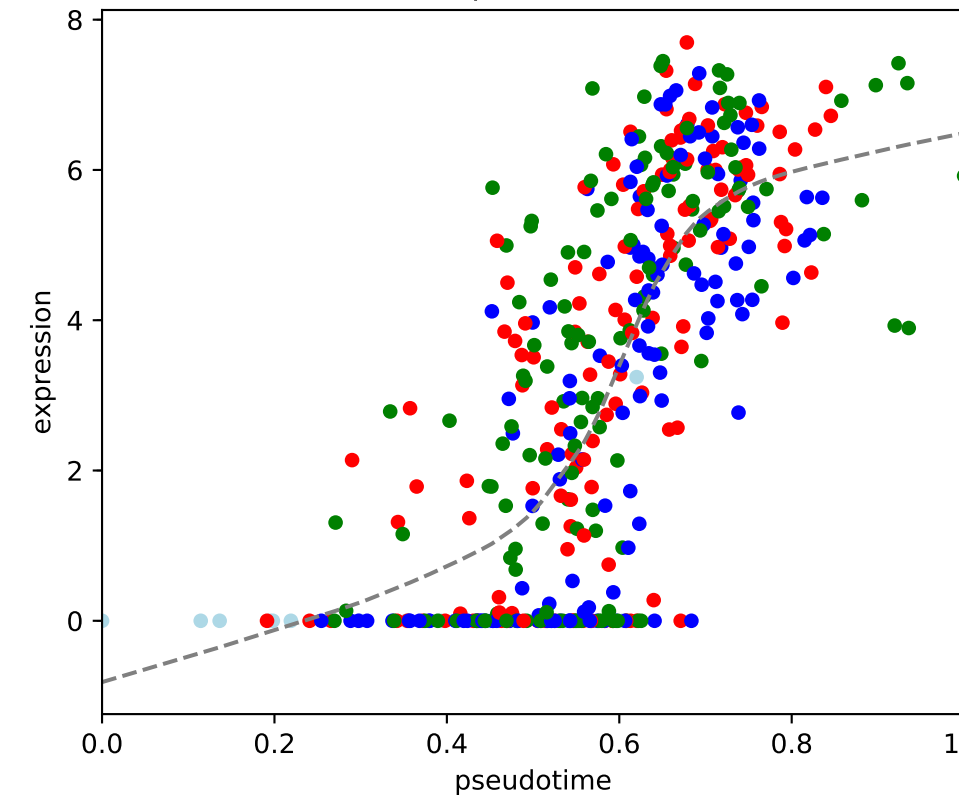


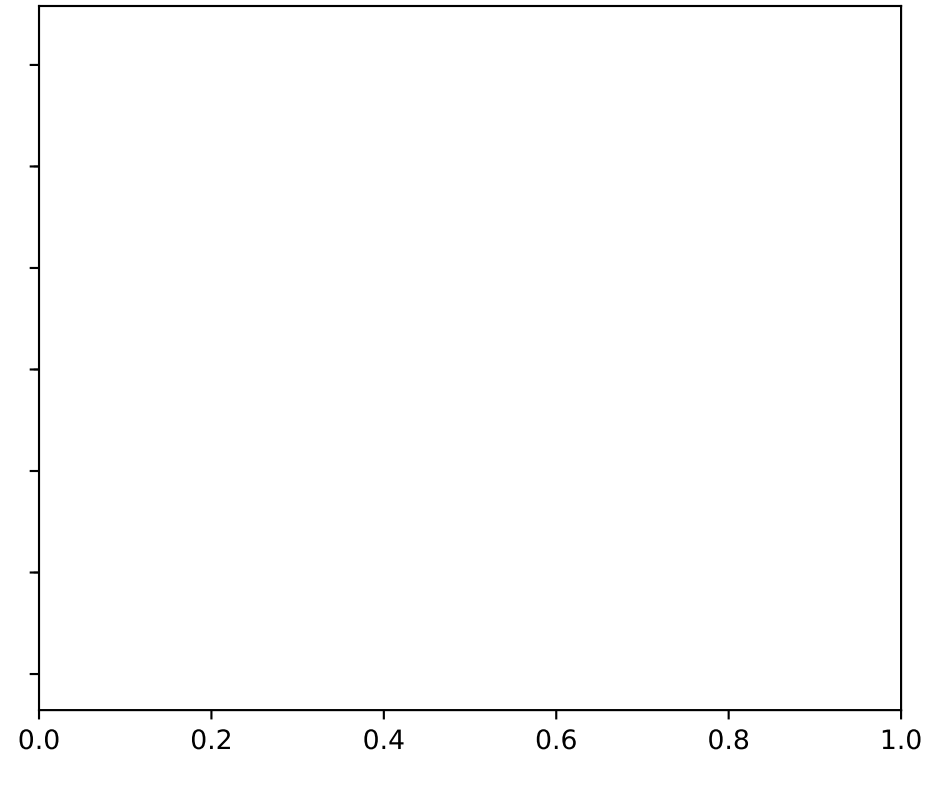
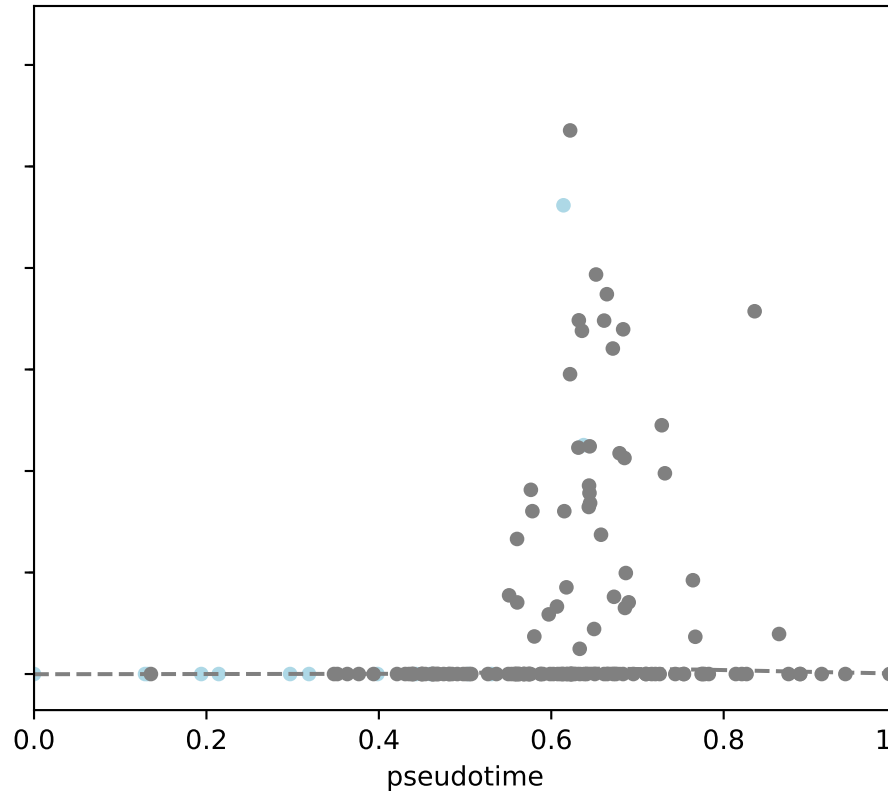
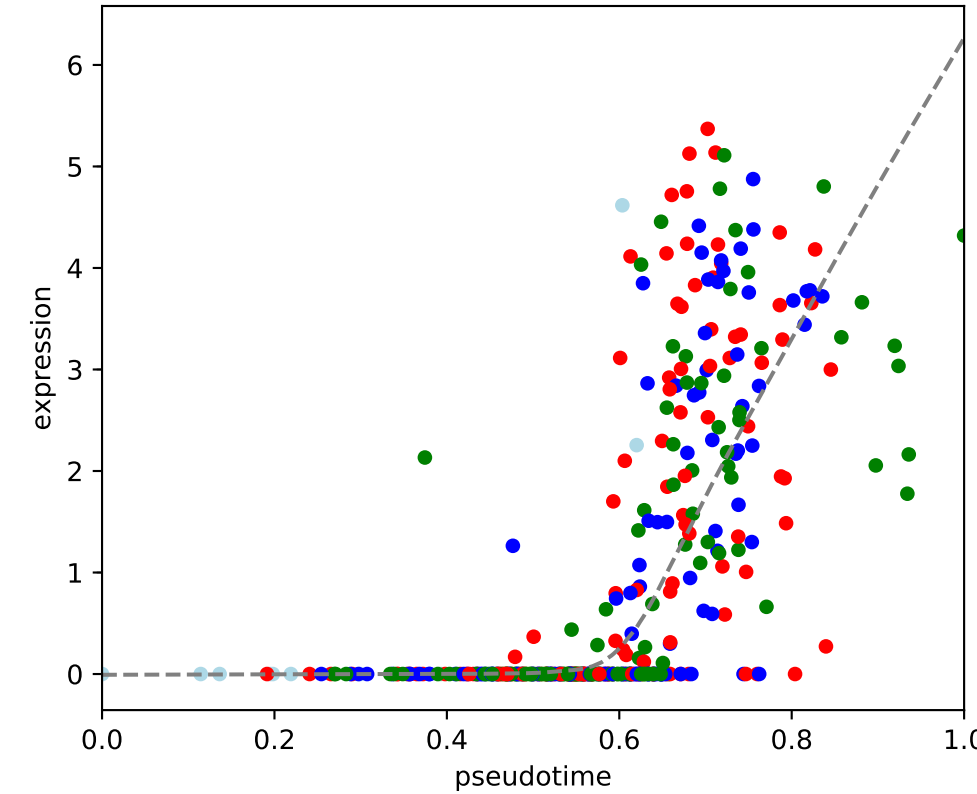
733 0spearman=0.74

733 1spearman=0.30



733 0spearman=0.74

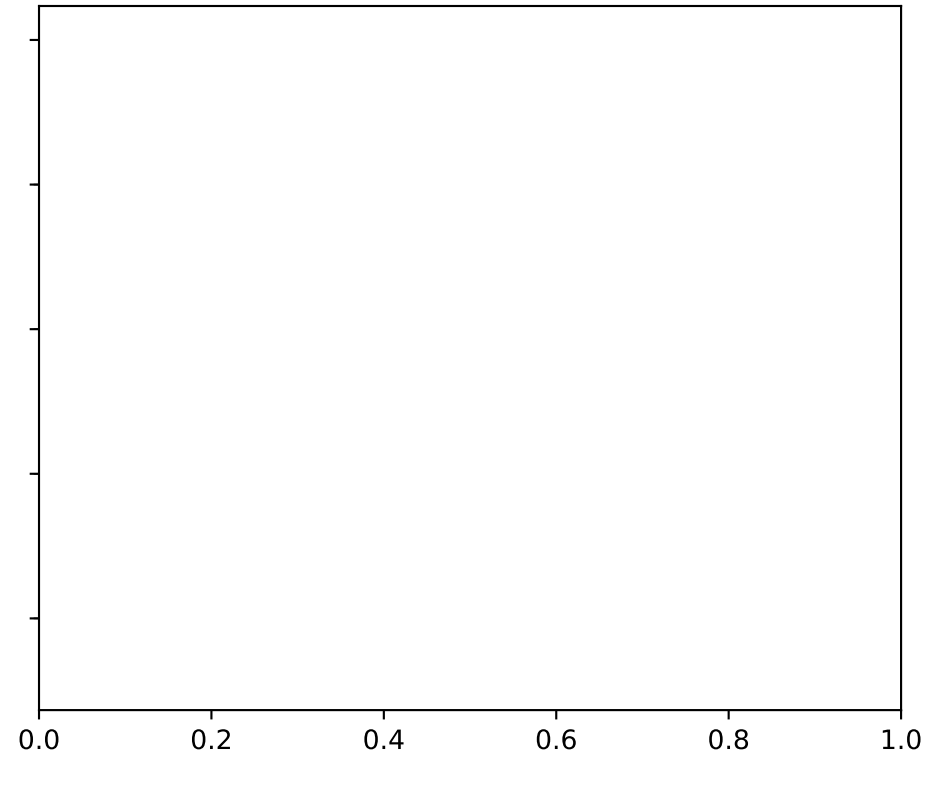
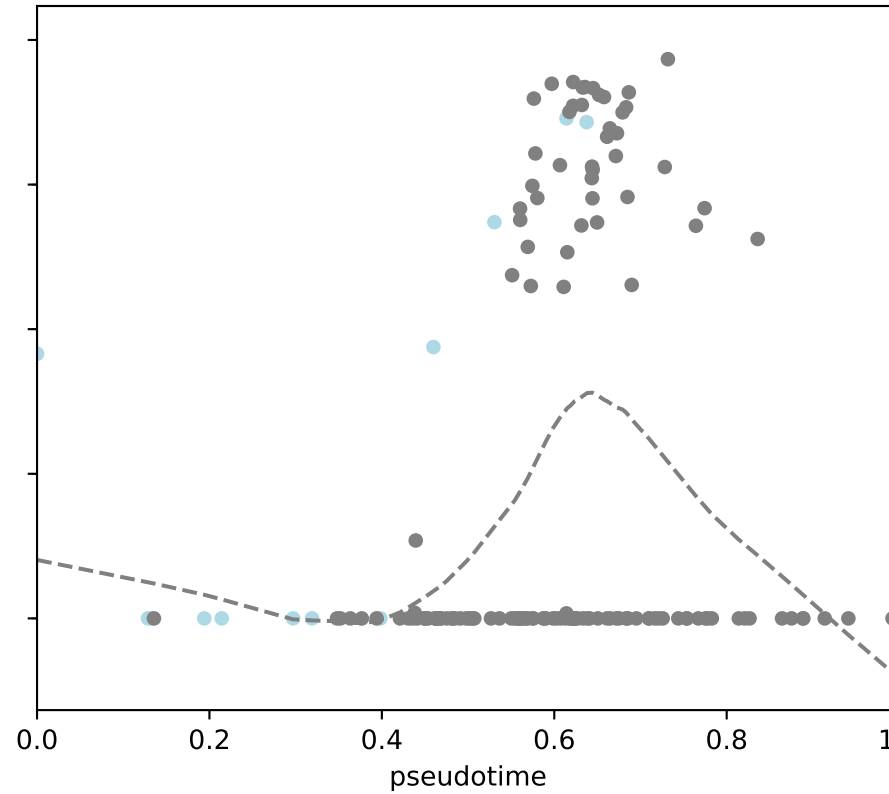
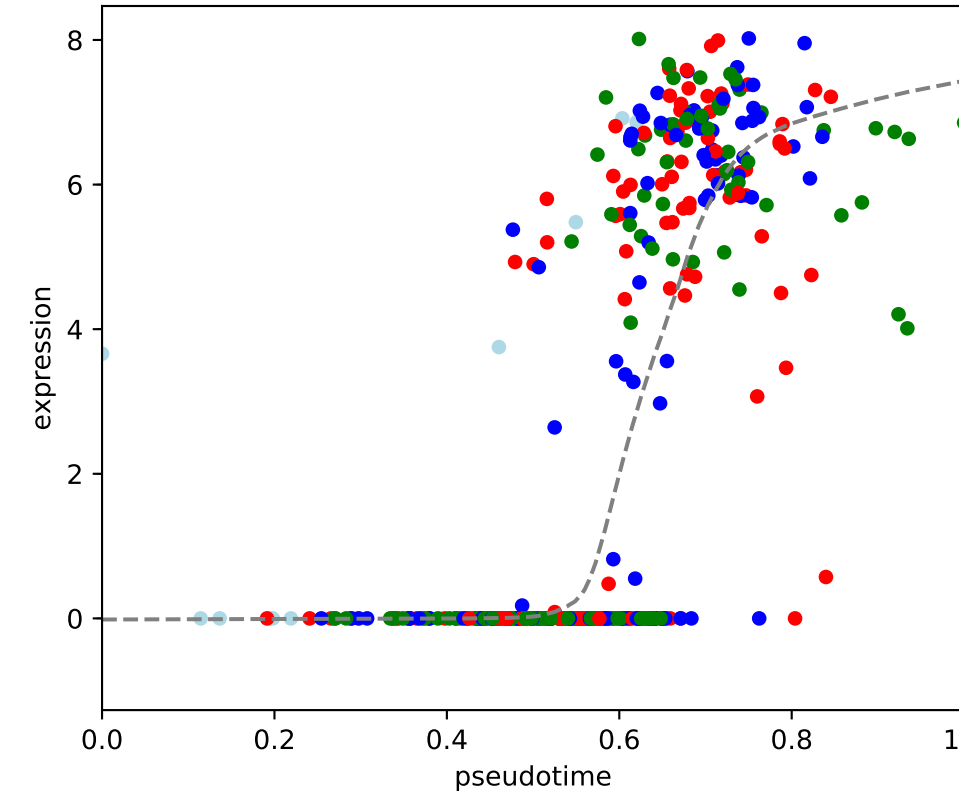
733 1spearman=0.33



ENST00000617575 NR2E3 (nuclear receptor subfamily 2 group E member 3)

733 0spearman=0.73

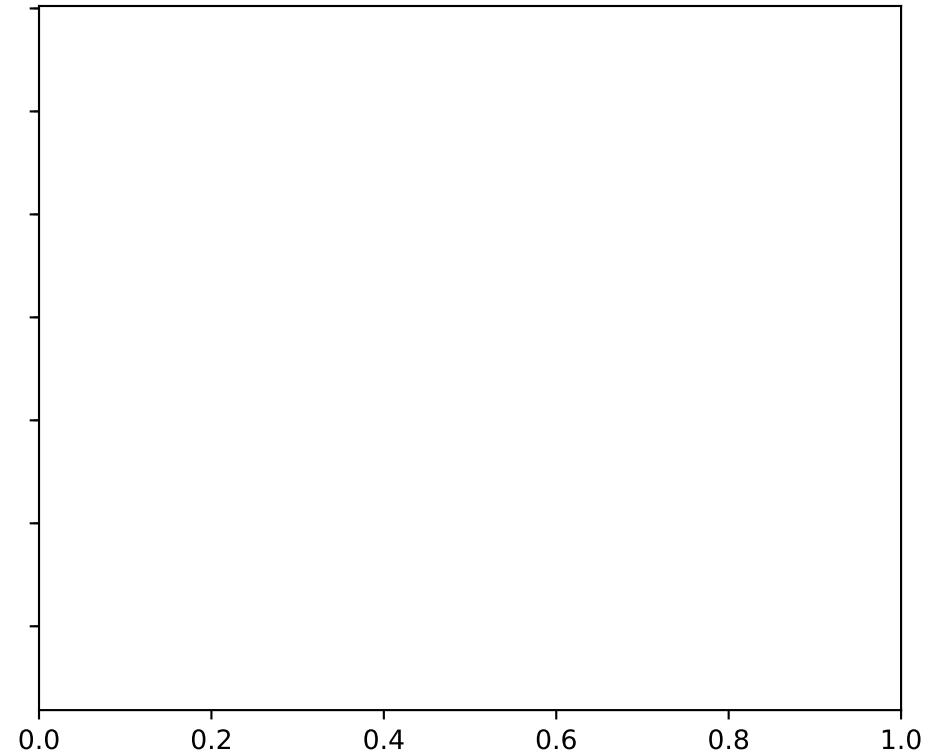
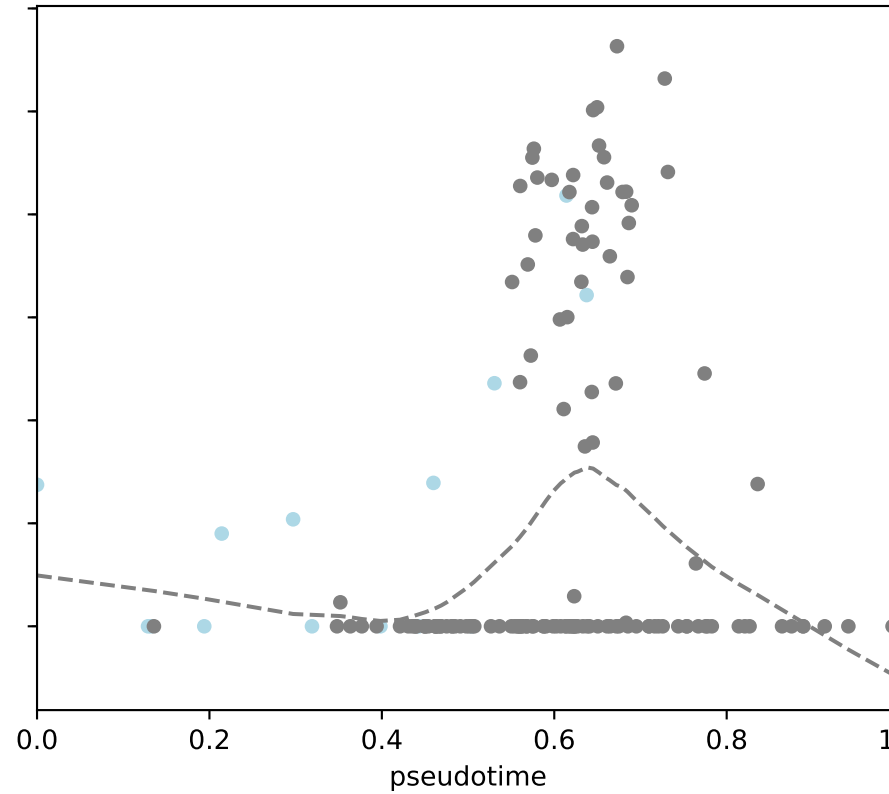
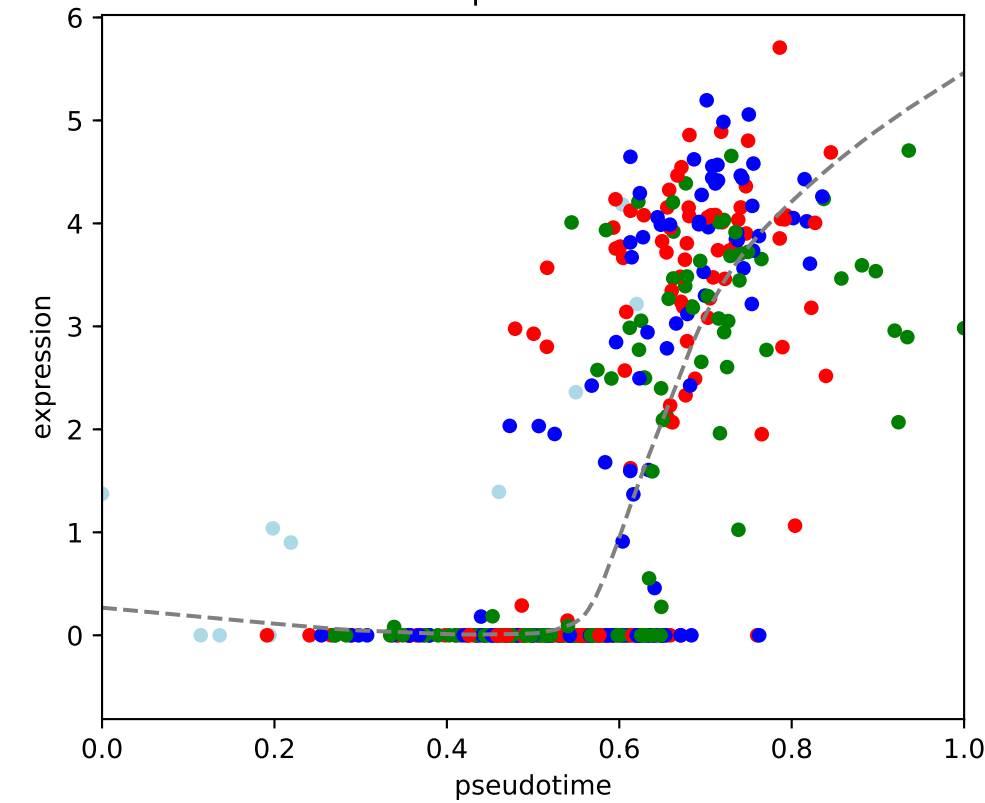
733 1spearman=0.23



ENST00000558280 NRL (neural retina leucine zipper)

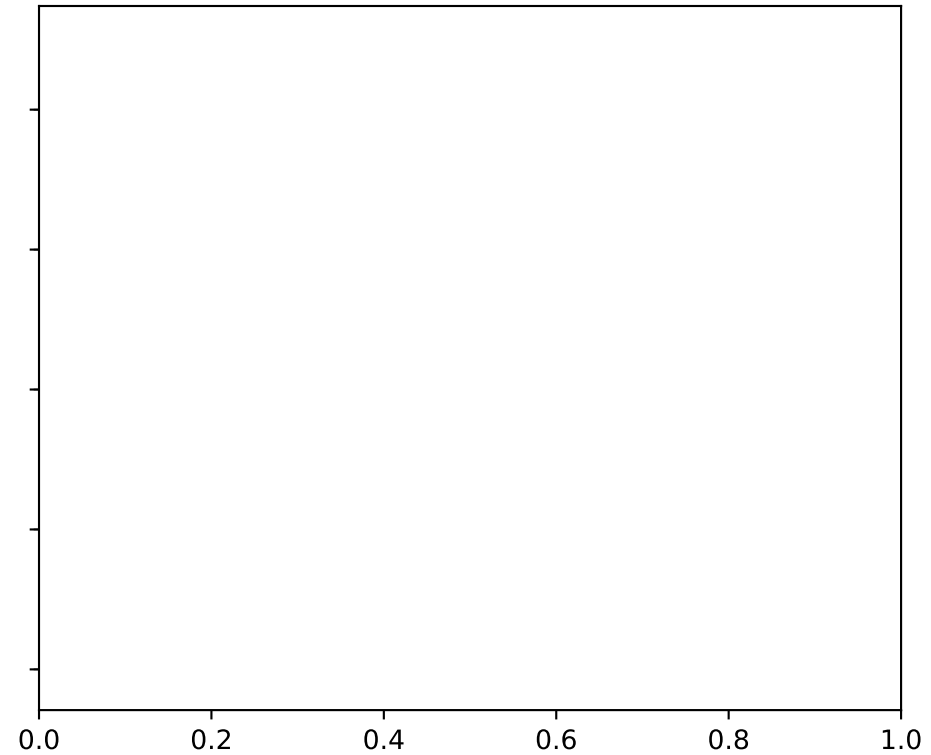
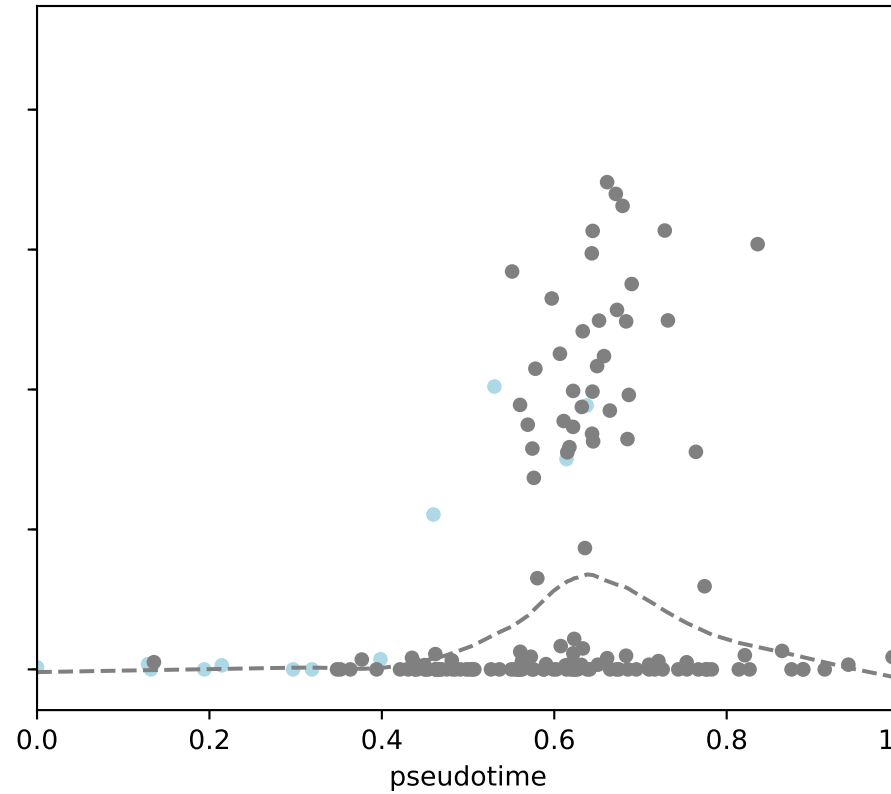
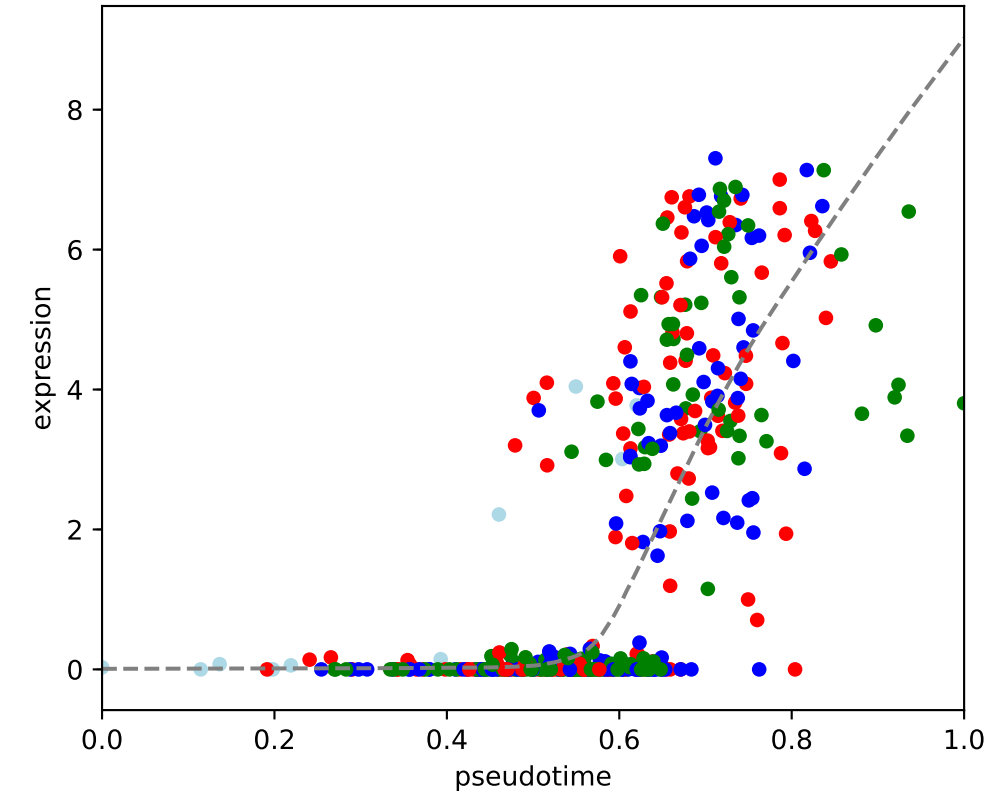
733 0spearman=0.72

733 1spearman=0.20



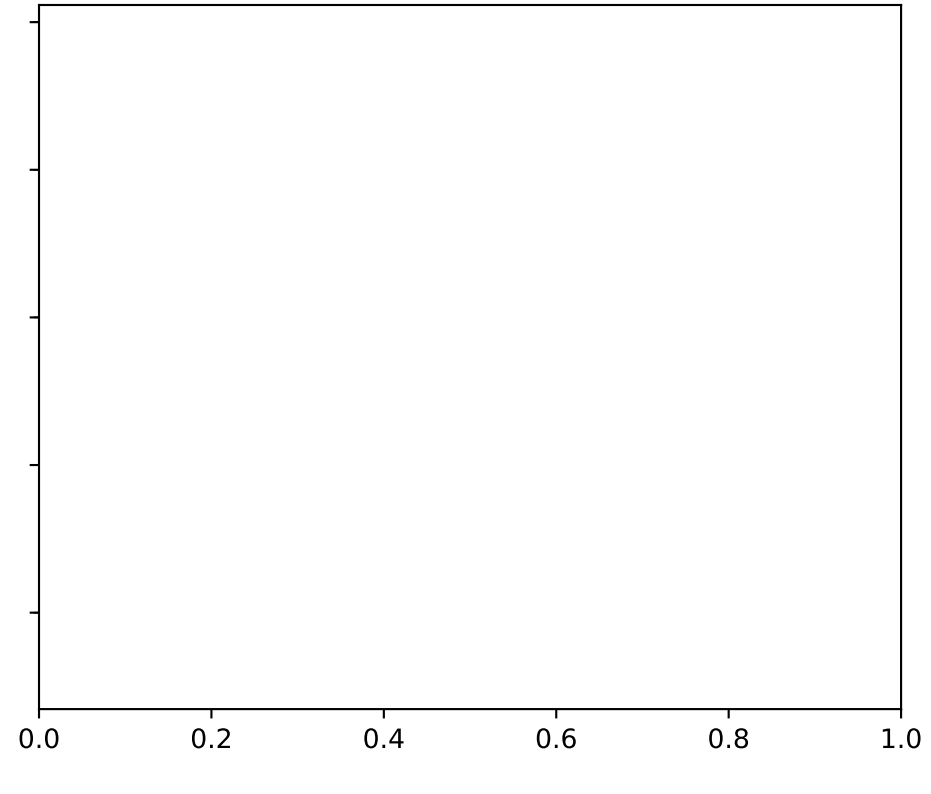
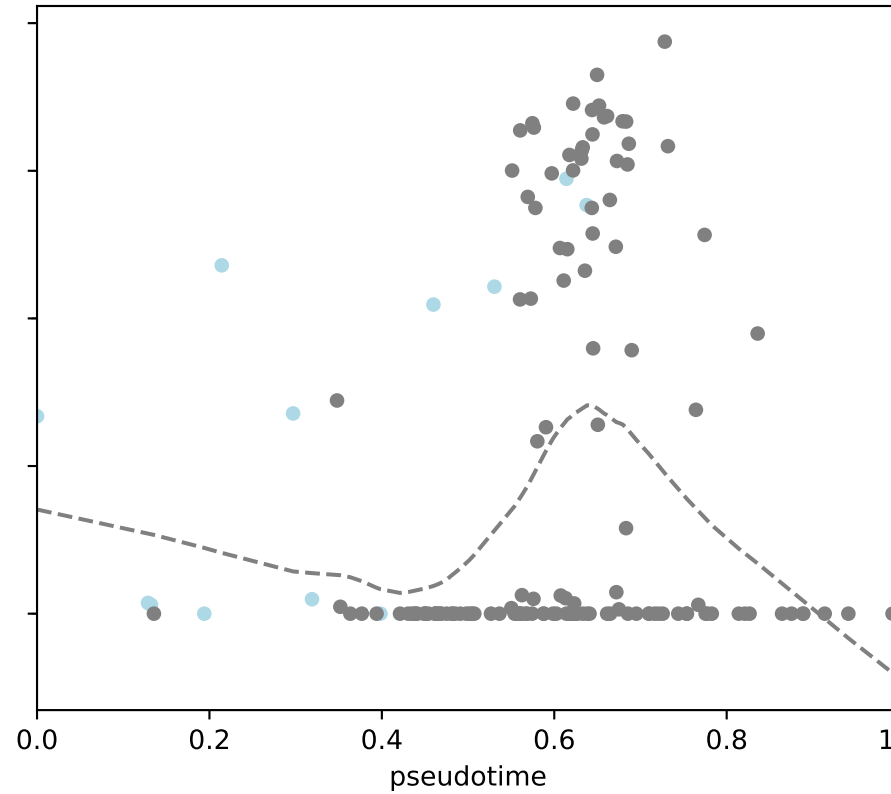
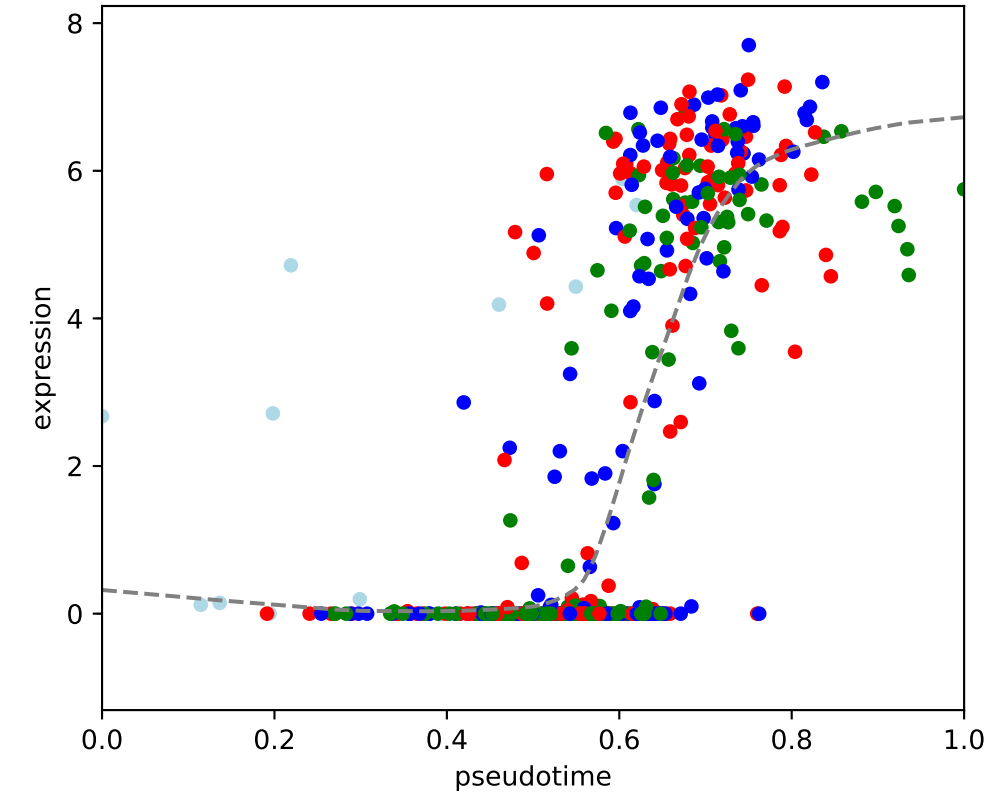
733 0spearman=0.71

733 1spearman=0.28



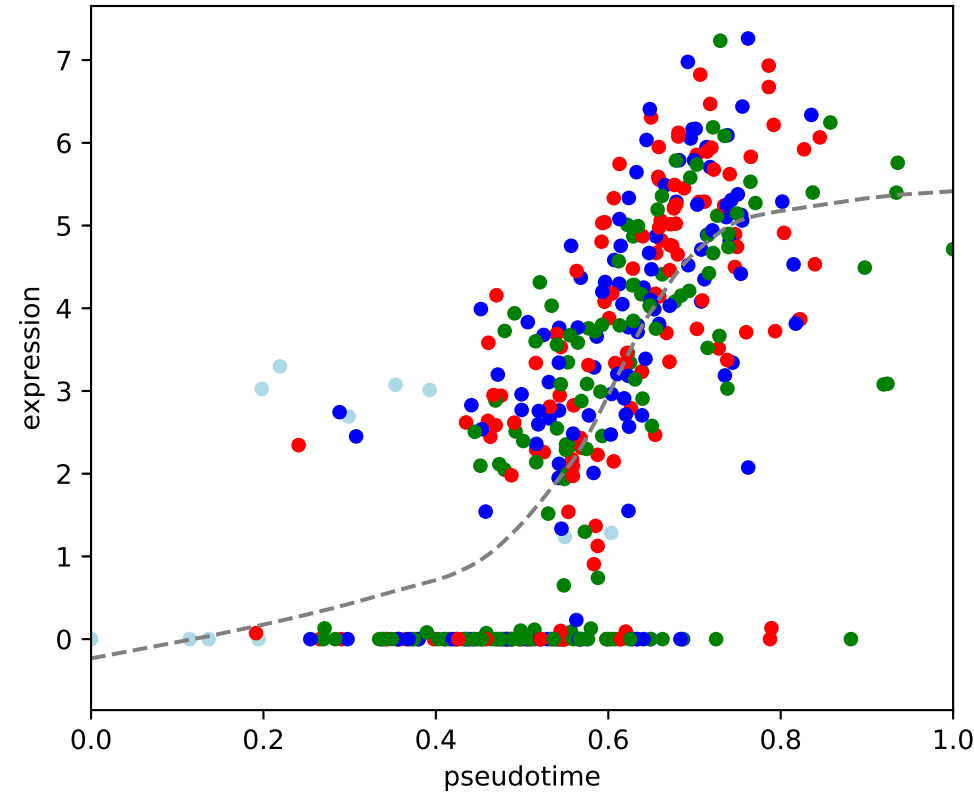
733 0spearman=0.71

733 1spearman=0.16

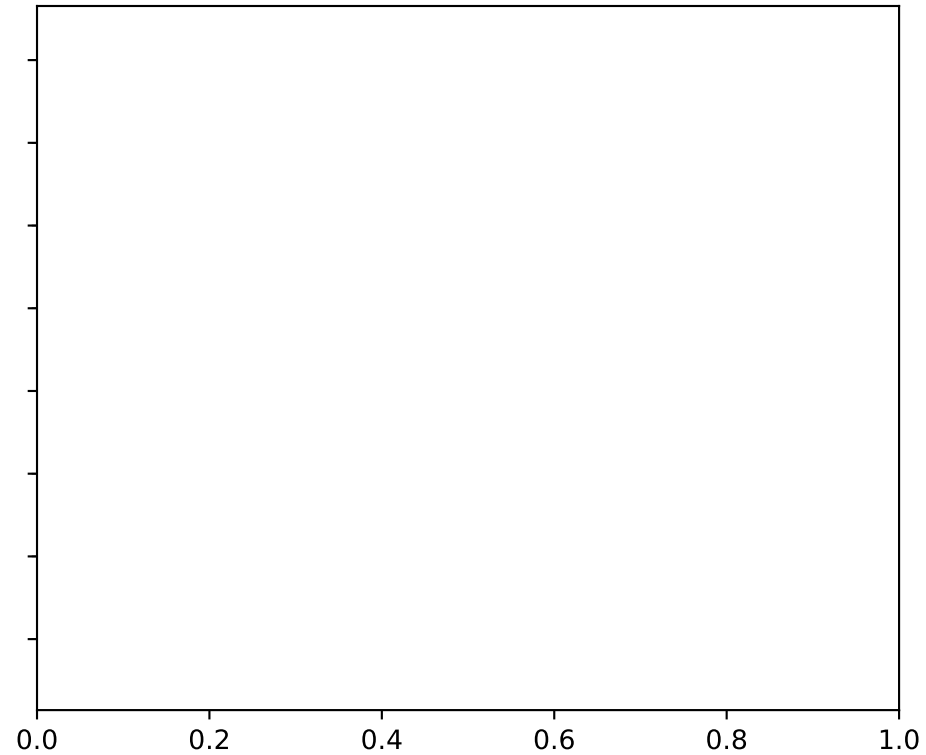
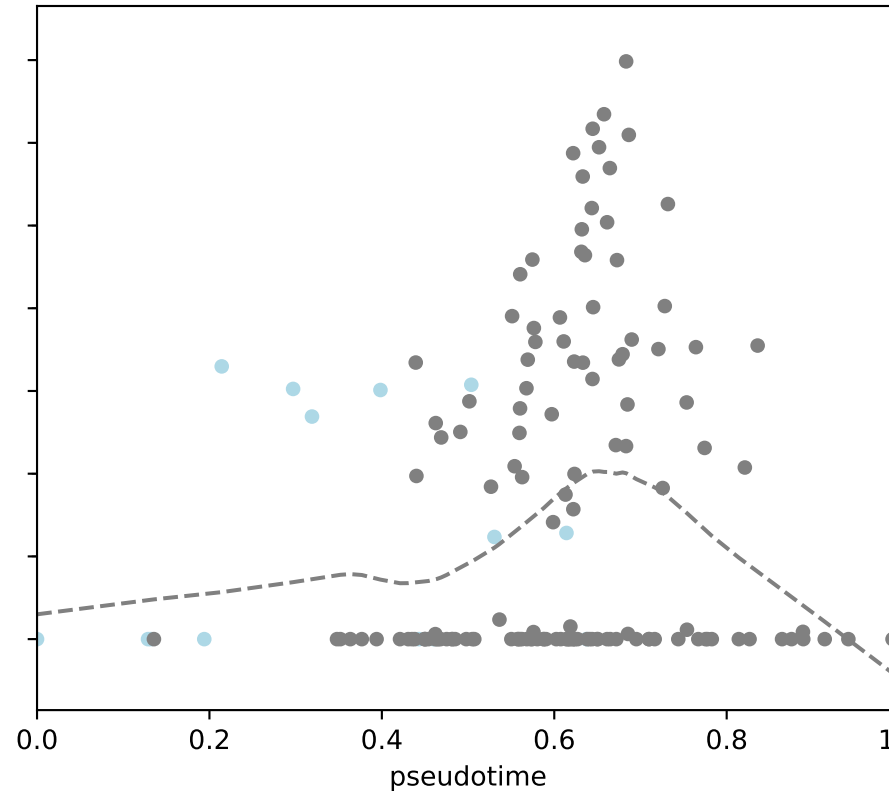


ENST00000311183 CNGB1 (cyclic nucleotide gated channel beta 1)

733 0spearman=0.71



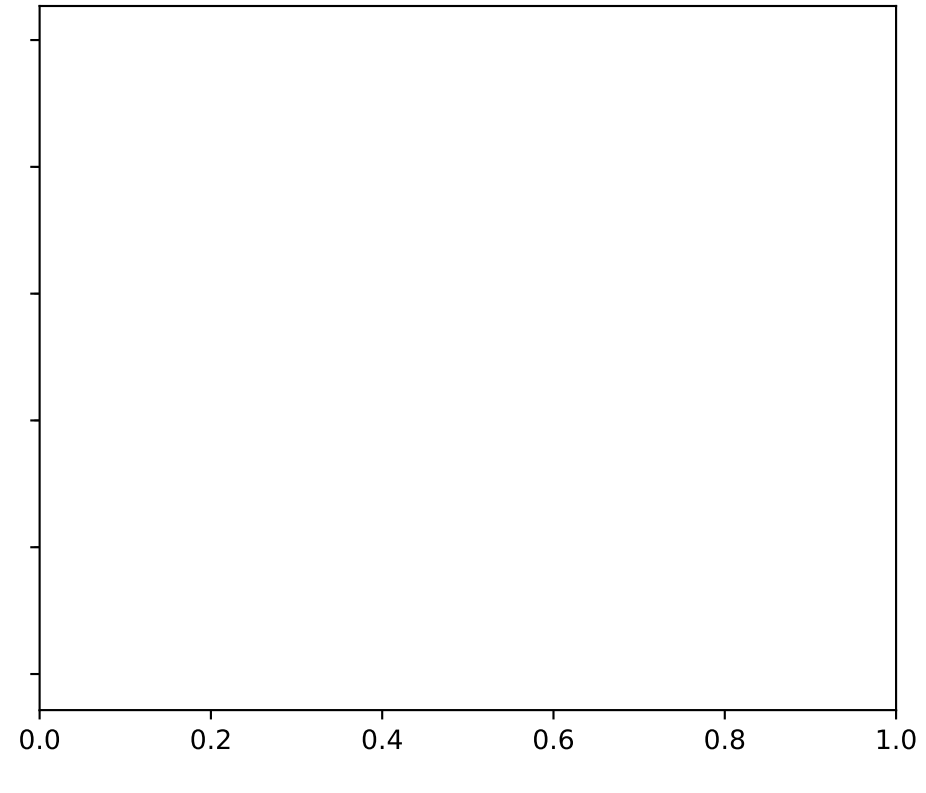
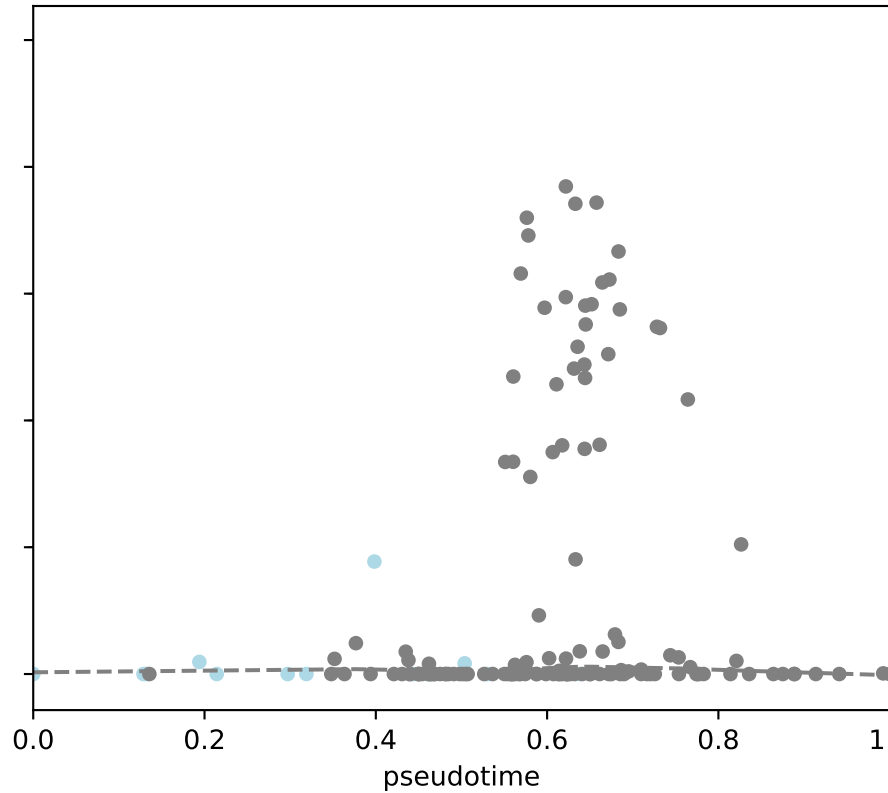
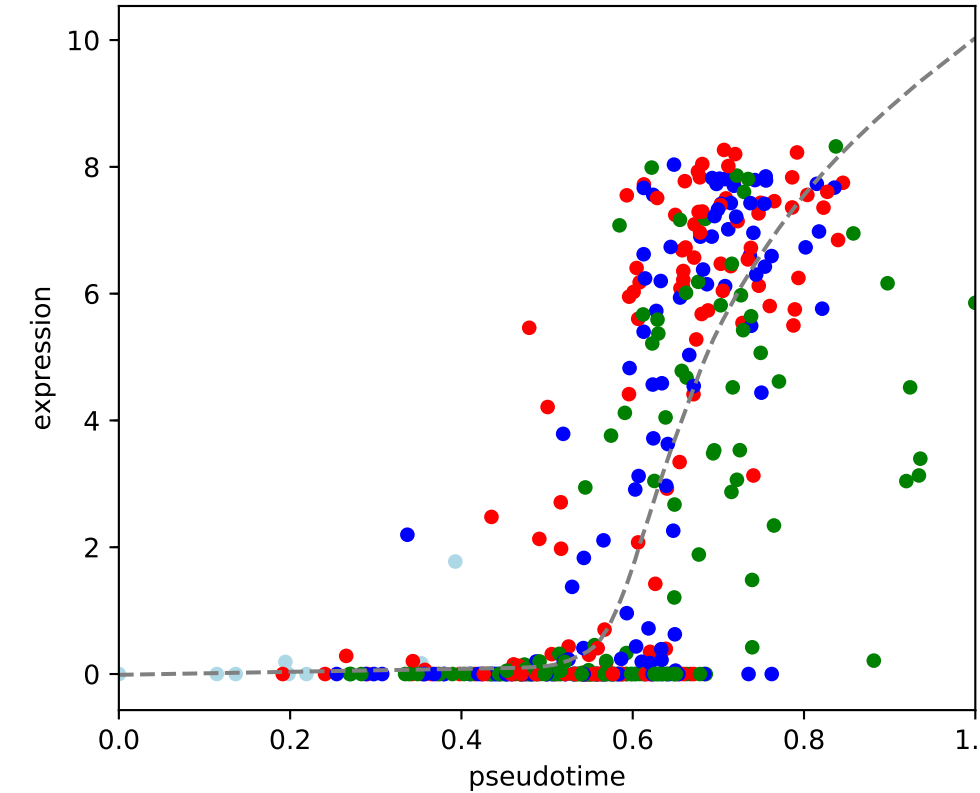
733 1spearman=0.18



ENST00000248572 GNGT1 (G protein subunit gamma transducin 1)

733 0spearman=0.70

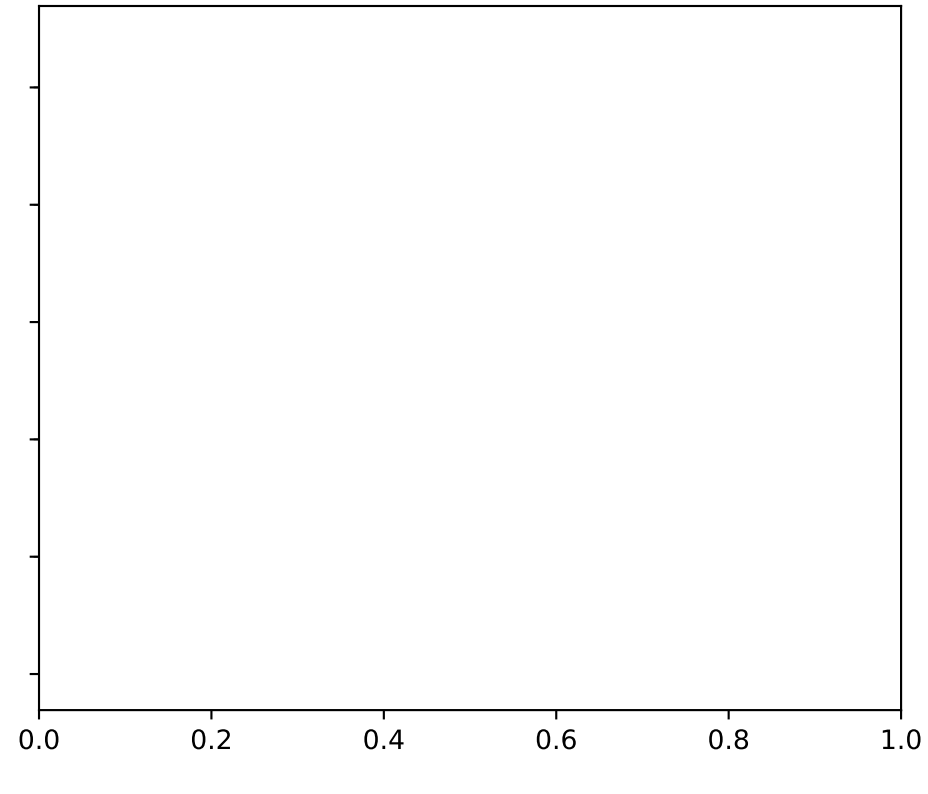
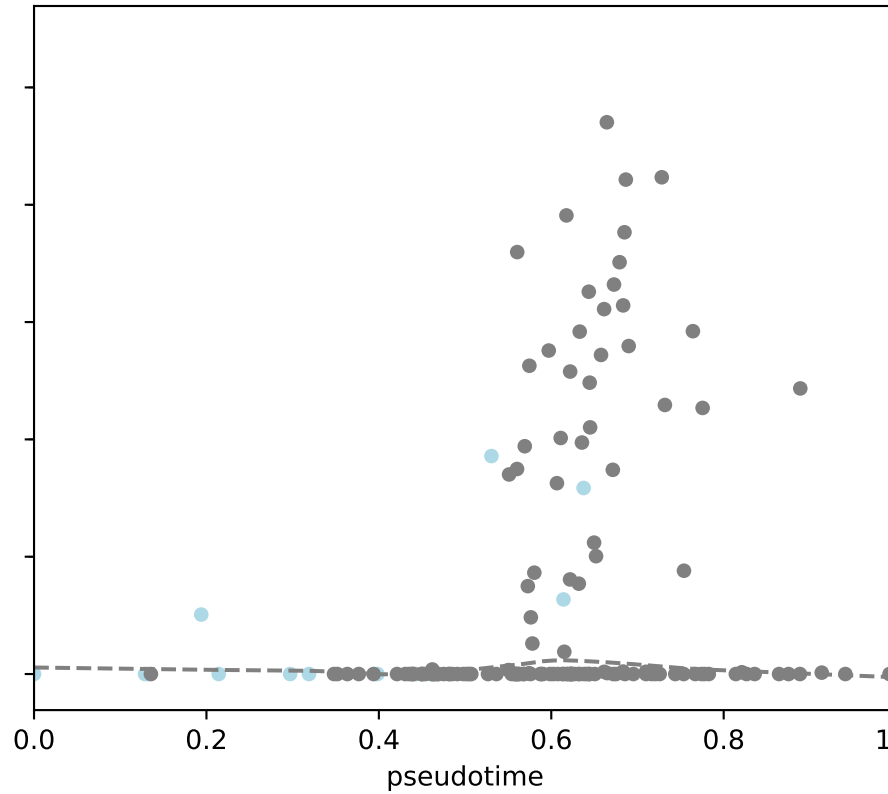
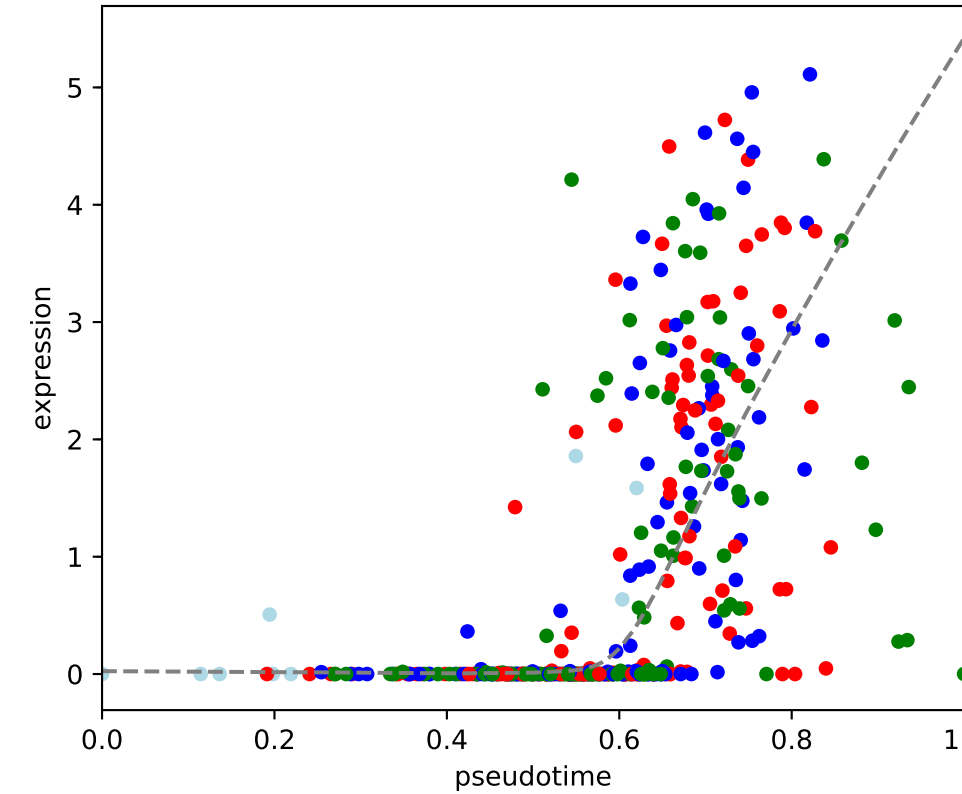
733 1spearman=0.22



ENST00000393386 PTPRZ1 (protein tyrosine phosphatase, receptor type Z1)

733 0spearman=0.70

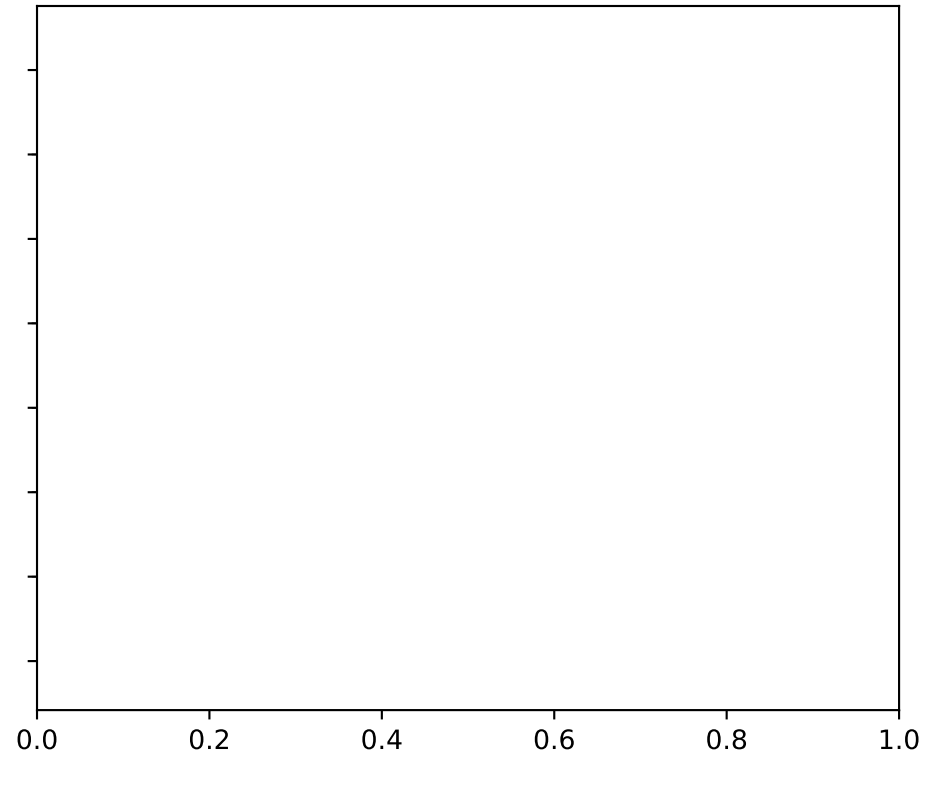
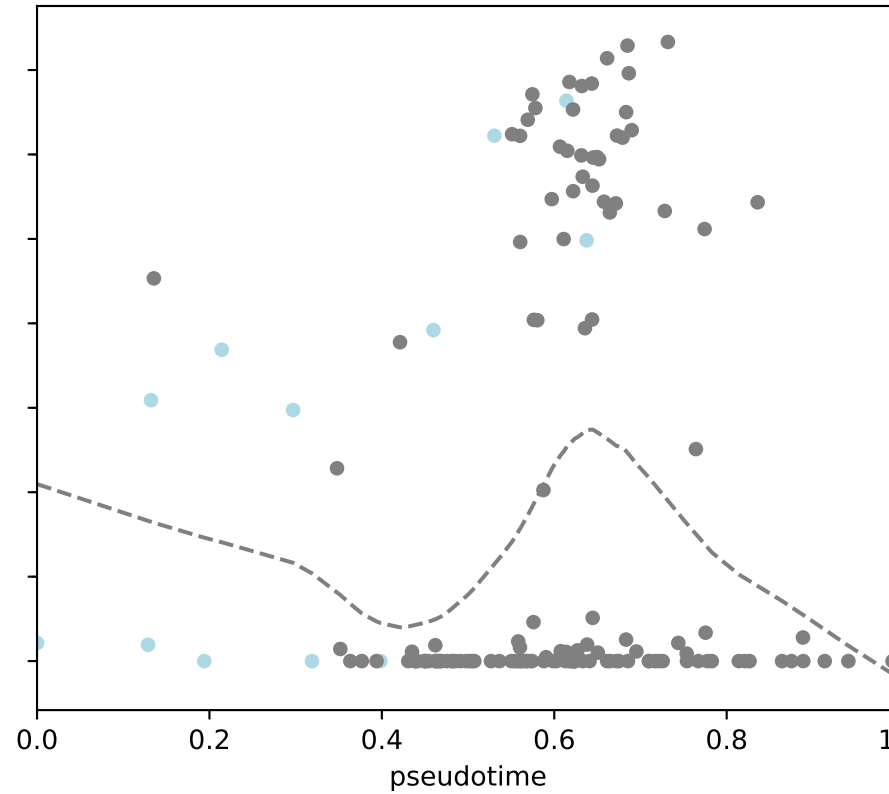
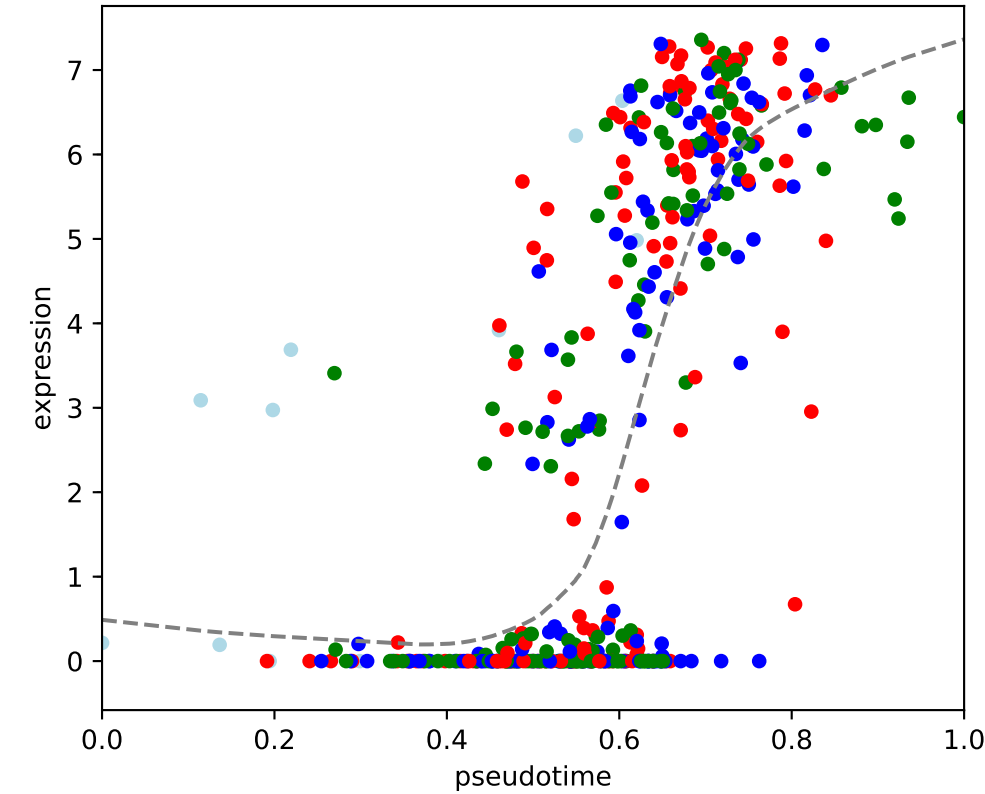
733 1spearman=0.31



ENST00000293255 CABP5 (calcium binding protein 5)

733 0spearman=0.69

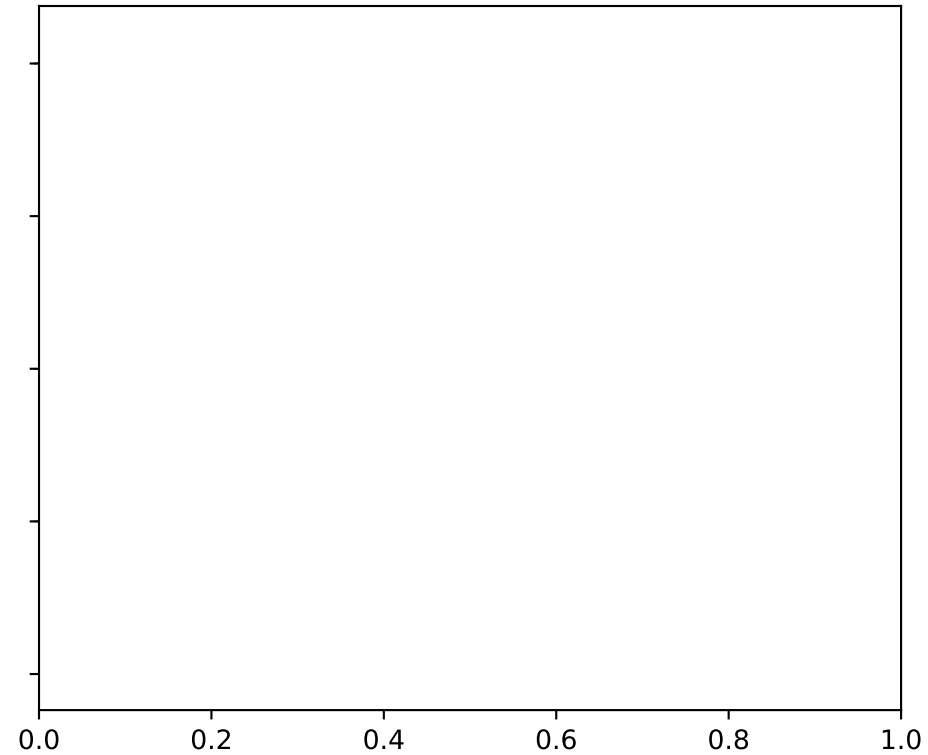
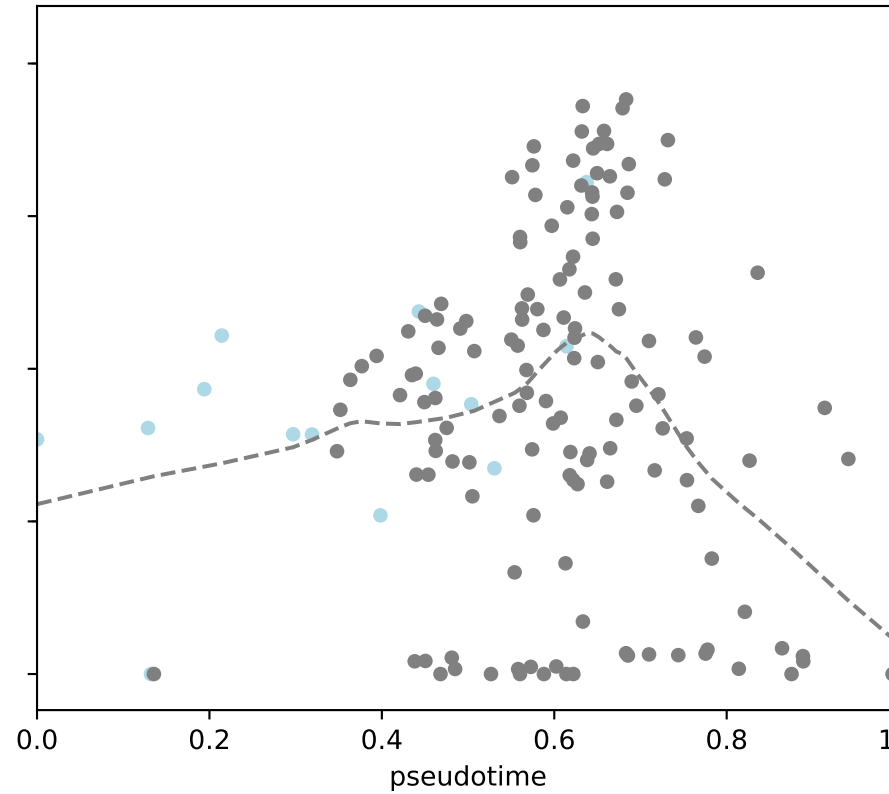
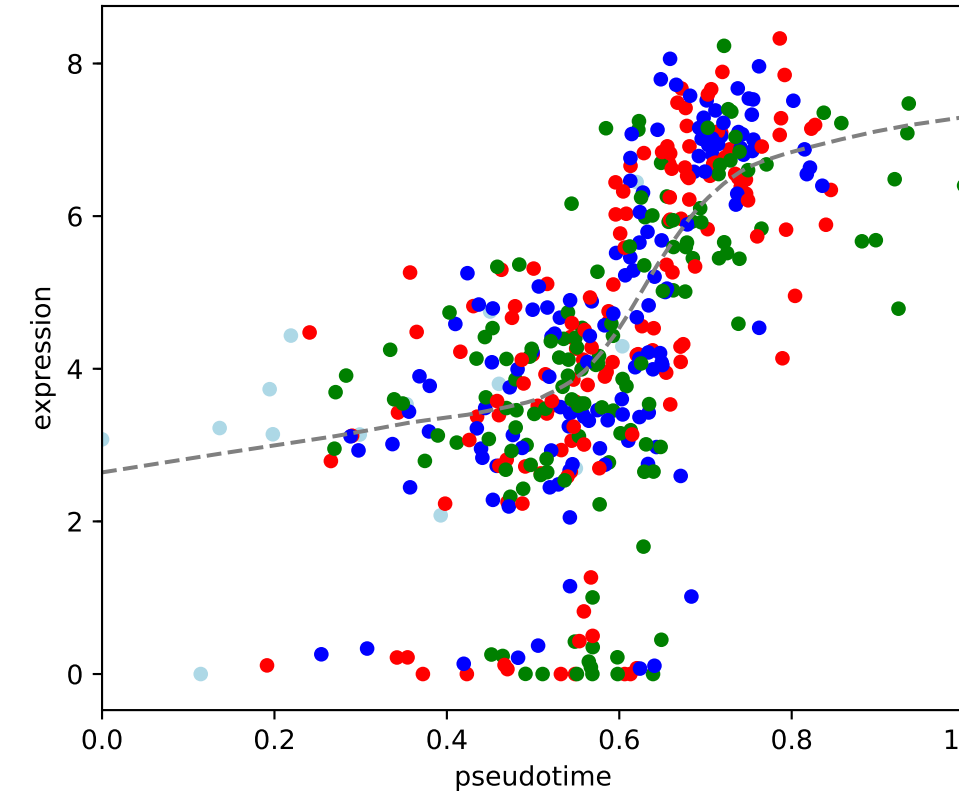
733 1spearman=0.17



ENST00000278833 ROM1 (retinal outer segment membrane protein 1)

733 0spearman=0.69

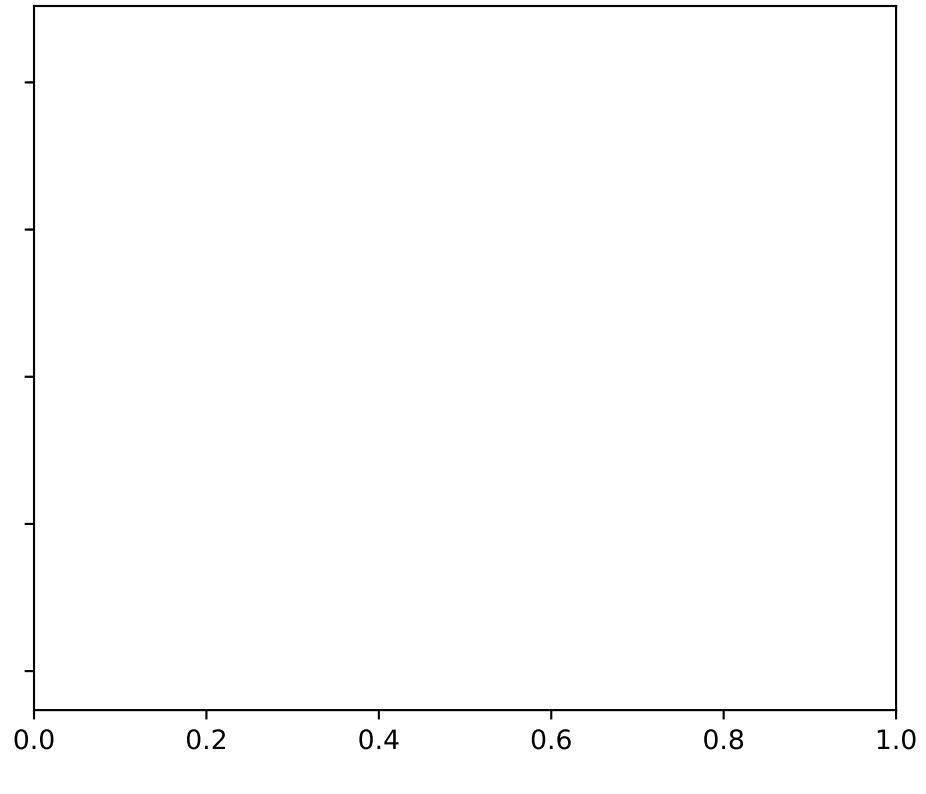
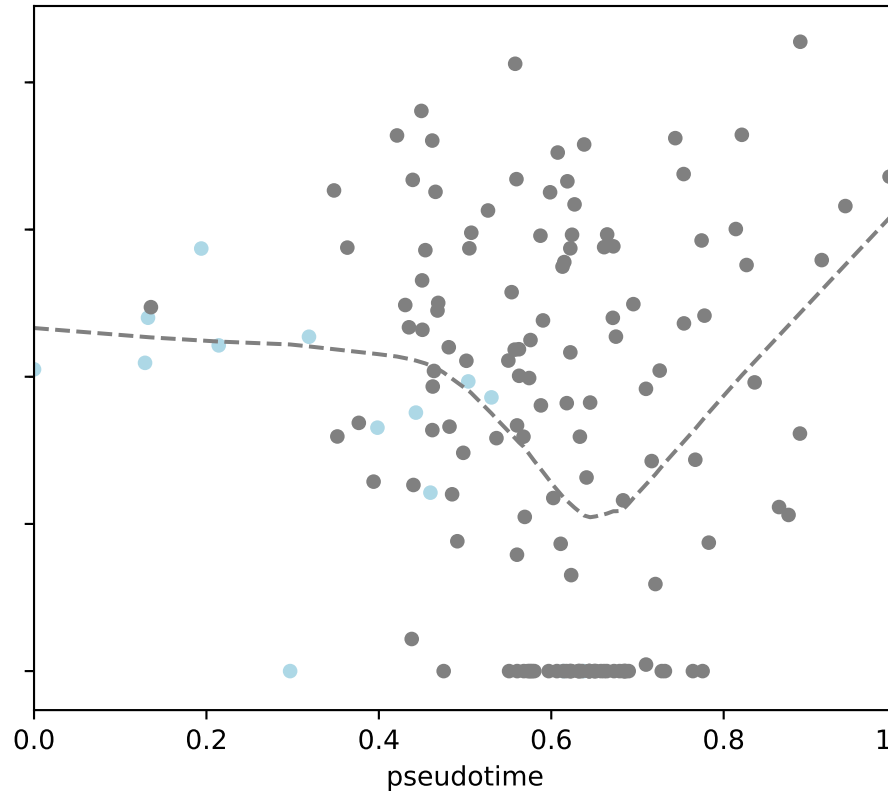
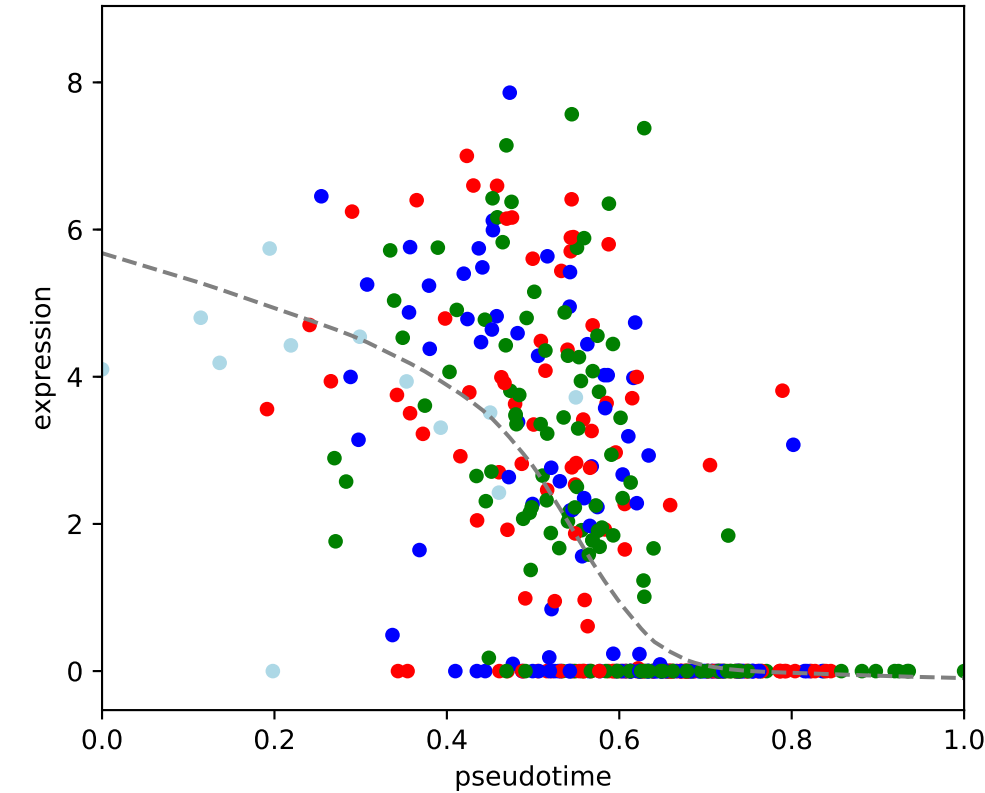
733 1spearman=0.06



ENST00000371464 RBP4 (retinol binding protein 4)

733 0spearman=-0.69

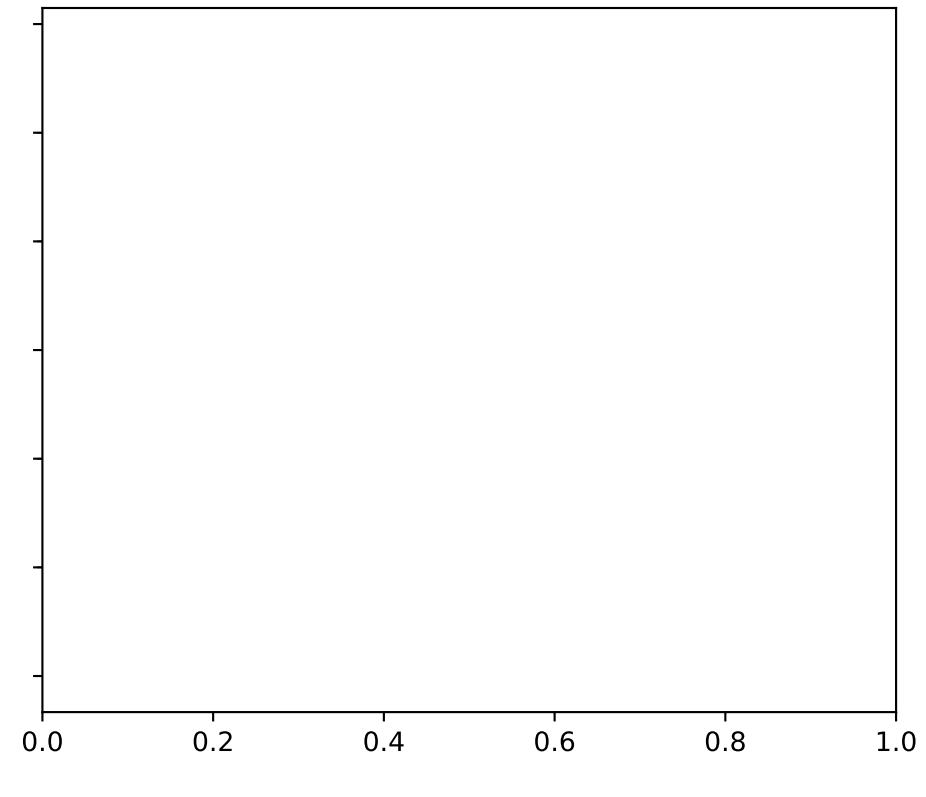
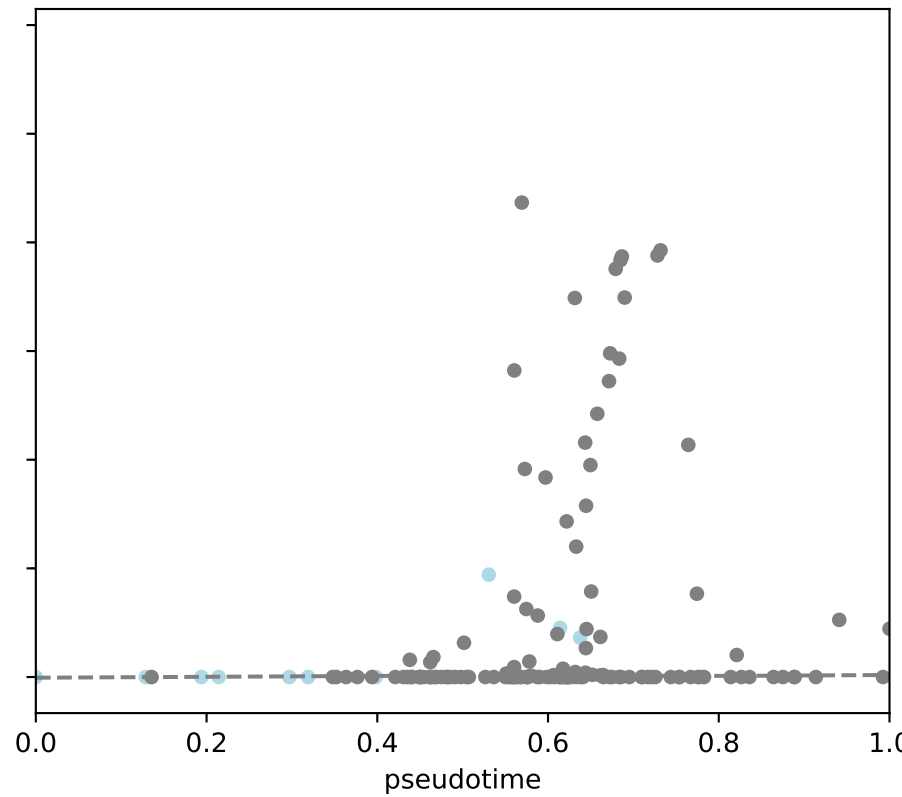
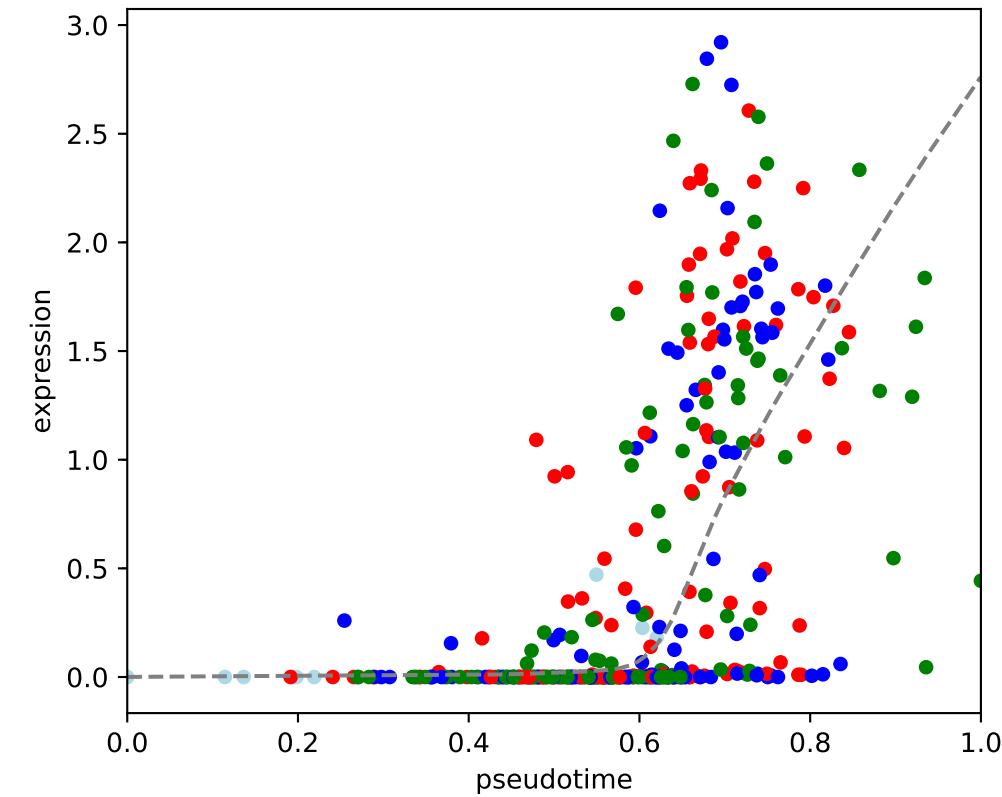
733 1spearman=-0.16



ENST00000628562 MRLN (myoregulin)

733 0spearman=0.69

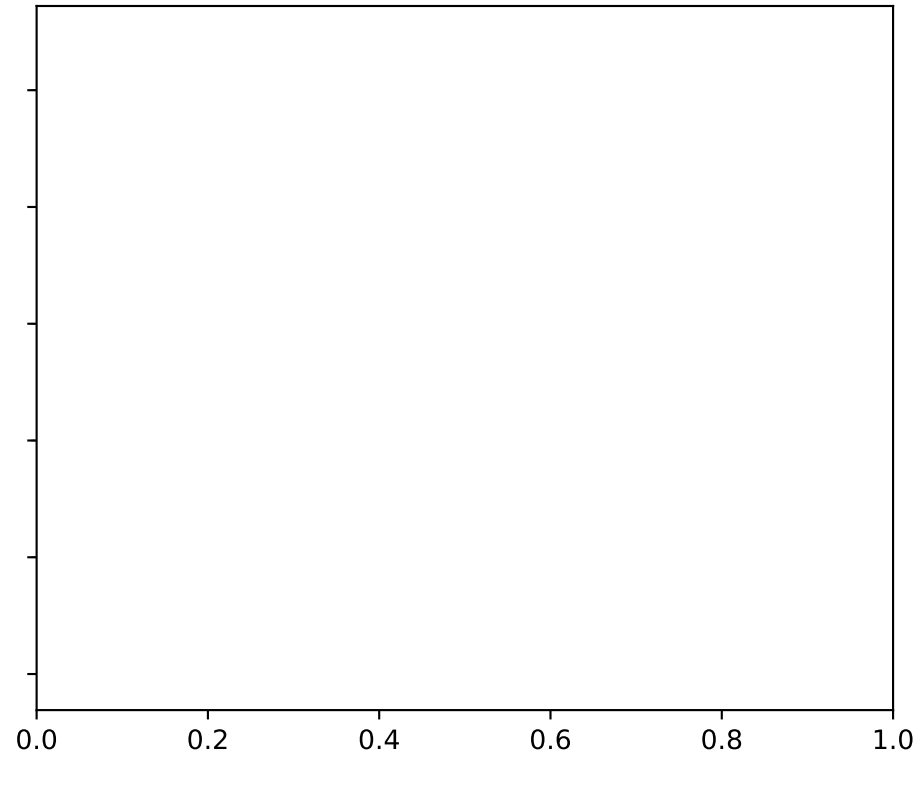
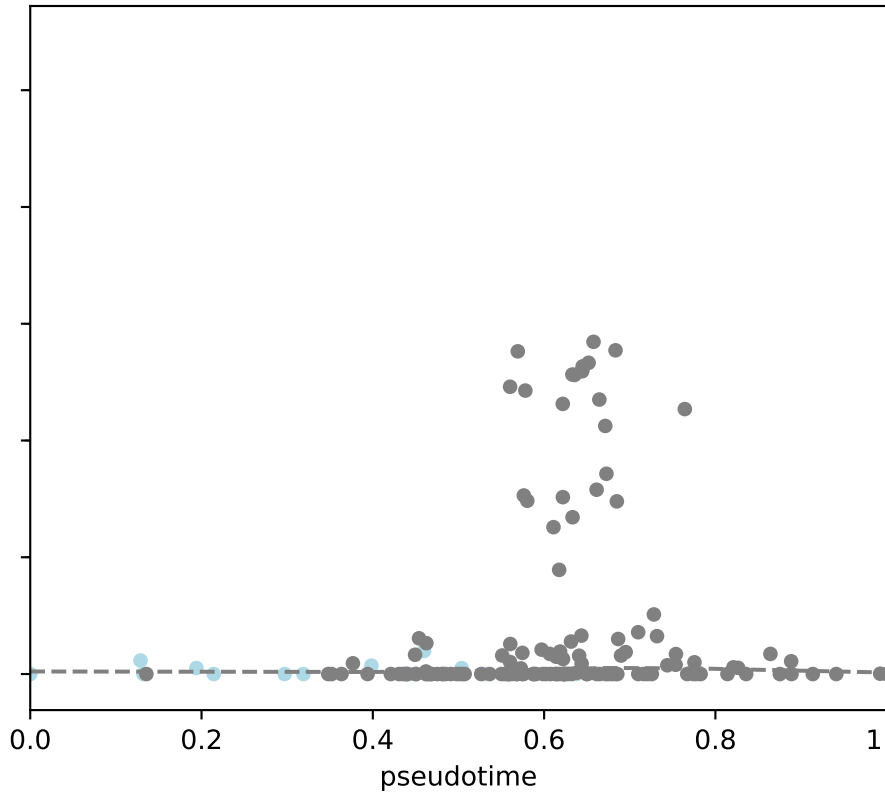
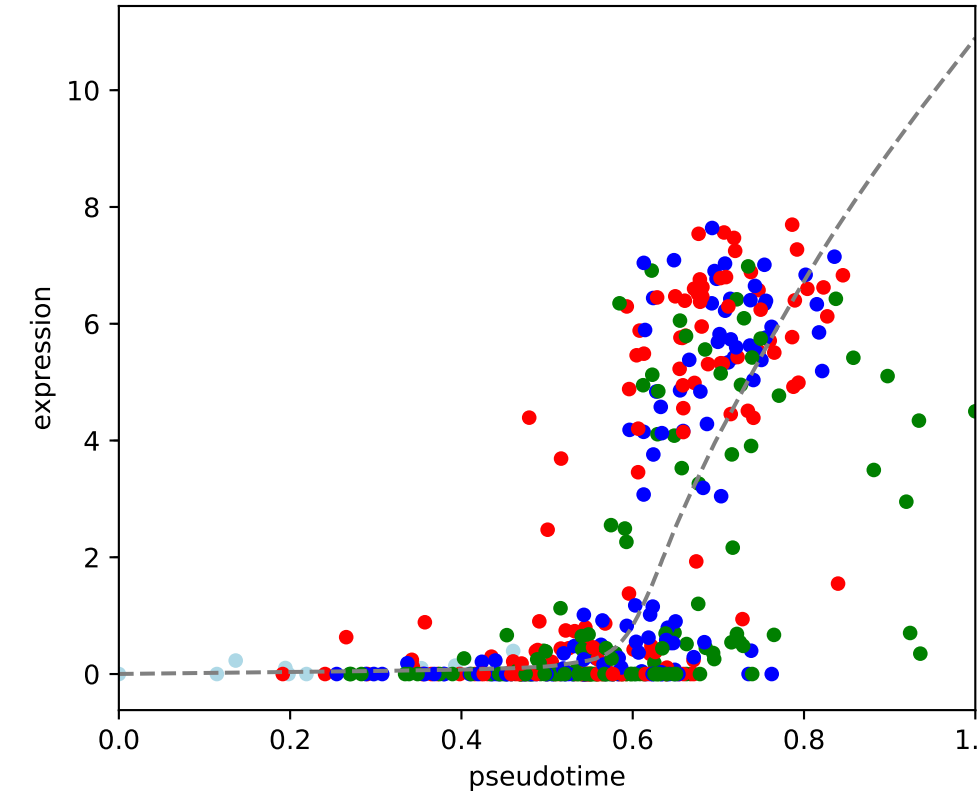
733 1spearman=0.28



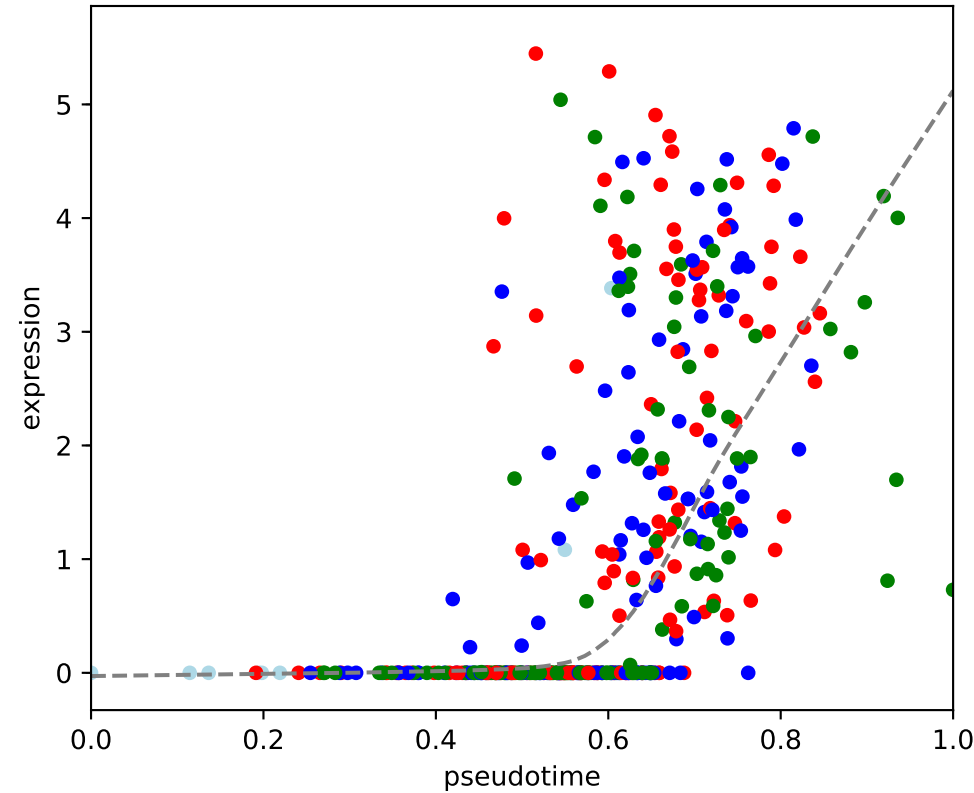
ENST00000429473 GNGT1 (G protein subunit gamma transducin 1)

733 0spearman=0.68

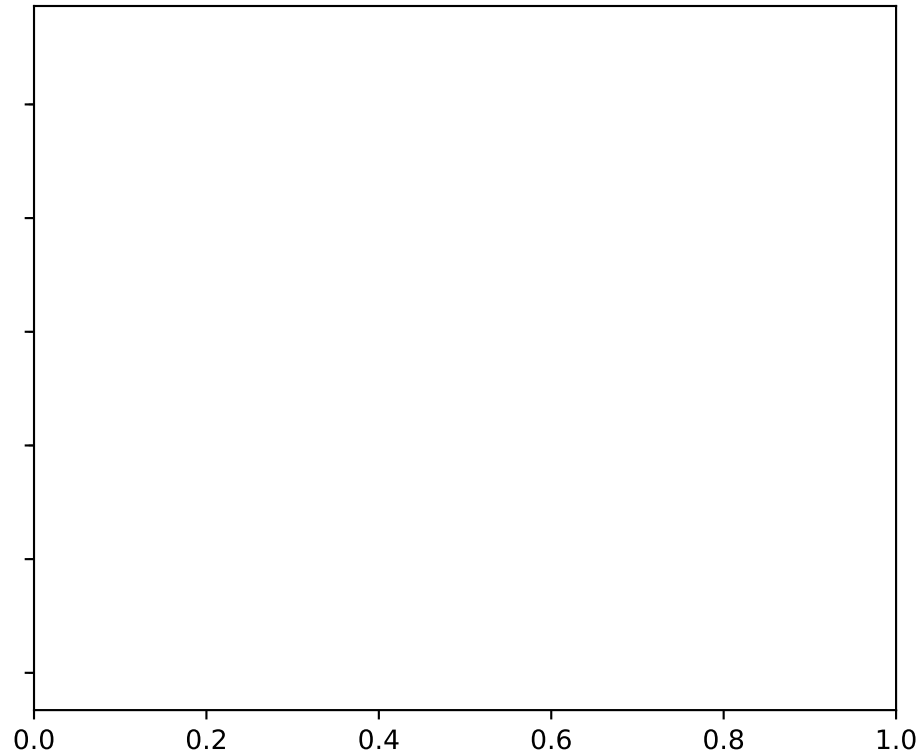
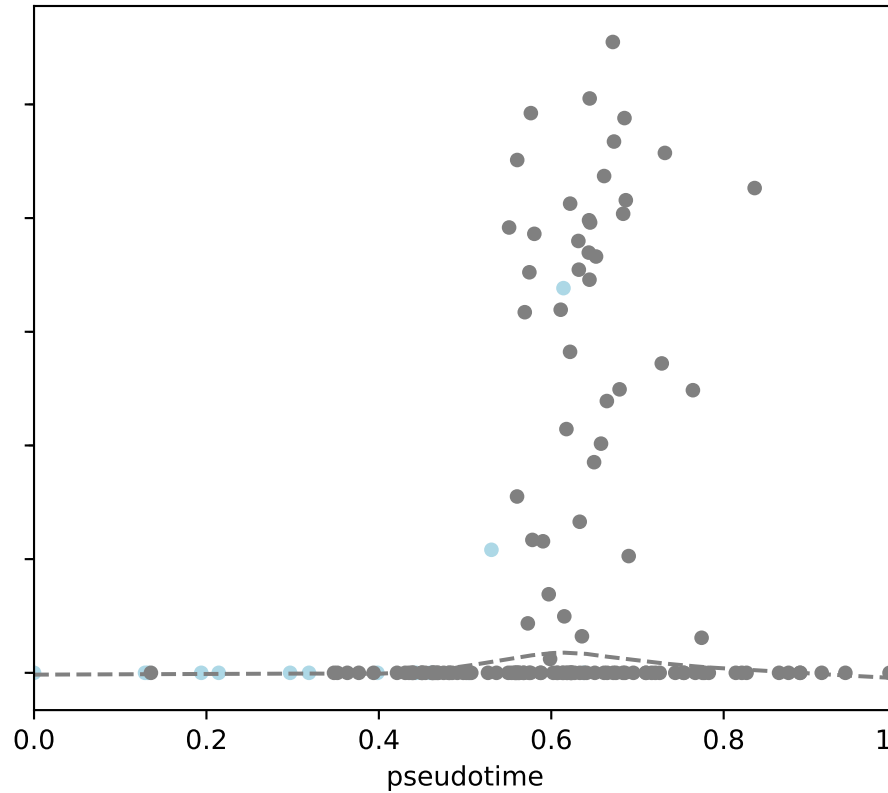
733 1spearman=0.24



733 0spearman=0.68

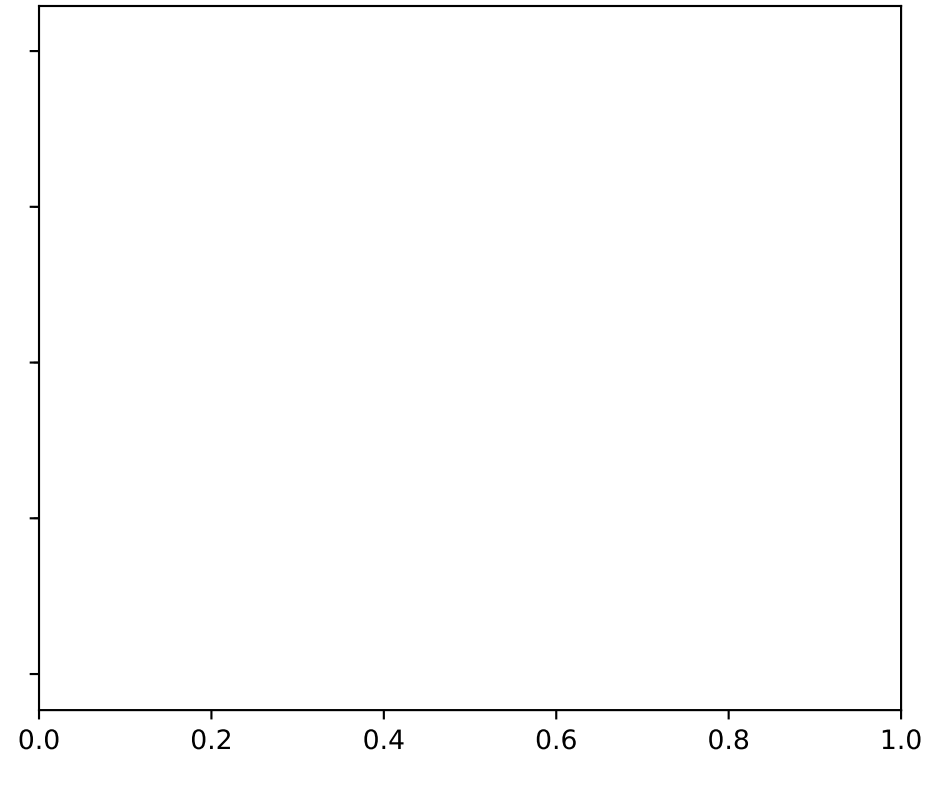
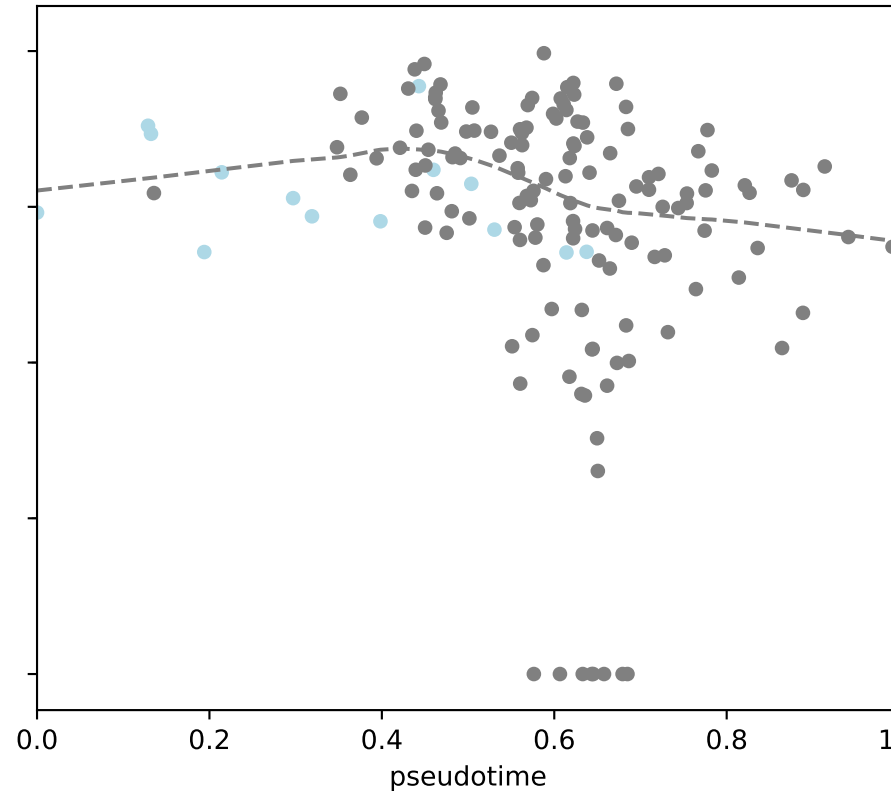
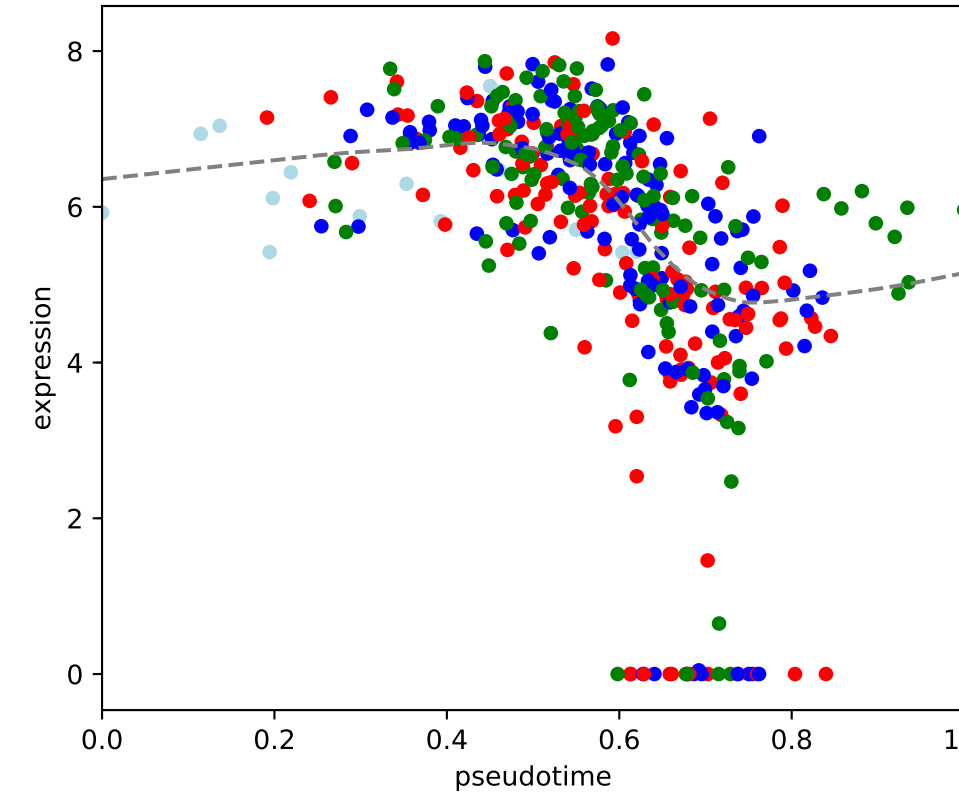


733 1spearman=0.26



733 0spearman=-0.68

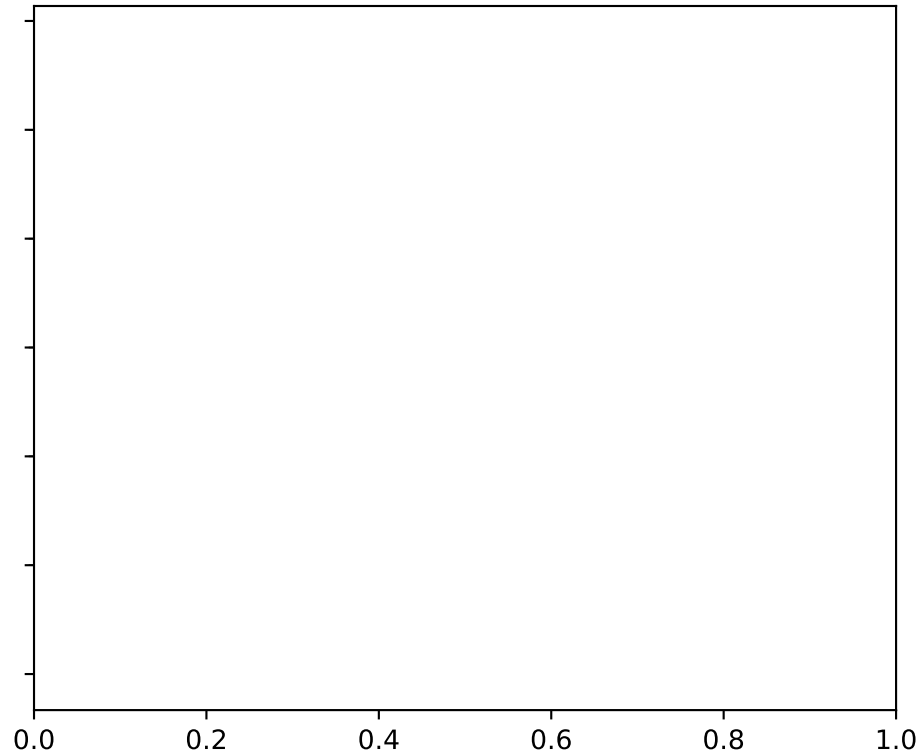
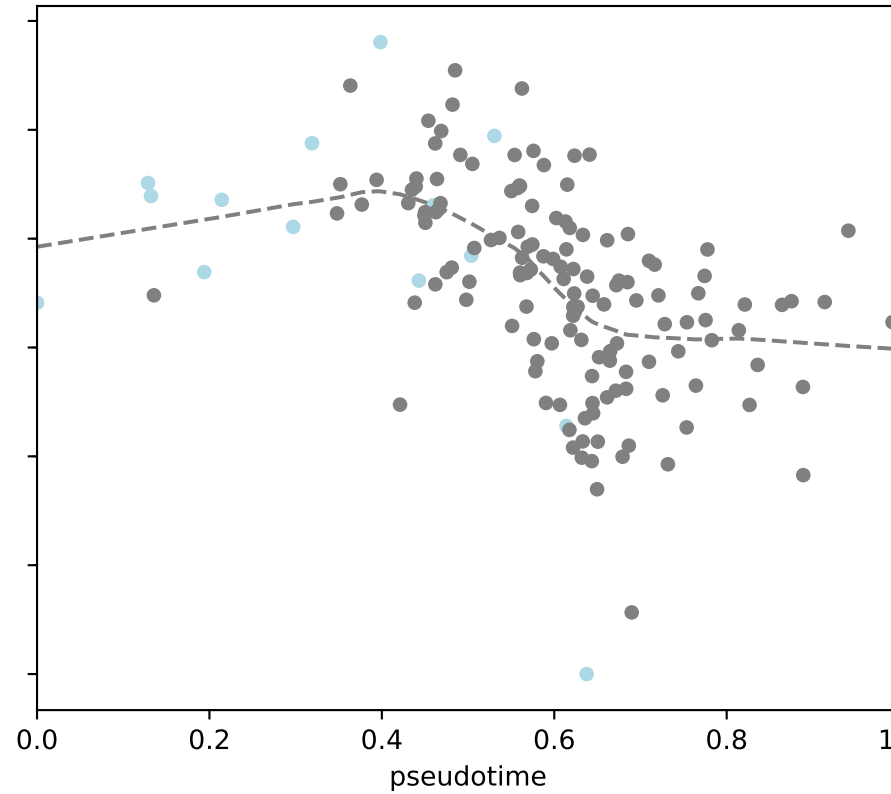
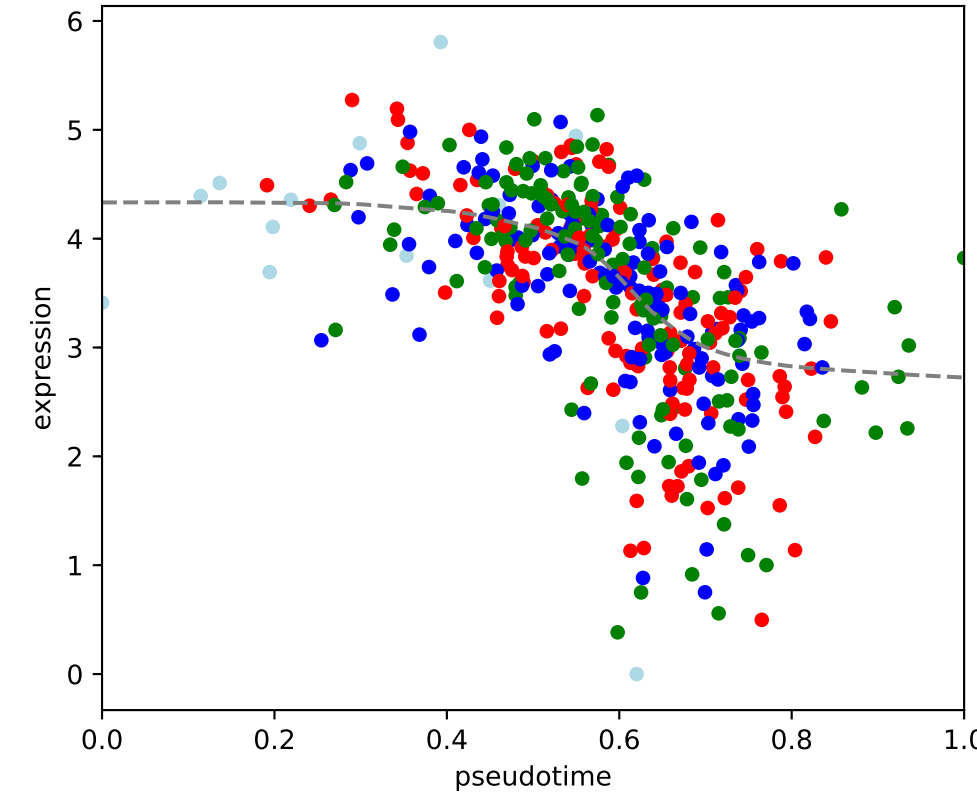
733 1spearman=-0.39



ENST00000391865 GUK1 (guanylate kinase 1)

733 0spearman=-0.67

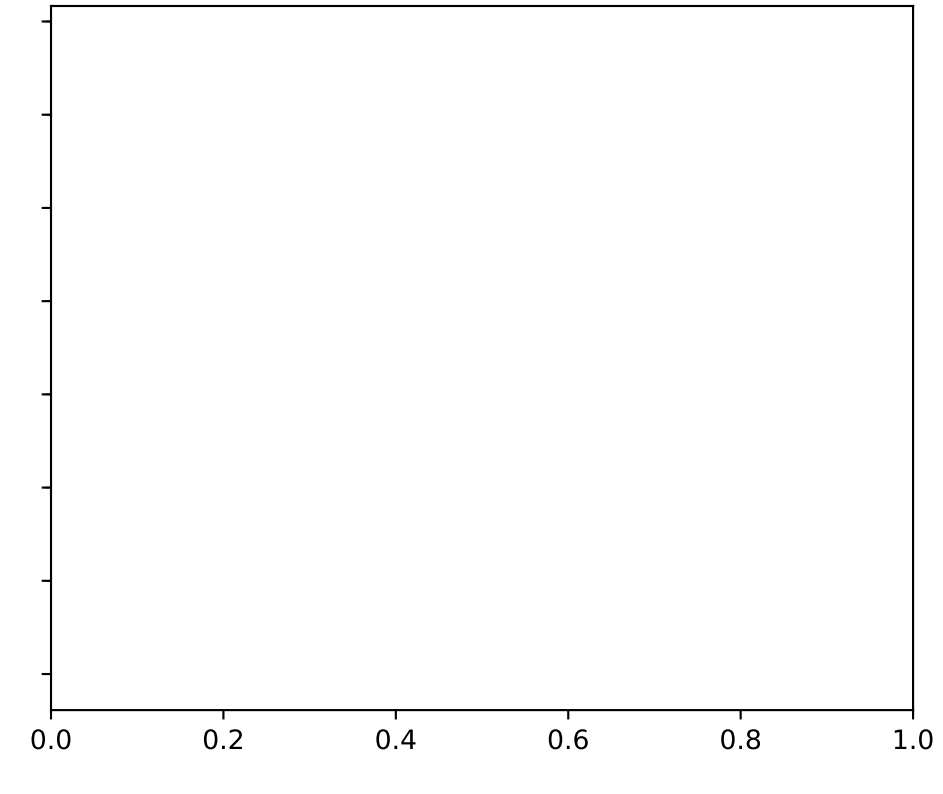
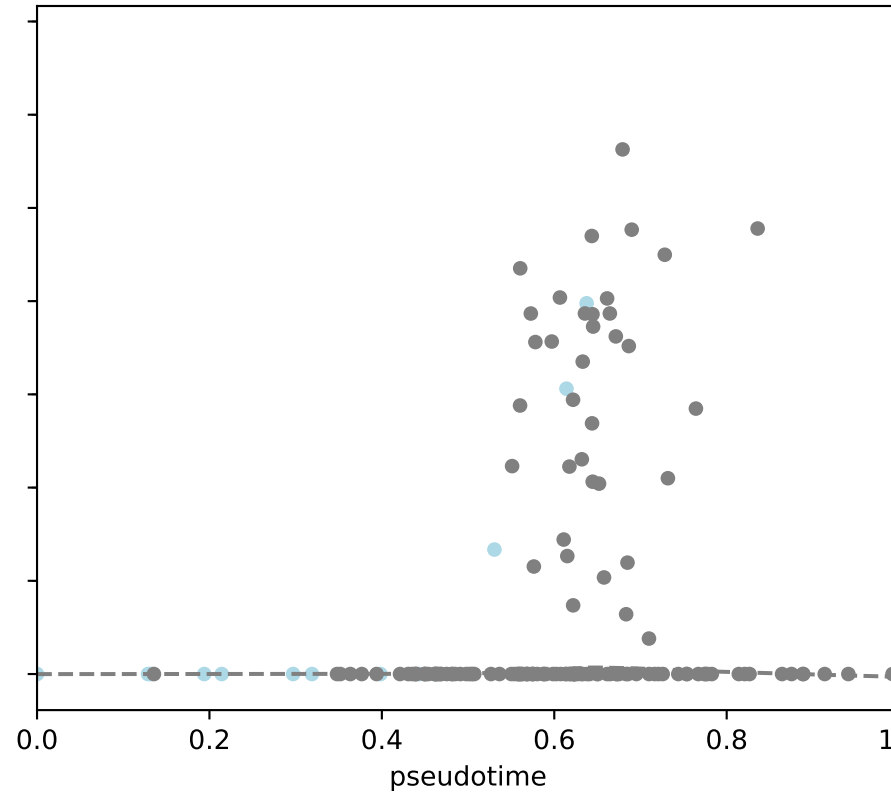
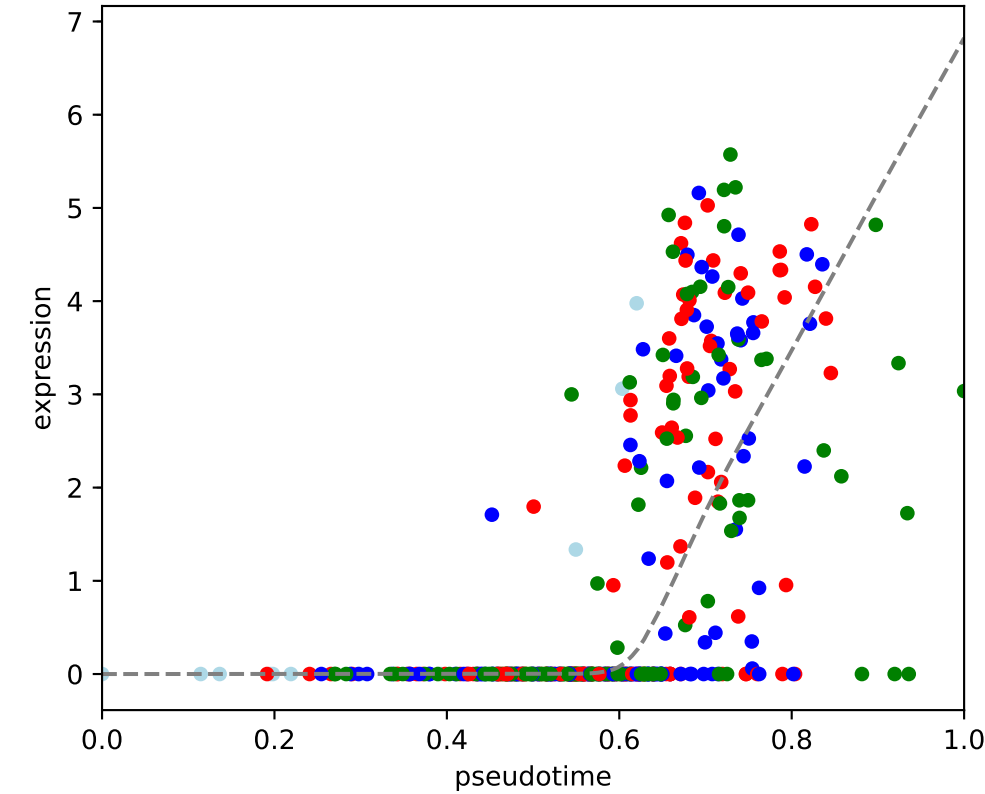
733 1spearman=-0.62



ENST00000562658 AC104938.1

733 0spearman=0.67

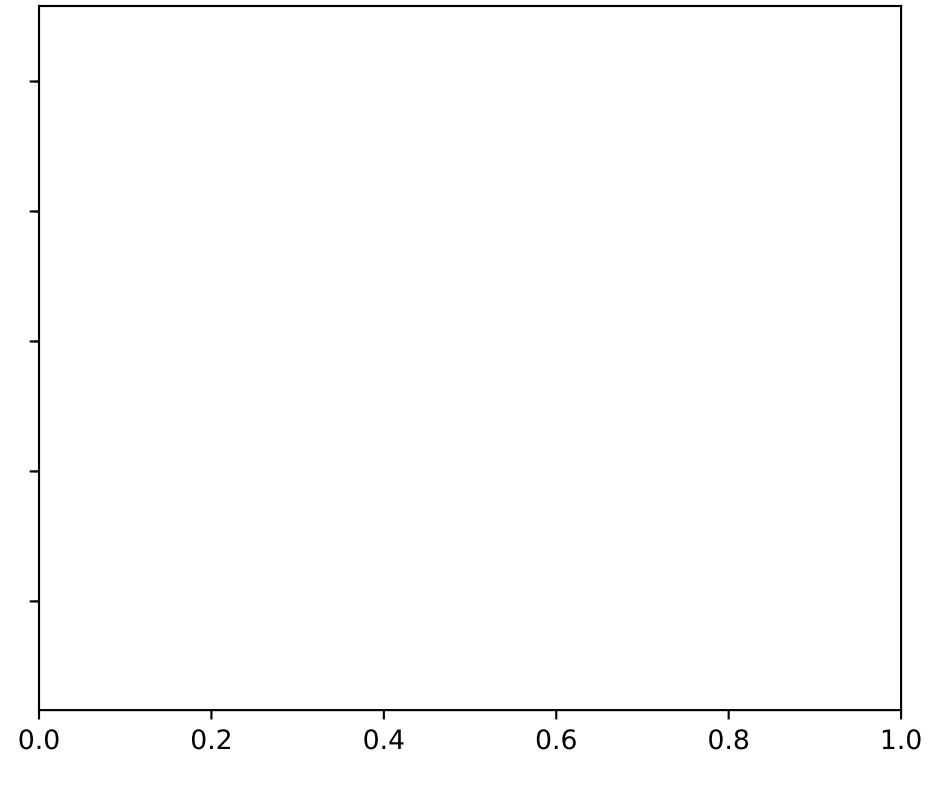
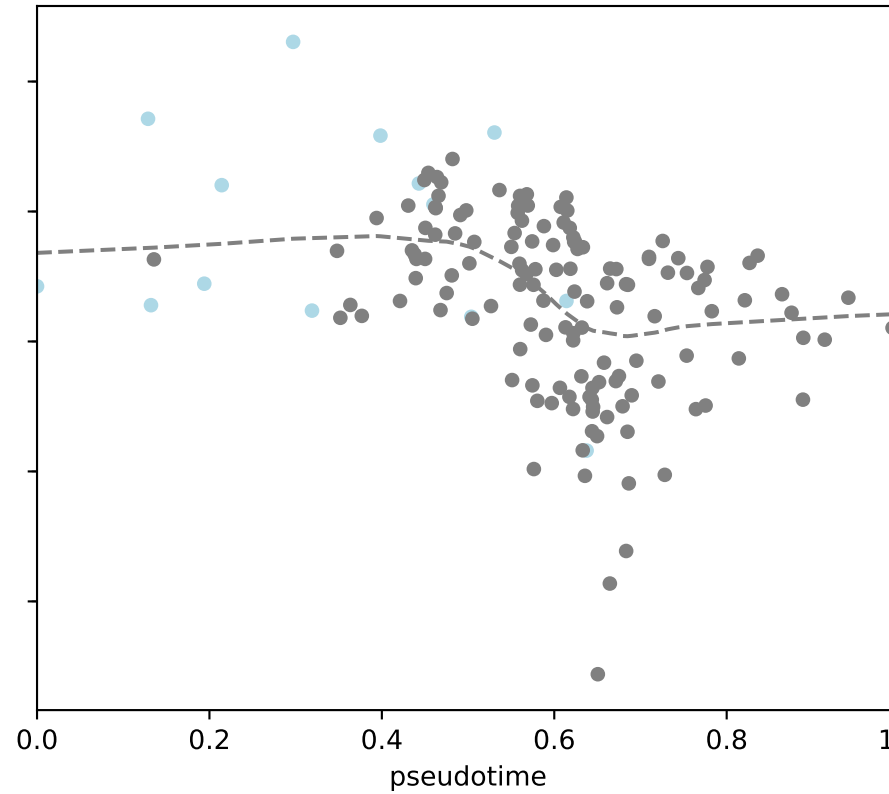
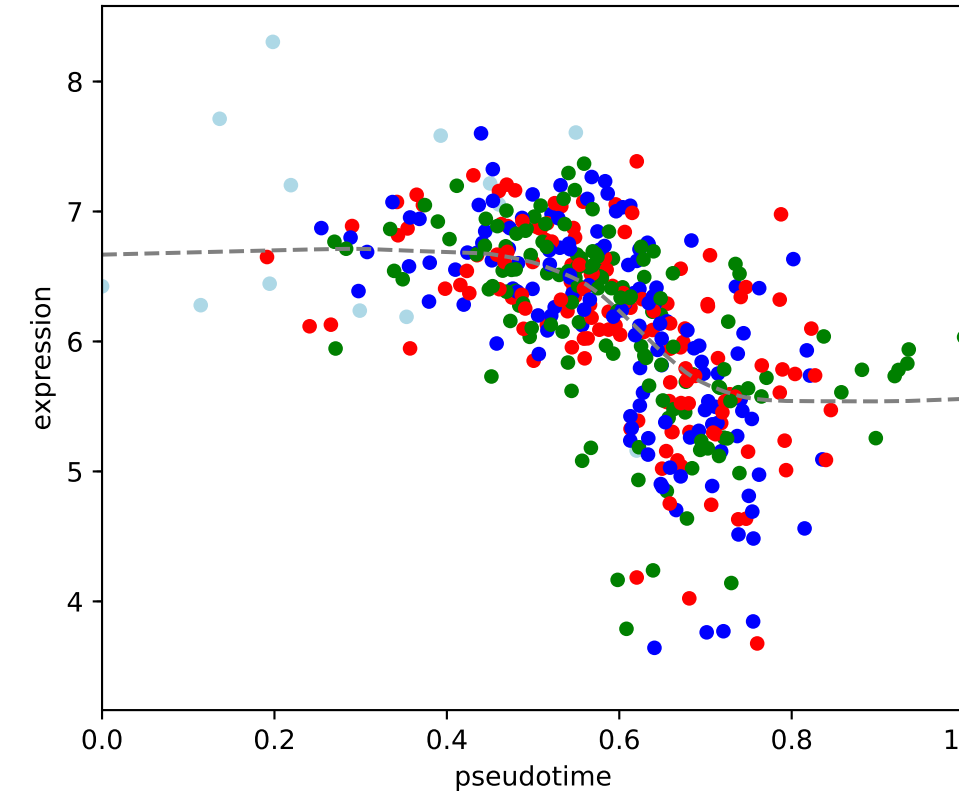
733 1spearman=0.26



ENST00000215754 MIF (macrophage migration inhibitory factor)

733 0spearman=-0.67

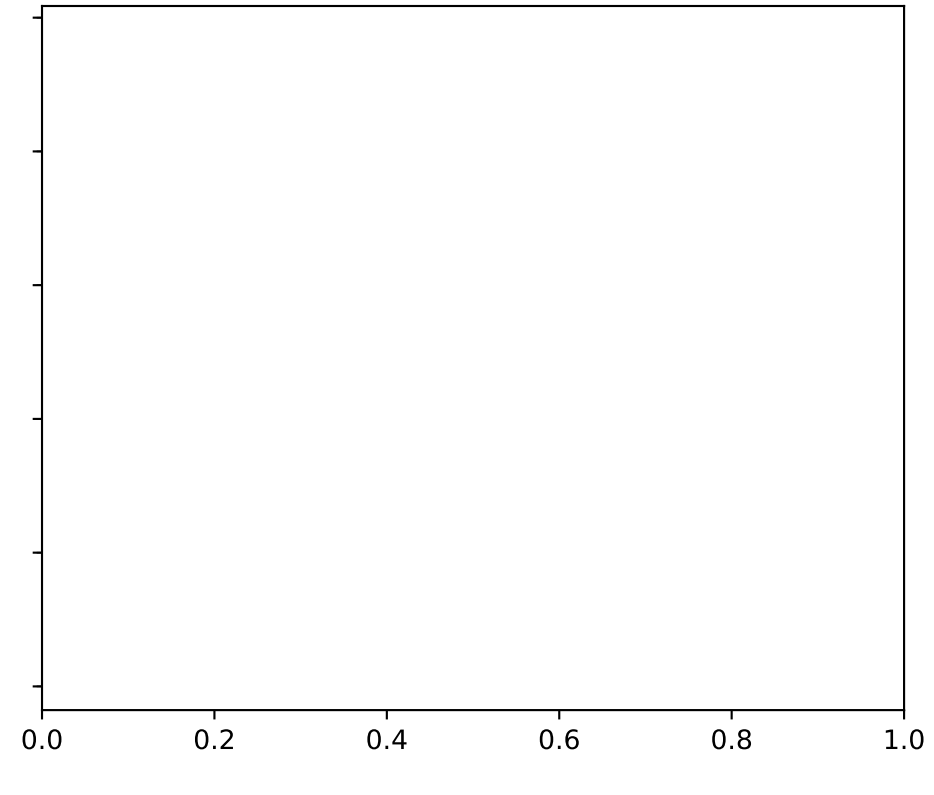
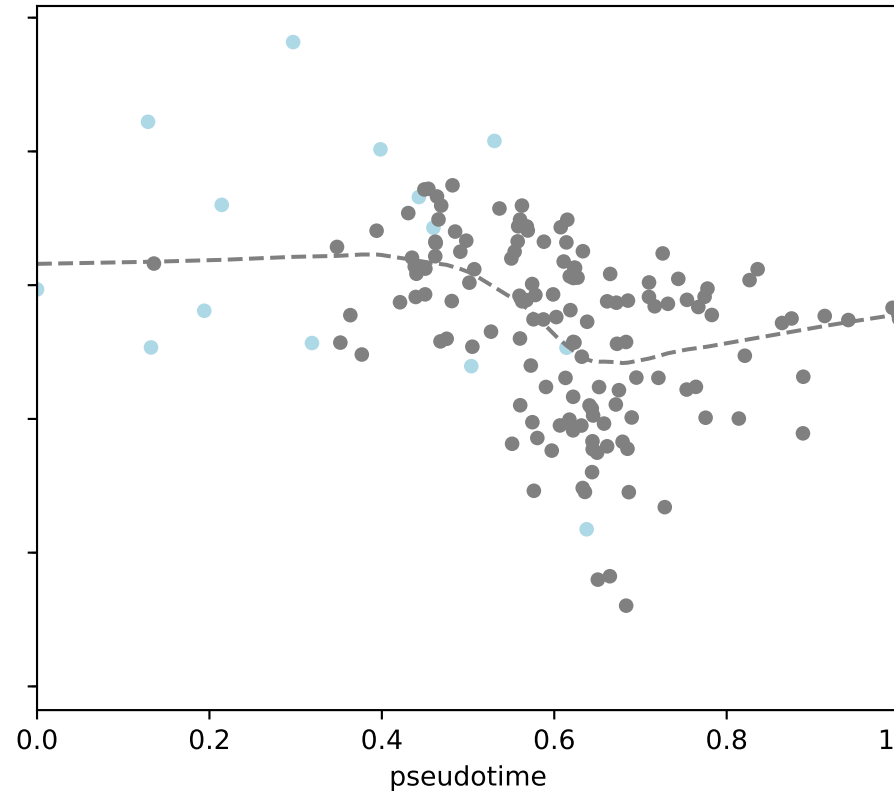
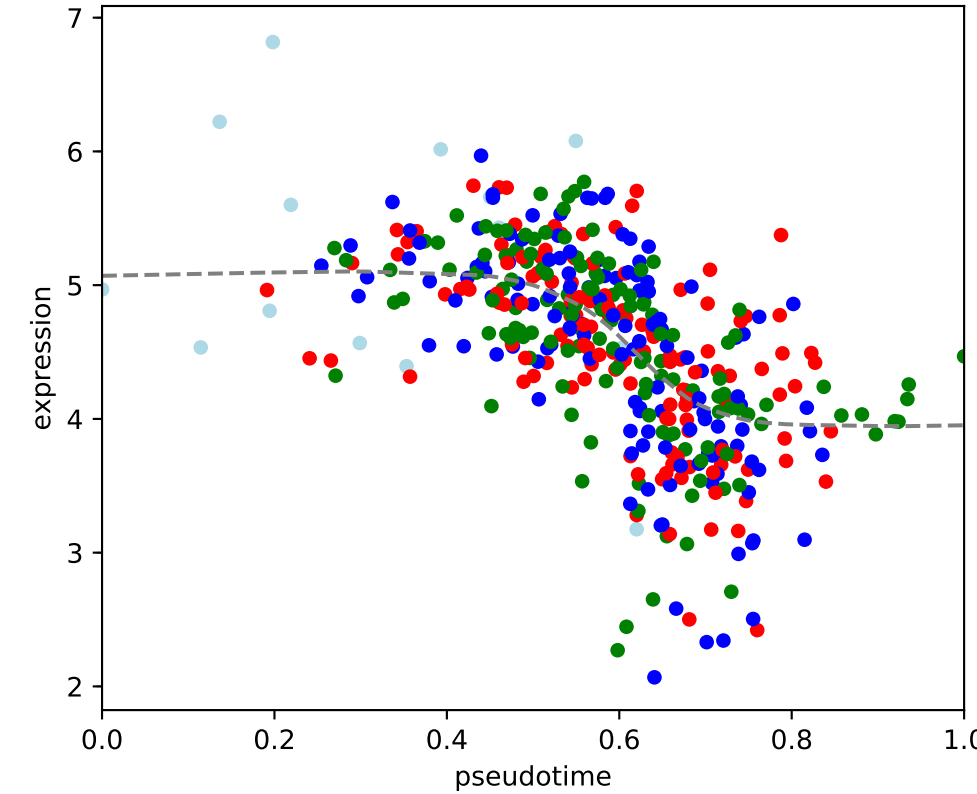
733 1spearman=-0.50



ENST00000406213 MIF-AS1 (MIF antisense RNA 1)

733 0spearman=-0.67

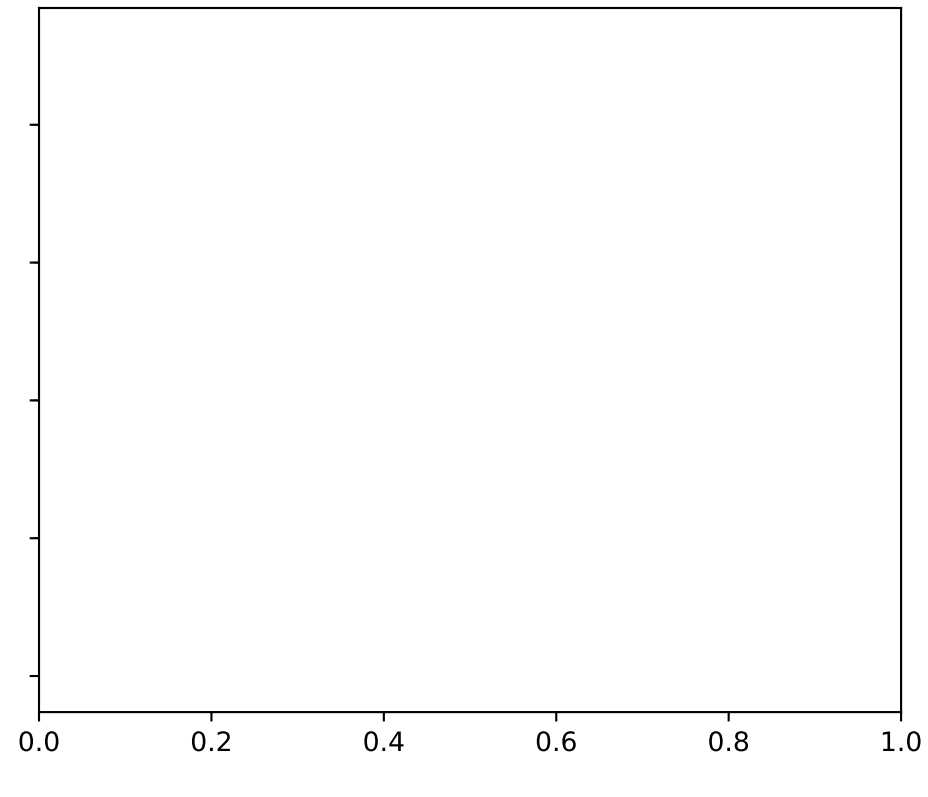
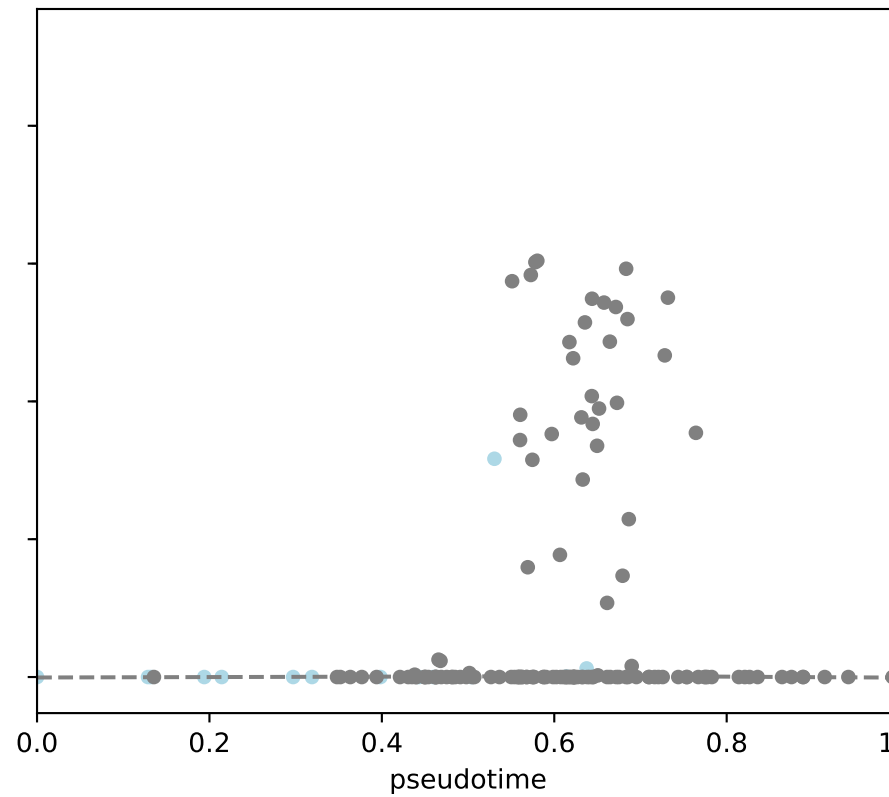
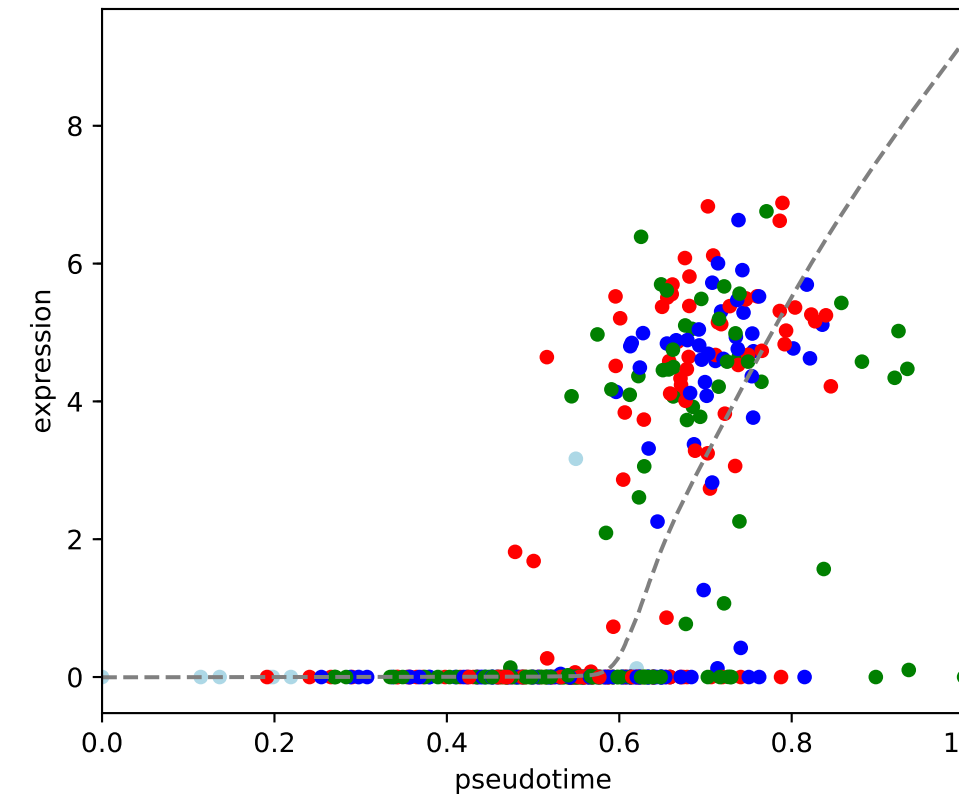
733 1spearman=-0.48



ENST00000430431 MRLN (myoregulin)

733 0spearman=0.67

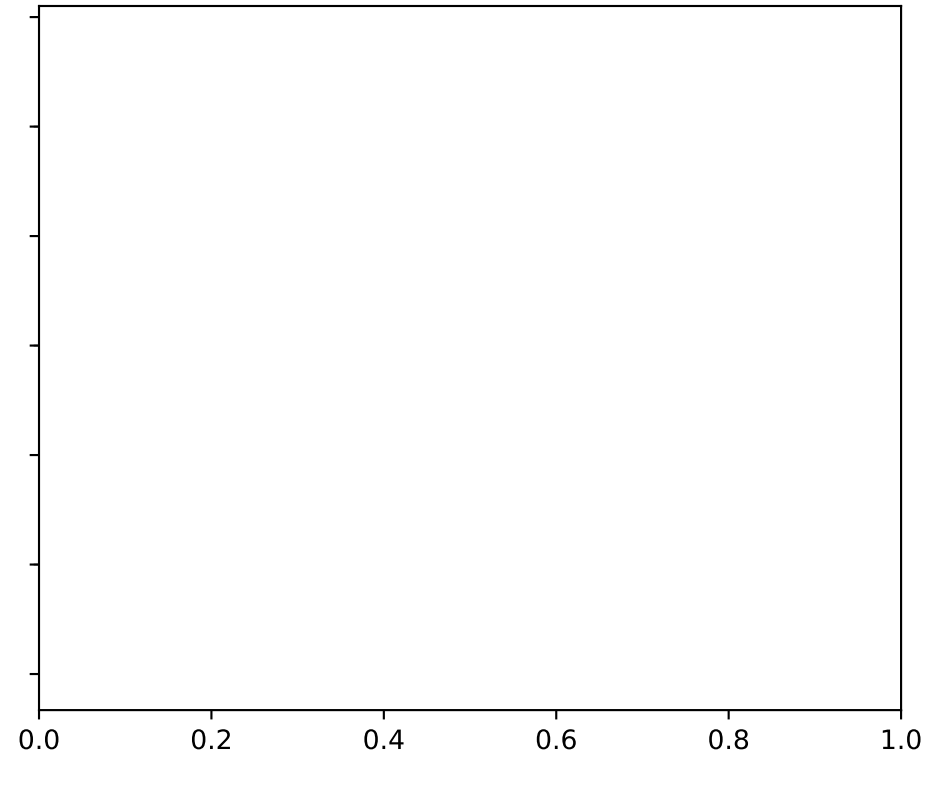
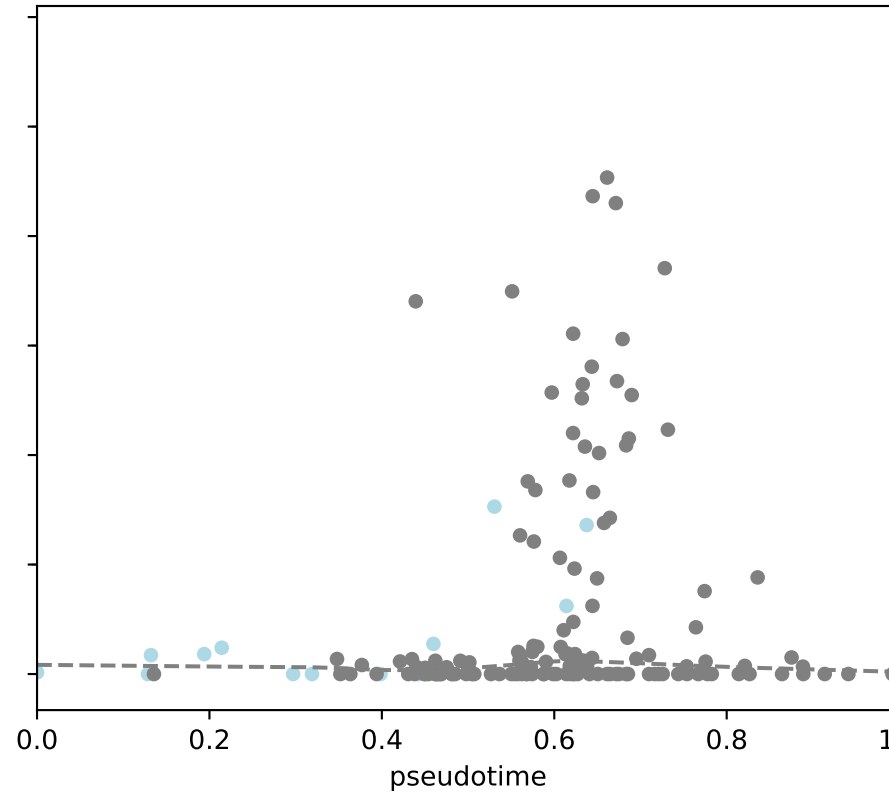
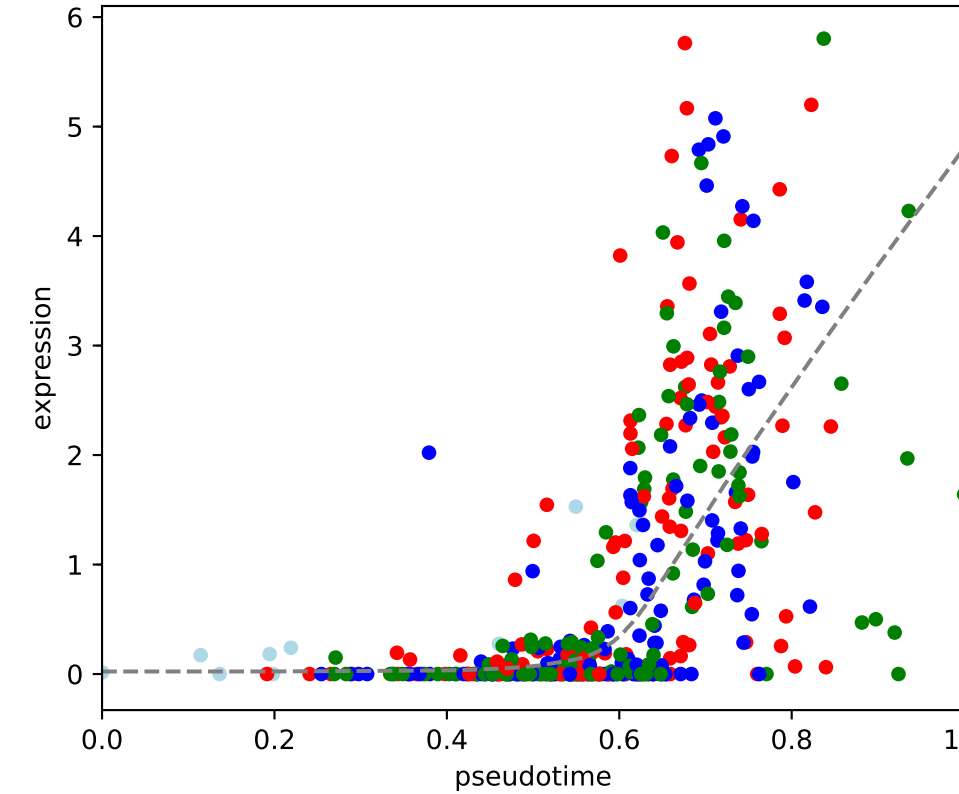
733 1spearman=0.16



ENST00000621736 NR2E3 (nuclear receptor subfamily 2 group E member 3)

733 0spearman=0.67

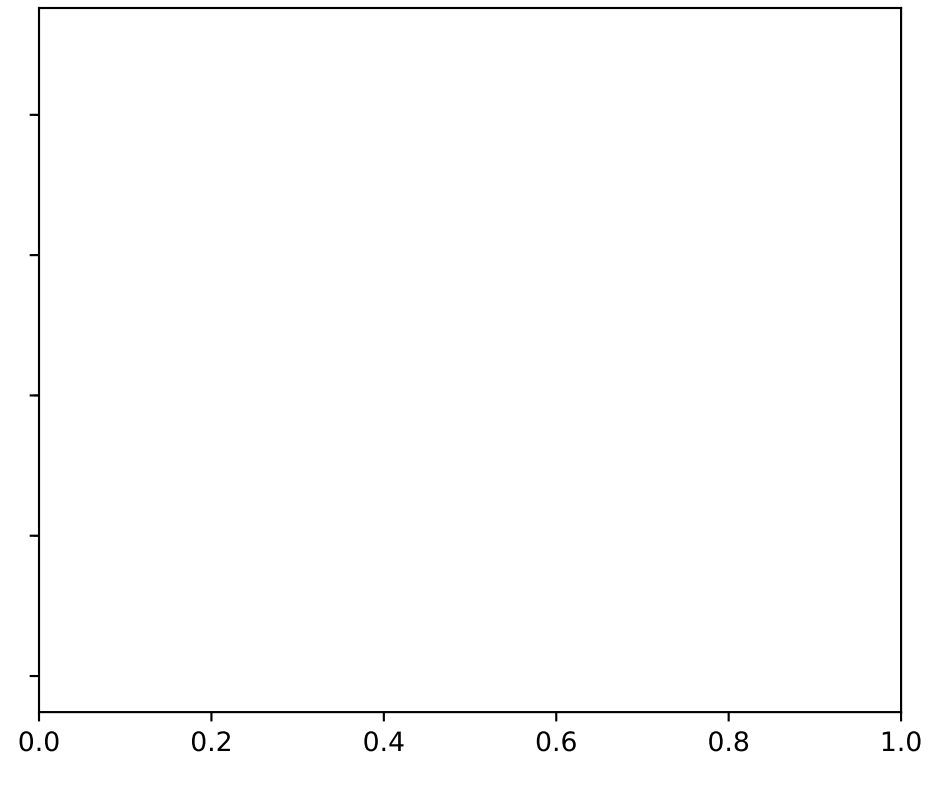
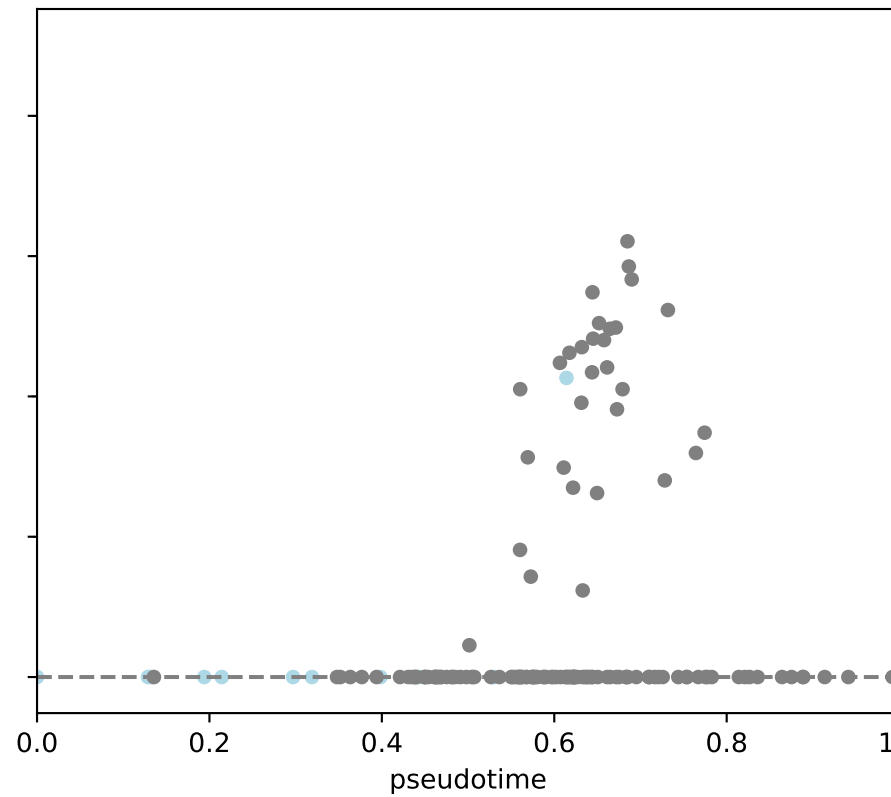
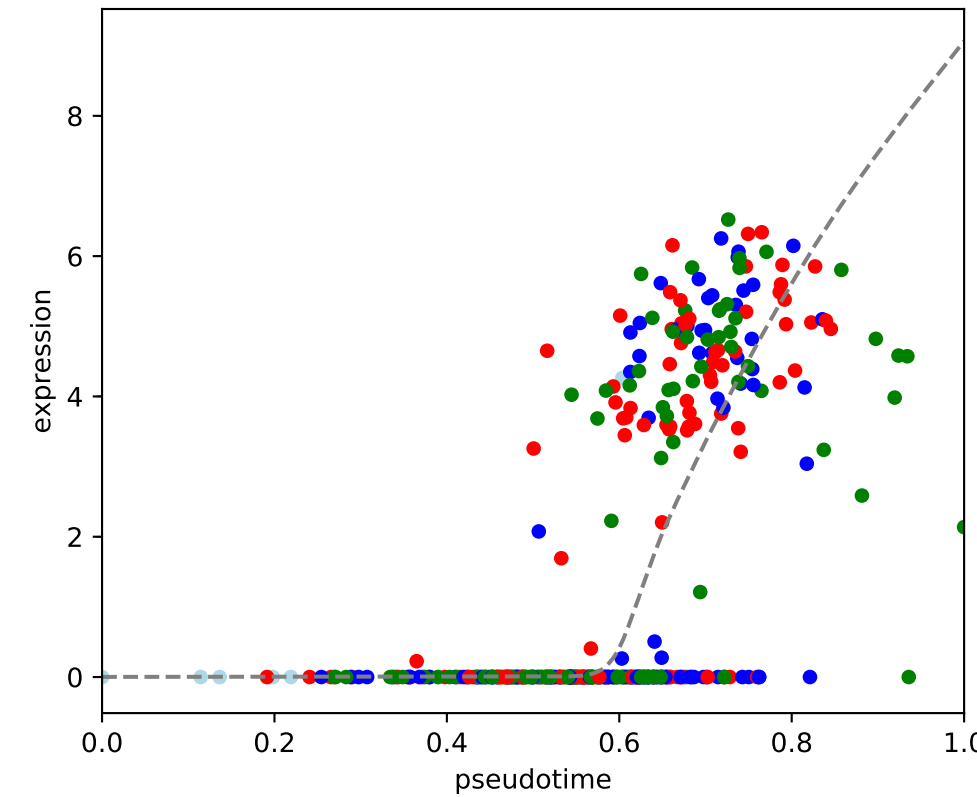
733 1spearman=0.17



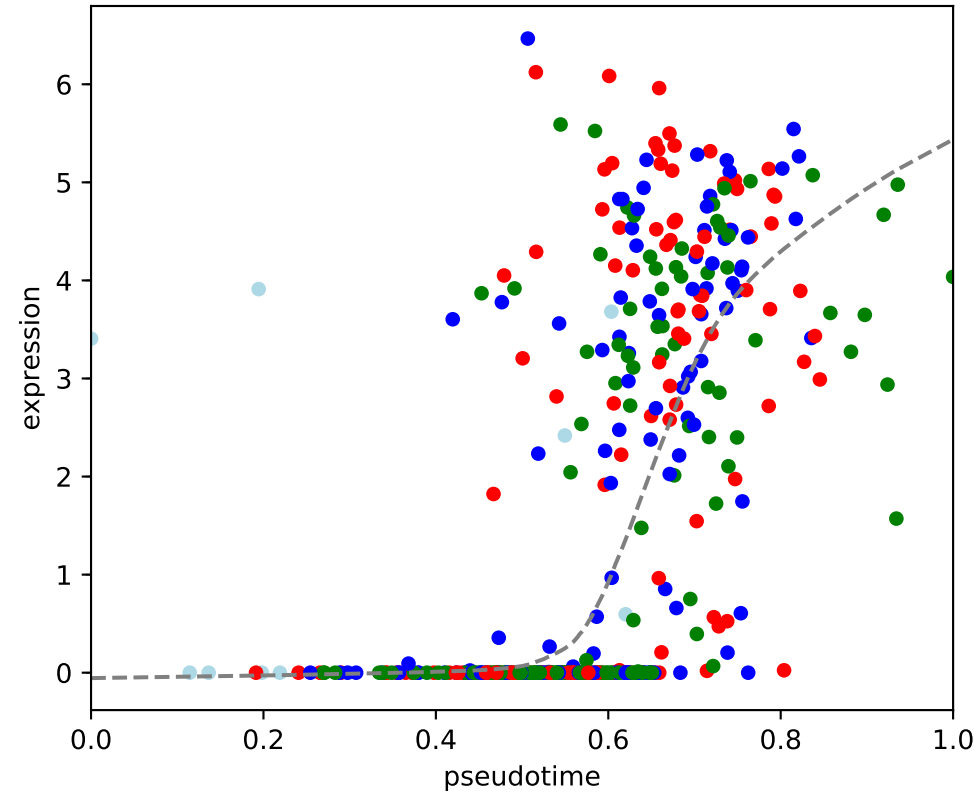
ENST00000414264 MRLN (myoregulin)

733 0spearman=0.67

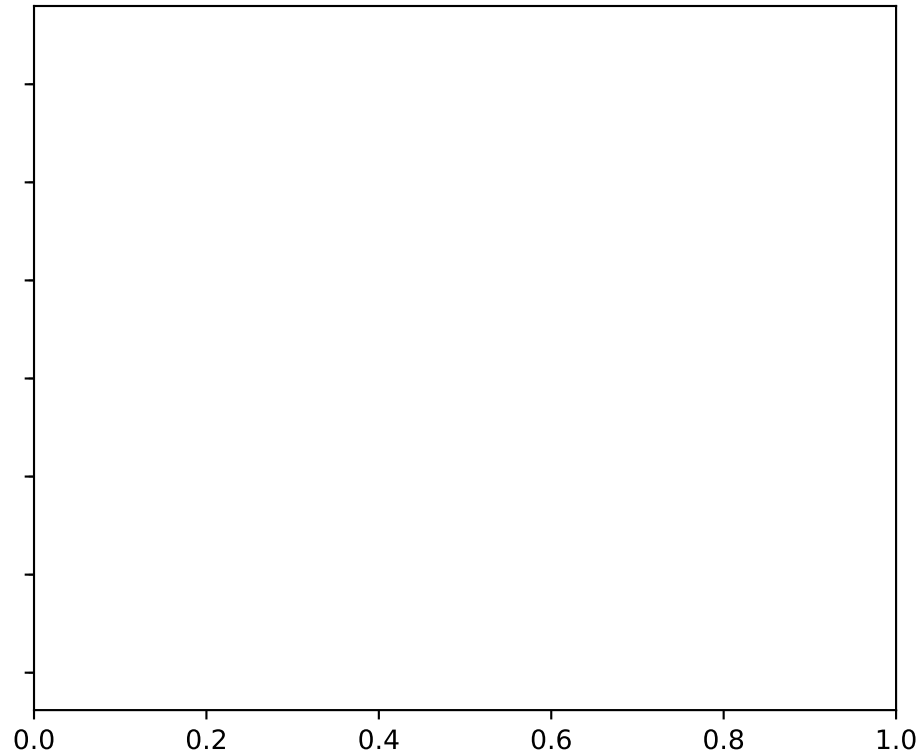
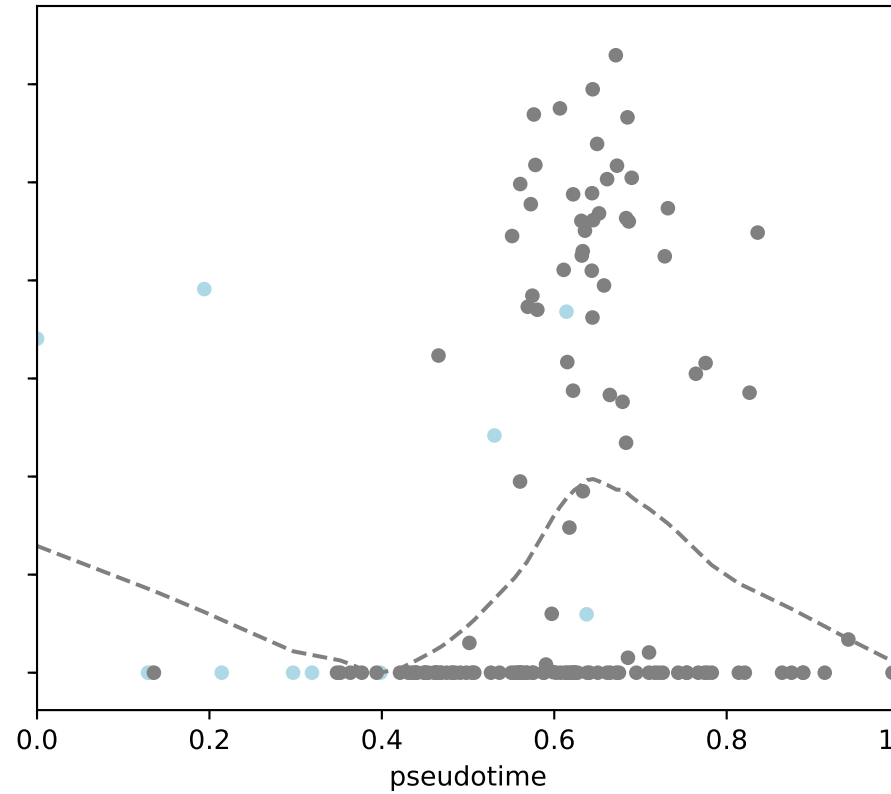
733 1spearman=0.26



733 0spearman=0.67



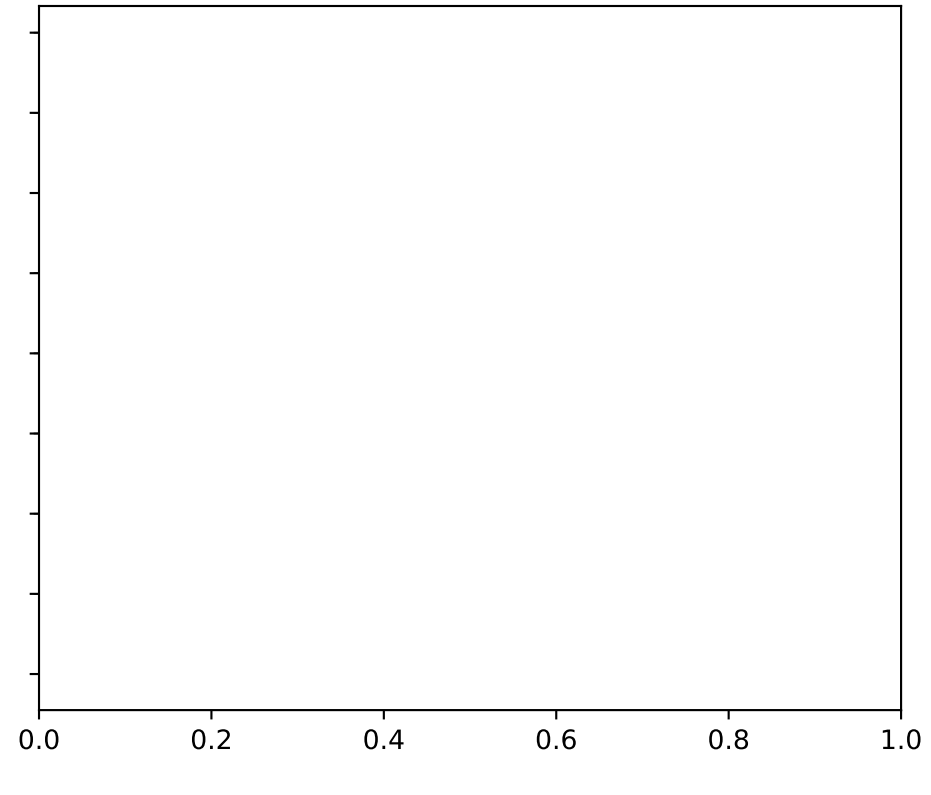
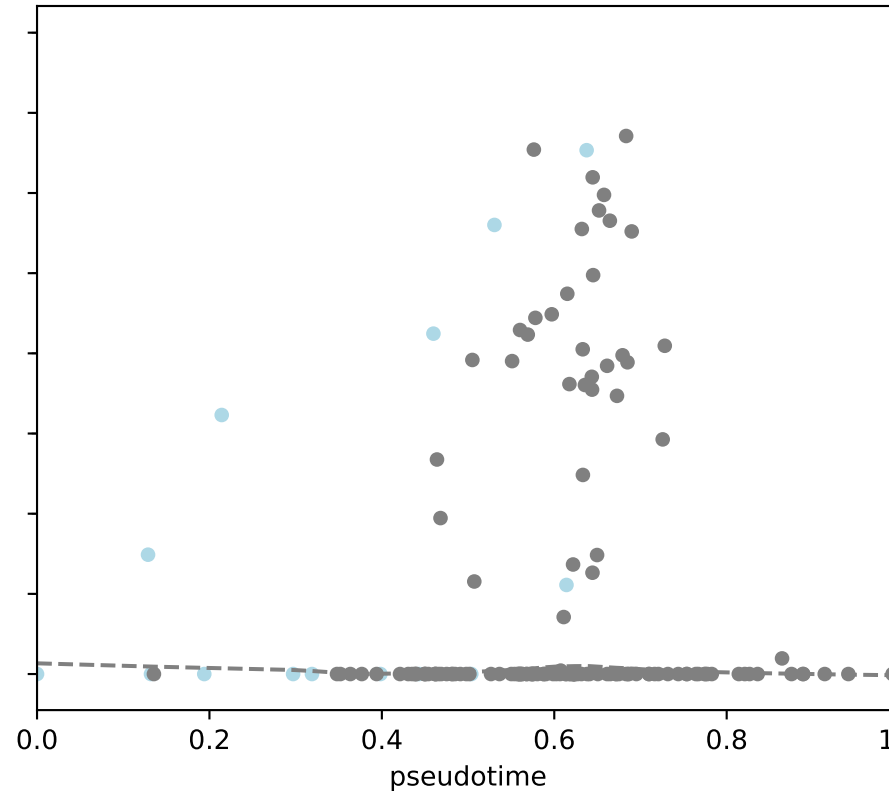
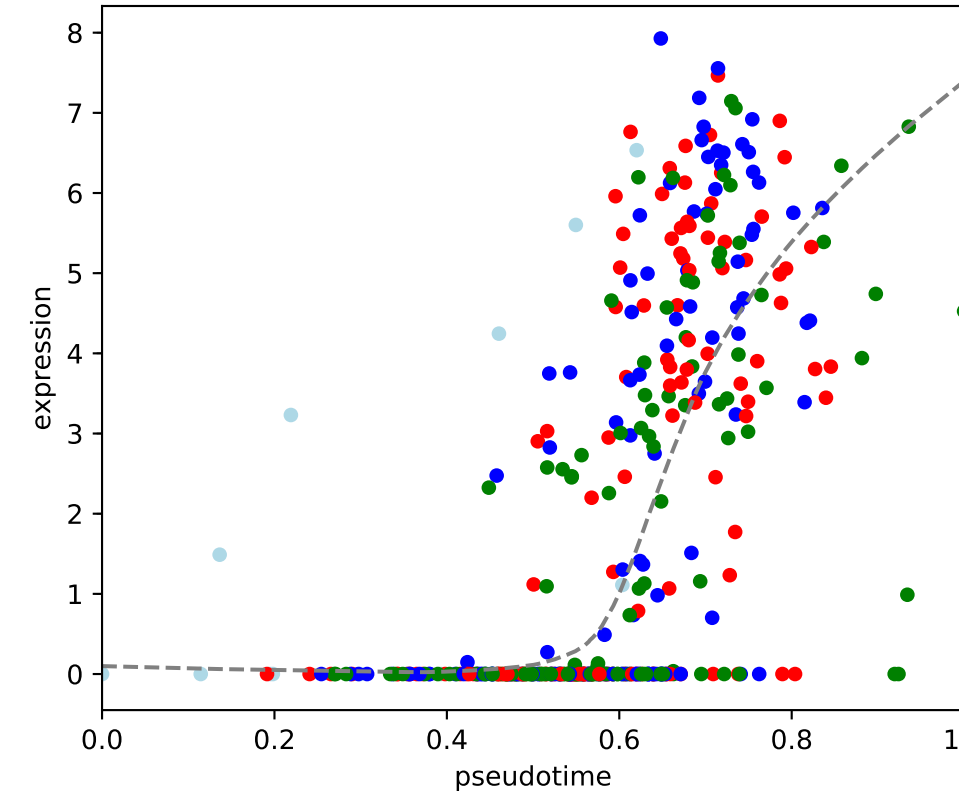
733 1spearman=0.28



ENST00000433068 GNAT1 (G protein subunit alpha transducin 1)

733 0spearman=0.66

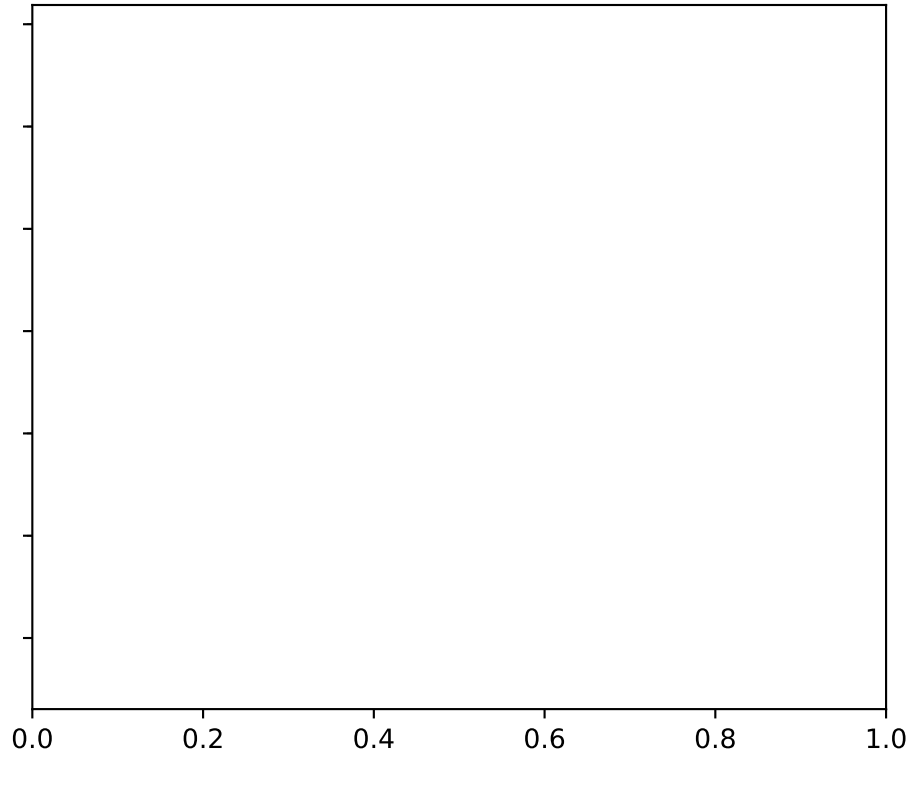
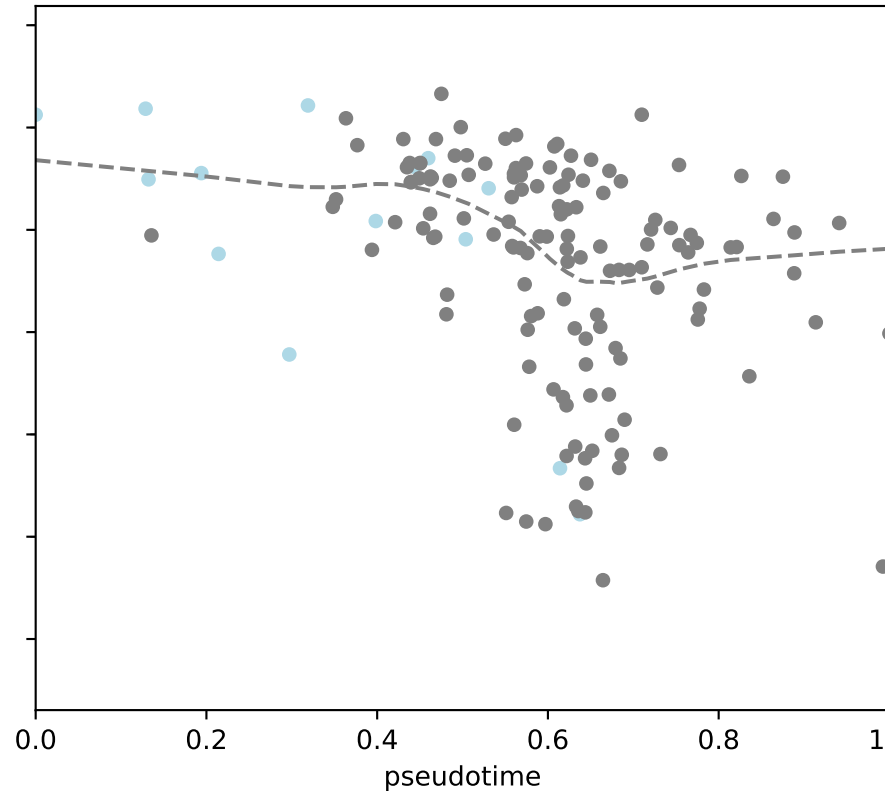
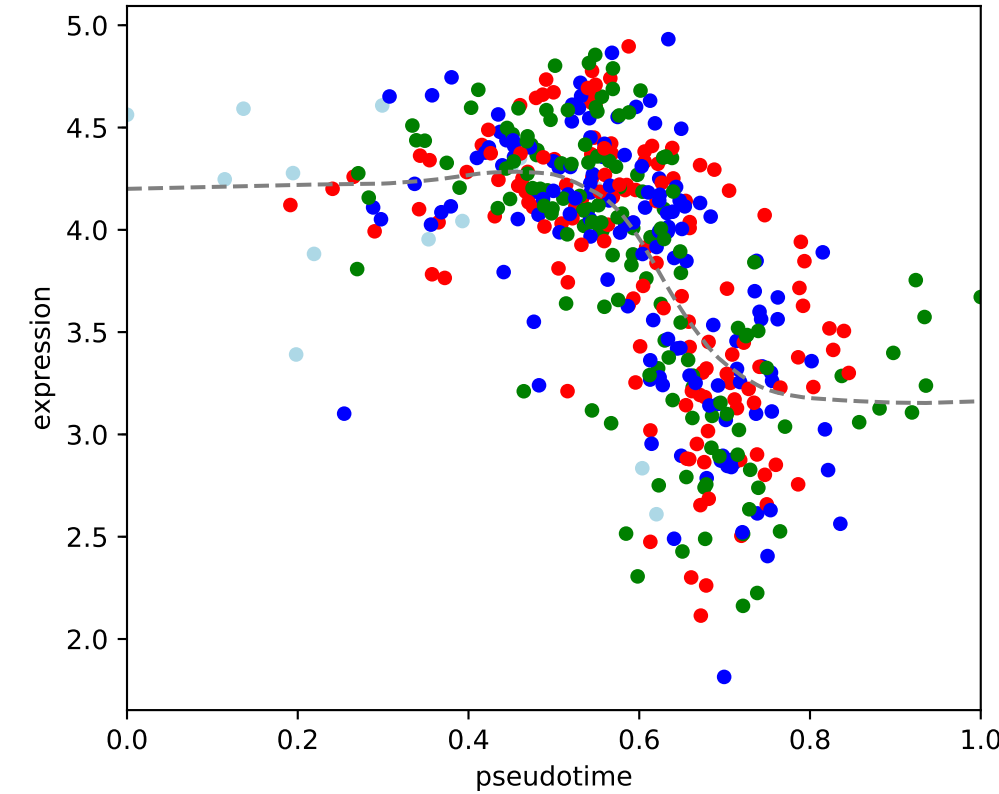
733 1spearman=0.11



ENST00000551020 ATP5F1B (ATP synthase F1 subunit beta)

733 0spearman=-0.66

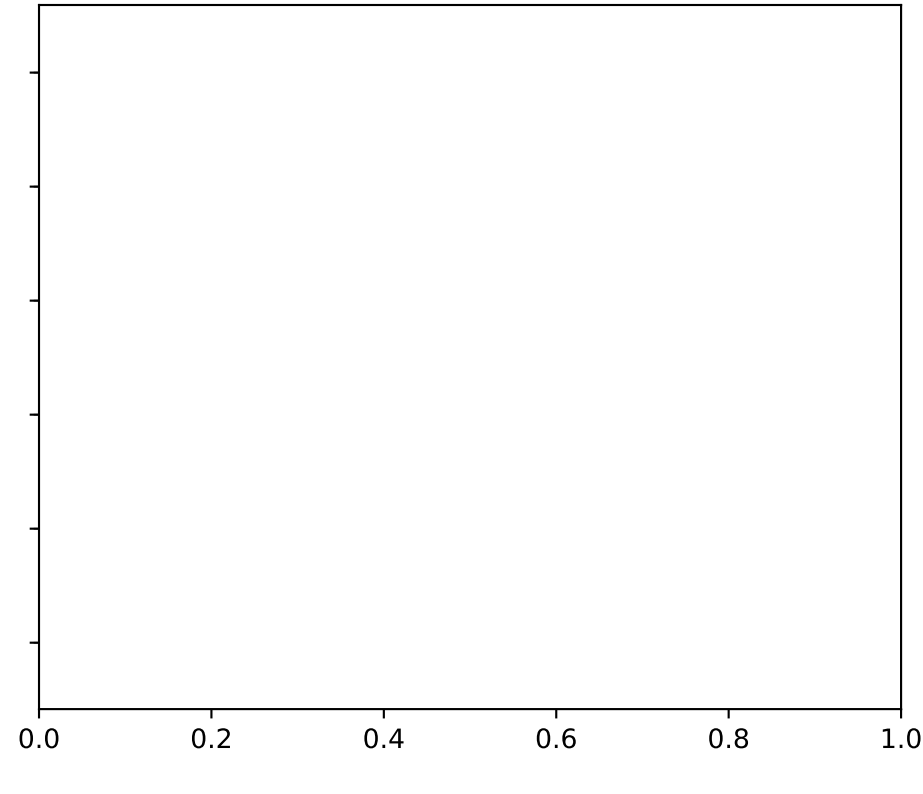
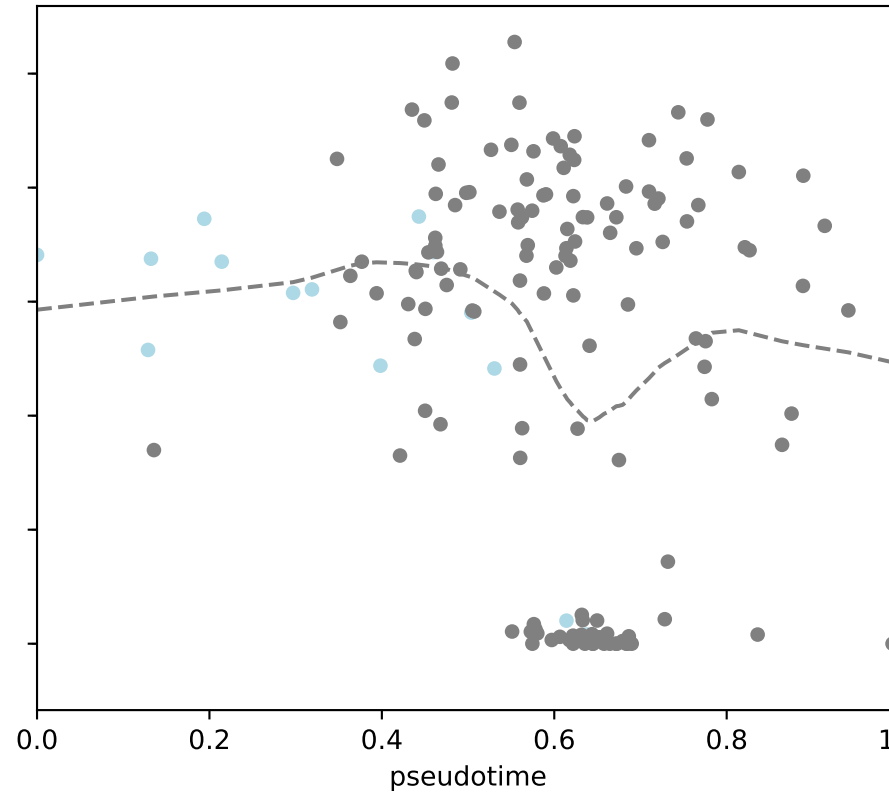
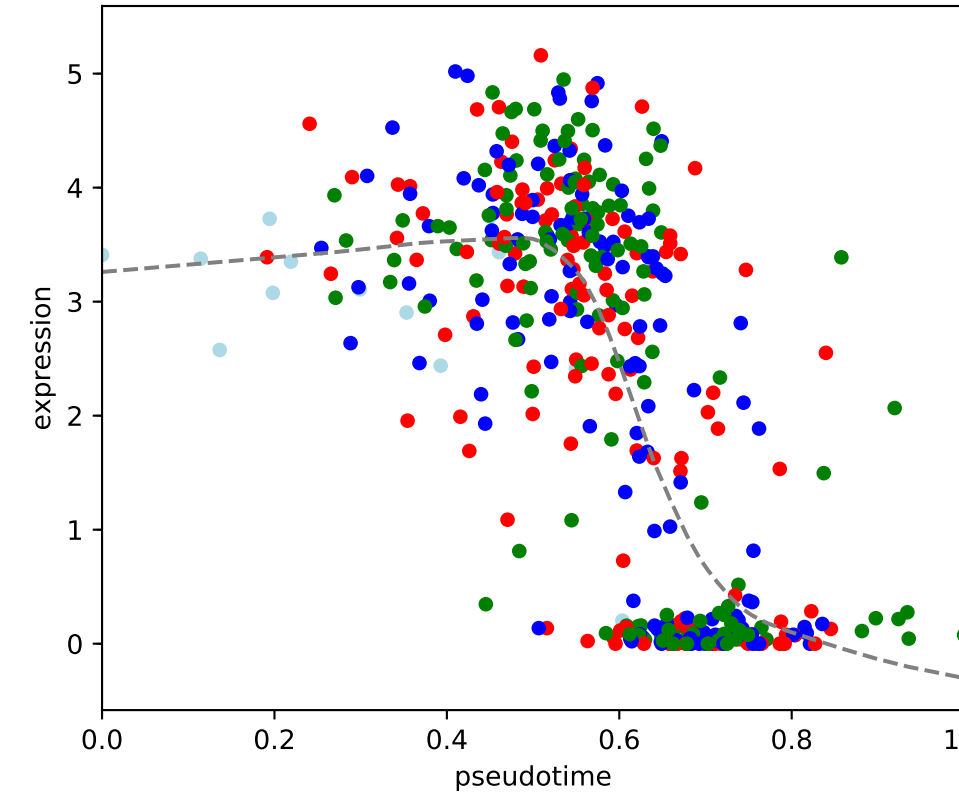
733 1spearman=-0.45



ENST00000251296 IGSF21 (immunoglobulin superfamily member 21)

733 0spearman=-0.66

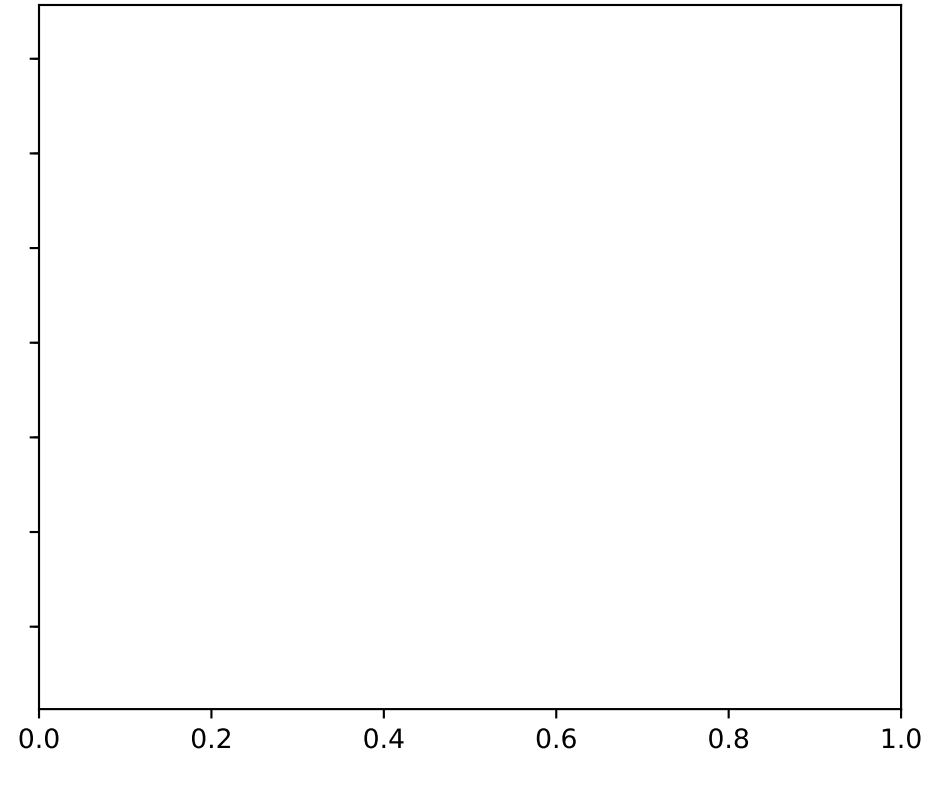
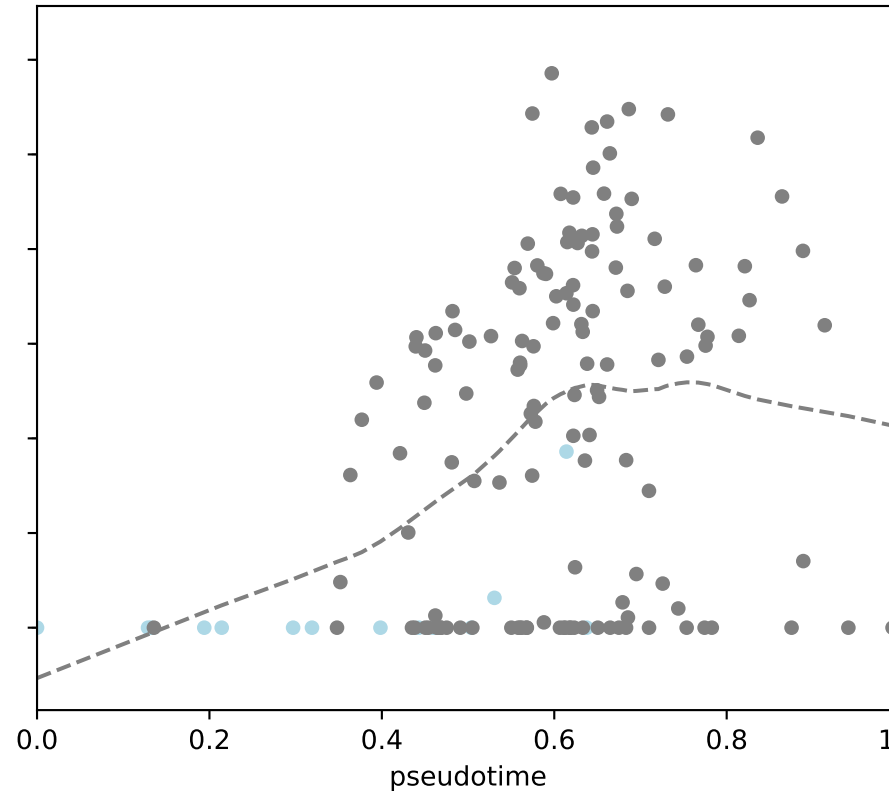
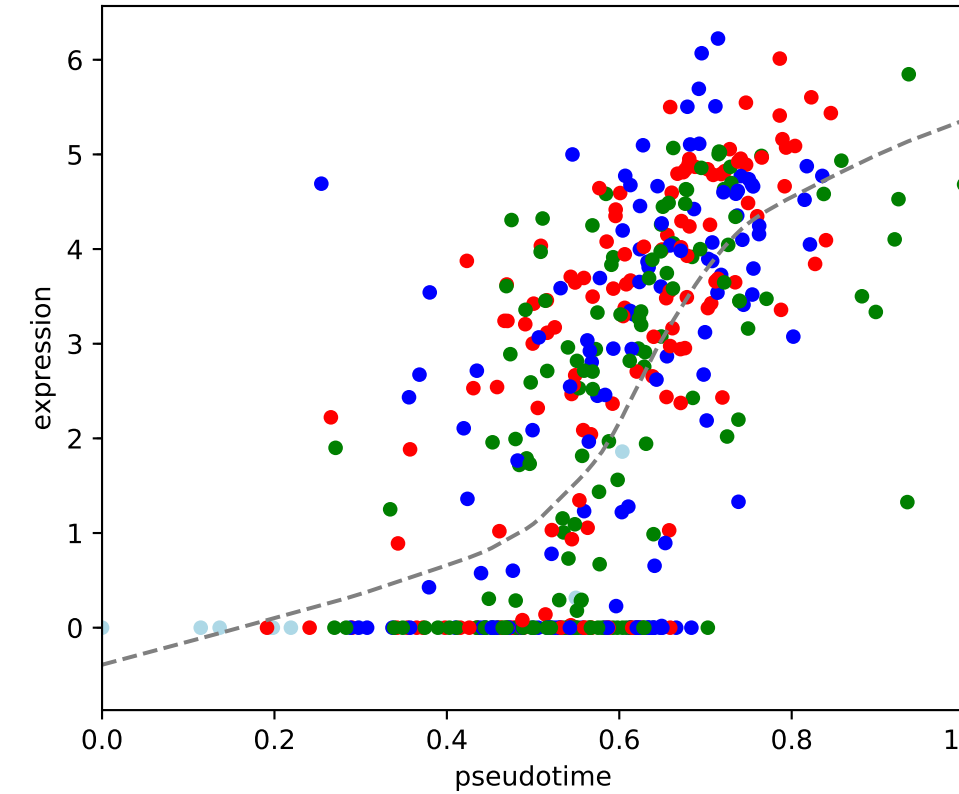
733 1spearman=-0.19



ENST00000341065 SAMD11 (sterile alpha motif domain containing 11)

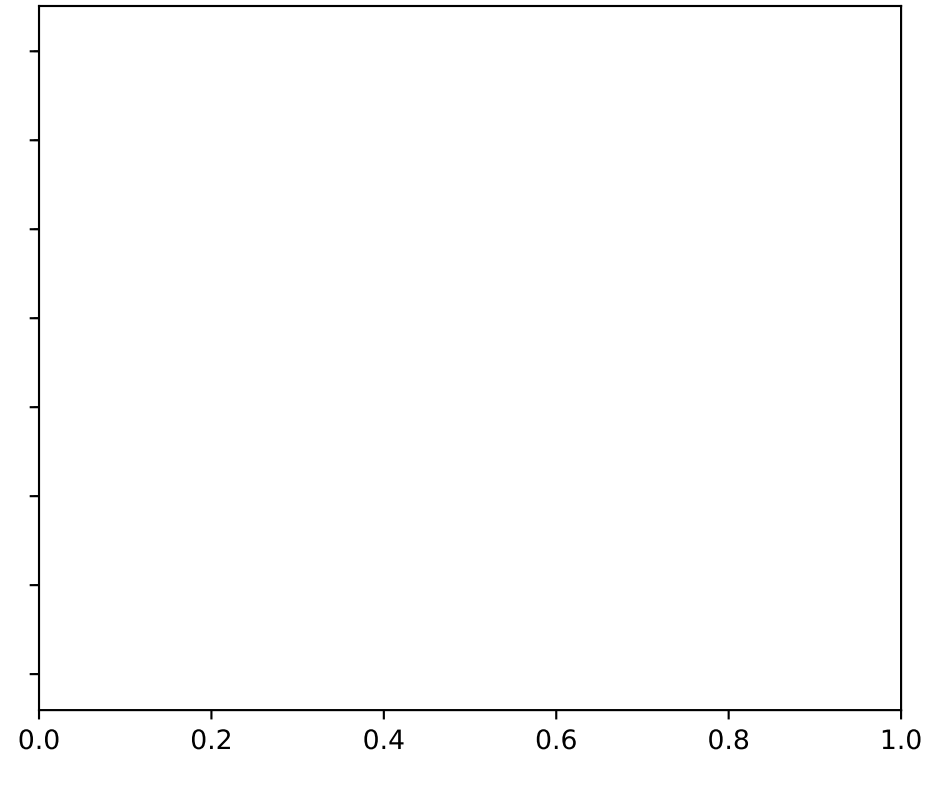
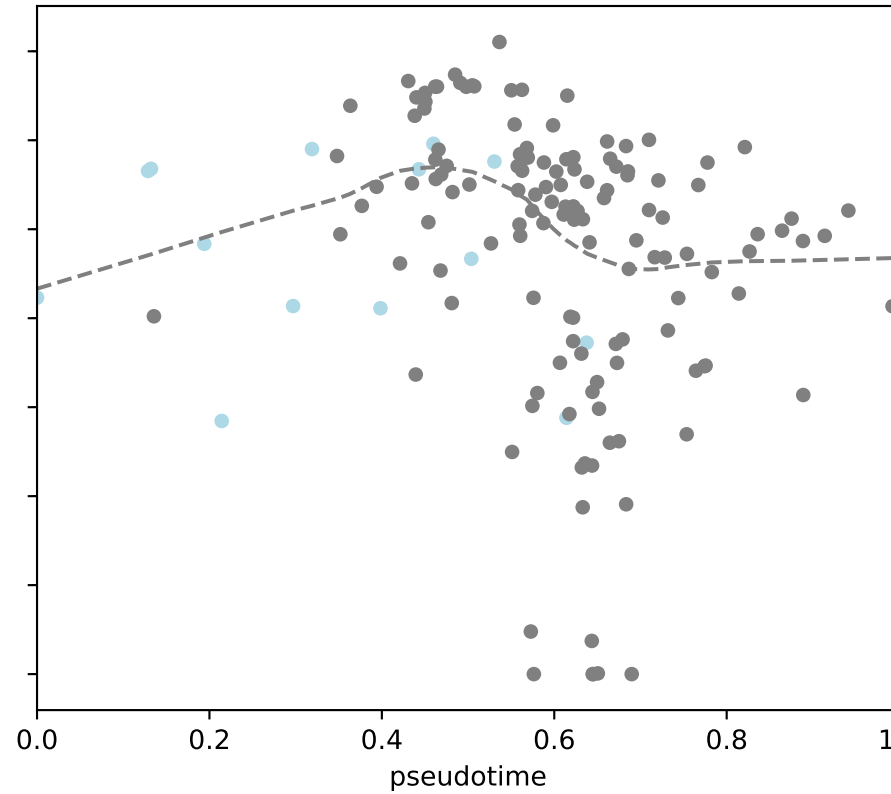
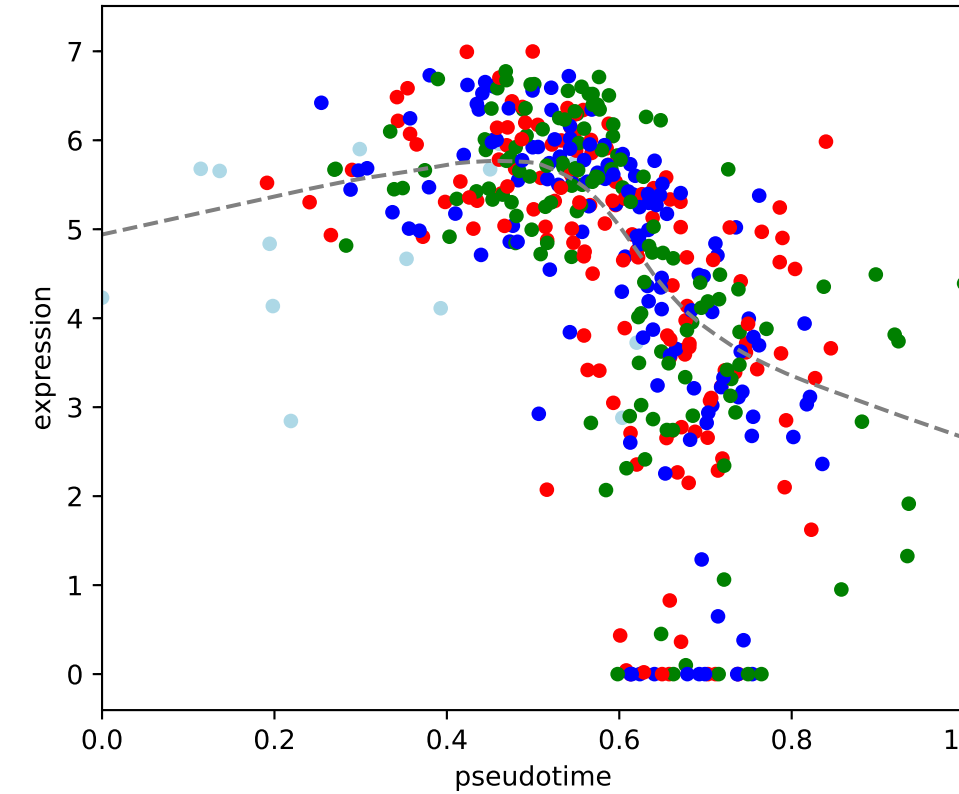
733 0spearman=0.66

733 1spearman=0.35



733 0spearman=-0.66

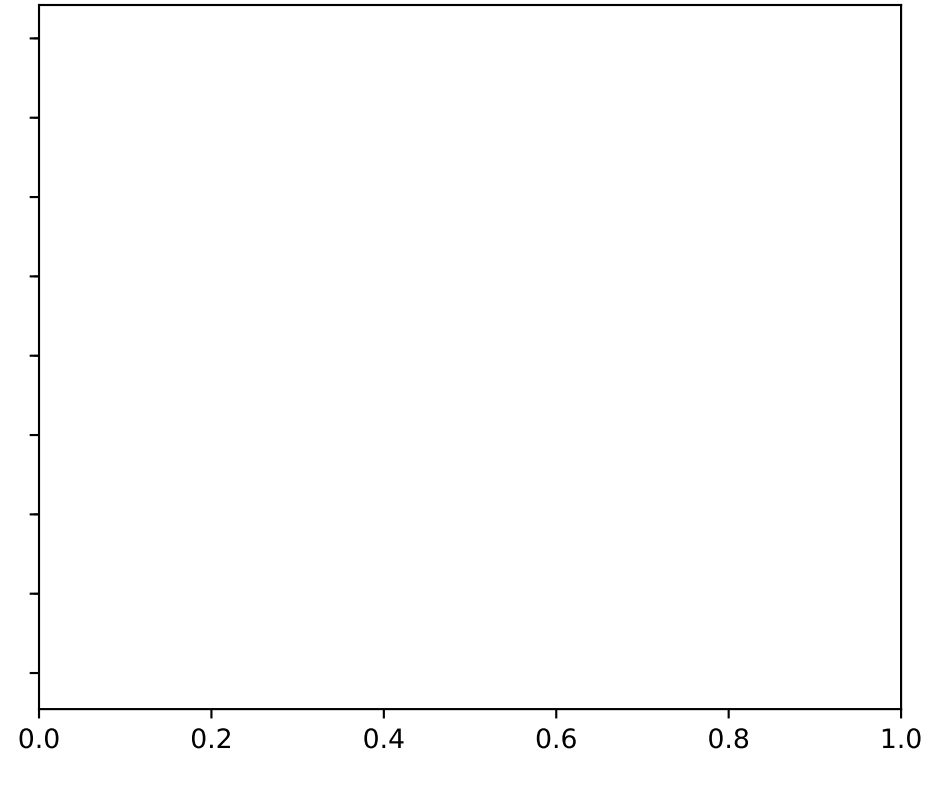
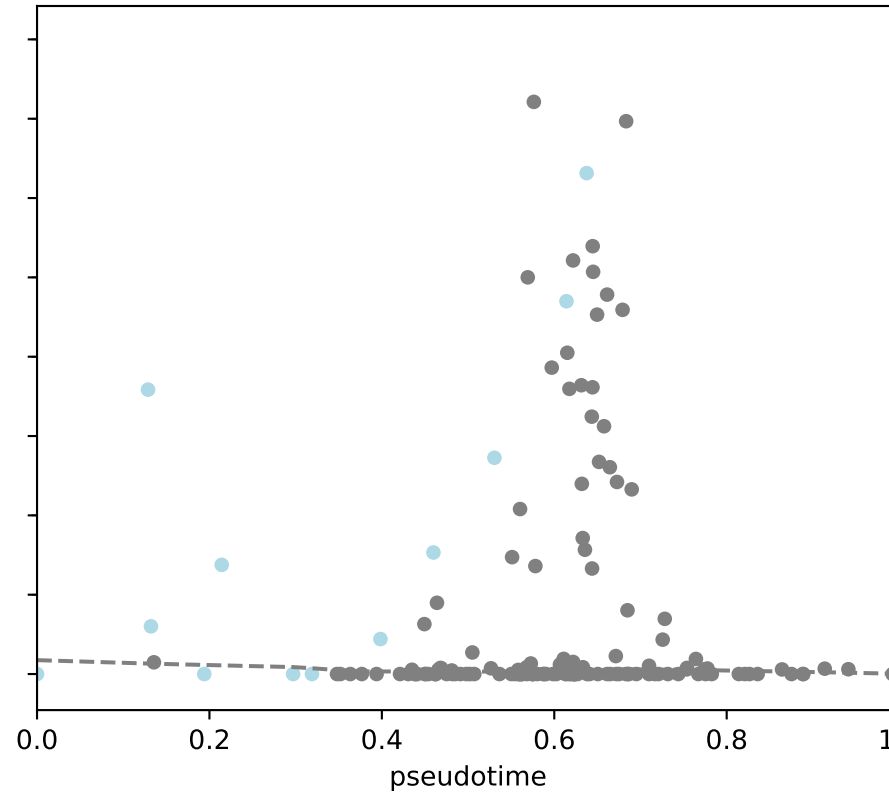
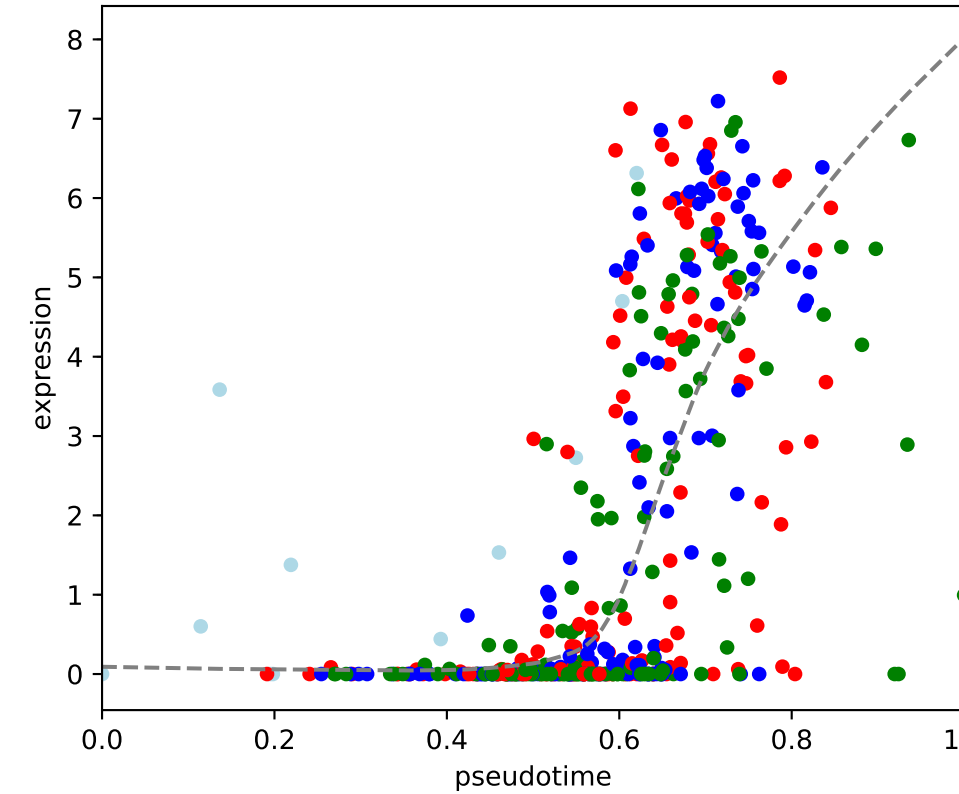
733 1spearman=-0.37



ENST00000232461 GNAT1 (G protein subunit alpha transducin 1)

733 0spearman=0.66

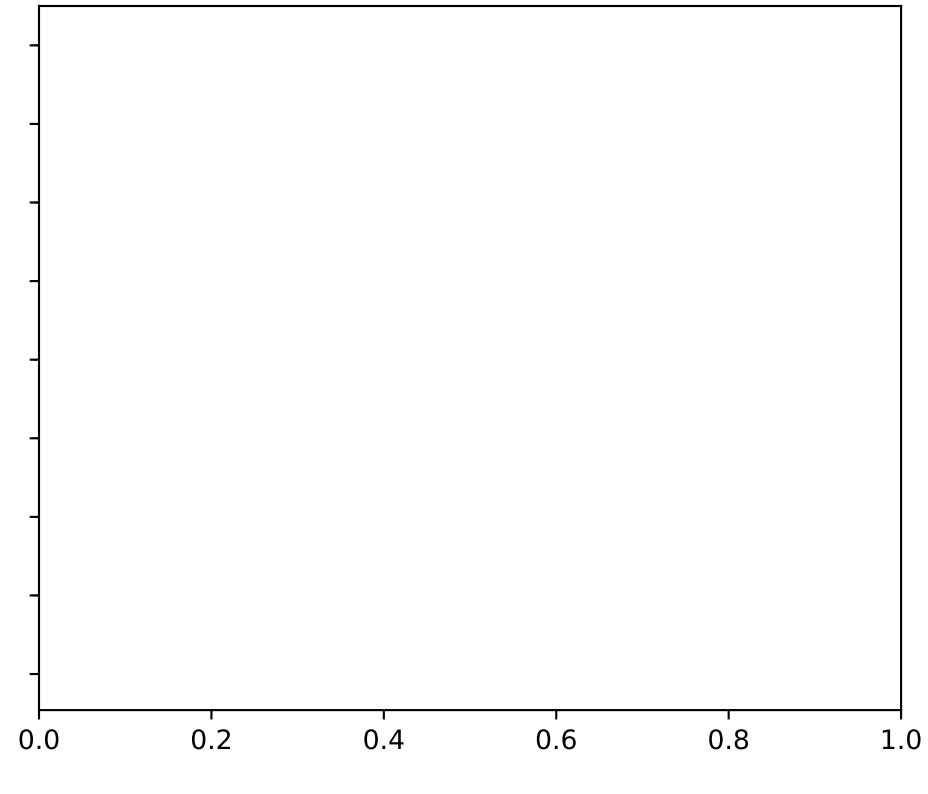
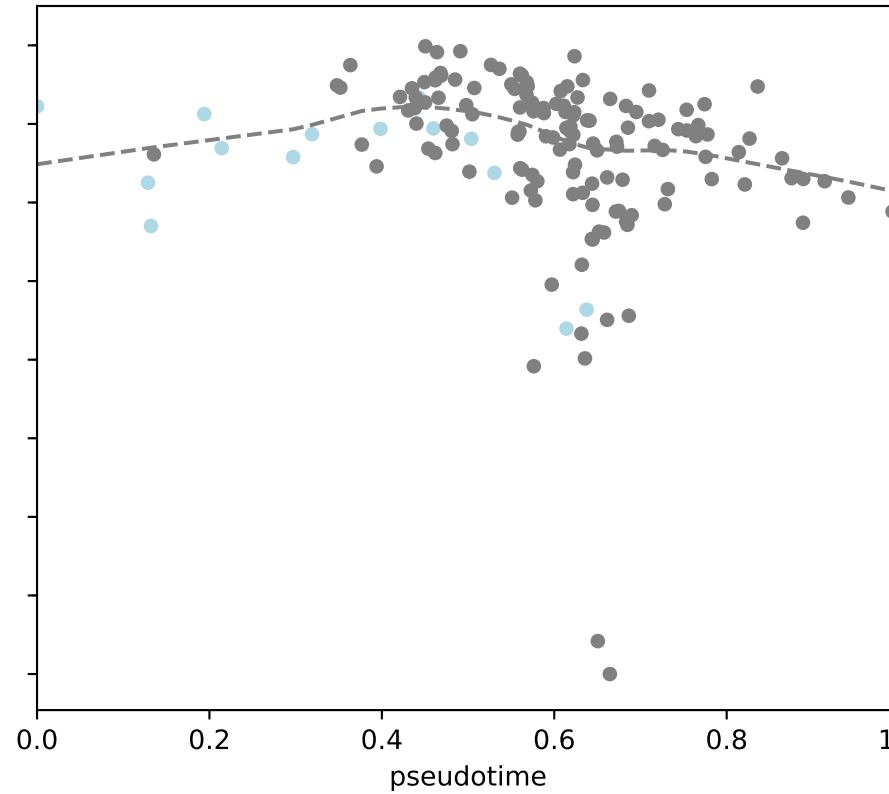
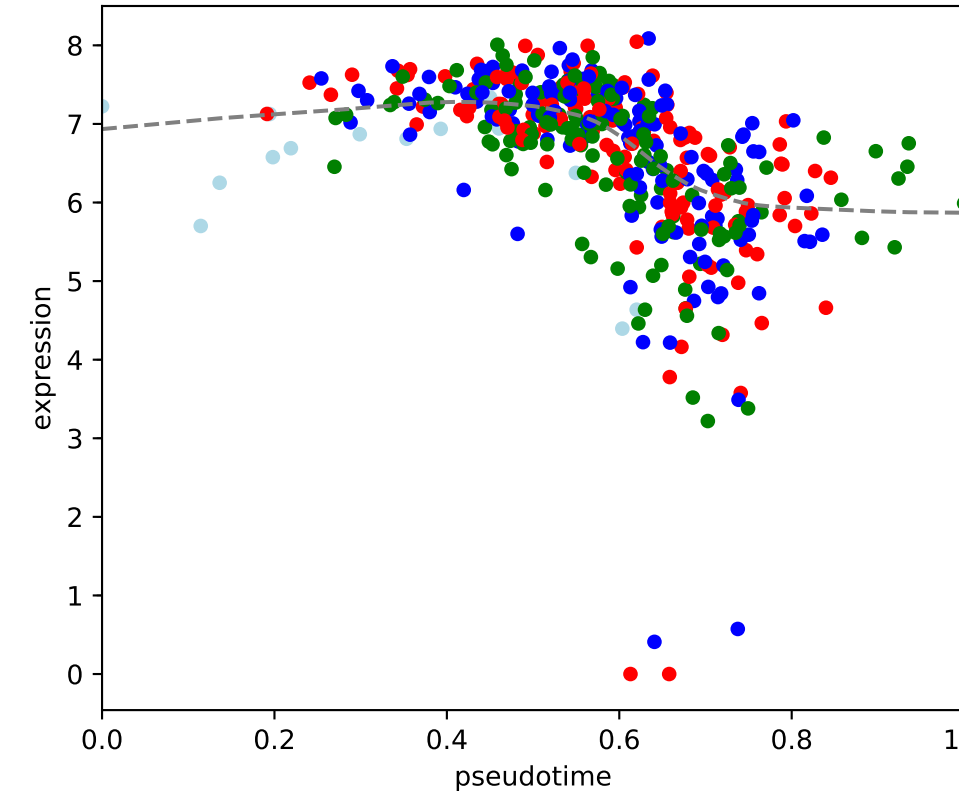
733 1spearman=0.11



ENST00000314133 COX8A (cytochrome c oxidase subunit 8A)

733 0spearman=-0.66

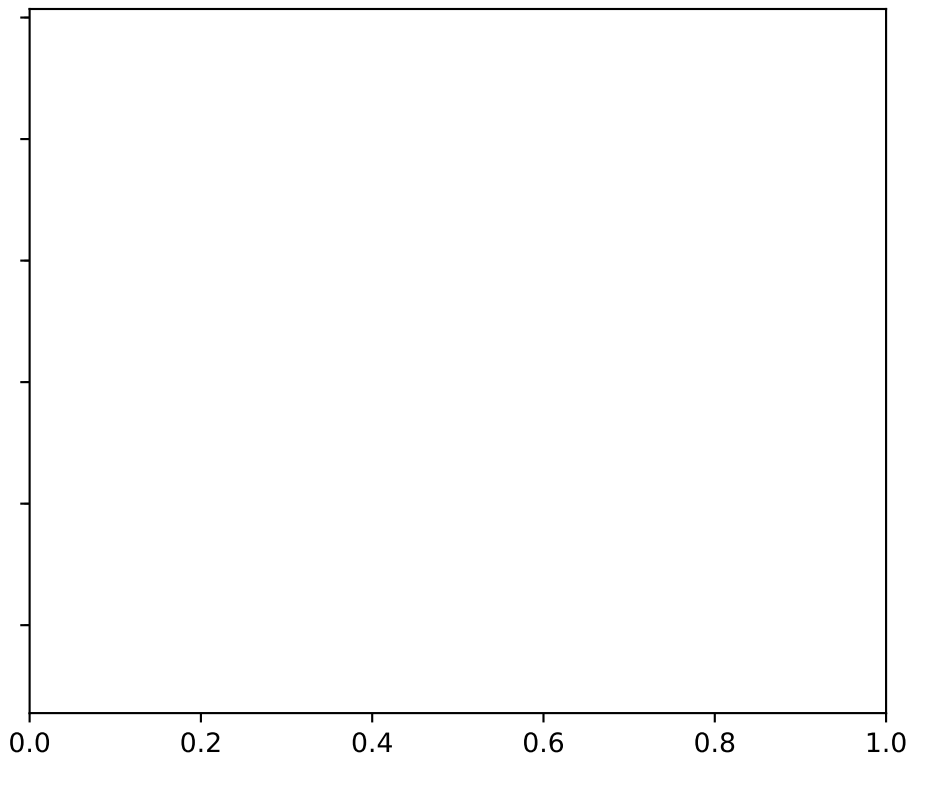
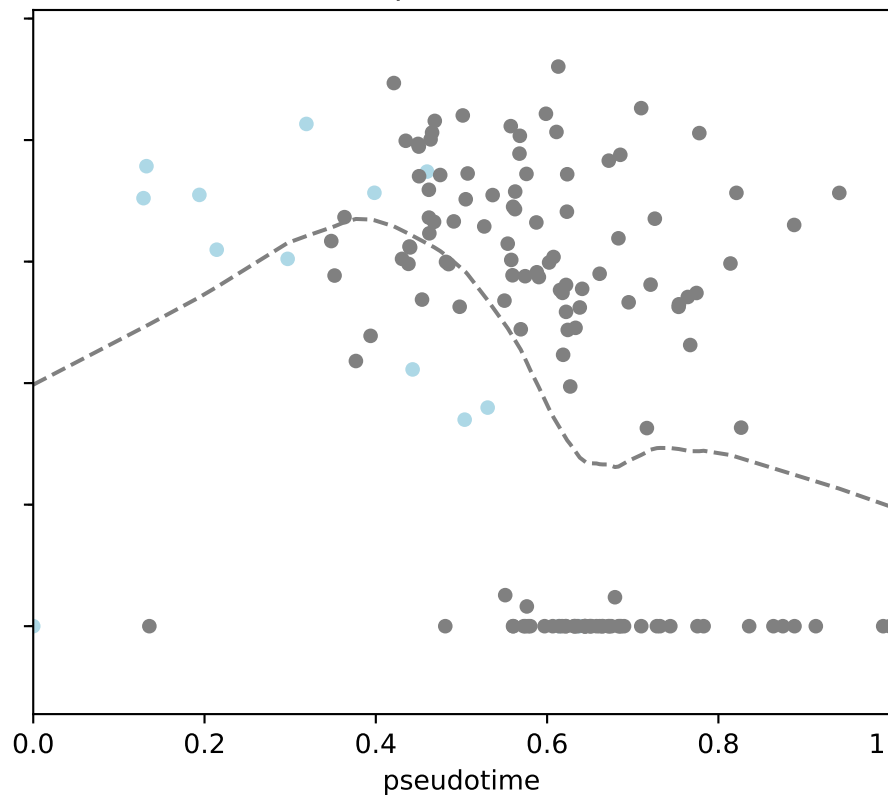
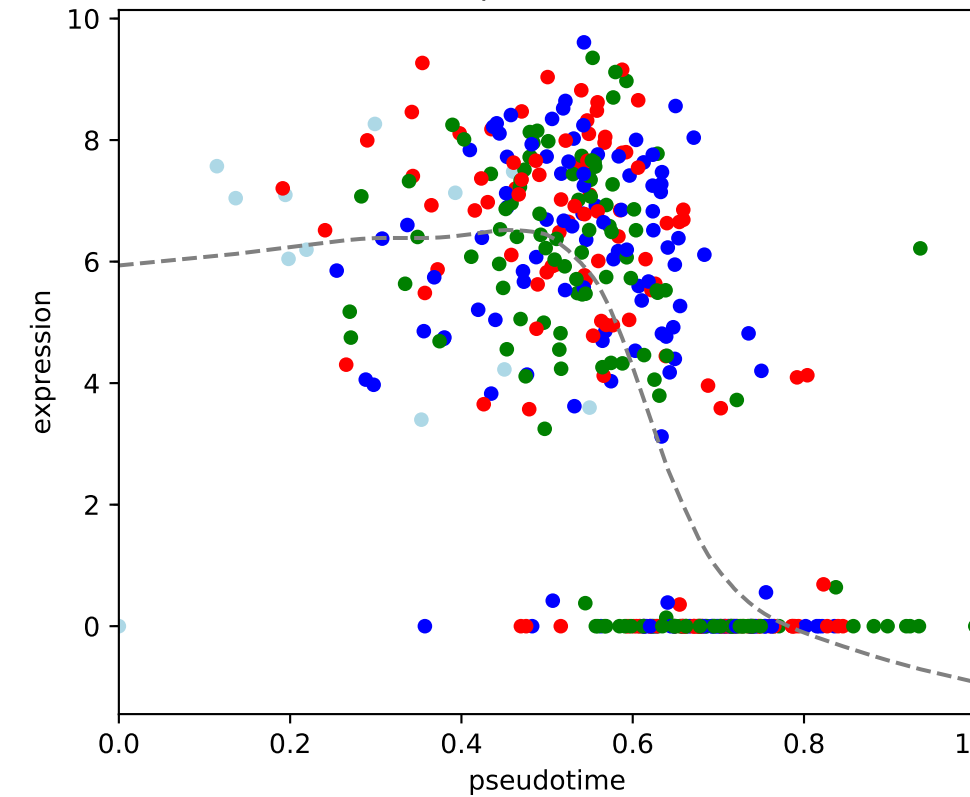
733 1spearman=-0.41



ENST00000372958 GUCA1A (guanylate cyclase activator 1A)

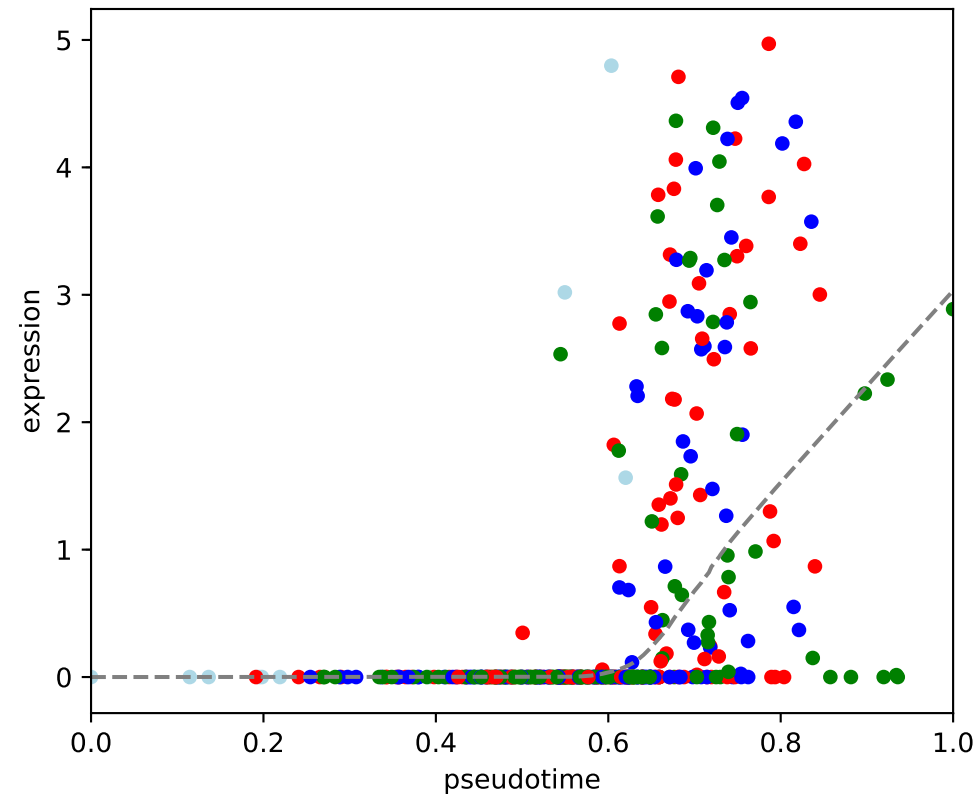
733 0spearman=-0.65

733 1spearman=-0.45

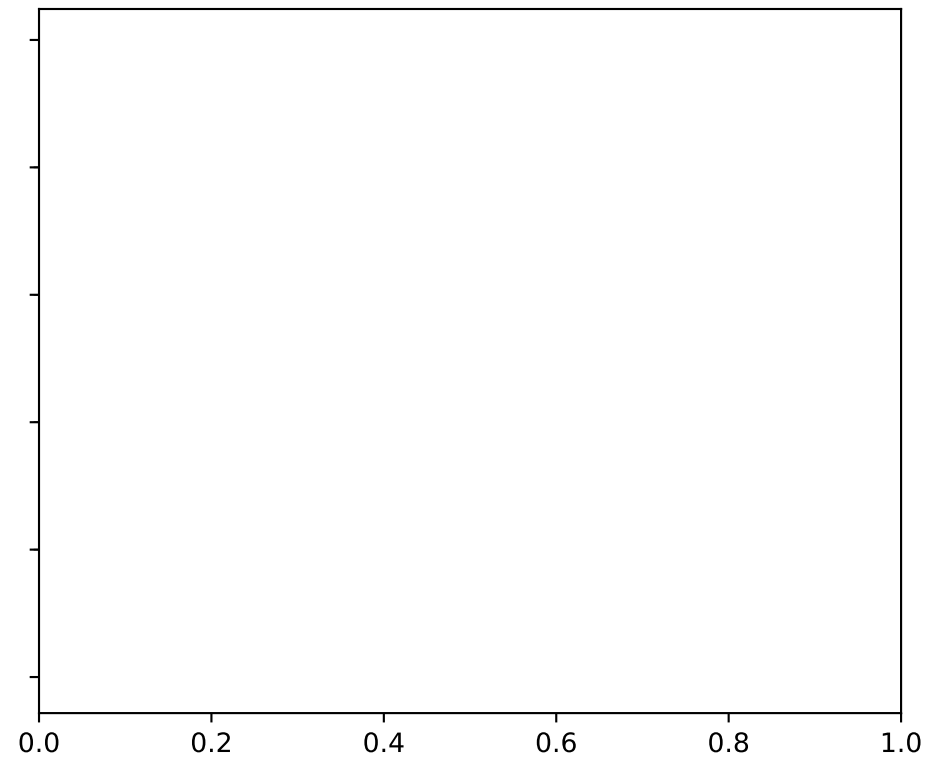
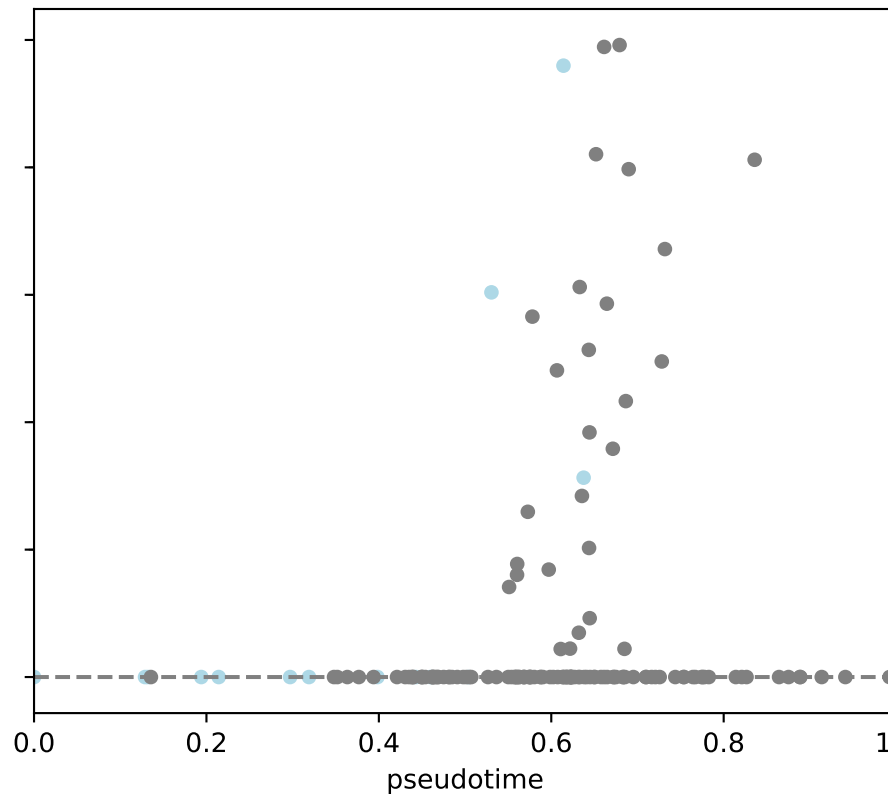


ENST00000561834 AC104938.1

733 0spearman=0.64



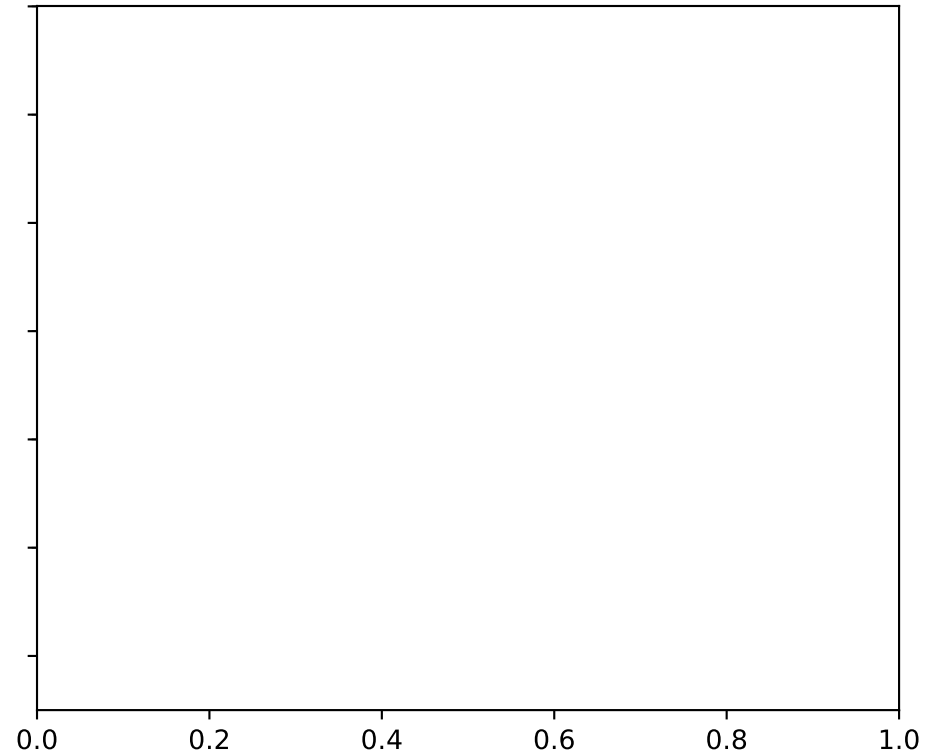
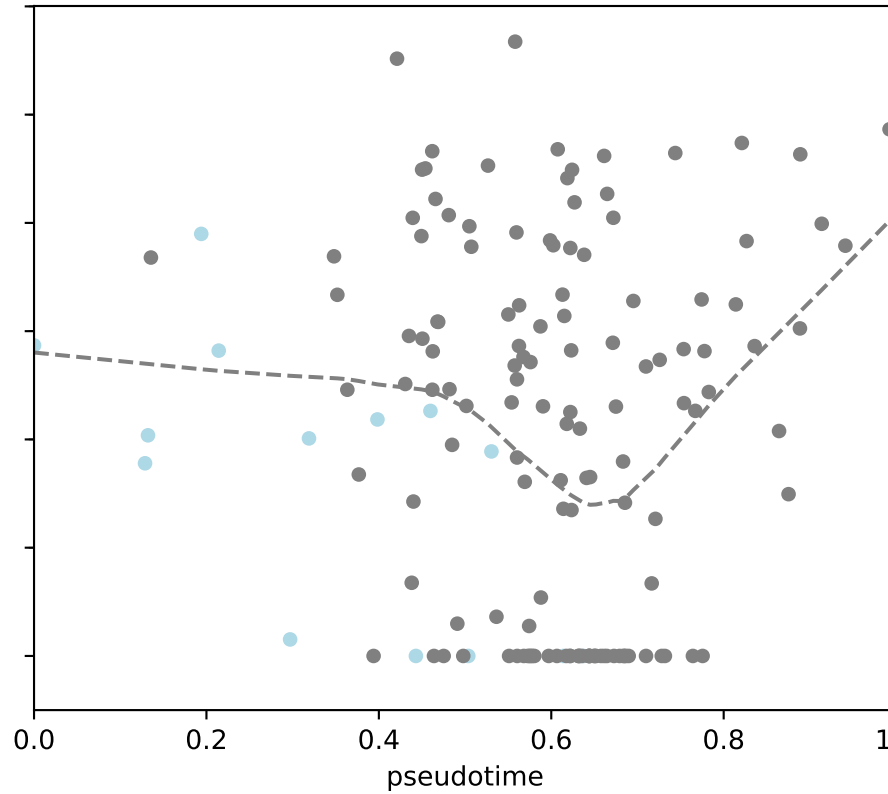
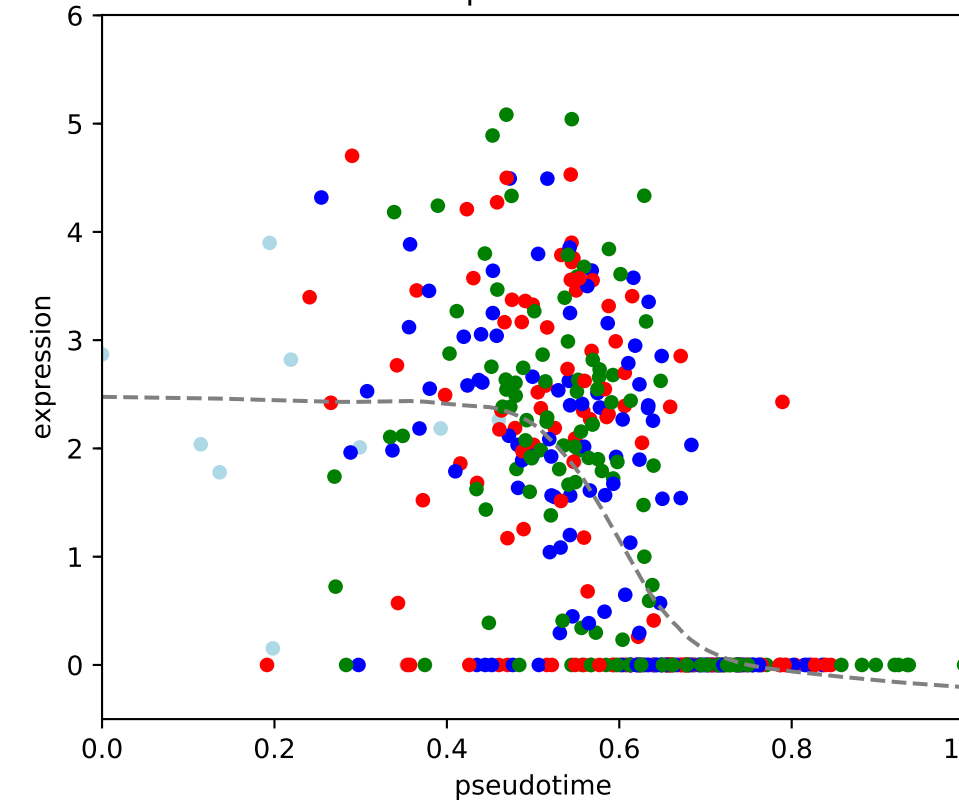
733 1spearman=0.21



ENST00000371469 RBP4 (retinol binding protein 4)

733 0spearman=-0.64

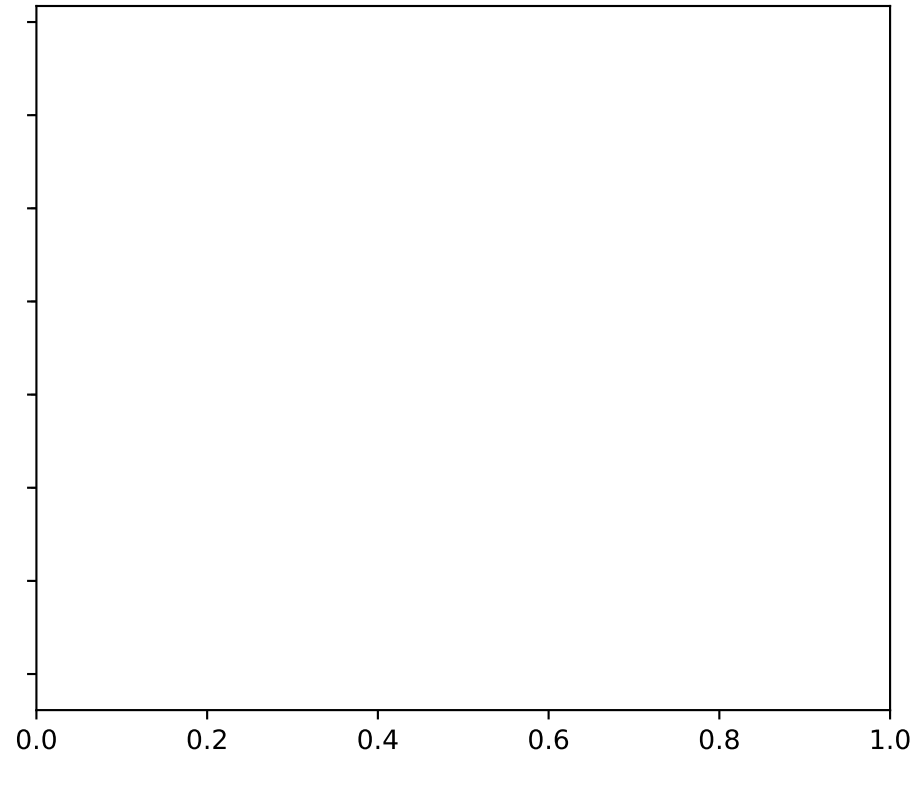
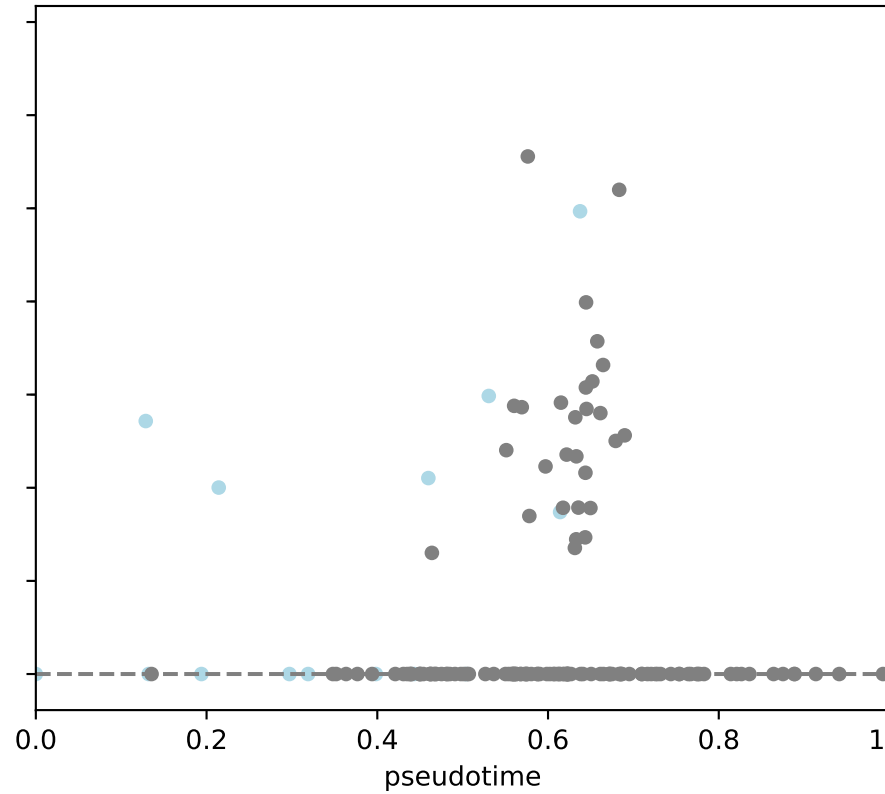
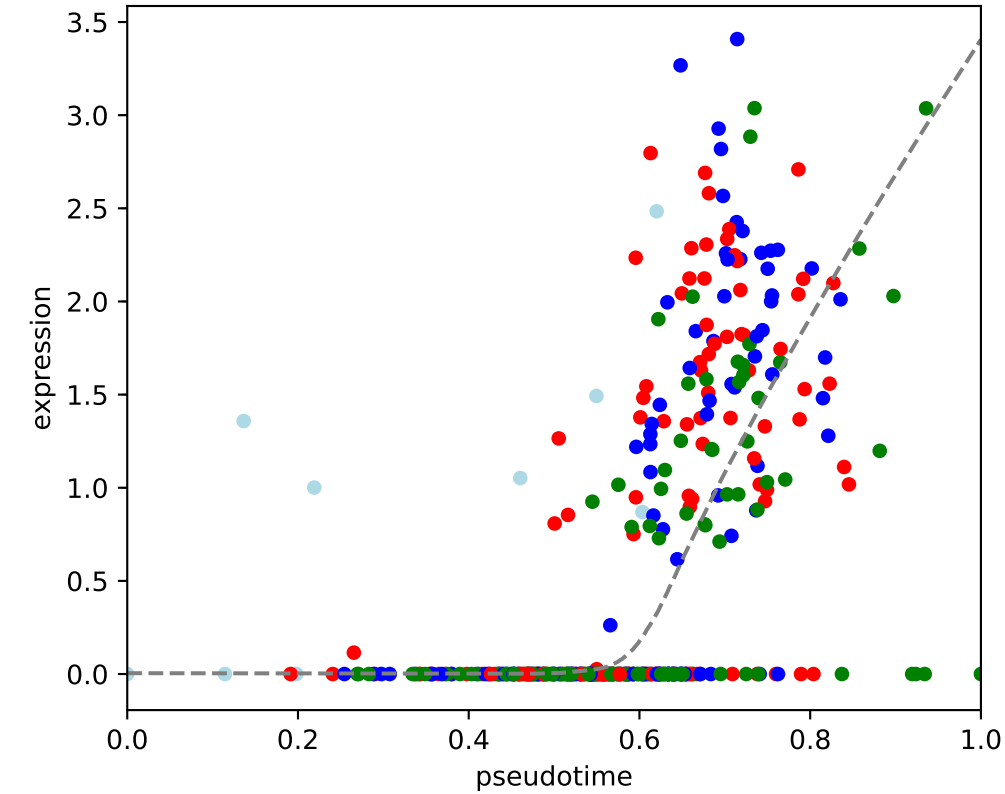
733 1spearman=-0.11



ENST00000440836 GNAT1 (G protein subunit alpha transducin 1)

733 0spearman=0.64

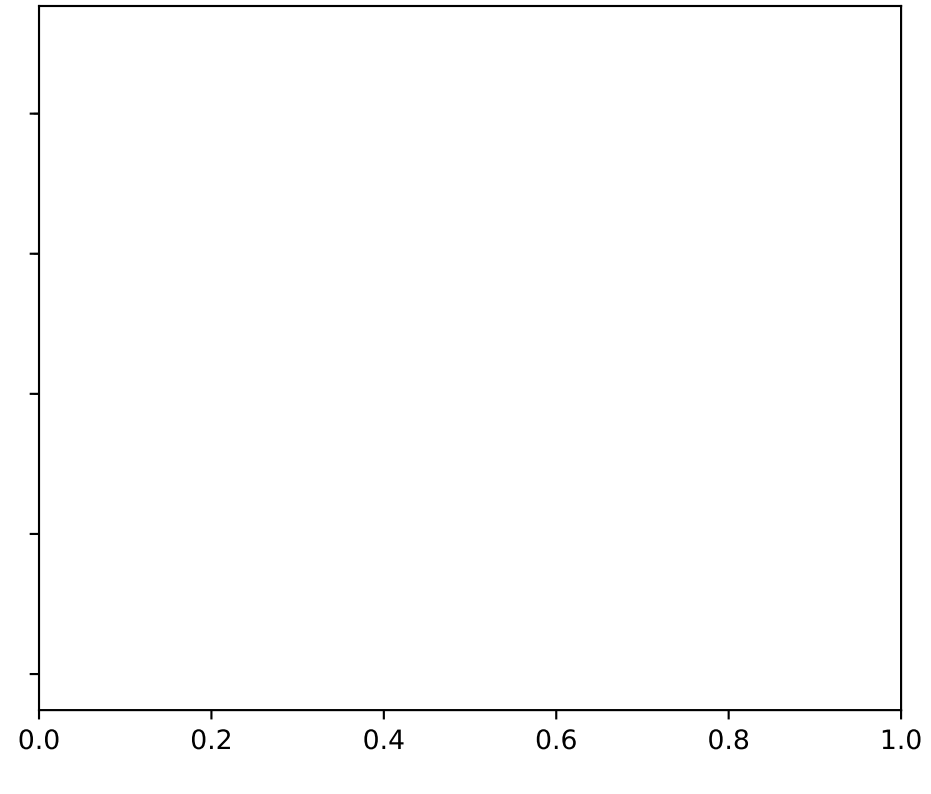
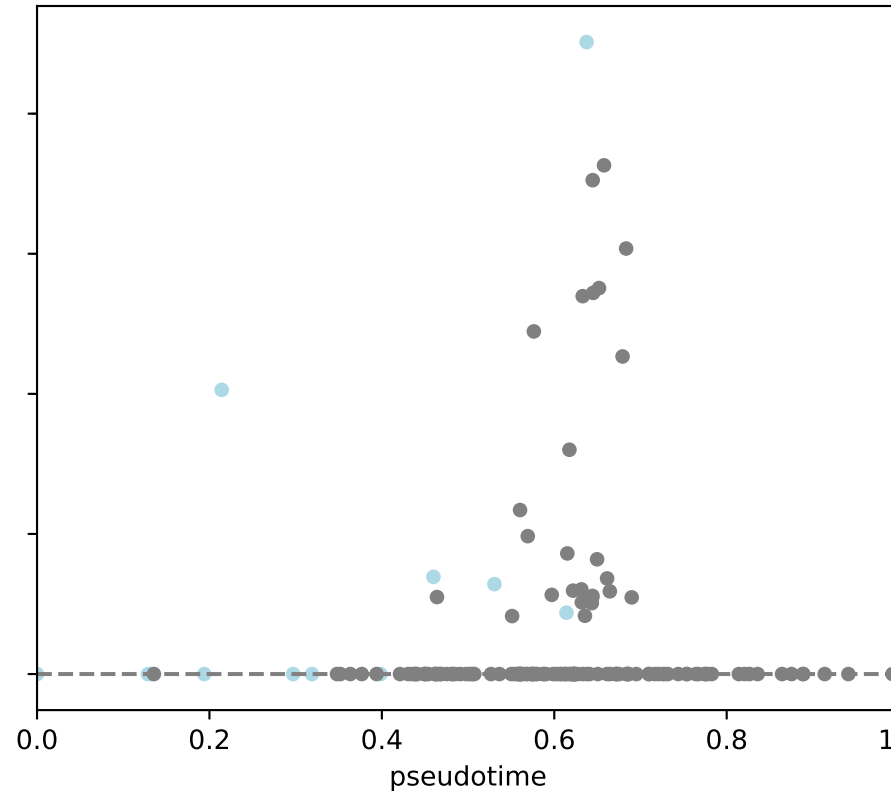
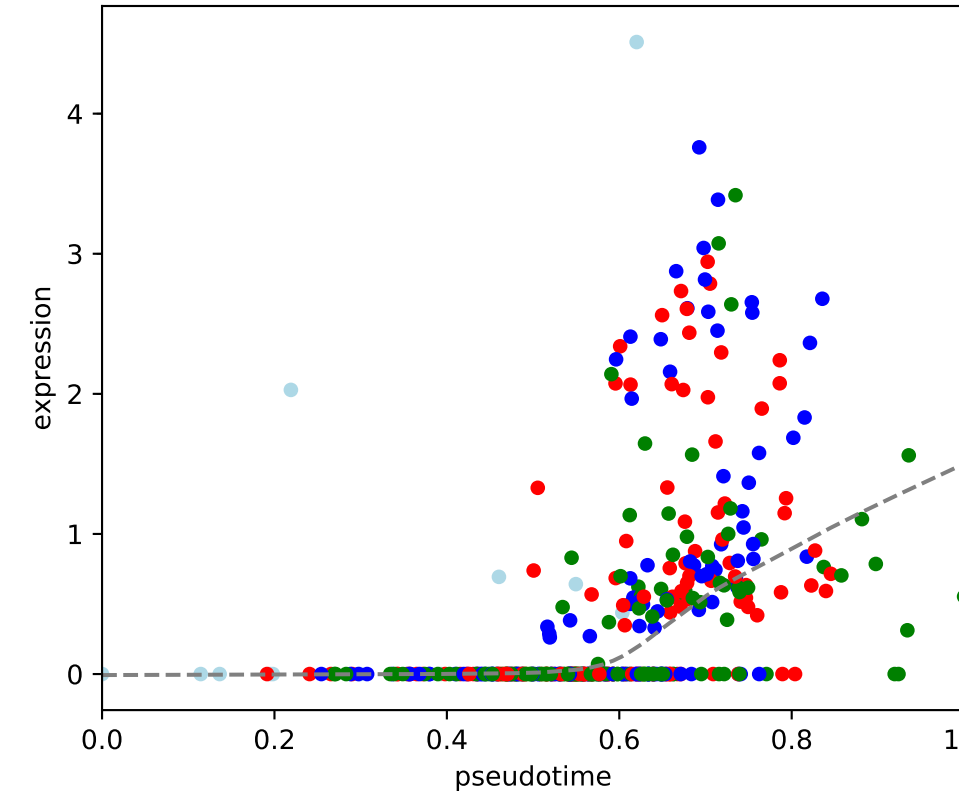
733 1spearman=0.07



ENST00000481246 GNAT1 (G protein subunit alpha transducin 1)

733 0spearman=0.64

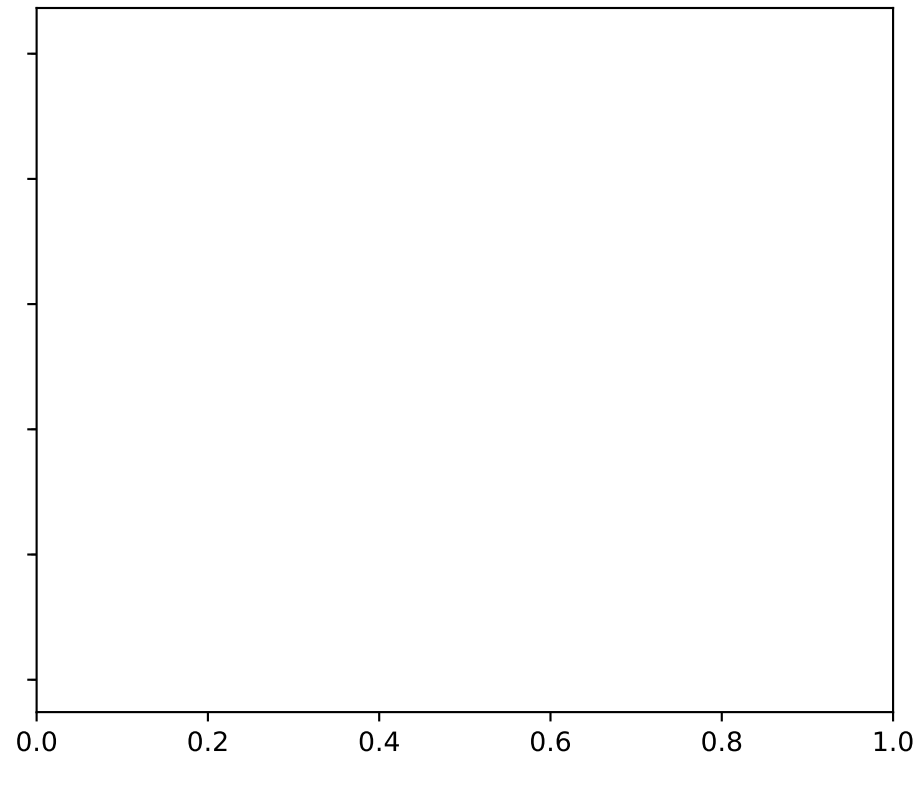
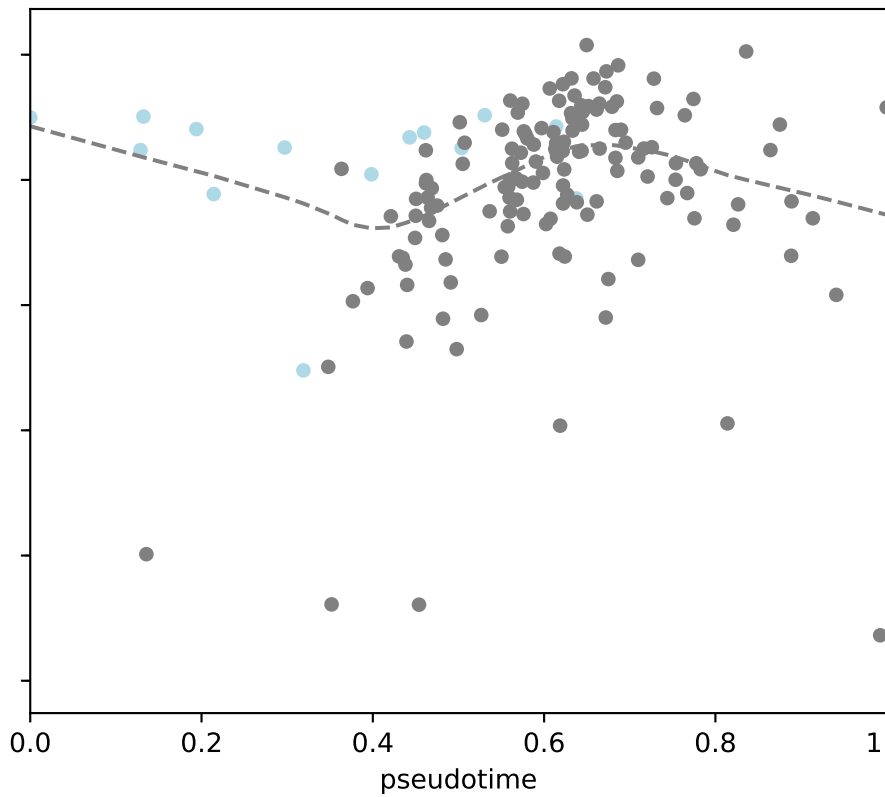
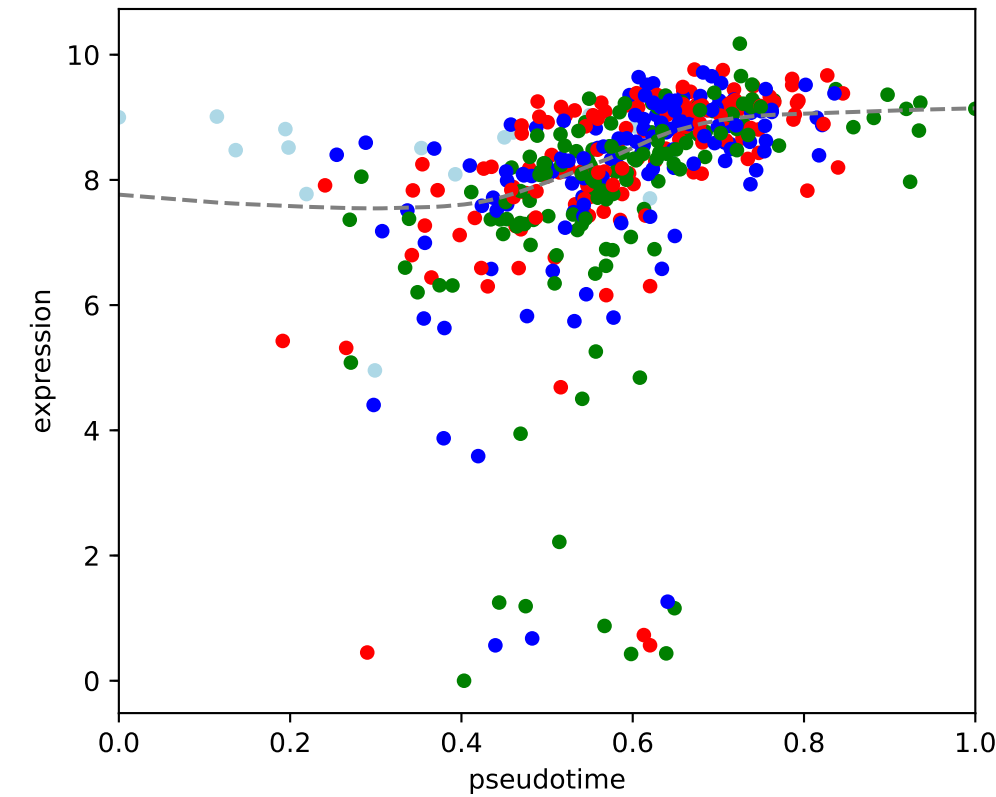
733 1spearman=0.09



ENST00000391997 PDC (phosducin)

733 0spearman=0.63

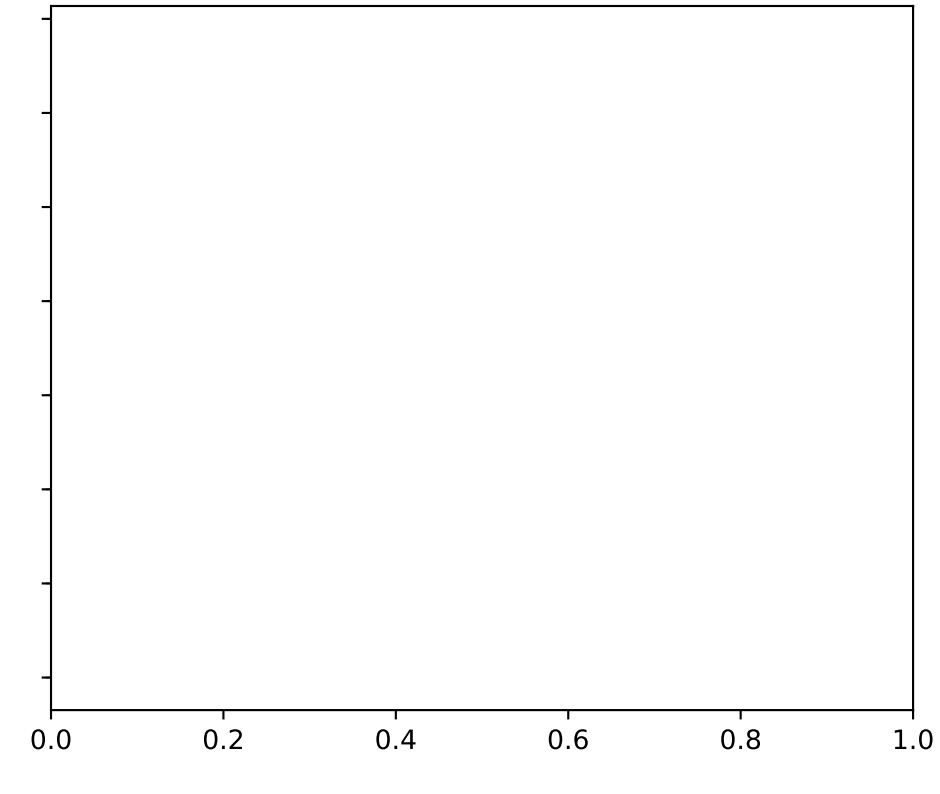
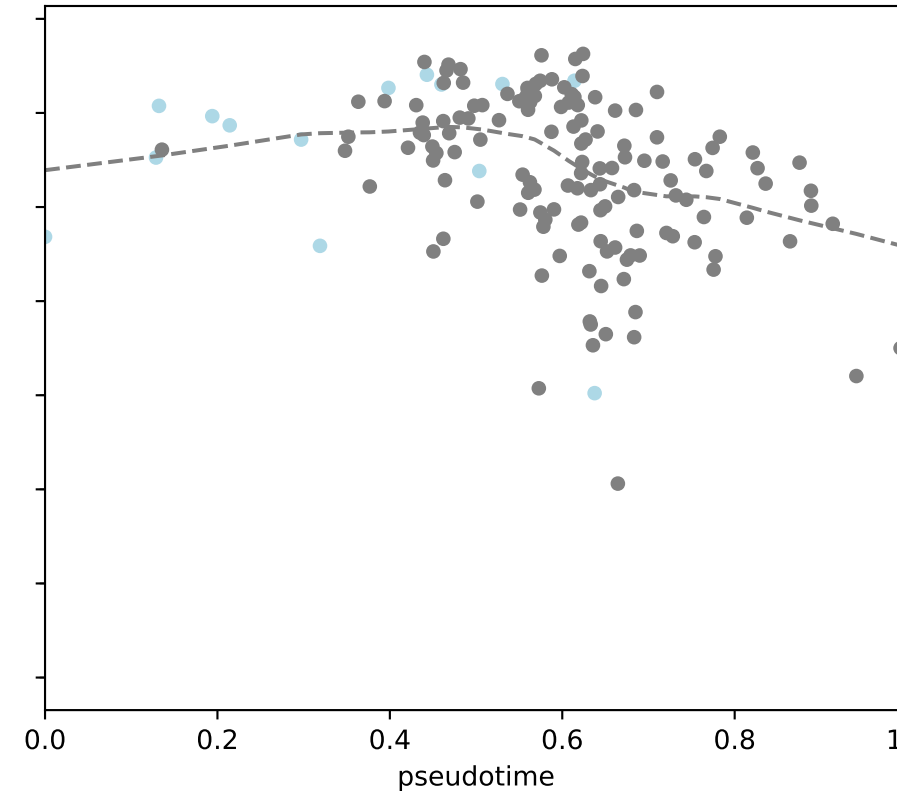
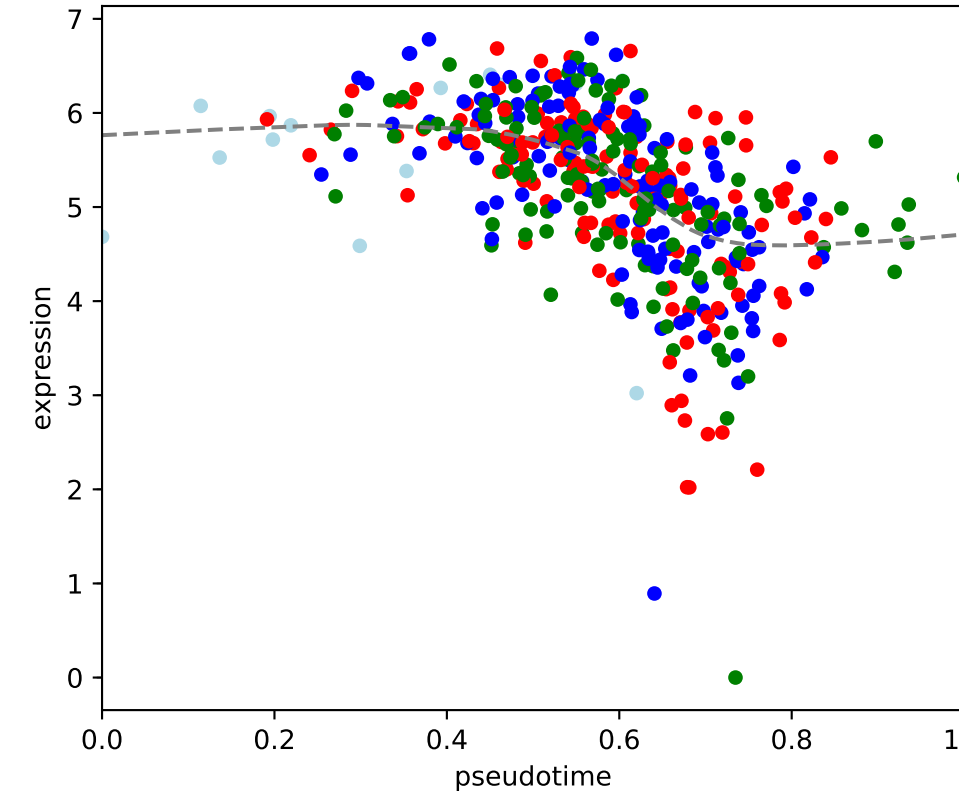
733 1spearman=0.28



ENST00000222247 RPL18A (ribosomal protein L18a)

733 0spearman=-0.63

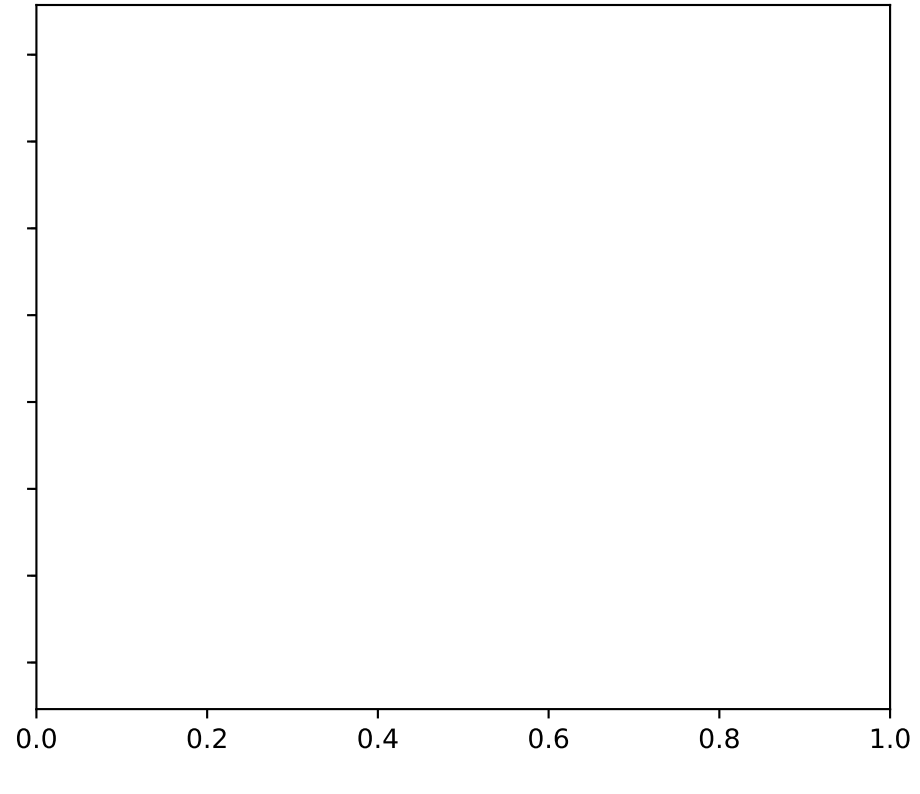
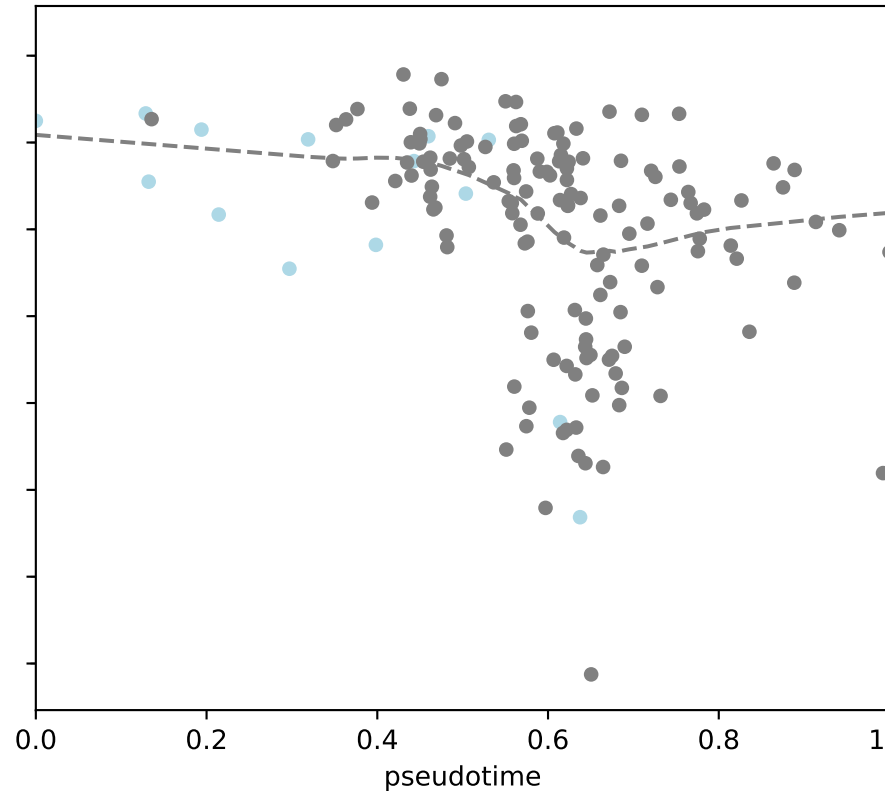
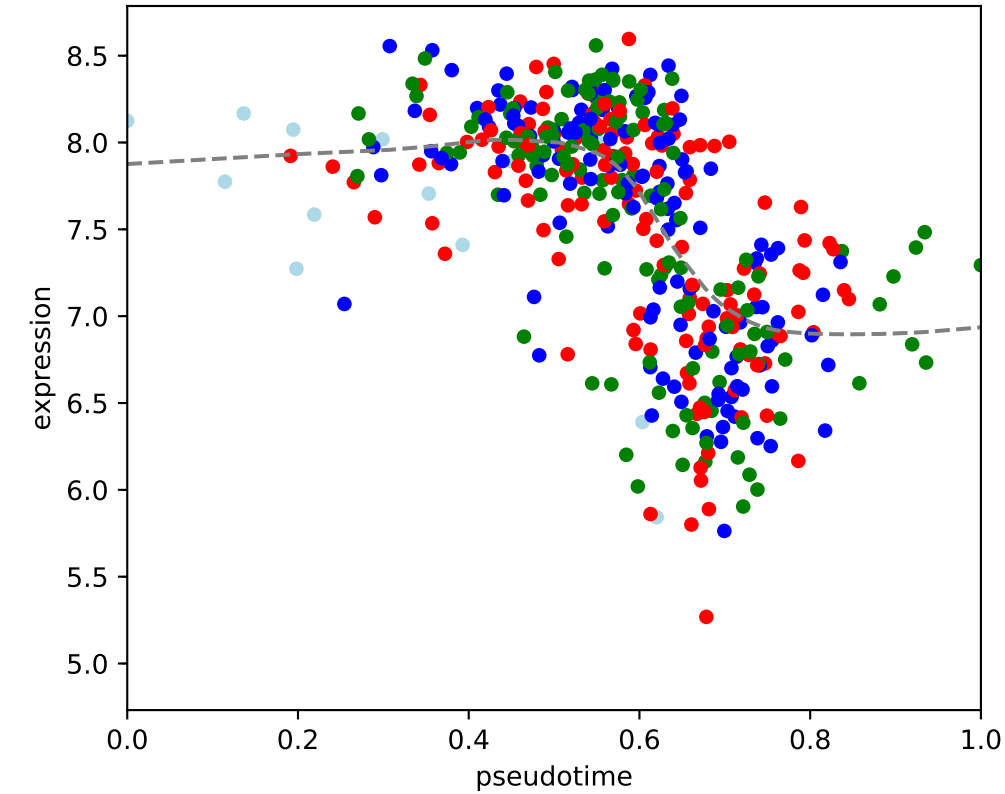
733 1spearman=-0.45



ENST00000262030 ATP5F1B (ATP synthase F1 subunit beta)

733 0spearman=-0.63

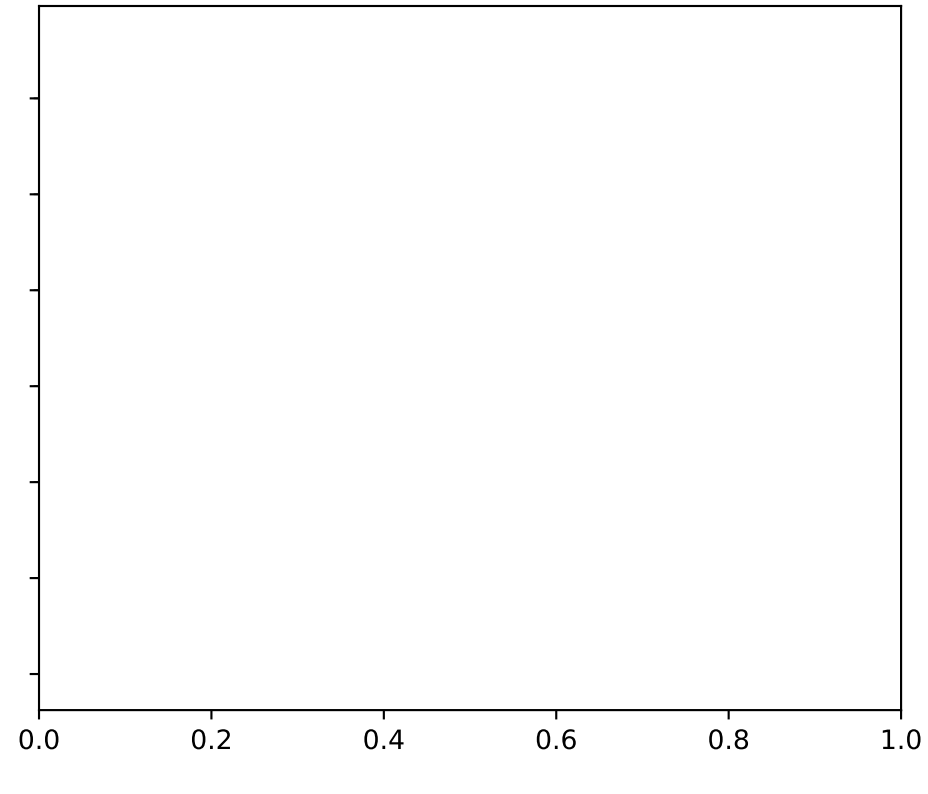
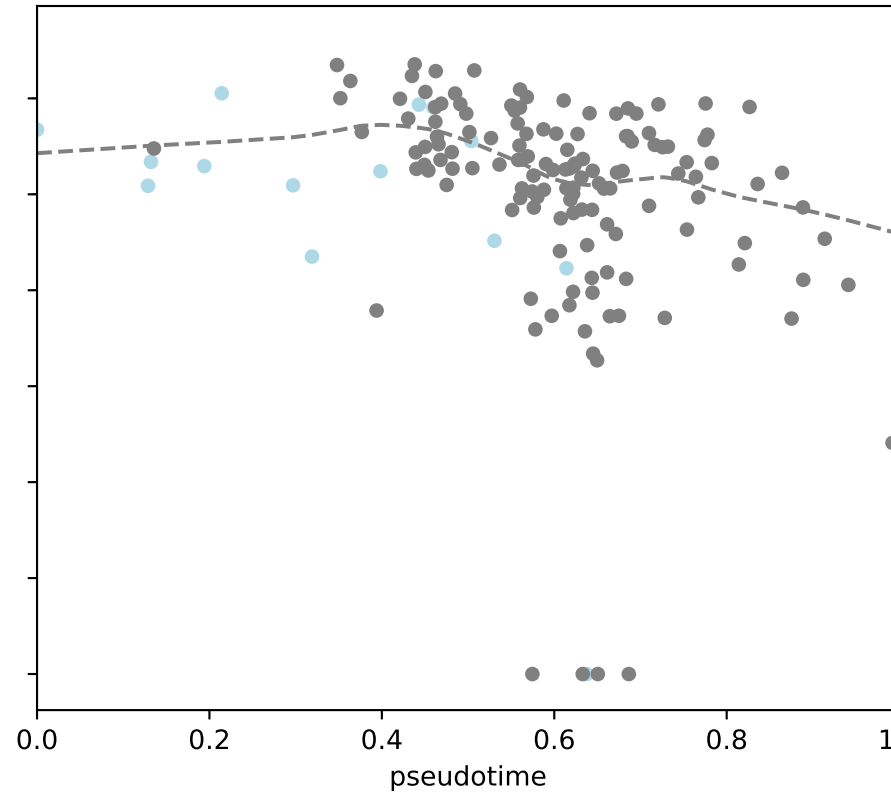
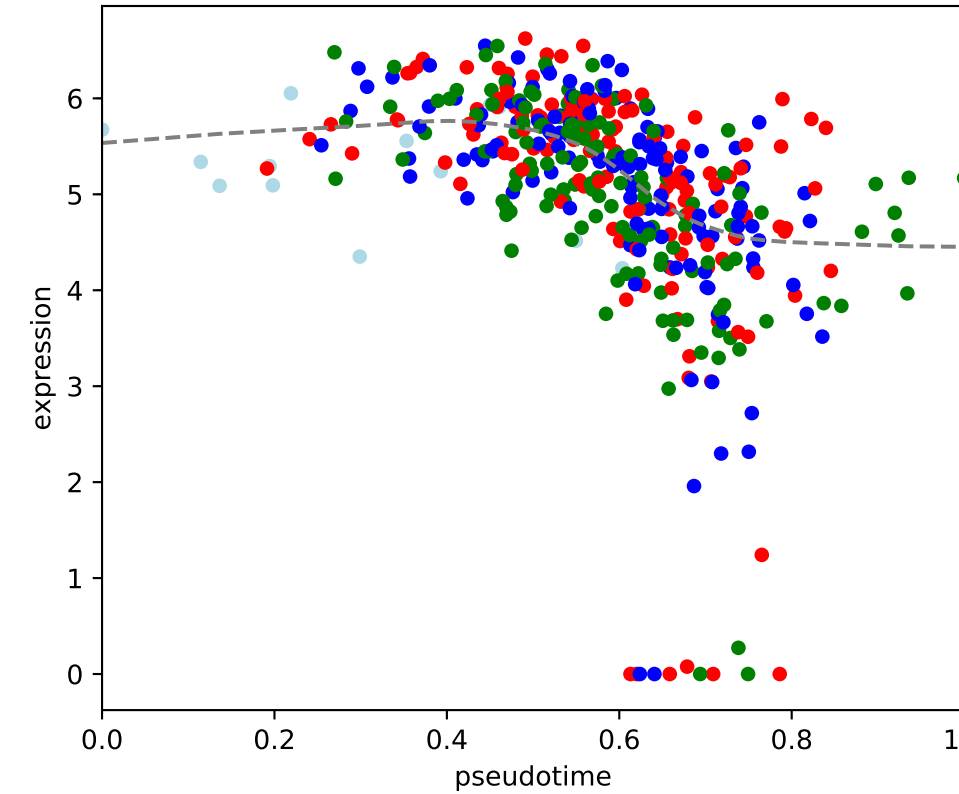
733 1spearman=-0.47



ENST00000313468 NDUFS8 (NADH:ubiquinone oxidoreductase core subunit S8)

733 0spearman=-0.63

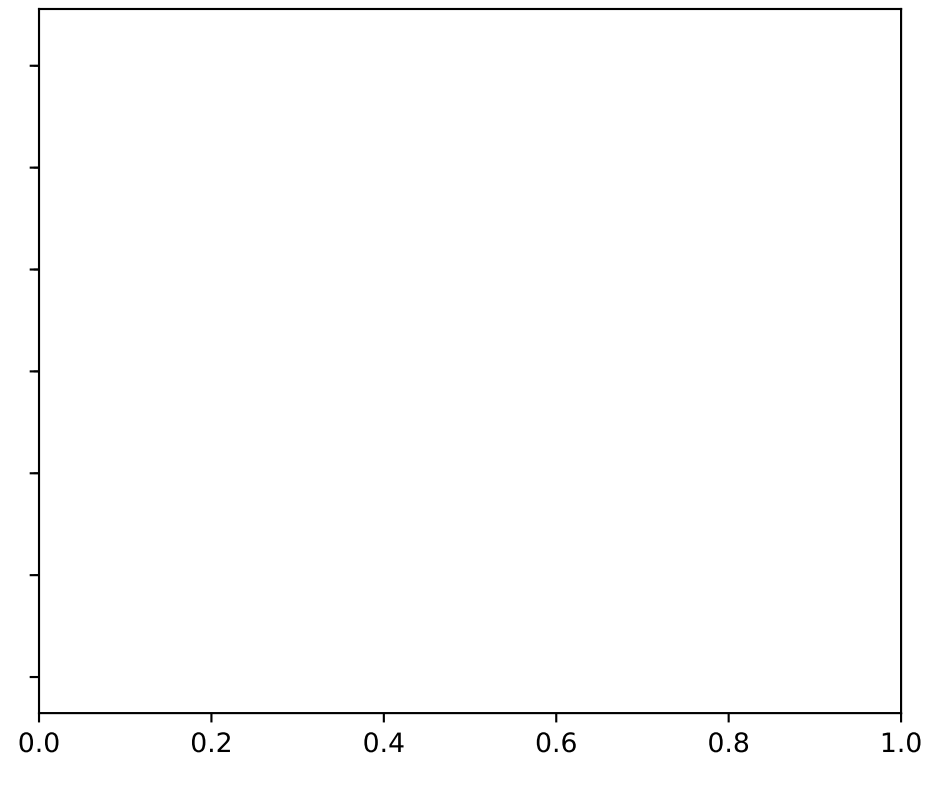
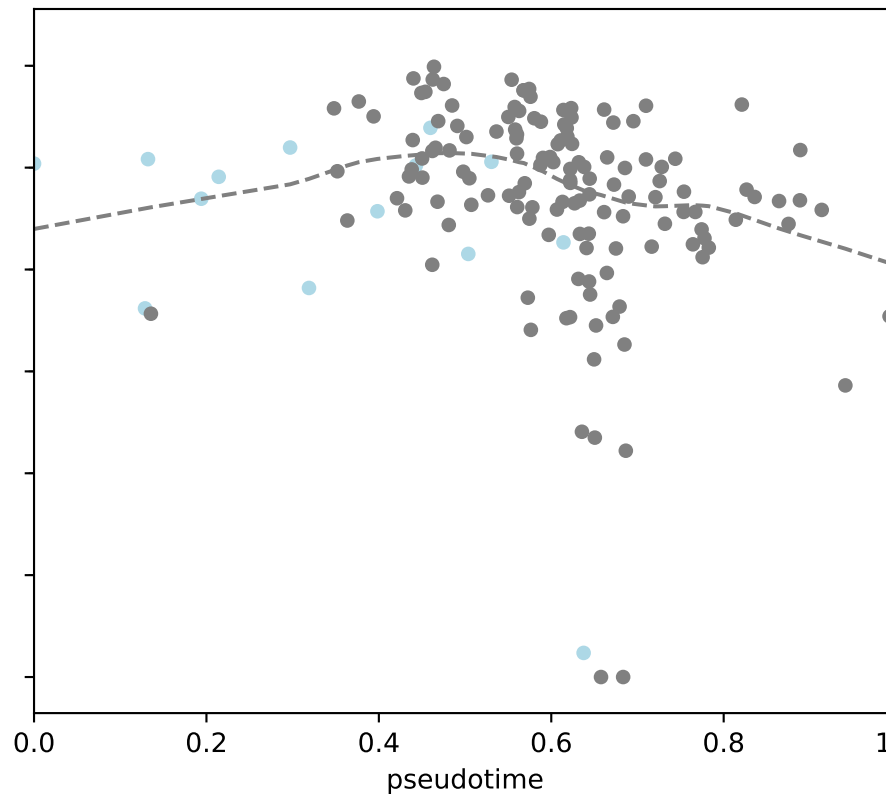
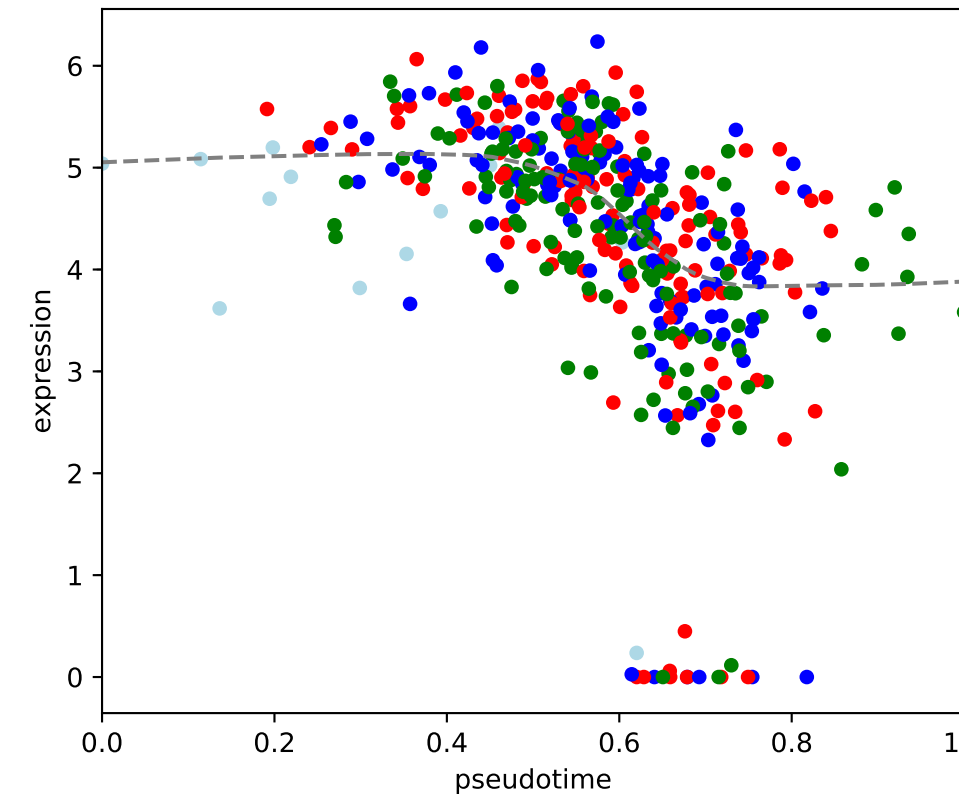
733 1spearman=-0.40



ENST00000321153 RPLP2 (ribosomal protein lateral stalk subunit P2)

733 0spearman=-0.63

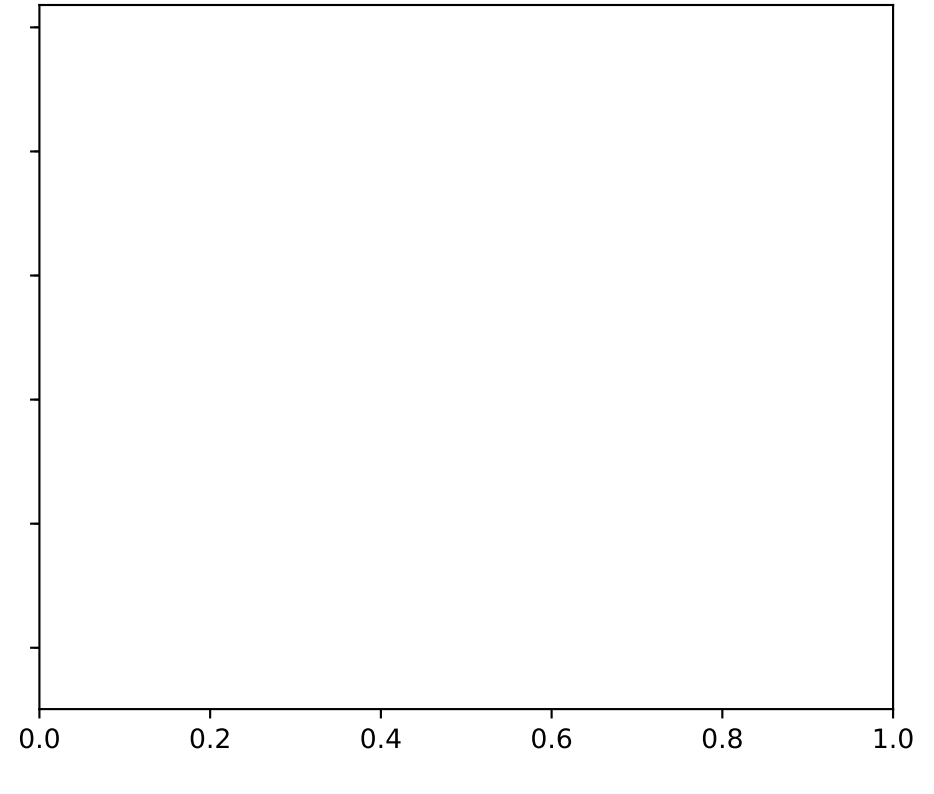
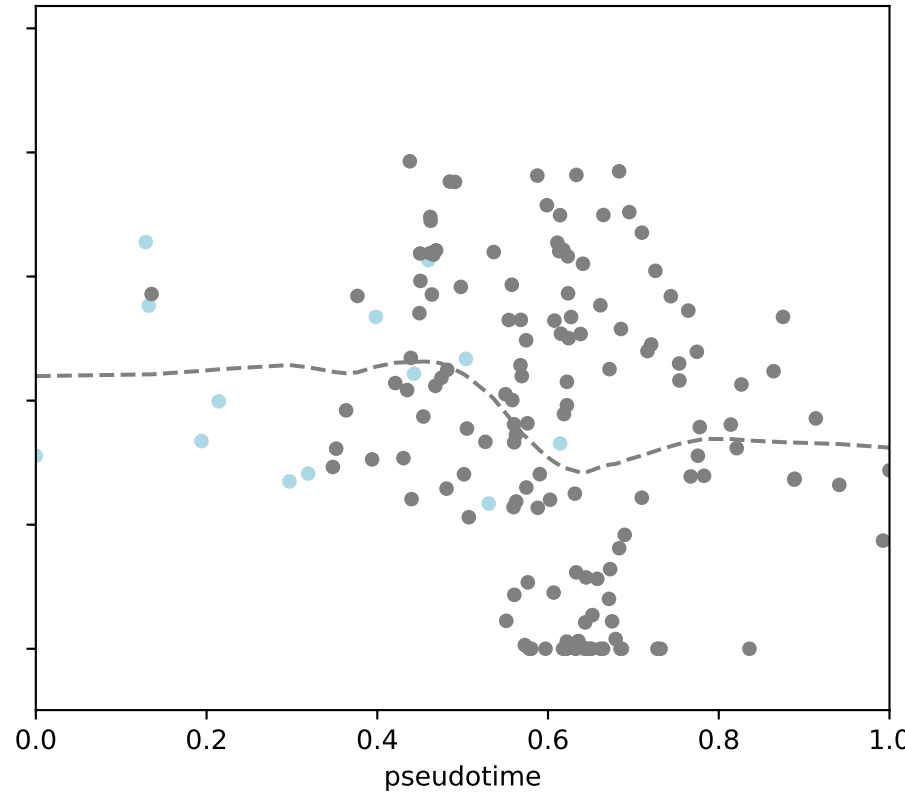
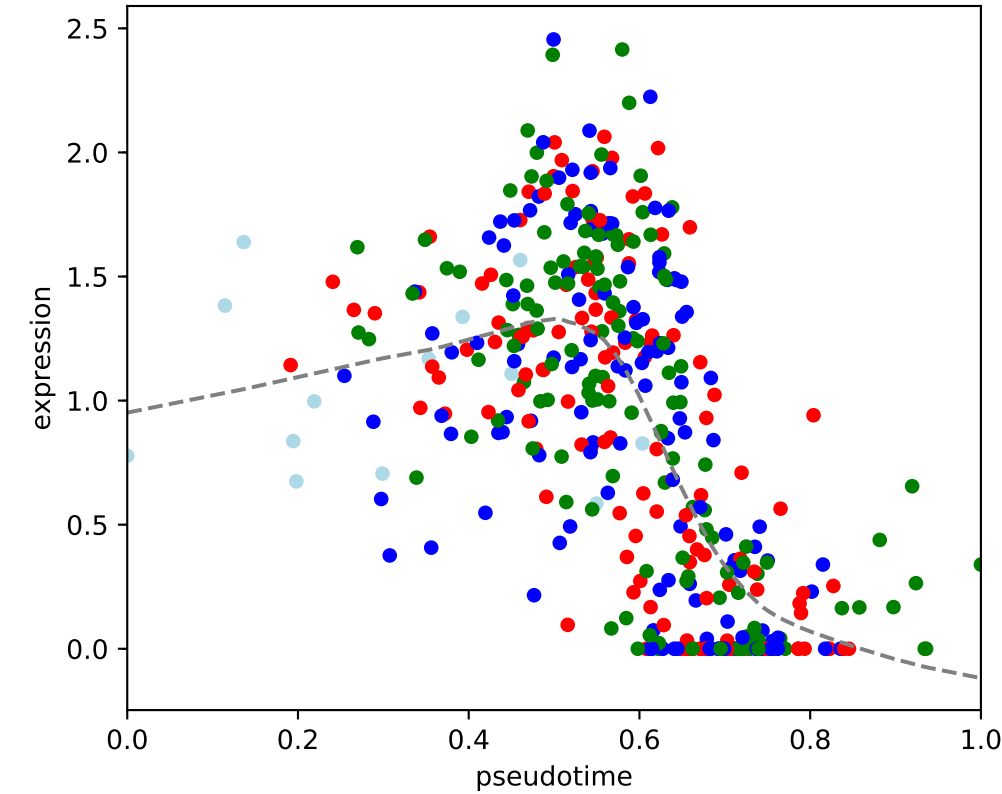
733 1spearman=-0.37



ENST00000523057 AC008429.2

733 0spearman=-0.63

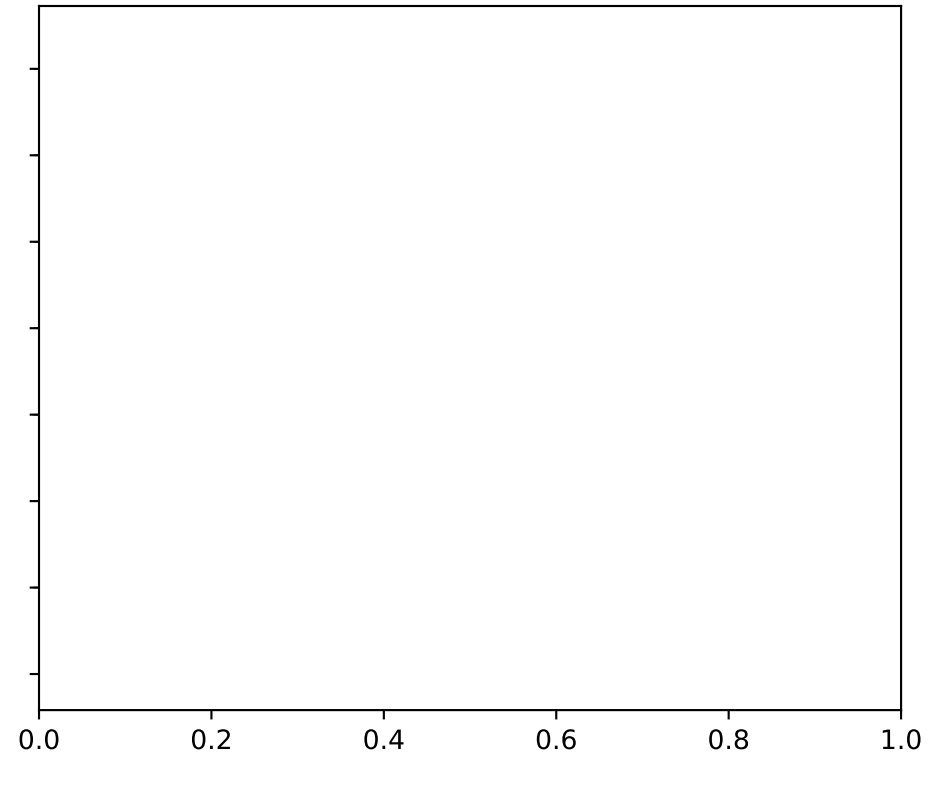
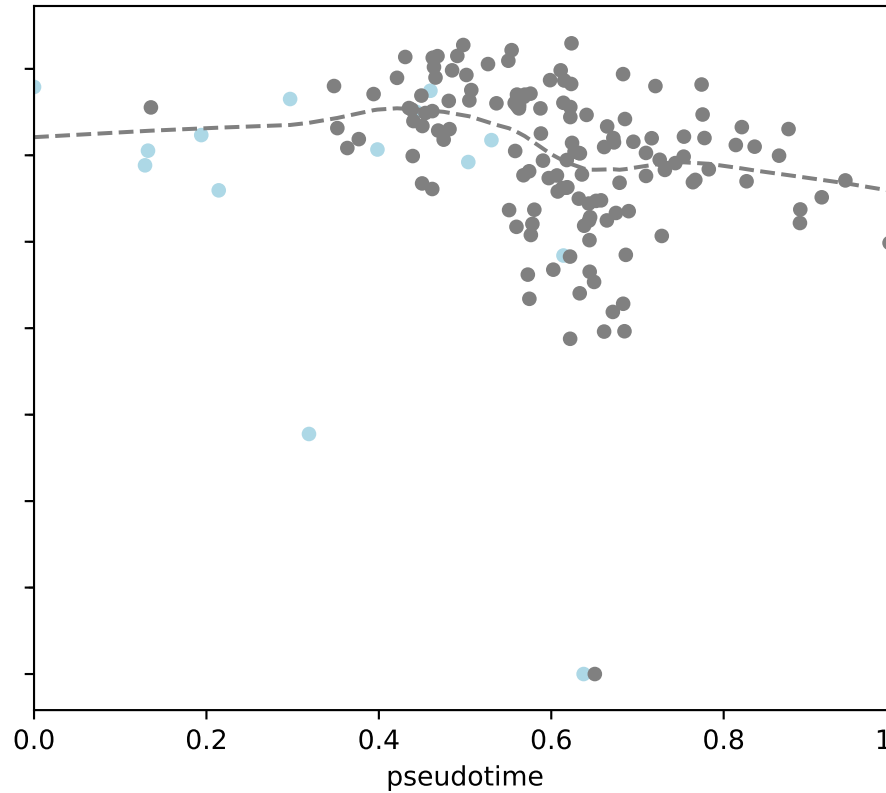
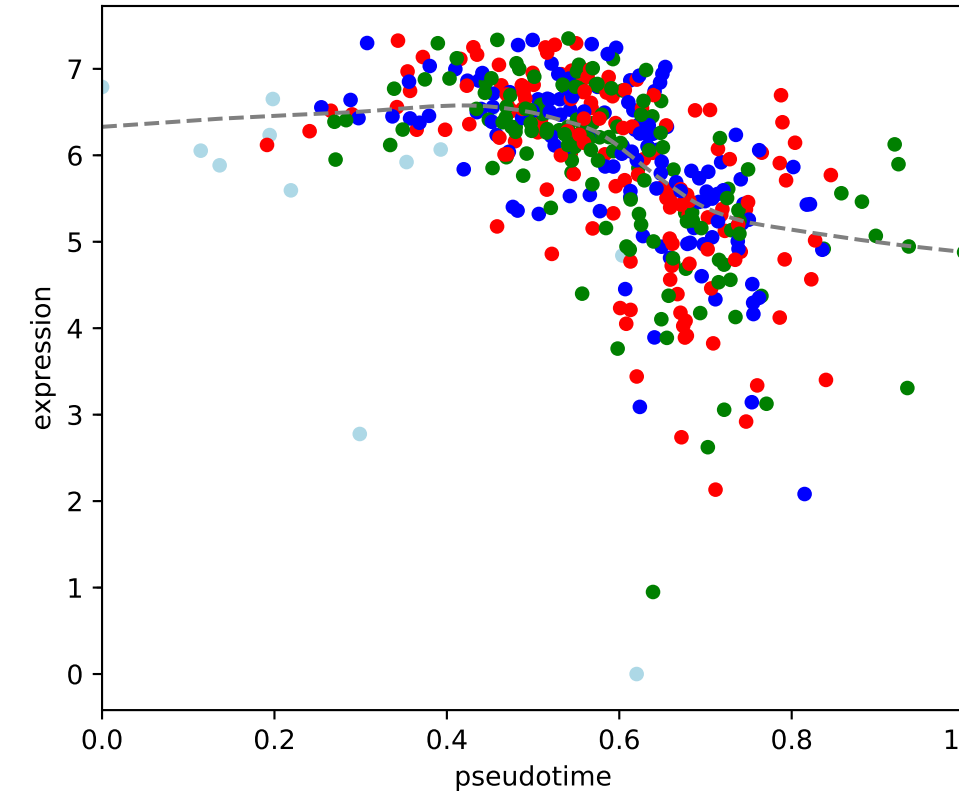
733 1spearman=-0.26



ENST00000247866 NDUFB2 (NADH:ubiquinone oxidoreductase subunit B2)

733 0spearman=-0.63

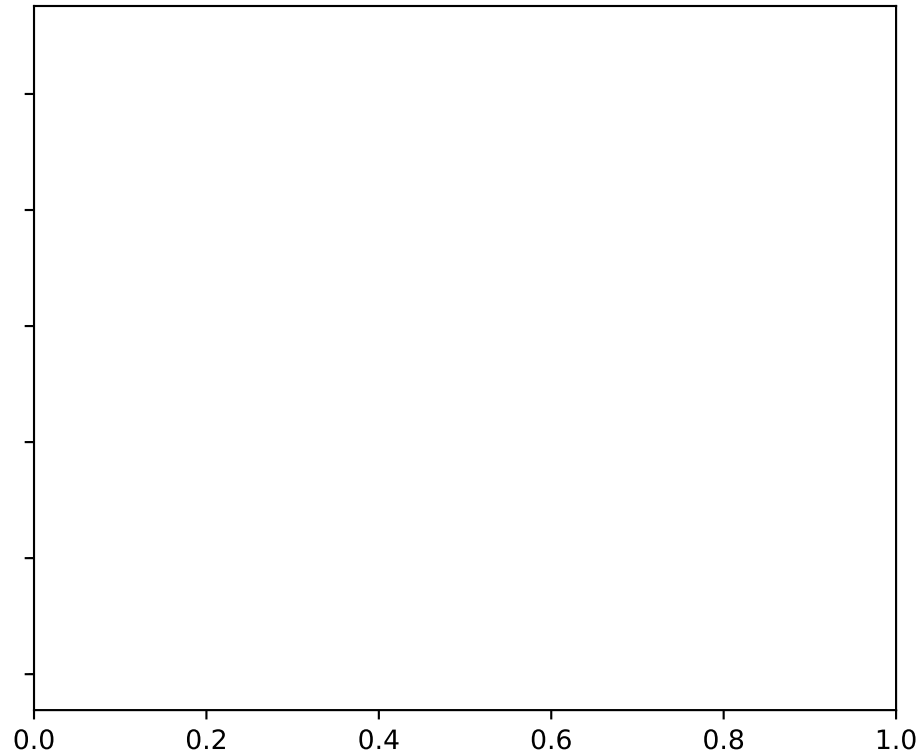
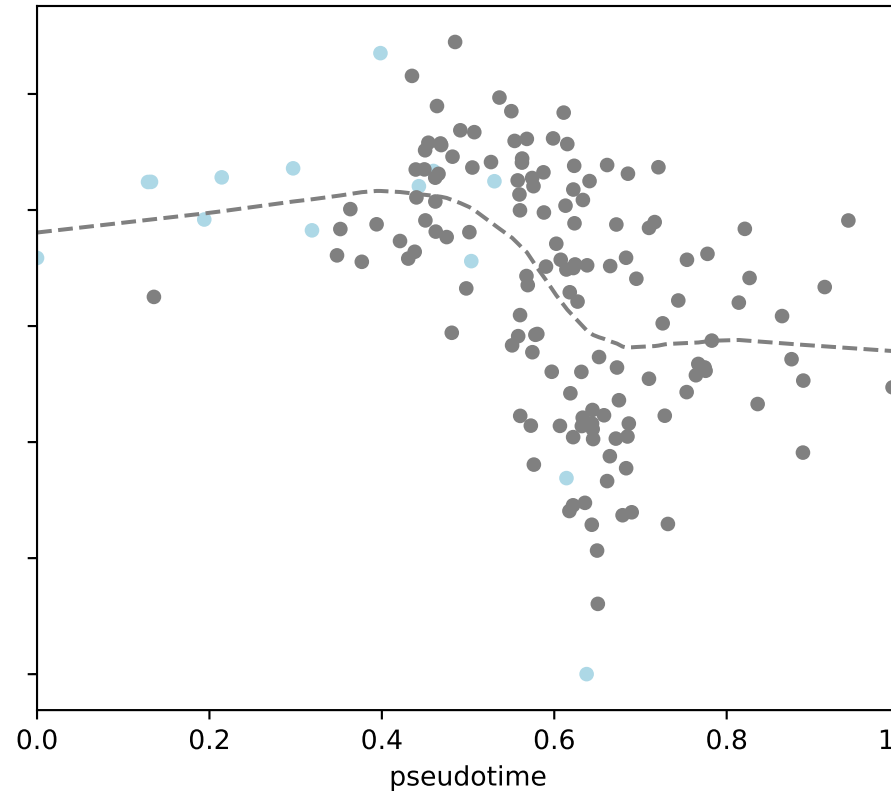
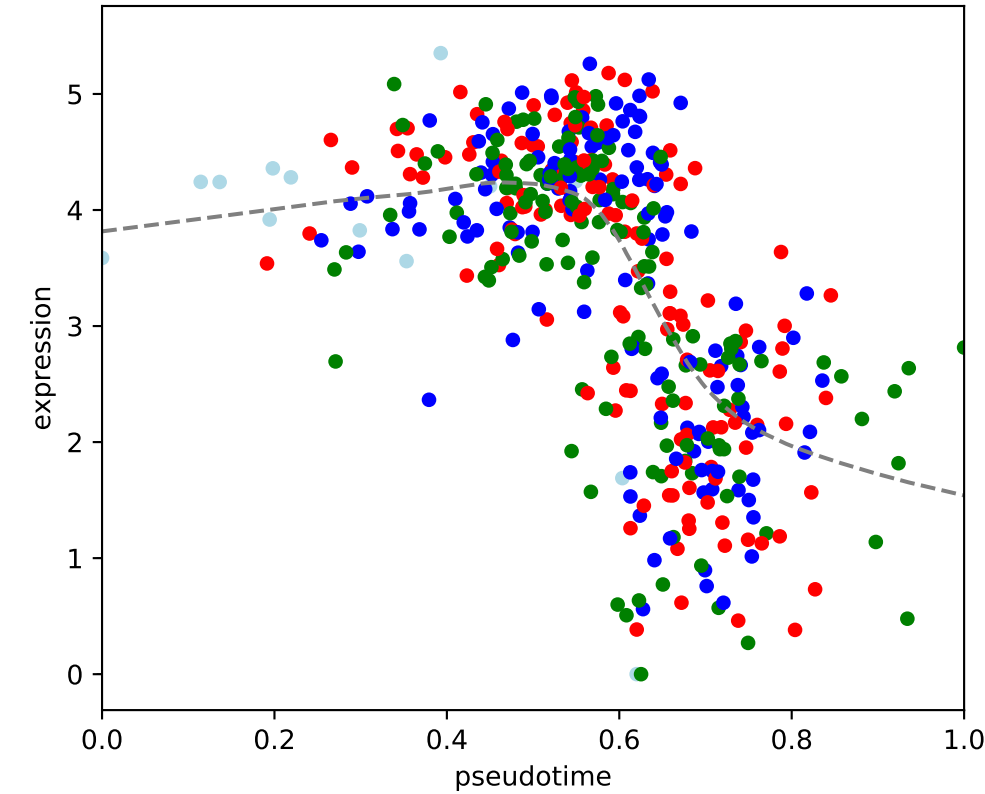
733 1spearman=-0.41



ENST00000366722 GUK1 (guanylate kinase 1)

733 0spearman=-0.63

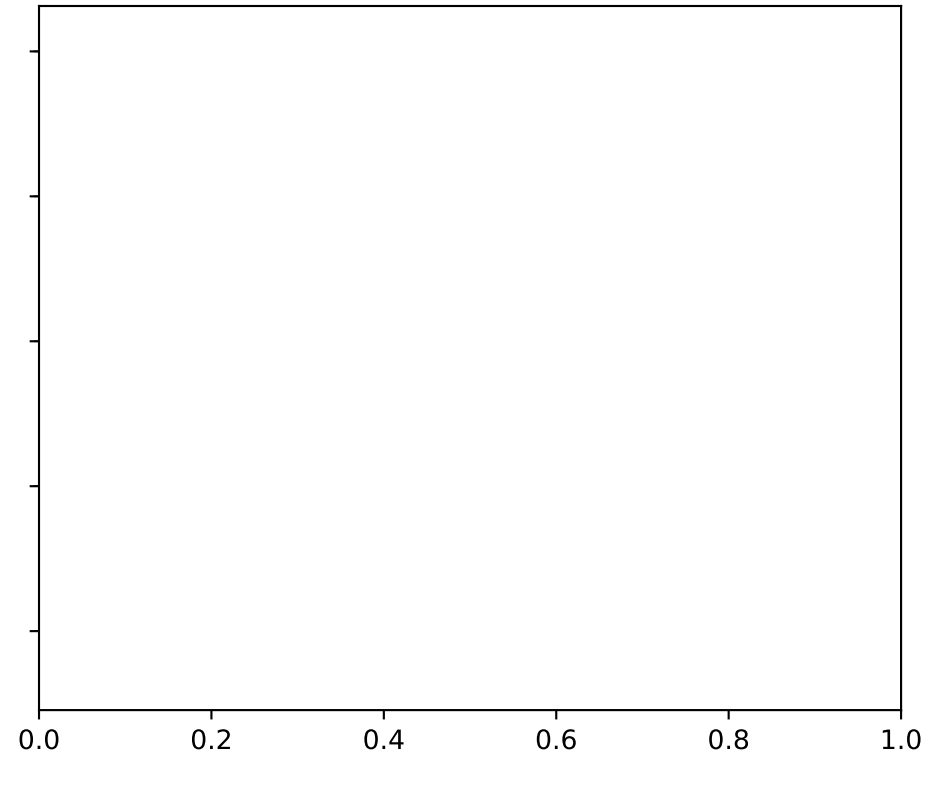
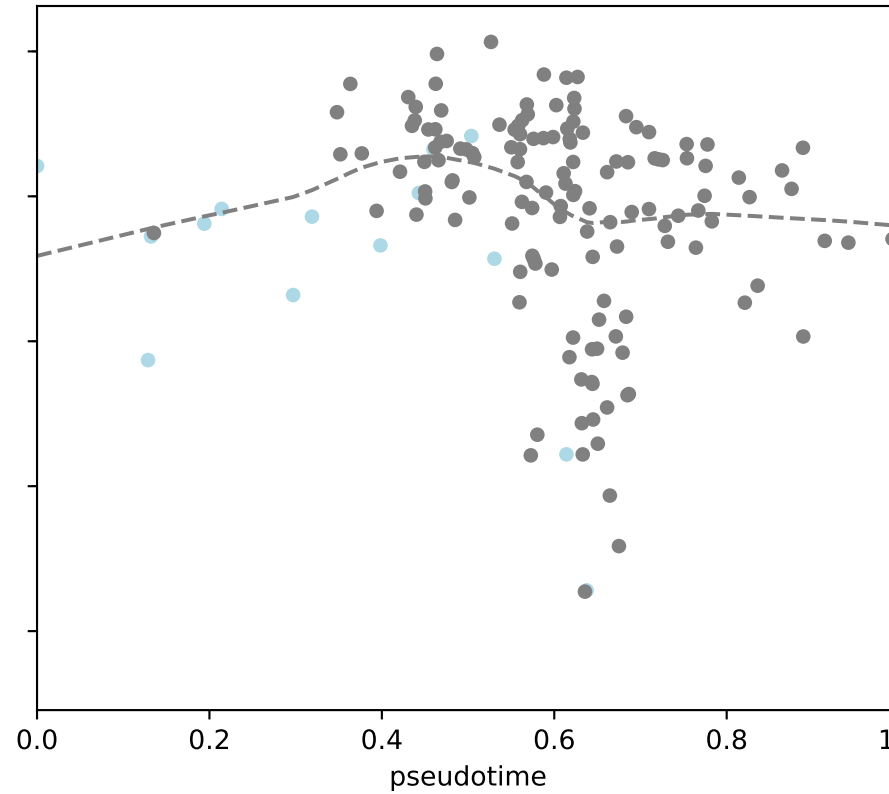
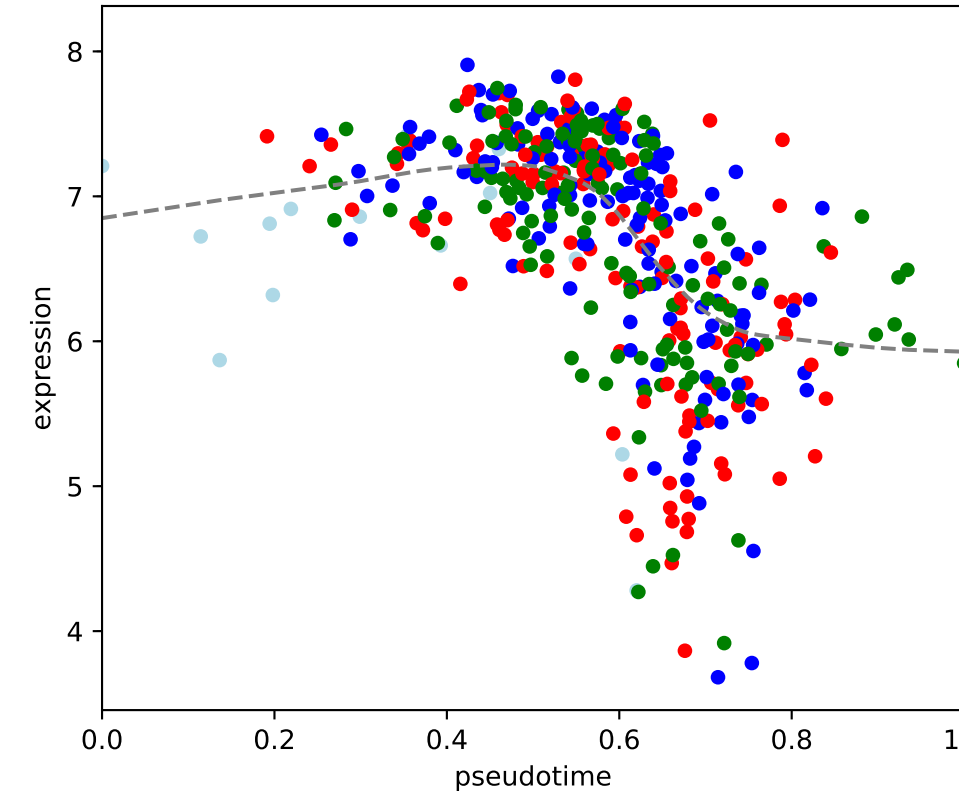
733 1spearman=-0.55



ENST00000247655 COX7C (cytochrome c oxidase subunit 7C)

733 0spearman=-0.63

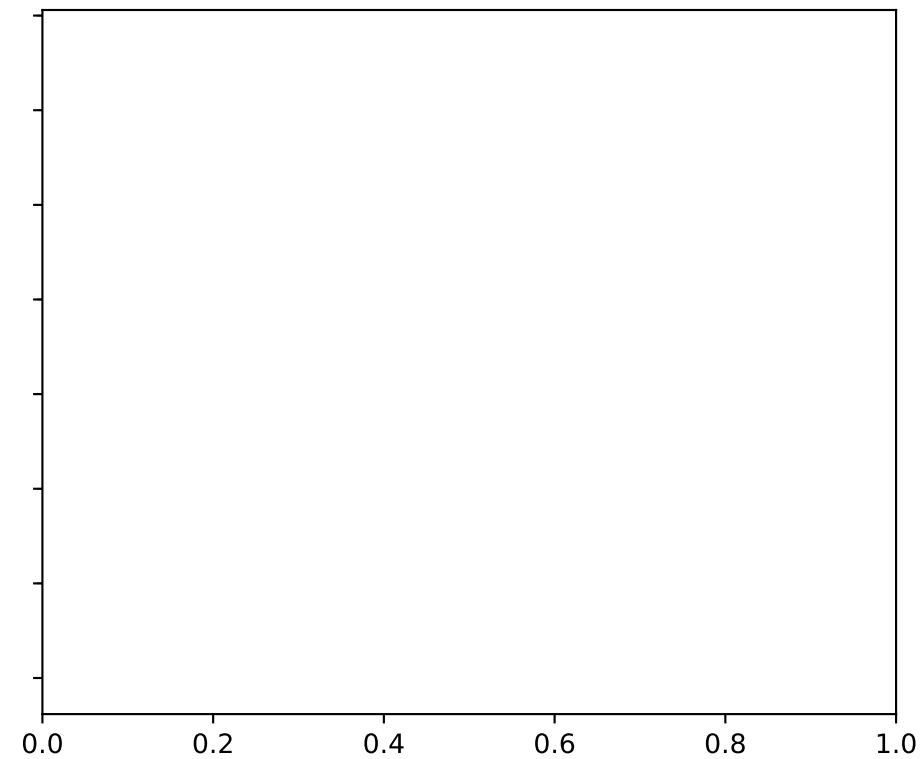
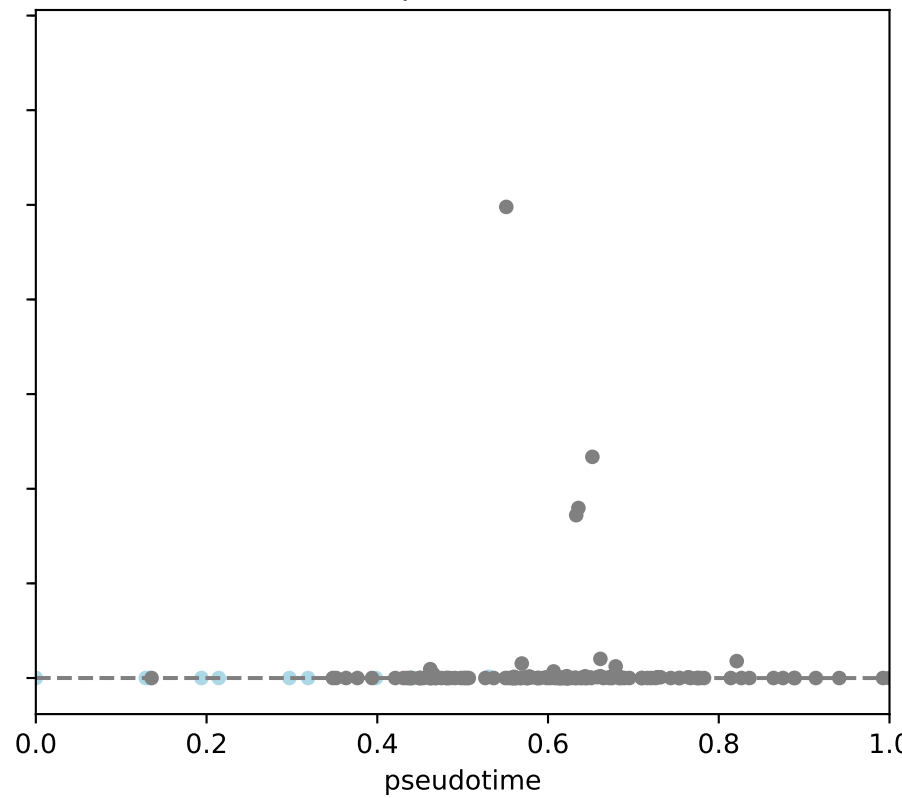
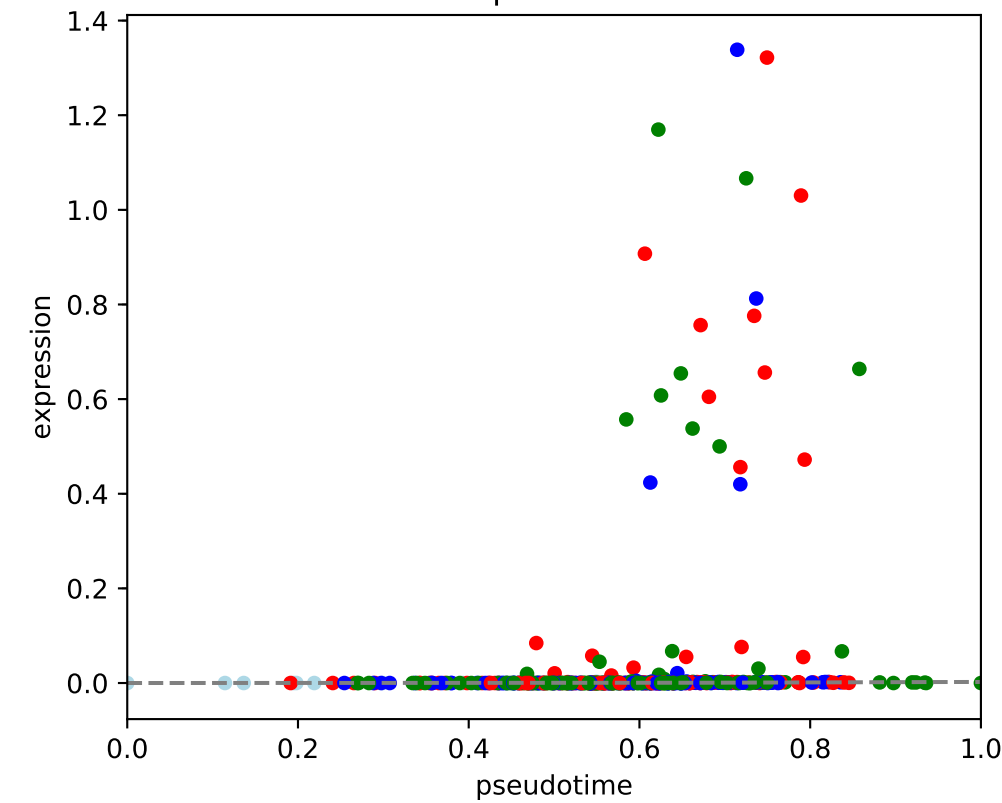
733 1spearman=-0.30



ENST00000598384 MRLN (myoregulin)

733 0spearman=0.63

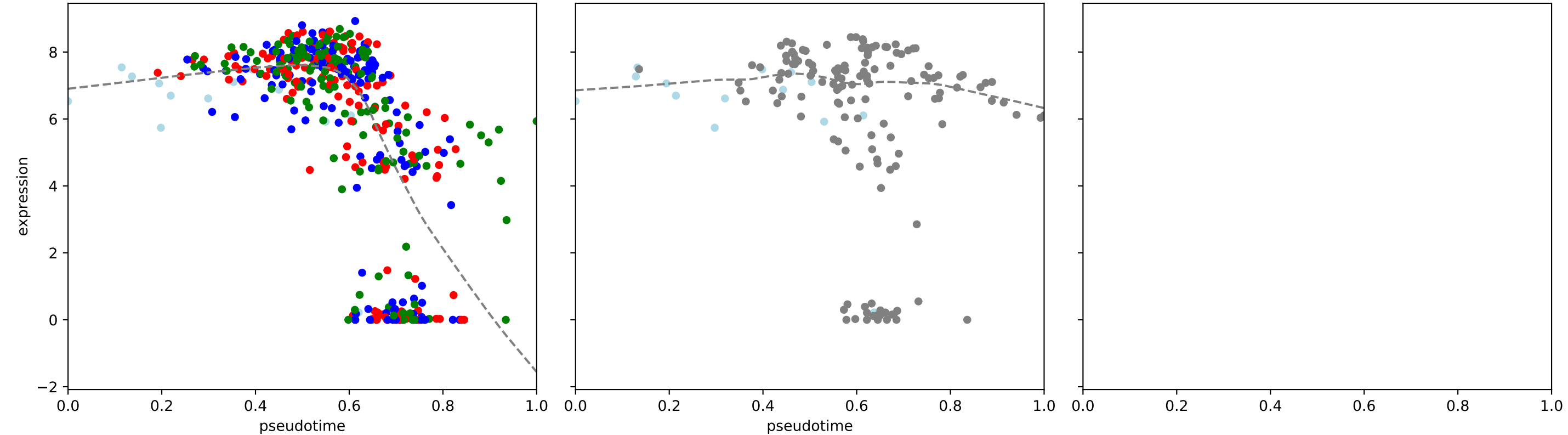
733 1spearman=0.18



ENST00000373232 PPA1 (pyrophosphatase (inorganic) 1)

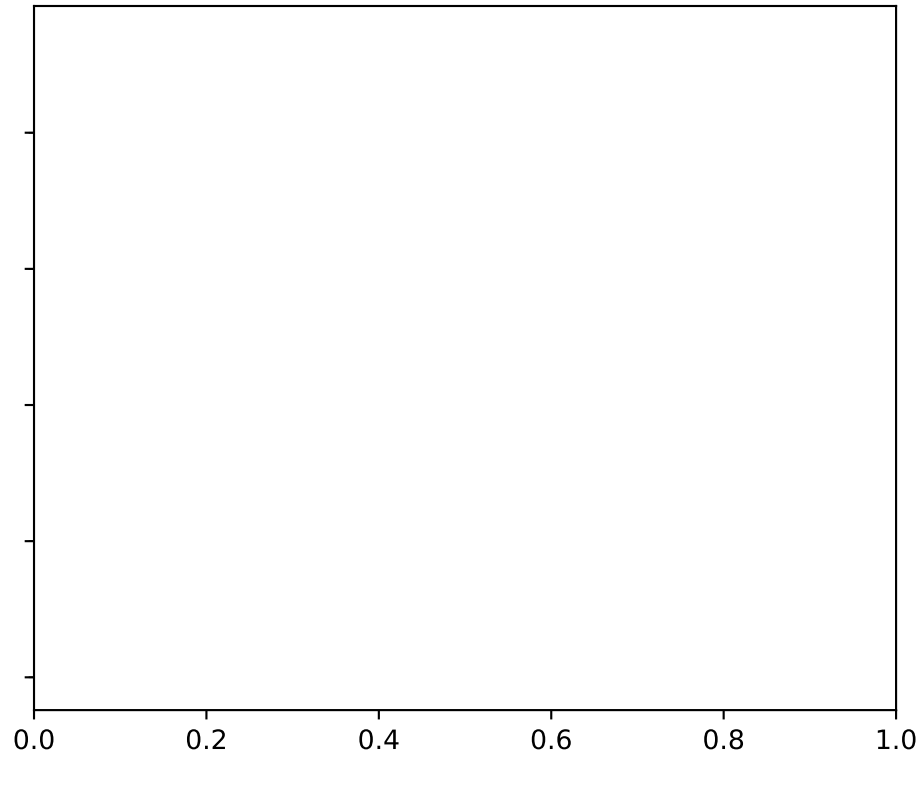
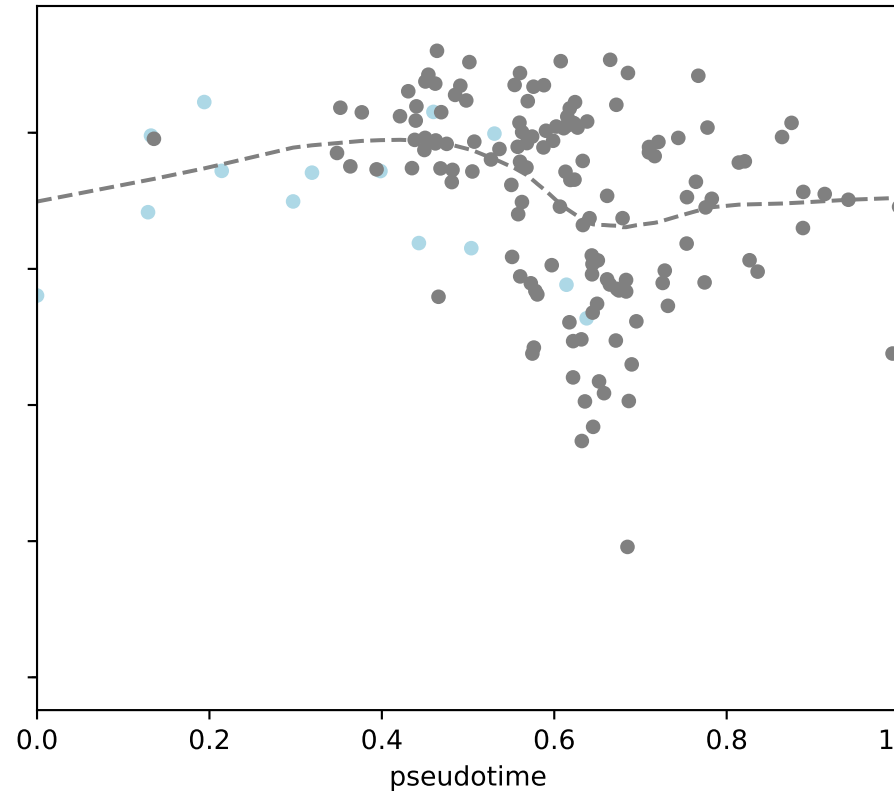
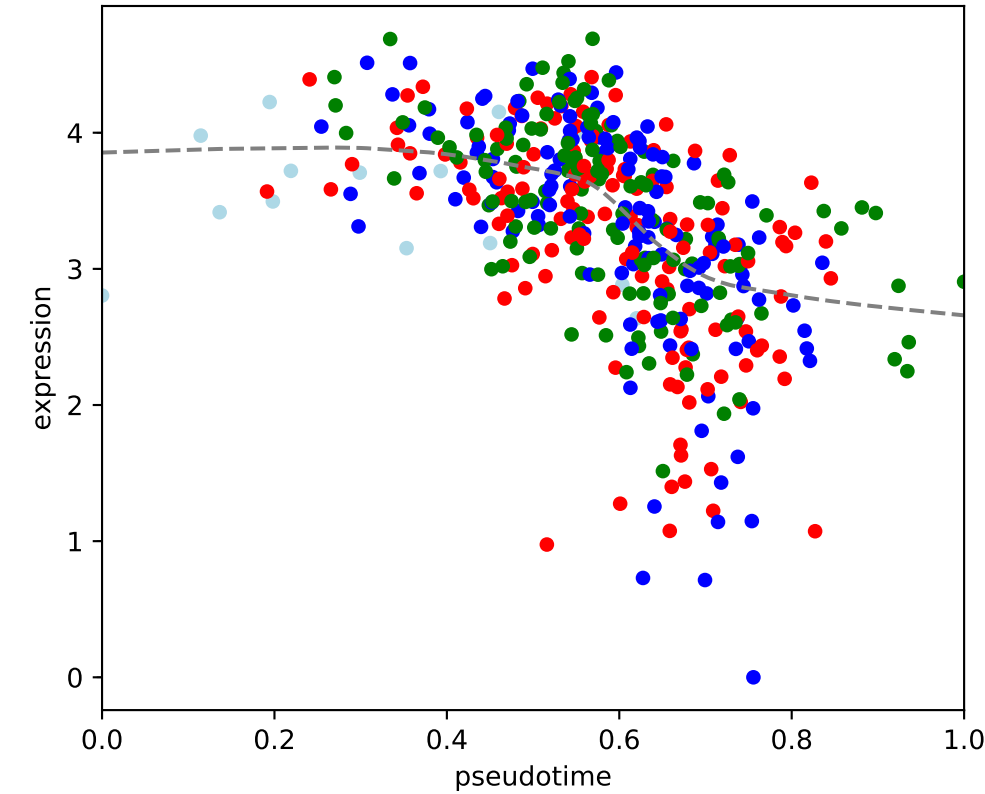
733 0spearman=-0.62

733 1spearman=-0.26



733 0spearman=-0.62

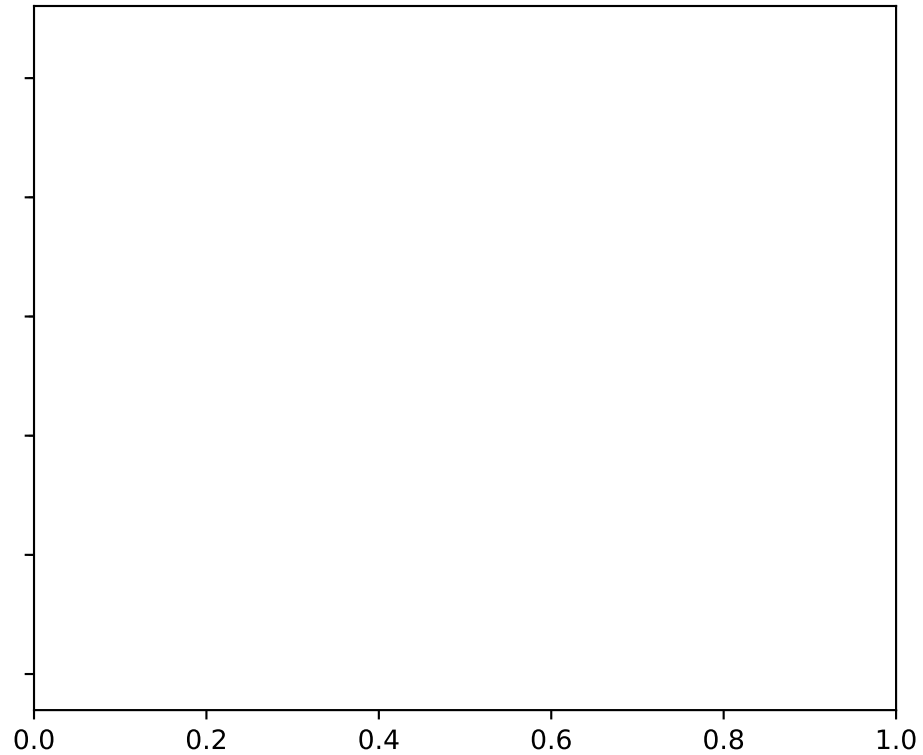
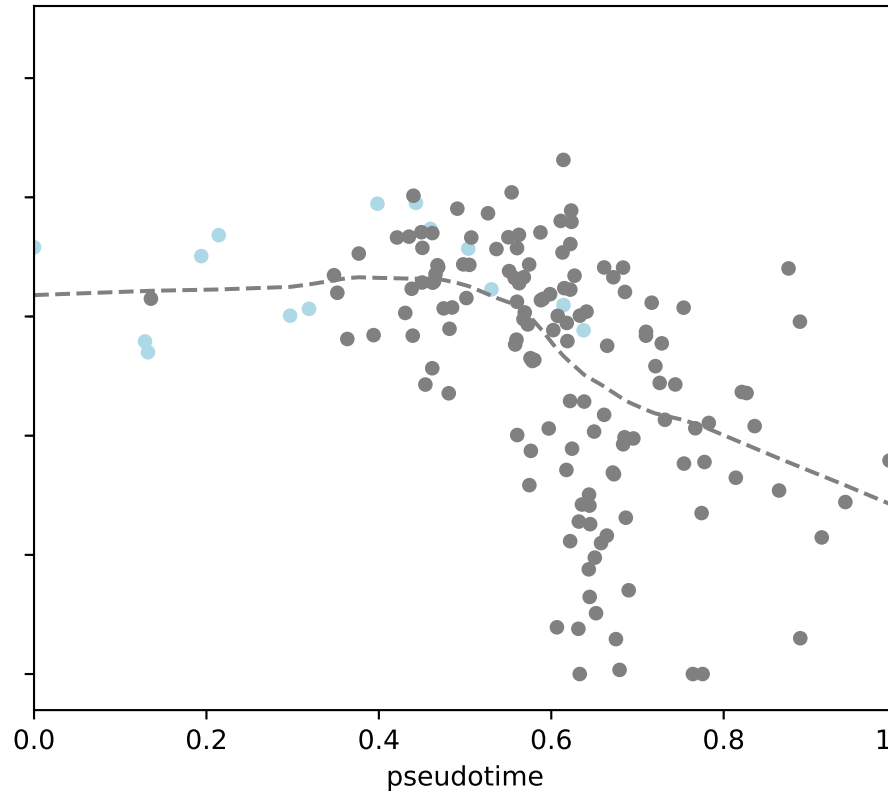
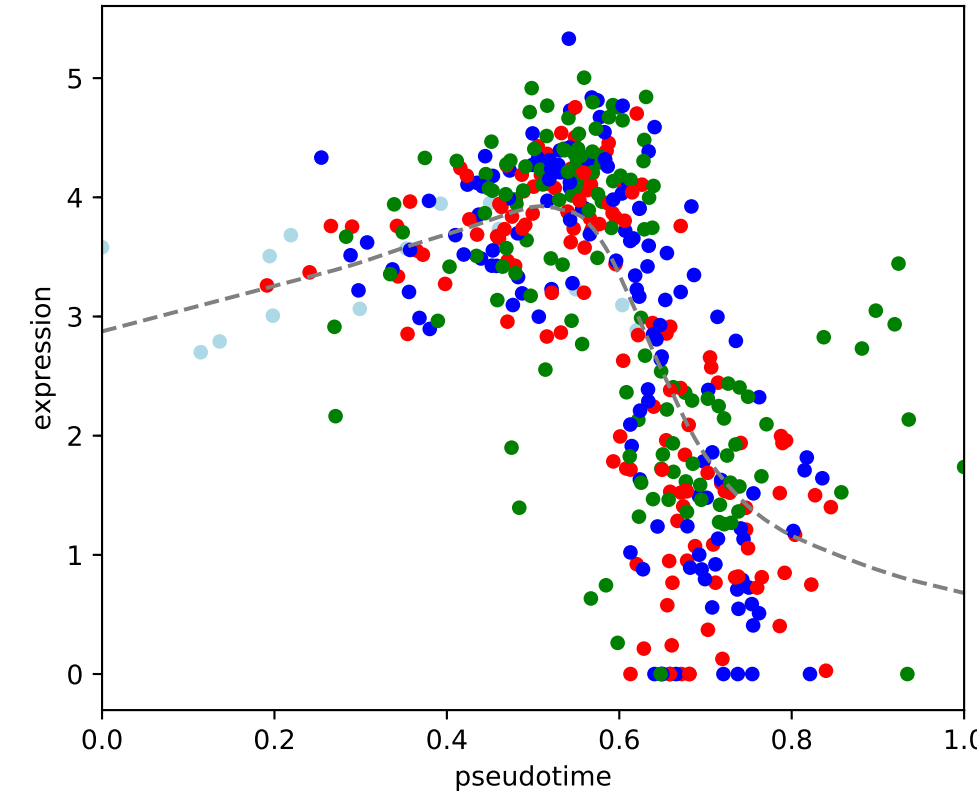
733 1spearman=-0.36



ENST00000552893 AC011603.2

733 0spearman=-0.62

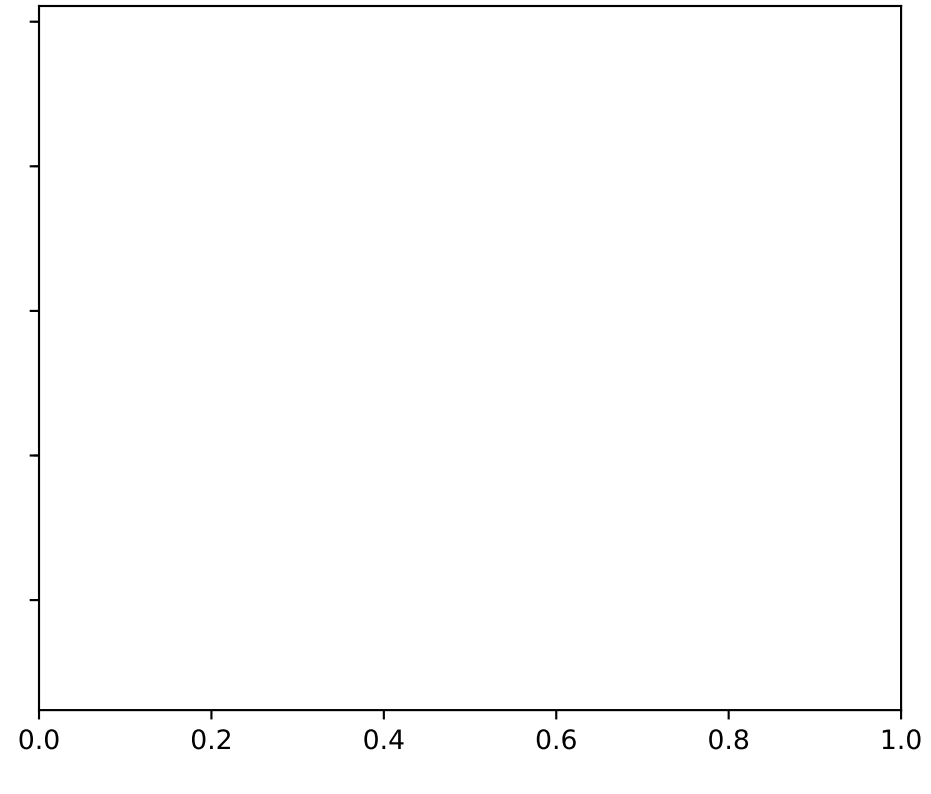
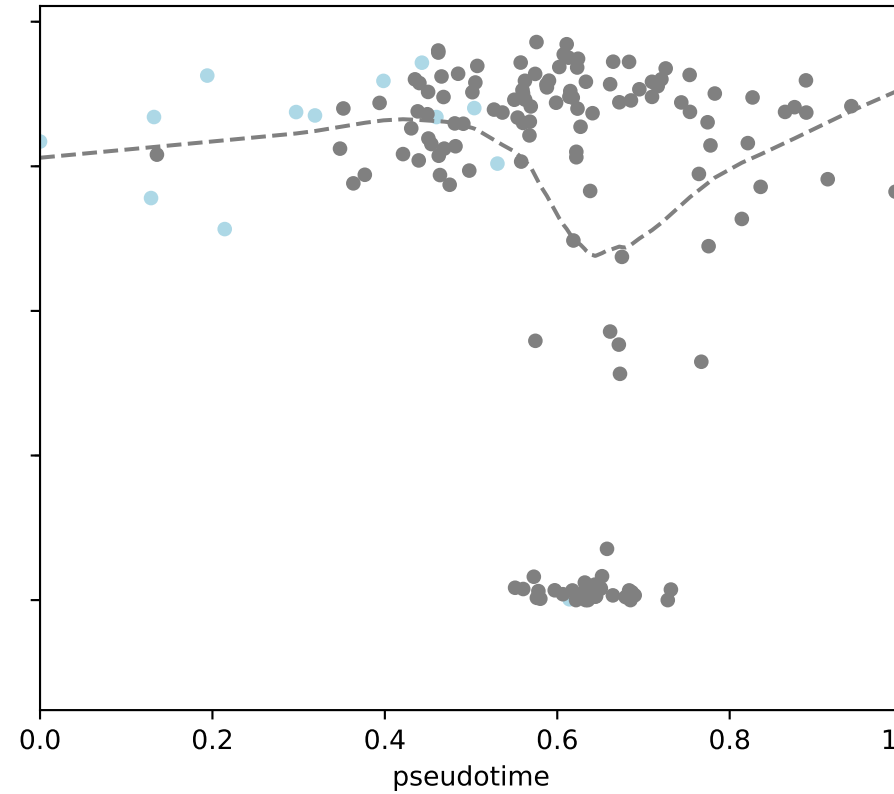
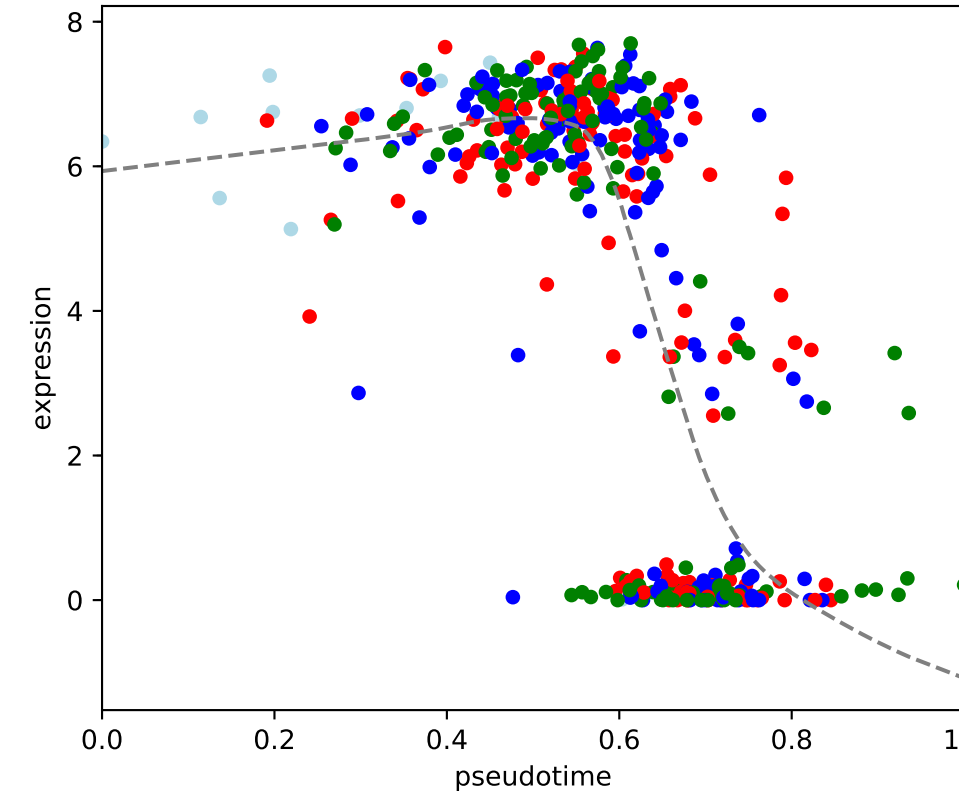
733 1spearman=-0.60



ENST00000351050 GNAT2 (G protein subunit alpha transducin 2)

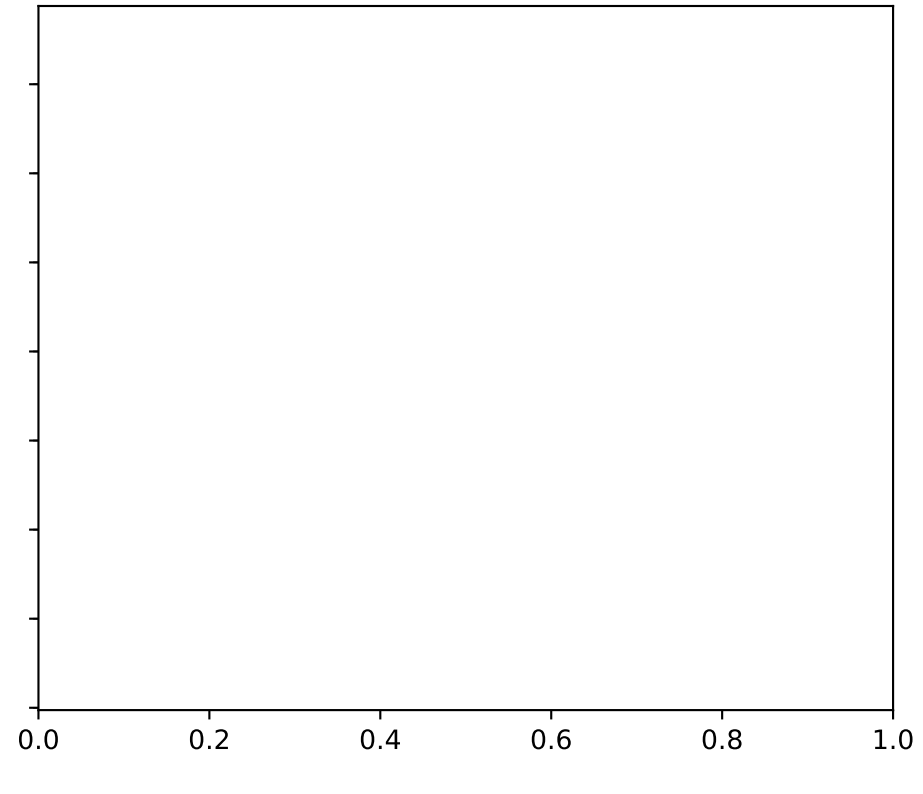
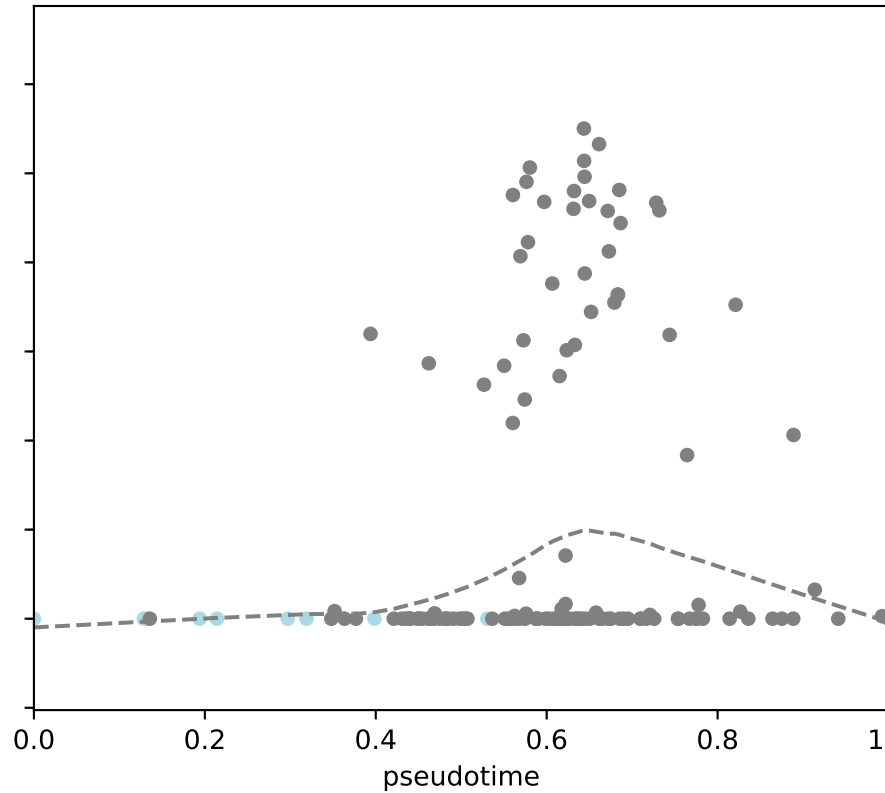
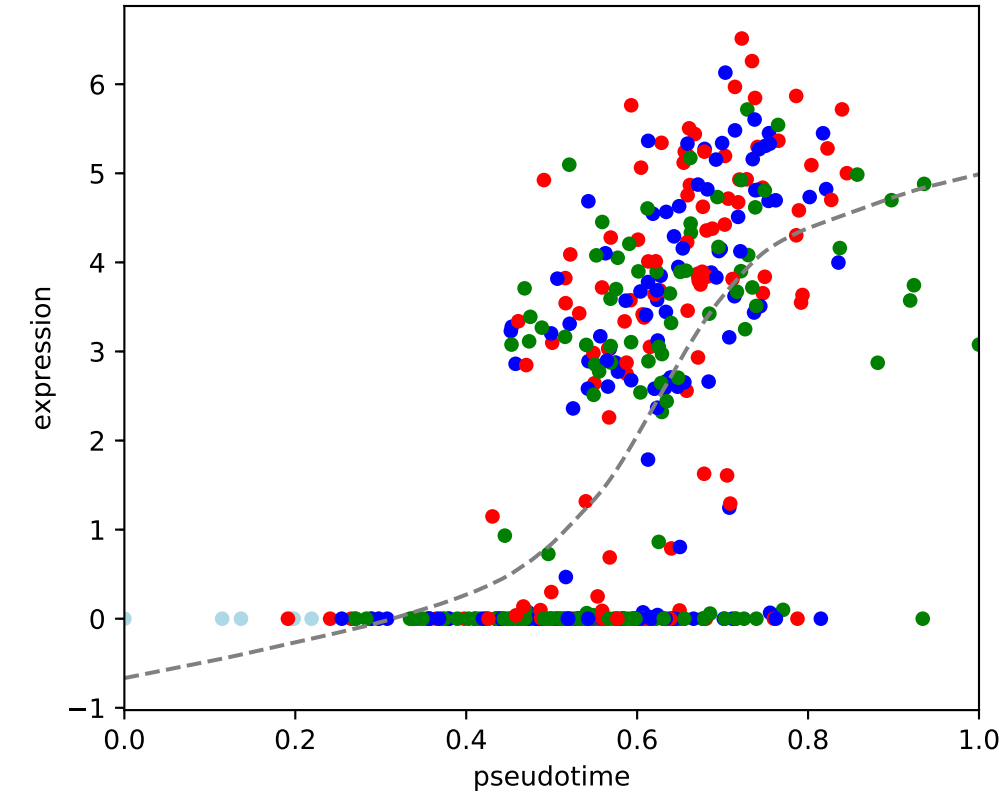
733 0spearman=-0.62

733 1spearman=-0.17



733 0spearman=0.62

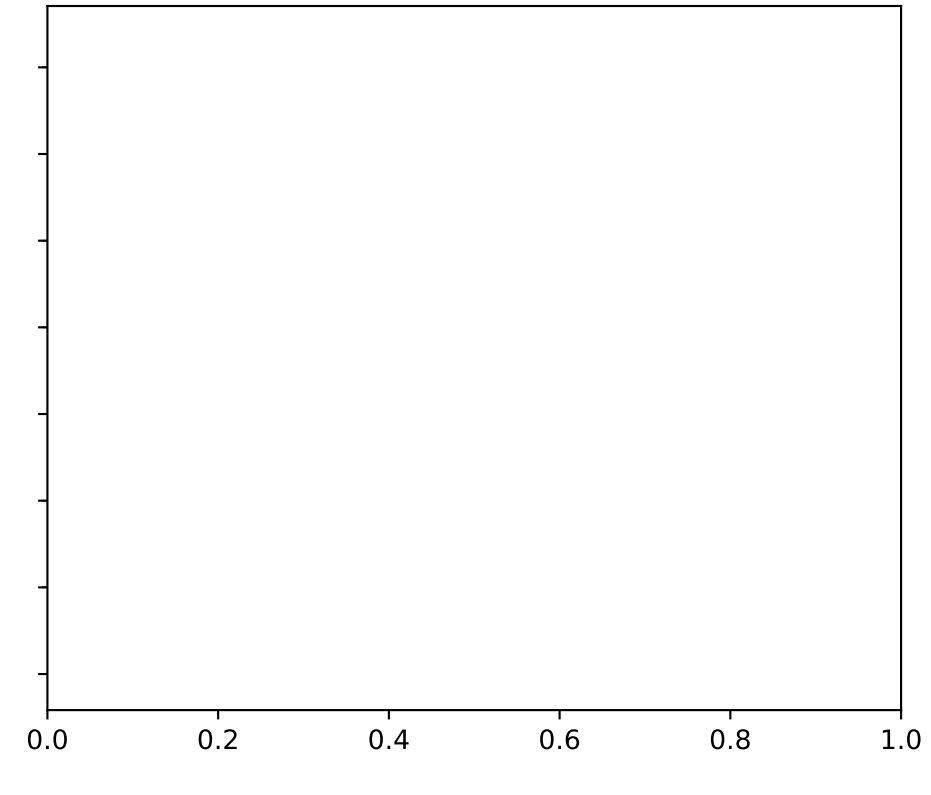
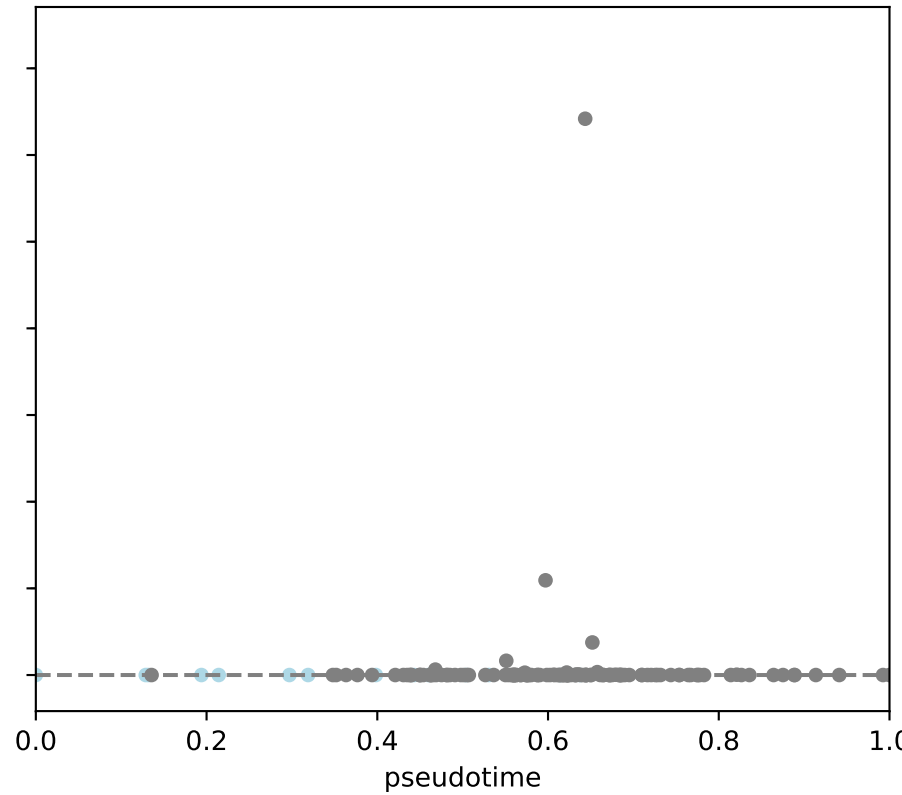
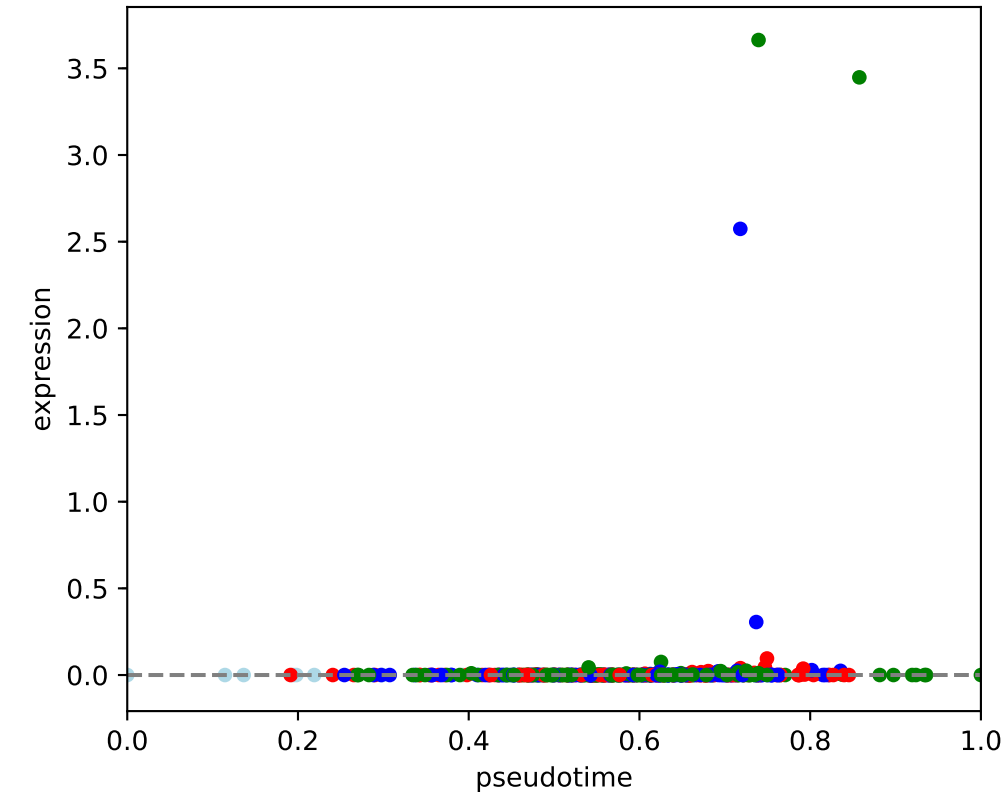
733 1spearman=0.26



ENST00000612853 MRLN (myoregulin)

733 0spearman=0.61

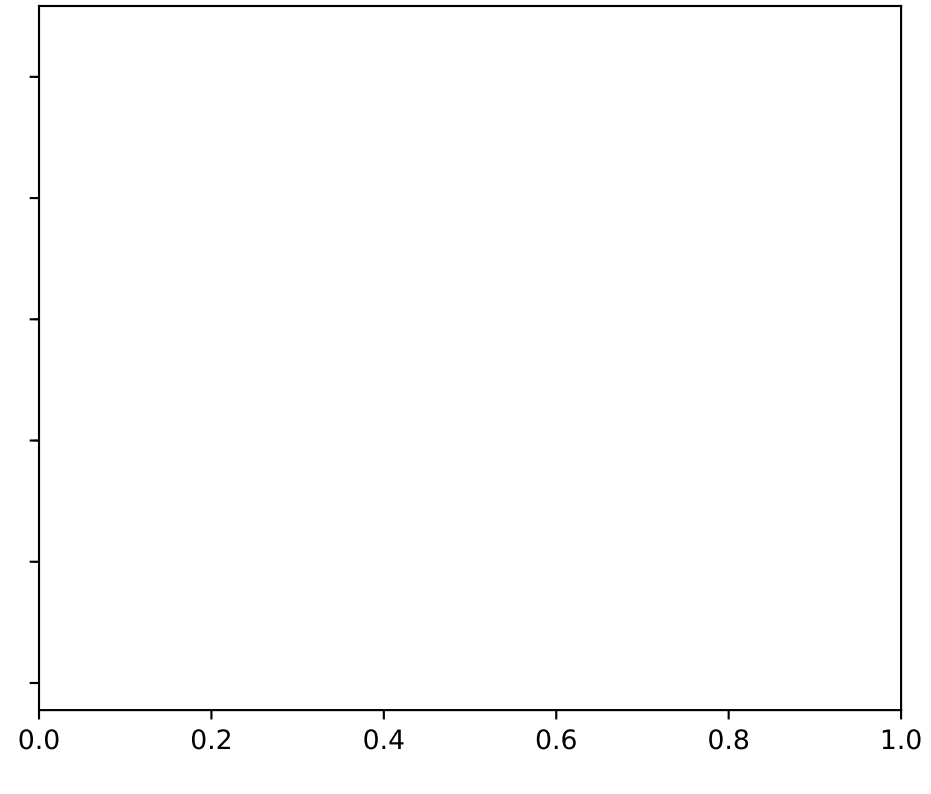
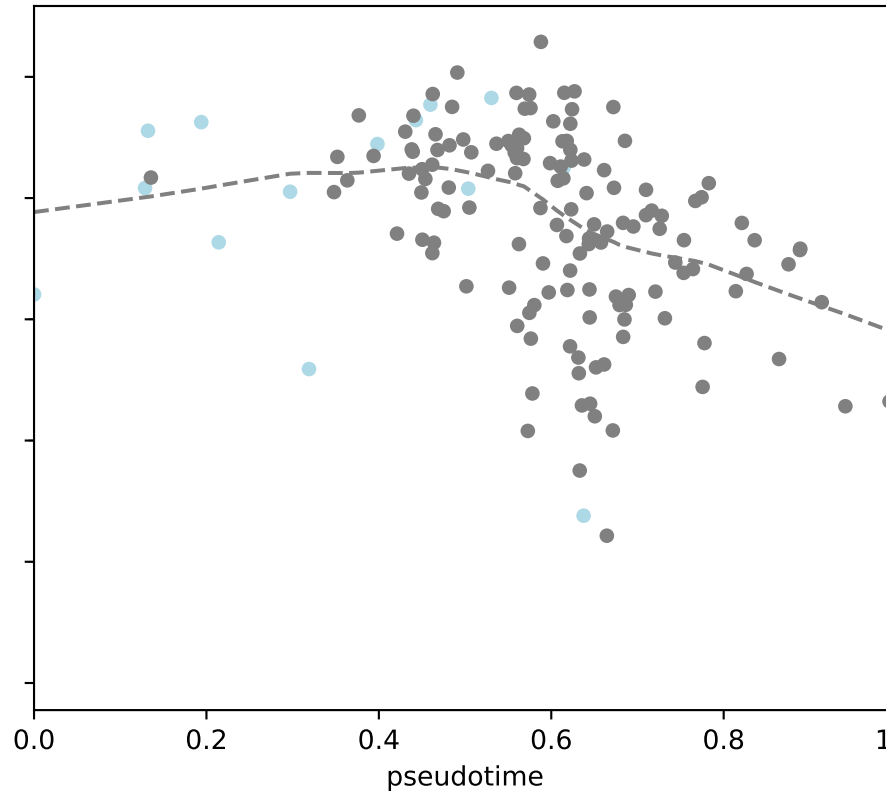
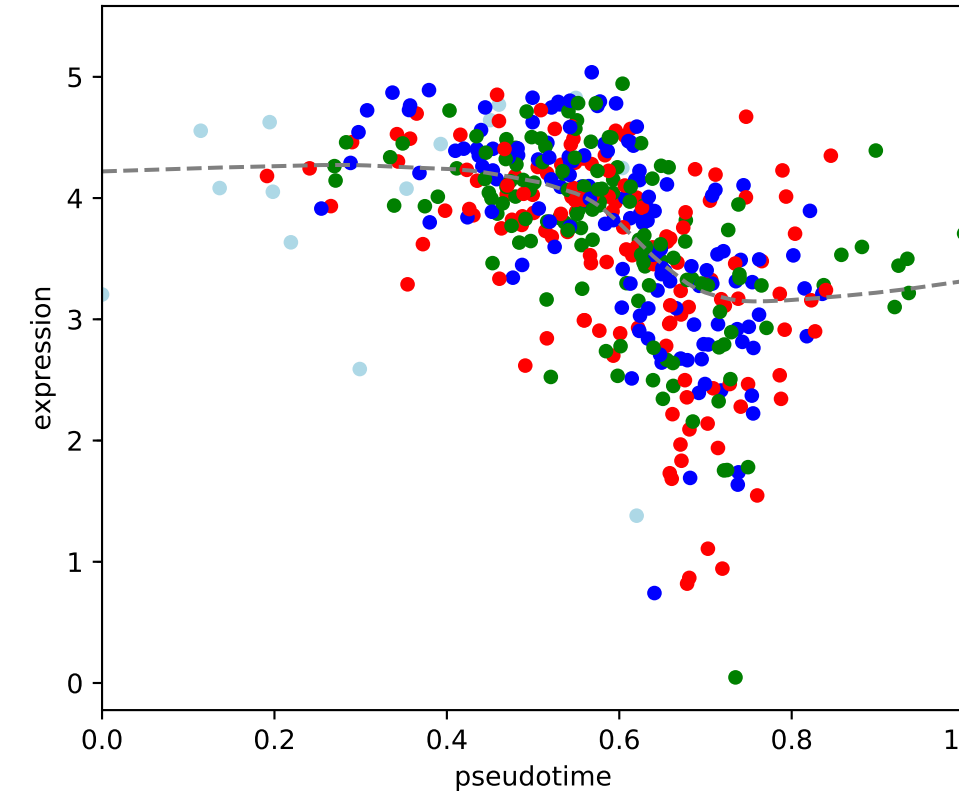
733 1spearman=0.16



ENST00000462885 RPL18AP3 (ribosomal protein L18a pseudogene 3)

733 0spearman=-0.61

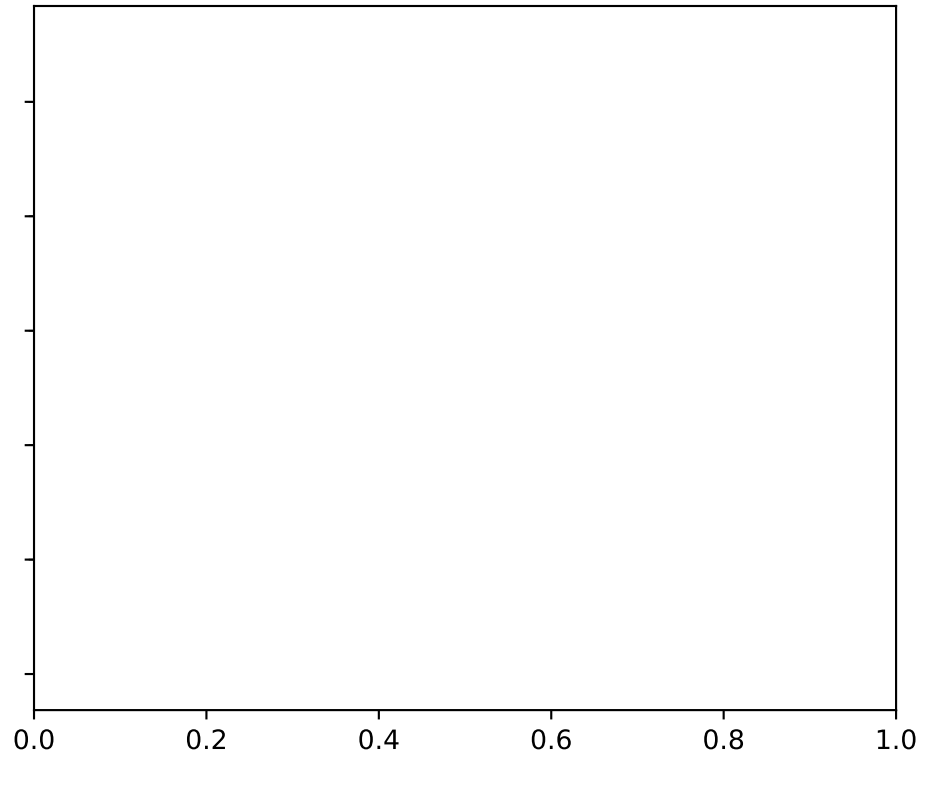
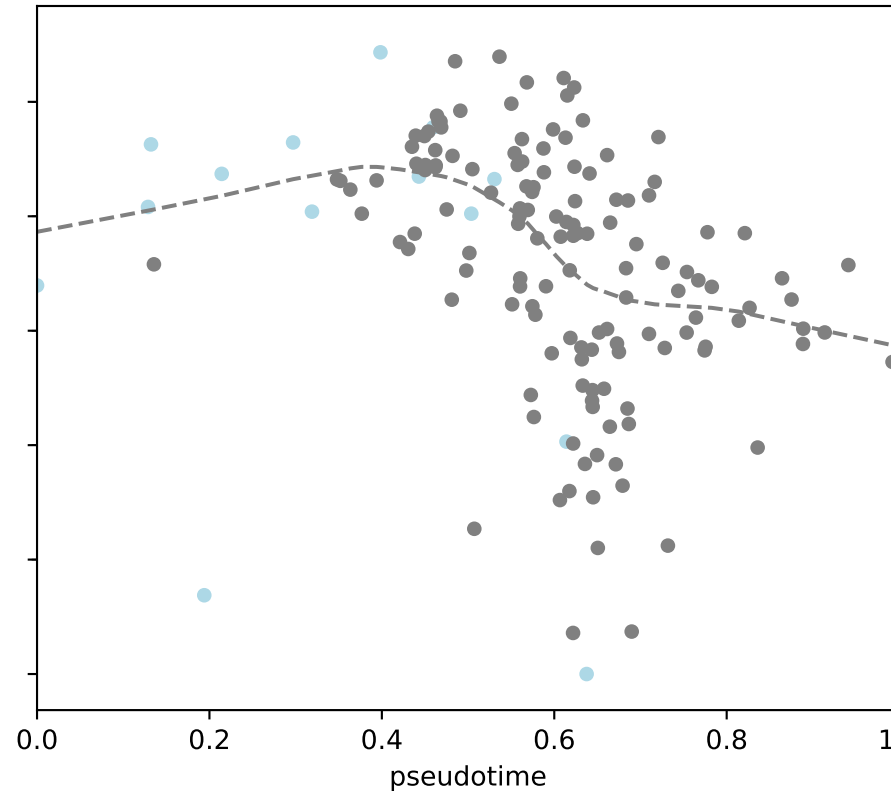
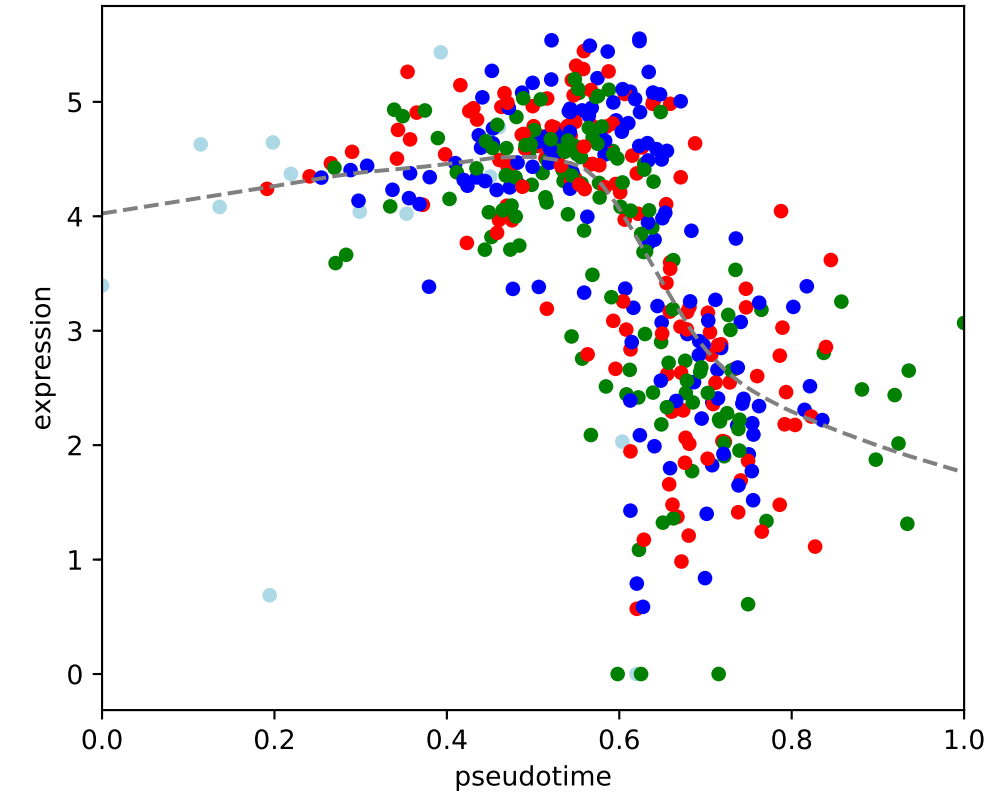
733 1spearman=-0.48



ENST00000477206 GUK1 (guanylate kinase 1)

733 0spearman=-0.61

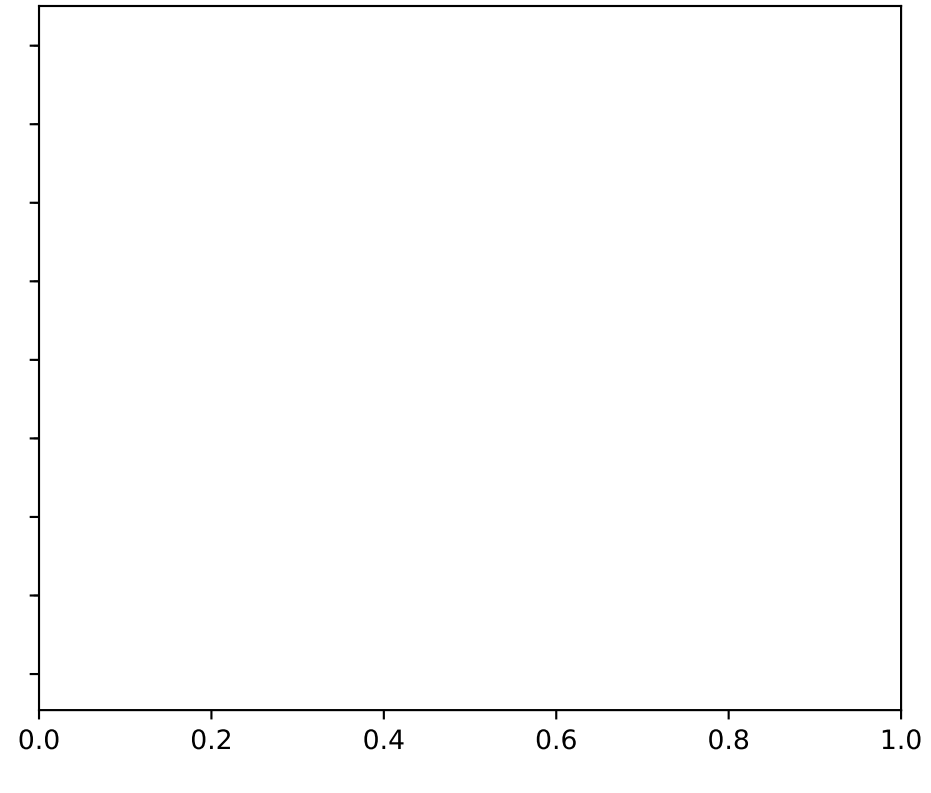
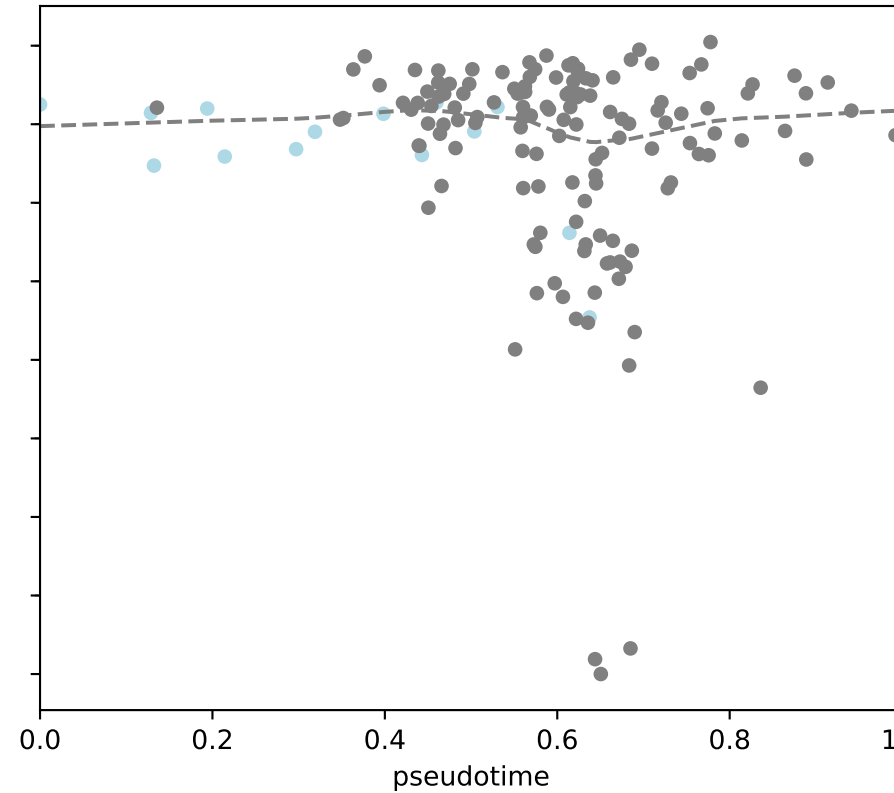
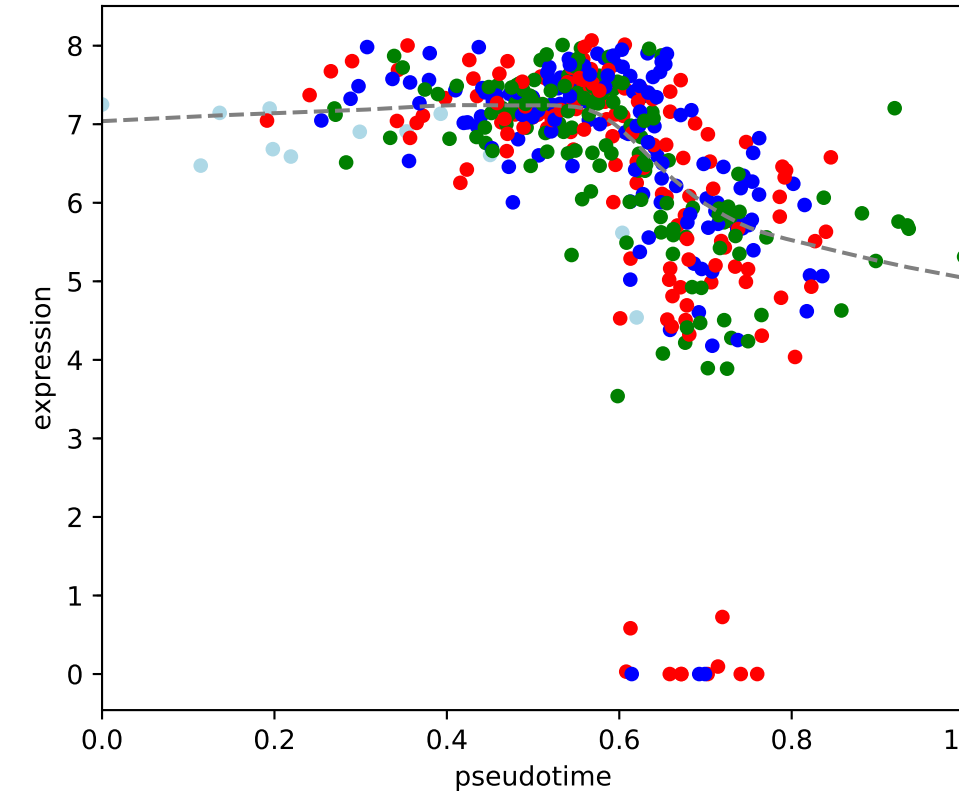
733 1spearman=-0.52



ENST00000318388 NDUFV2 (NADH:ubiquinone oxidoreductase core subunit V2)

733 0spearman=-0.61

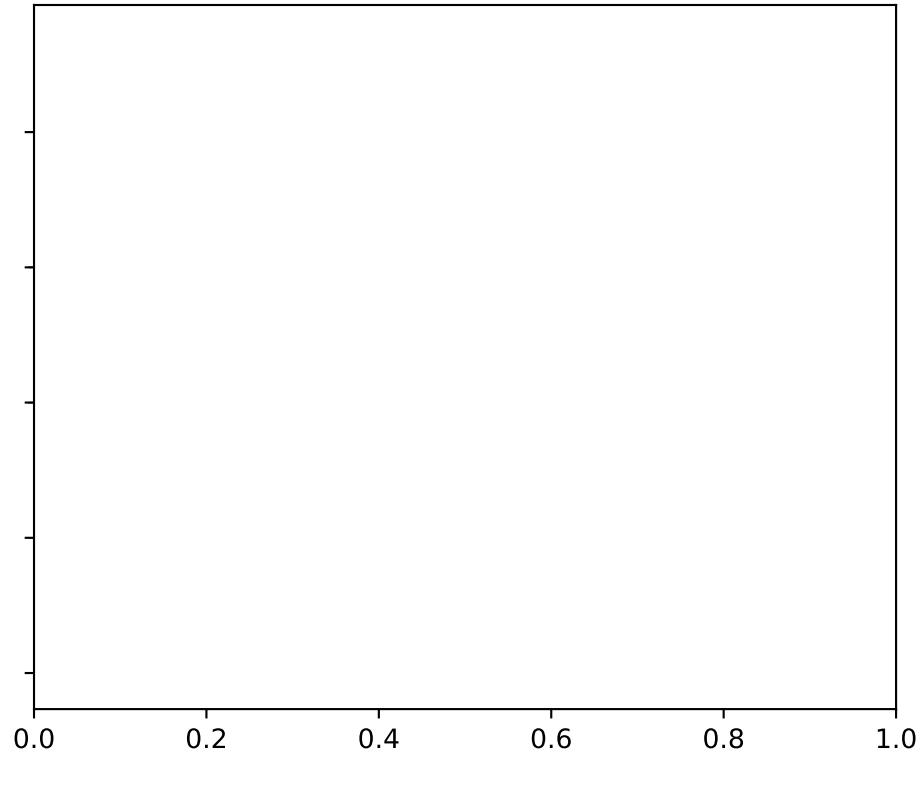
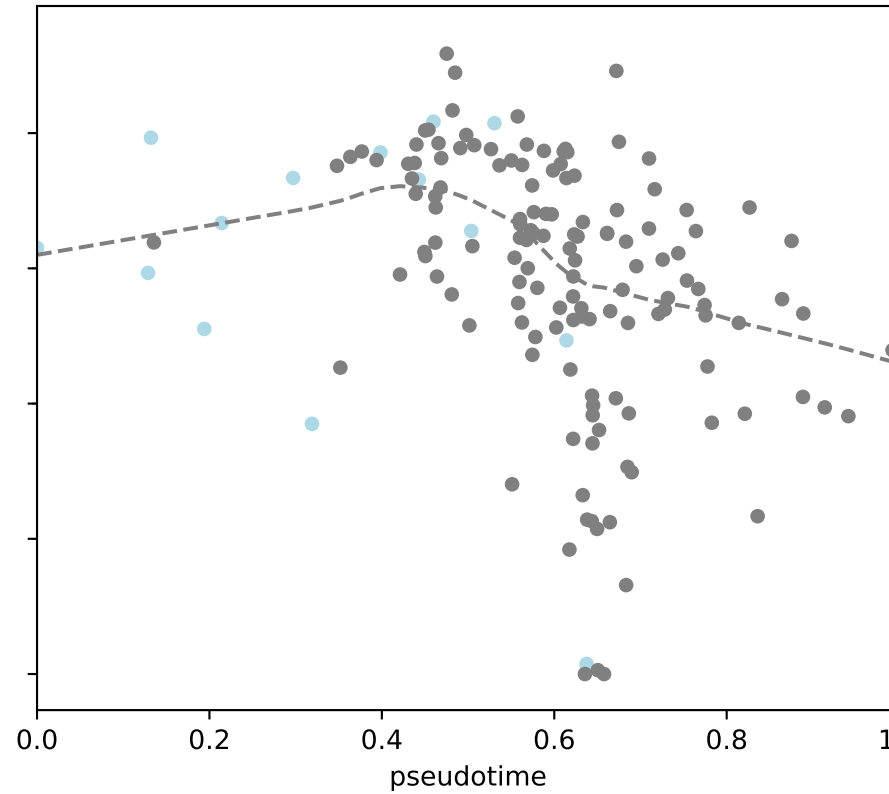
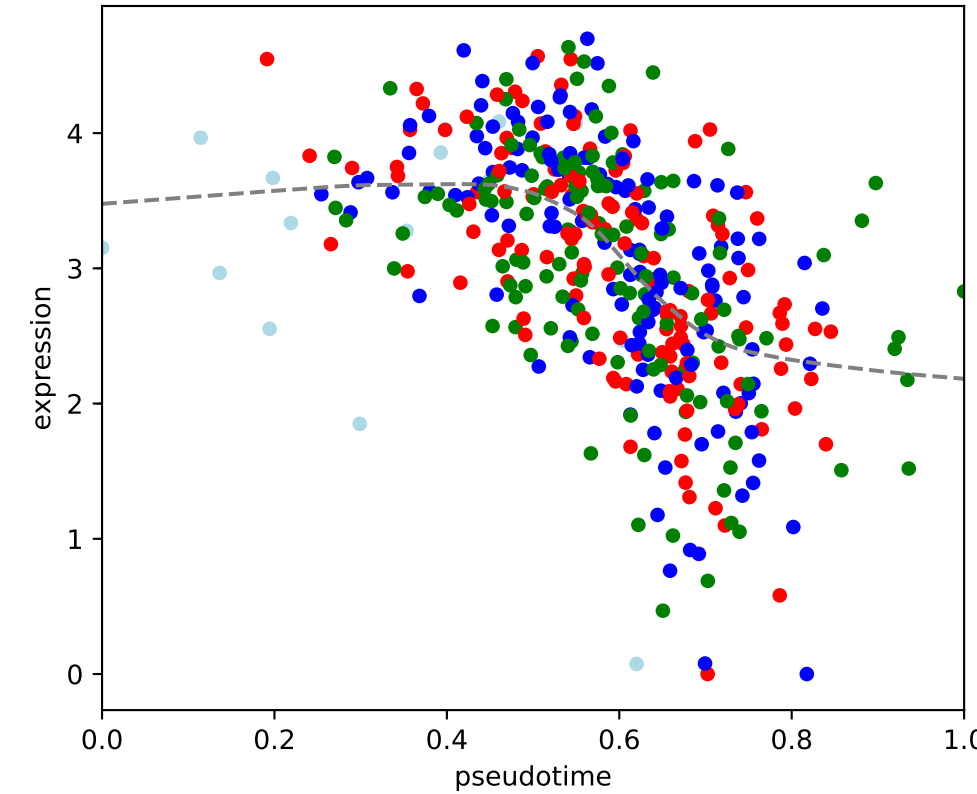
733 1spearman=-0.16



ENST00000518938 AC026403.1

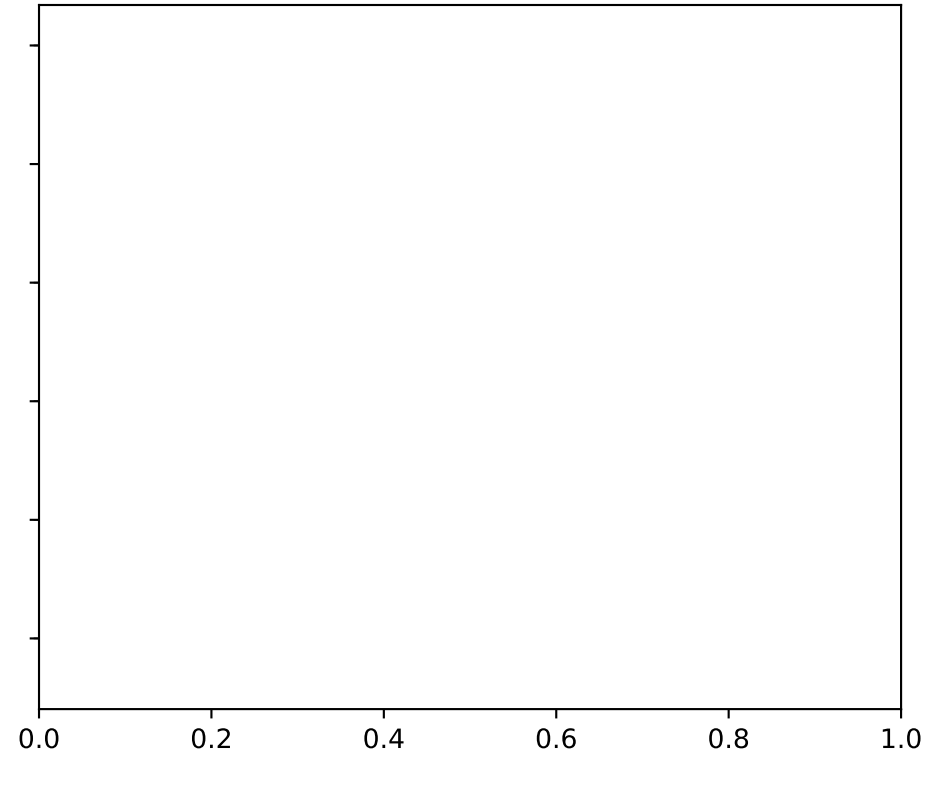
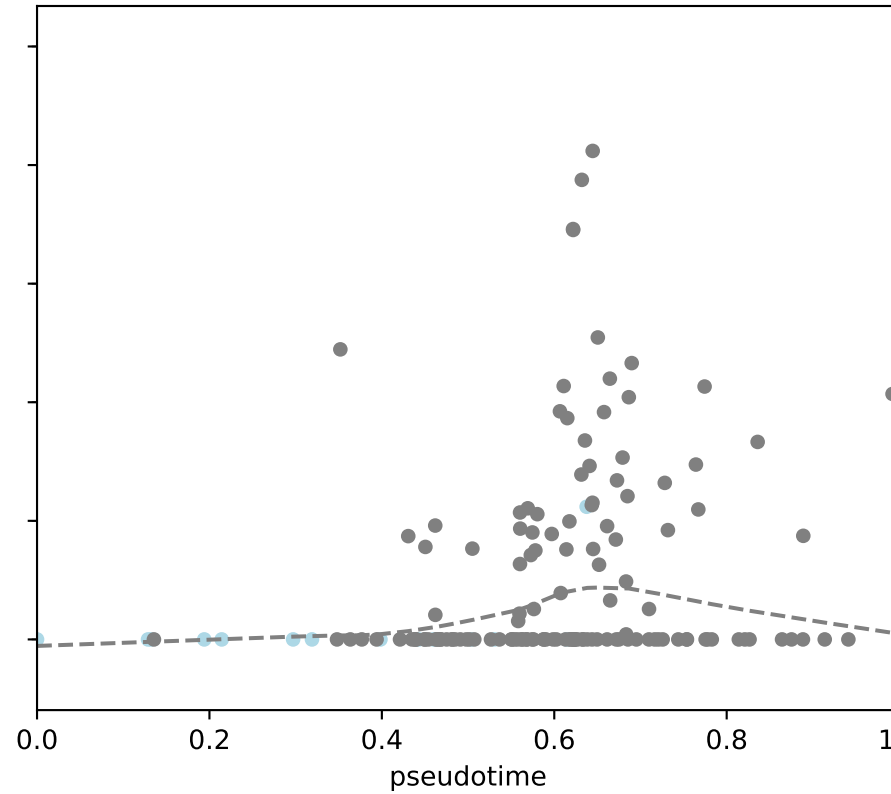
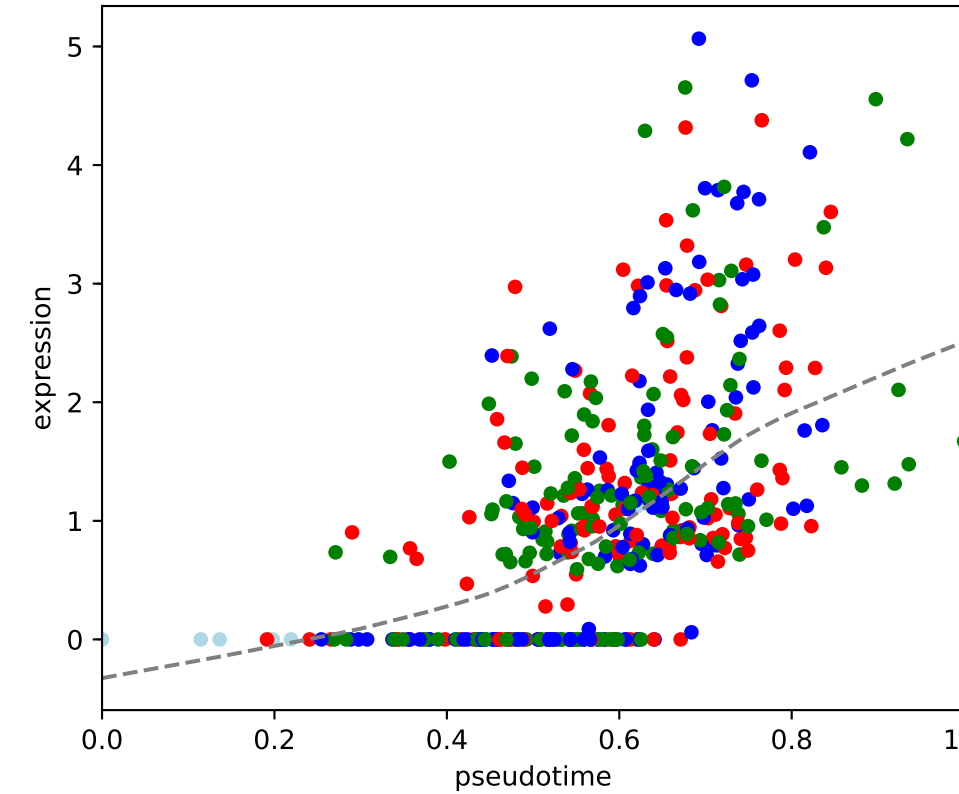
733 0spearman=-0.61

733 1spearman=-0.49



733 0spearman=0.61

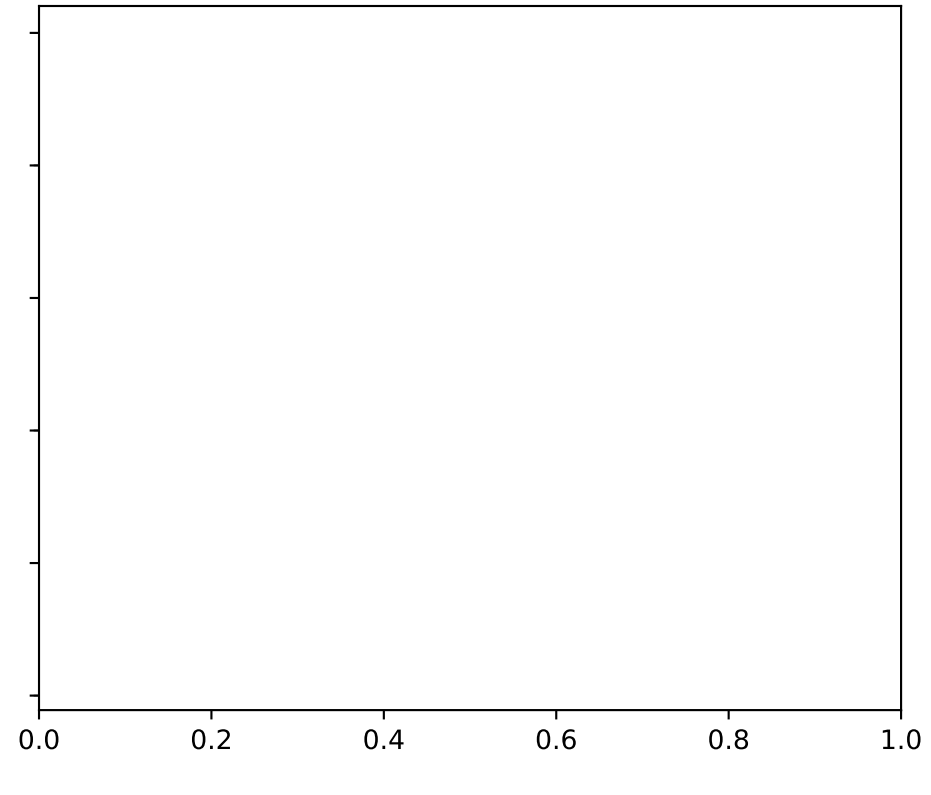
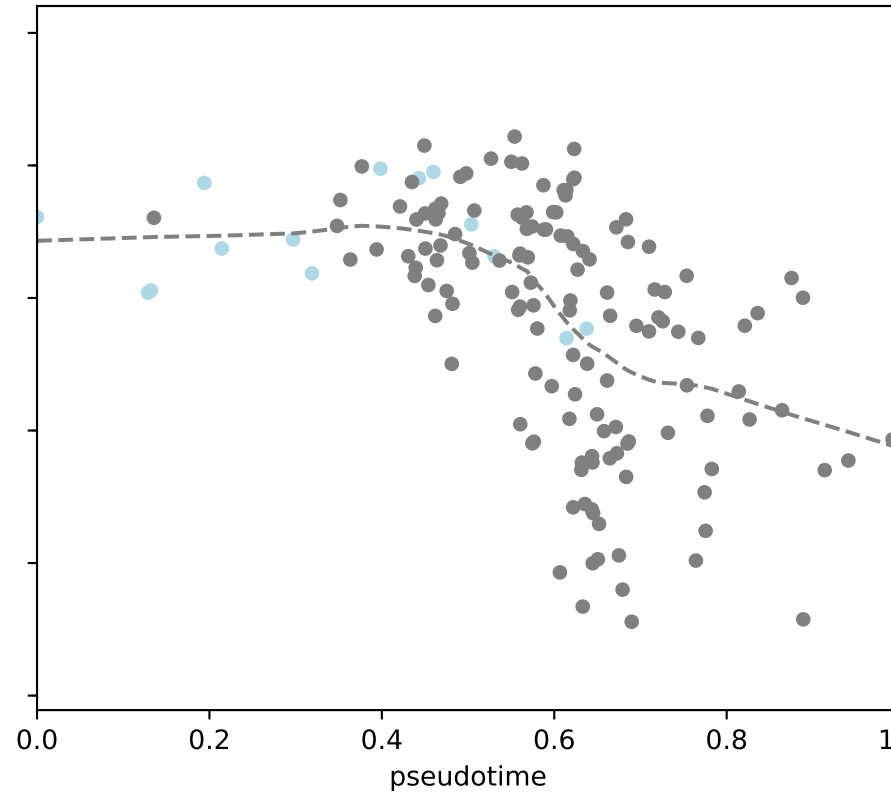
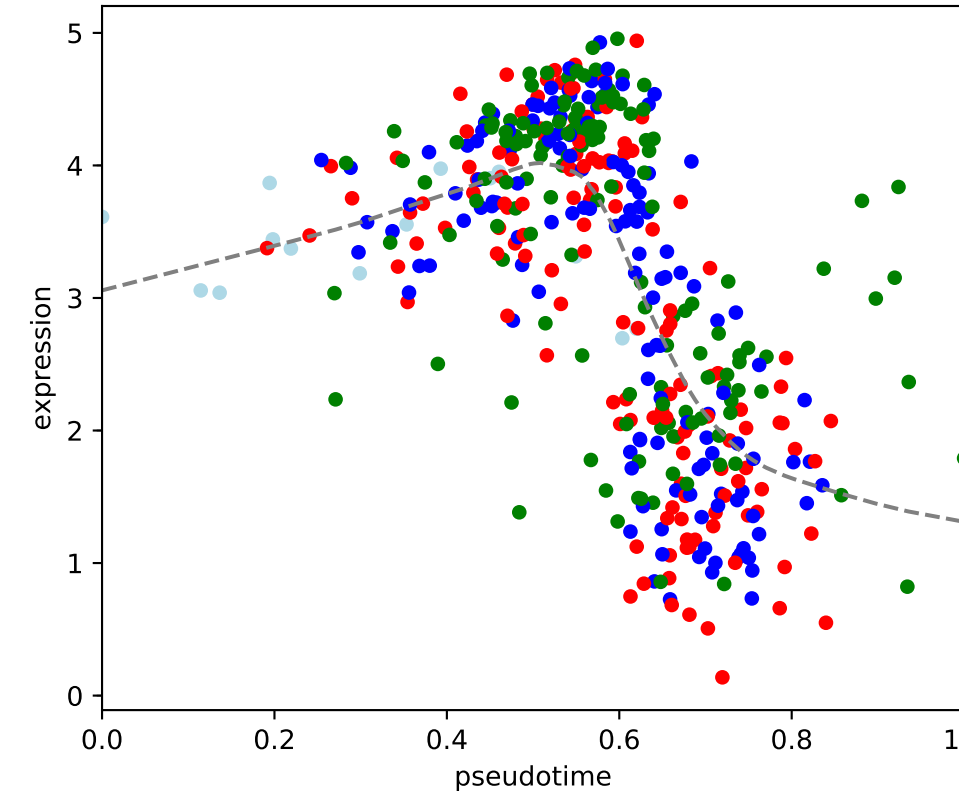
733 1spearman=0.30



ENST00000530835 TUBAP2 (tubulin alpha pseudogene 2)

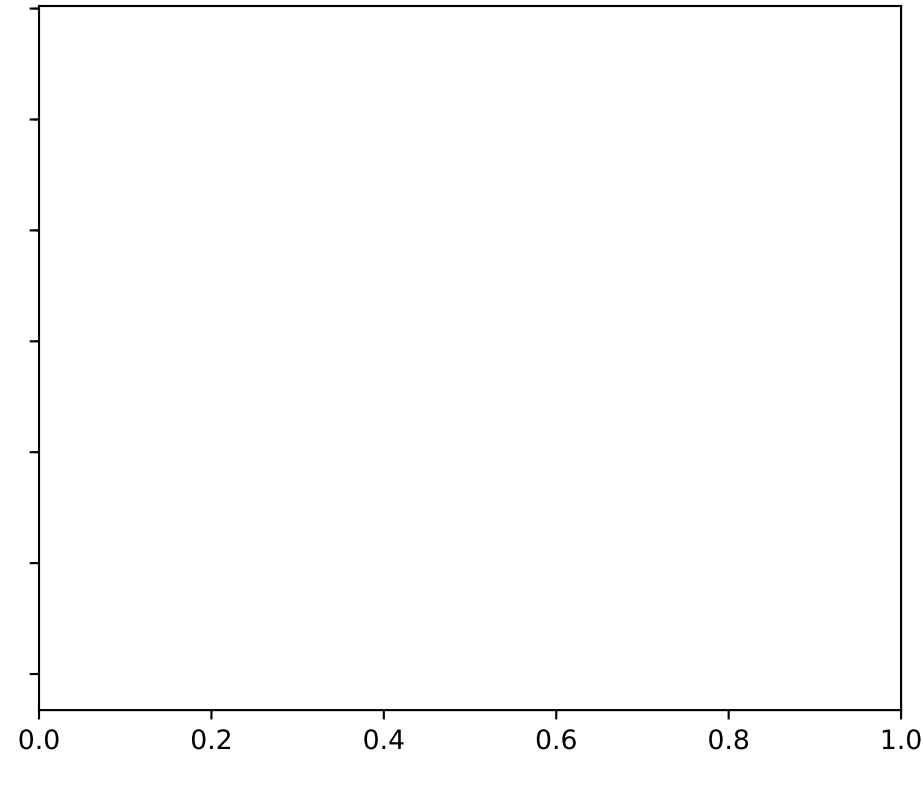
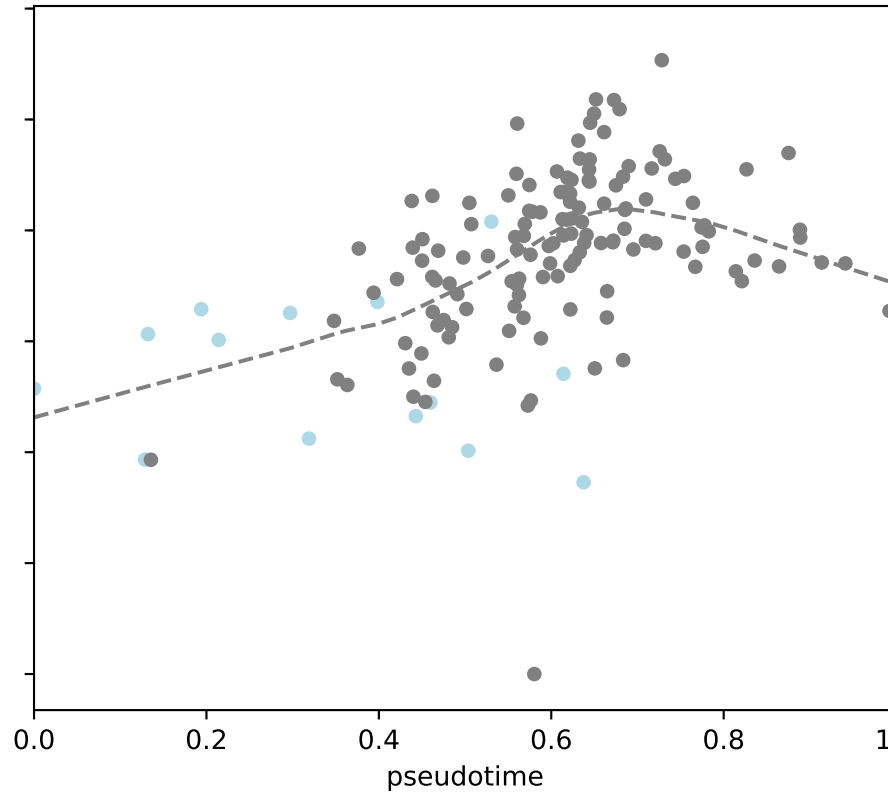
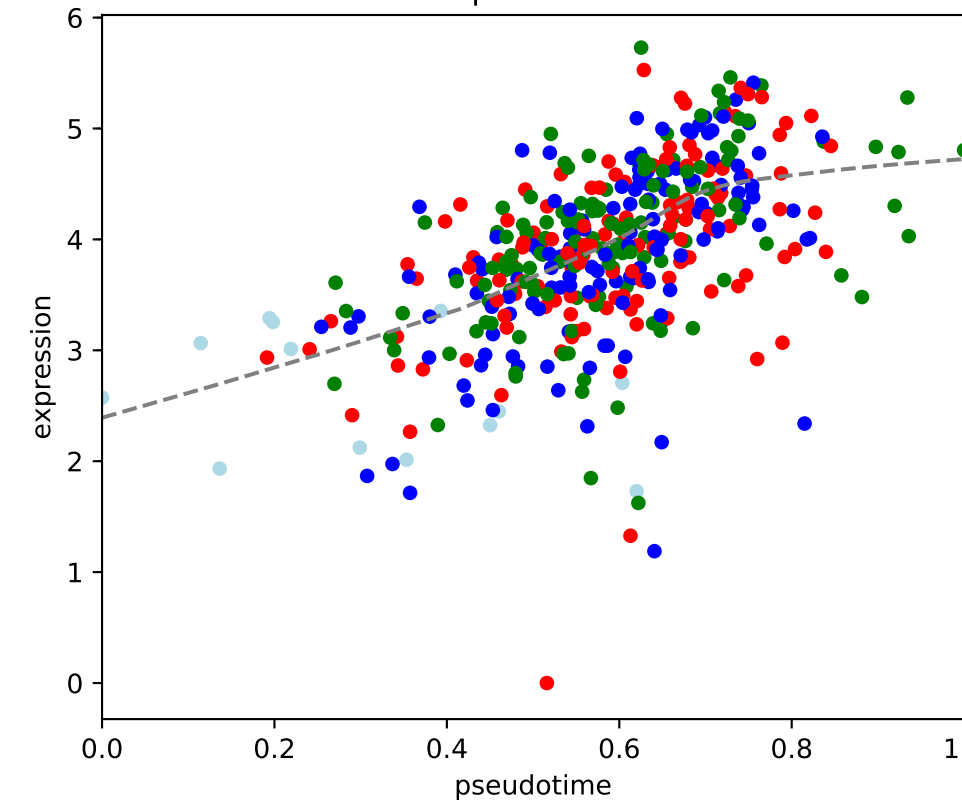
733 0spearman=-0.61

733 1spearman=-0.62



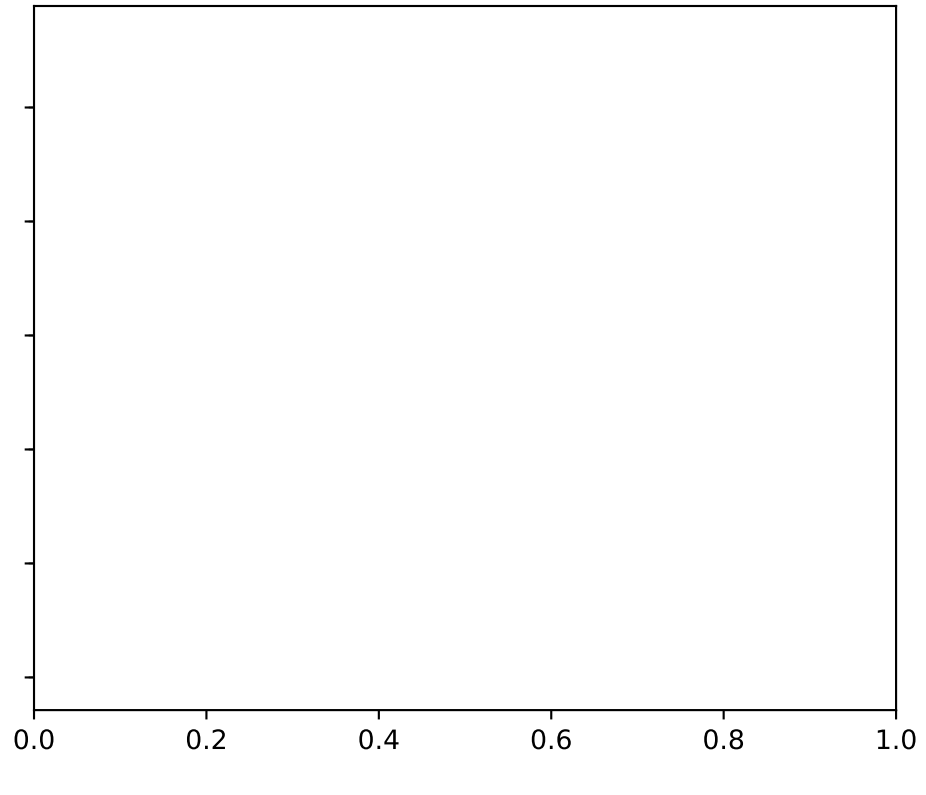
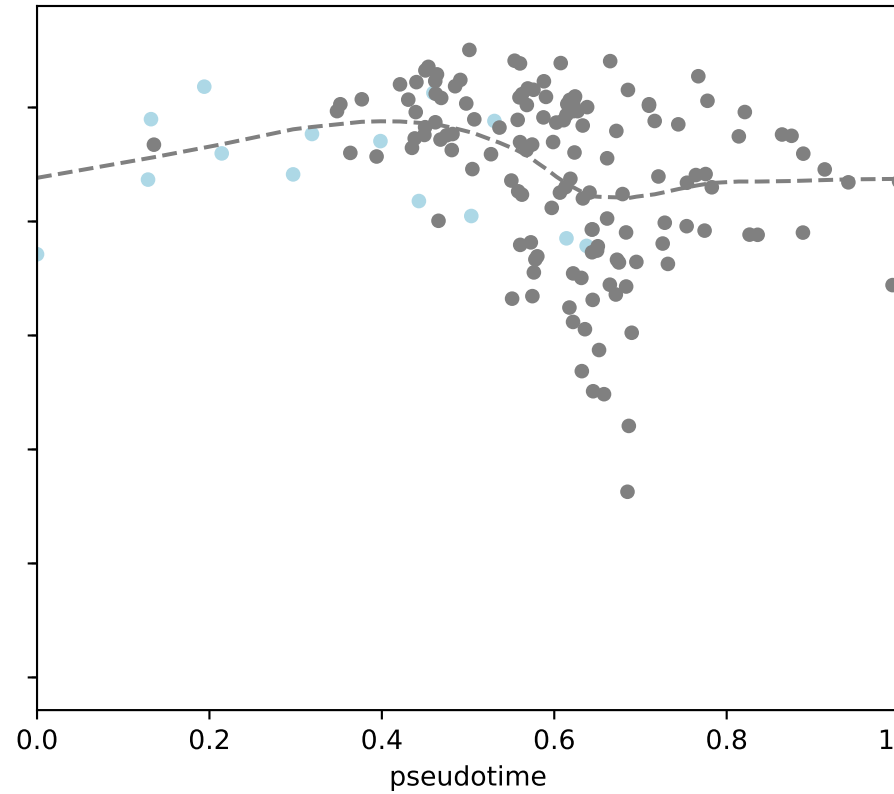
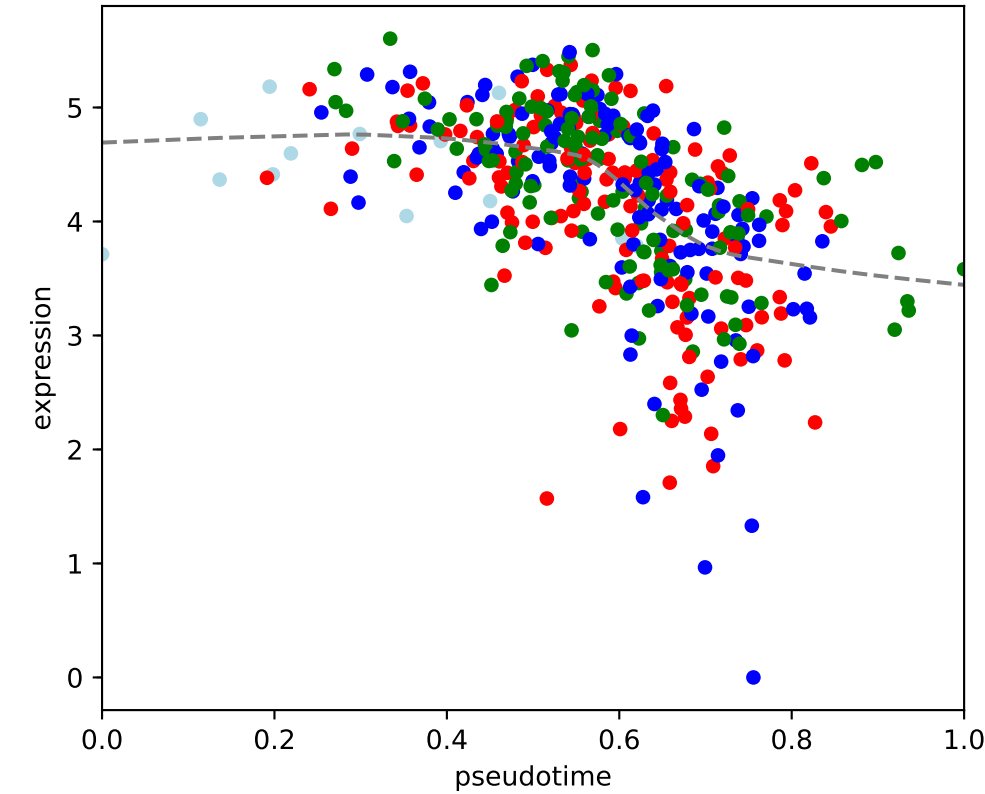
733 0spearman=0.61

733 1spearman=0.50



733 0spearman=-0.61

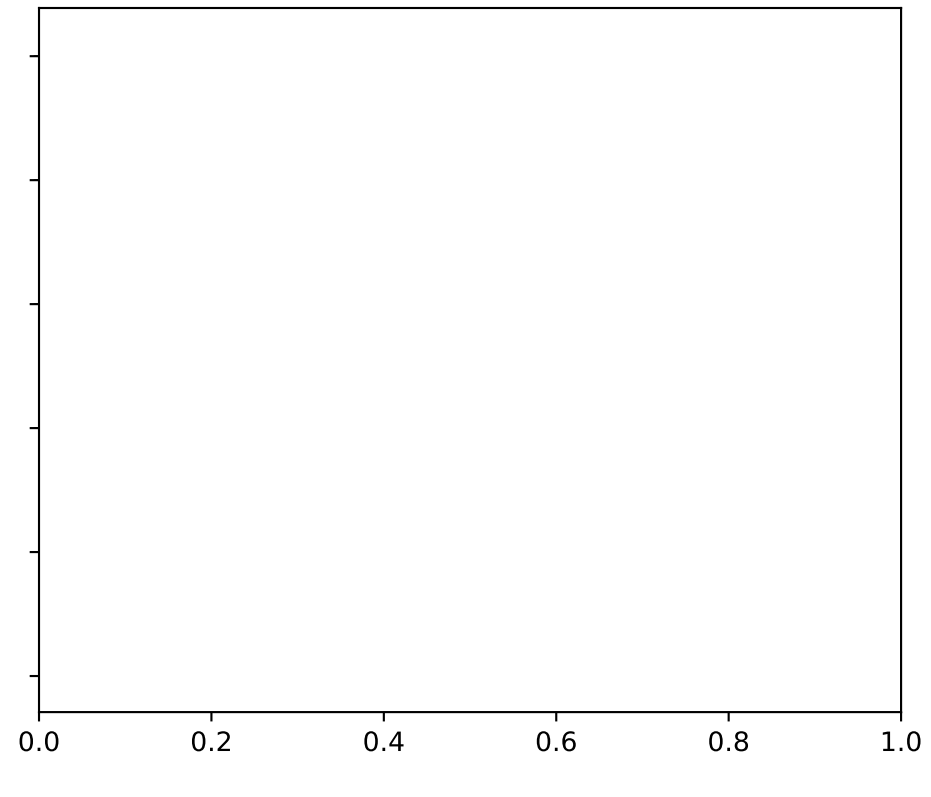
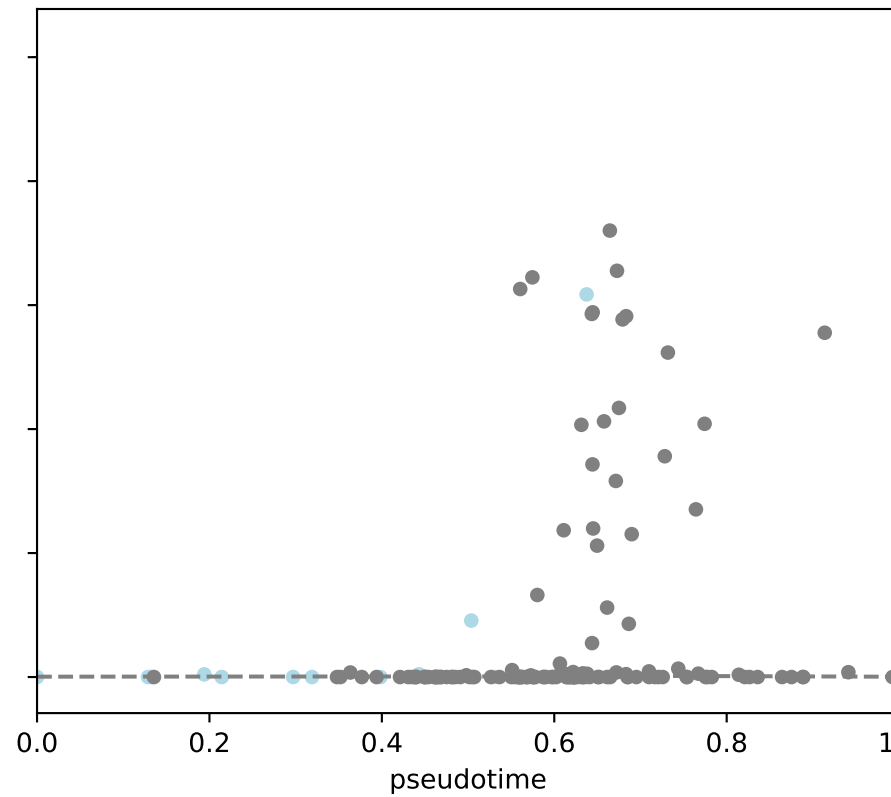
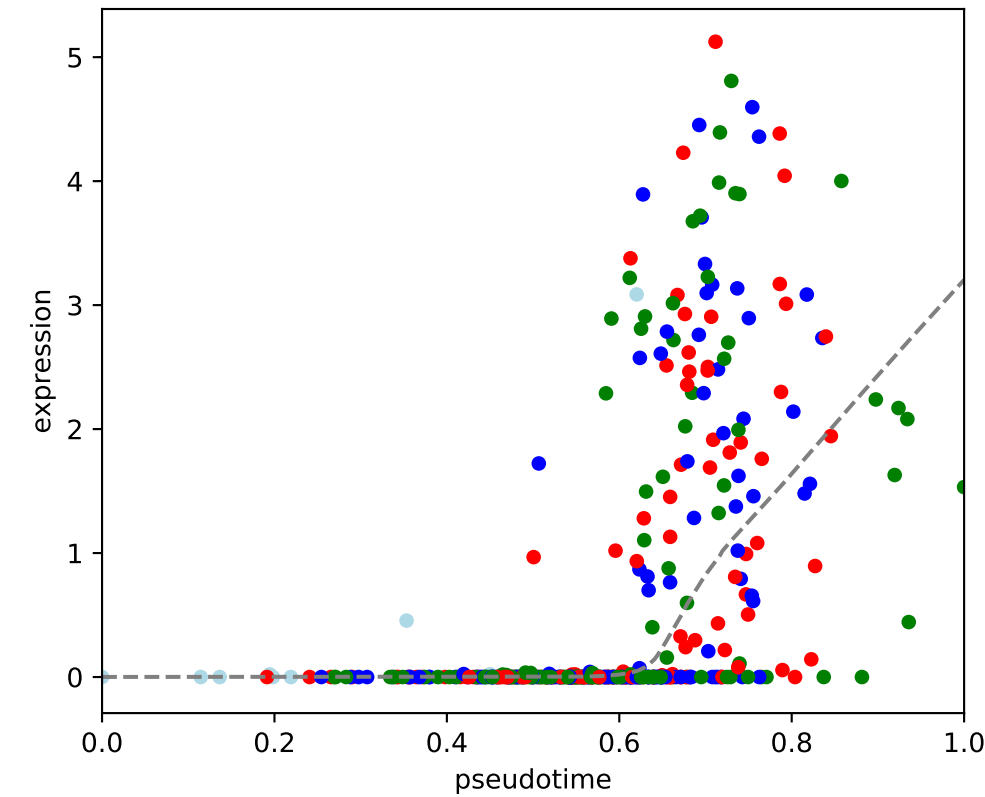
733 1spearman=-0.38



ENST00000281282 CGNL1 (cingulin like 1)

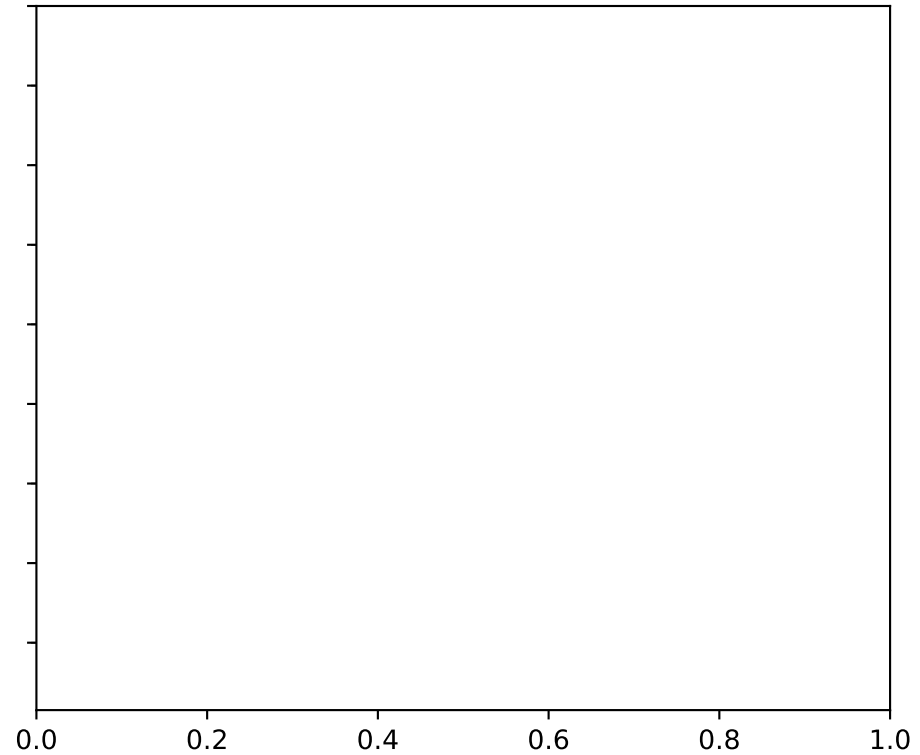
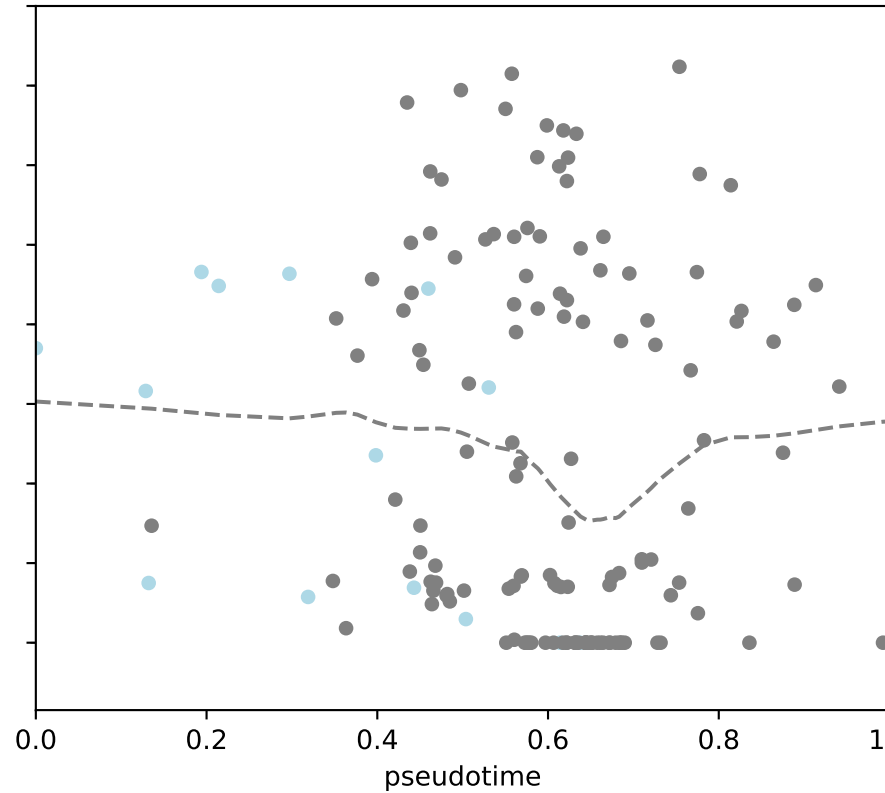
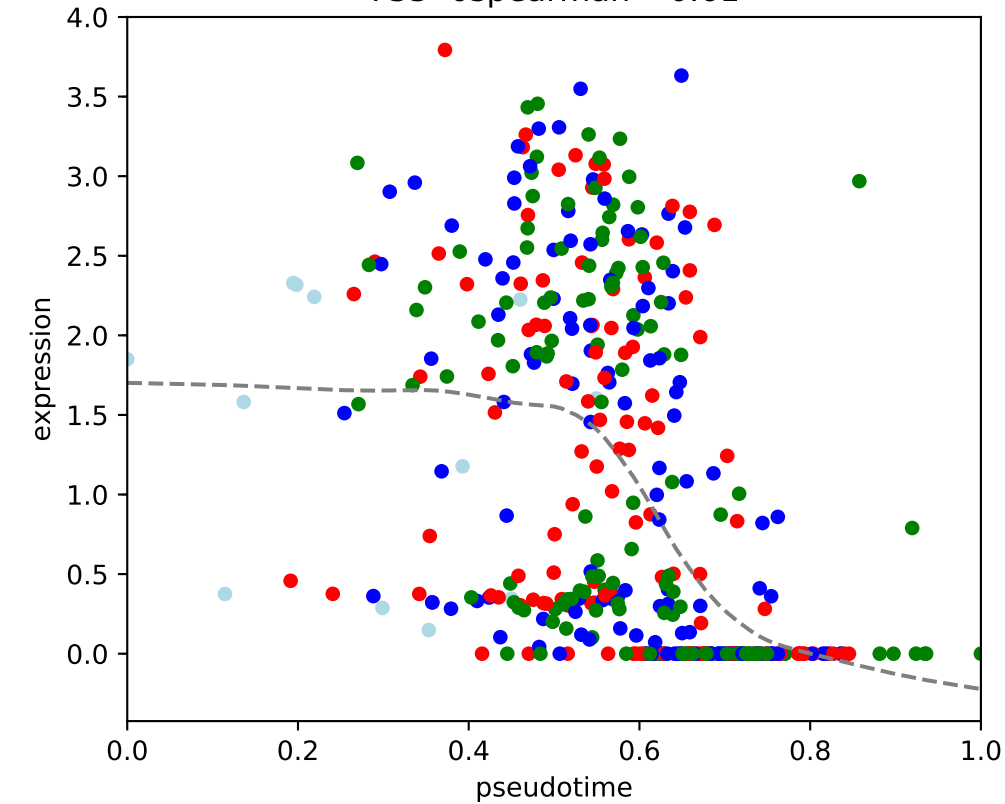
733 0spearman=0.61

733 1spearman=0.33



733 0spearman=-0.61

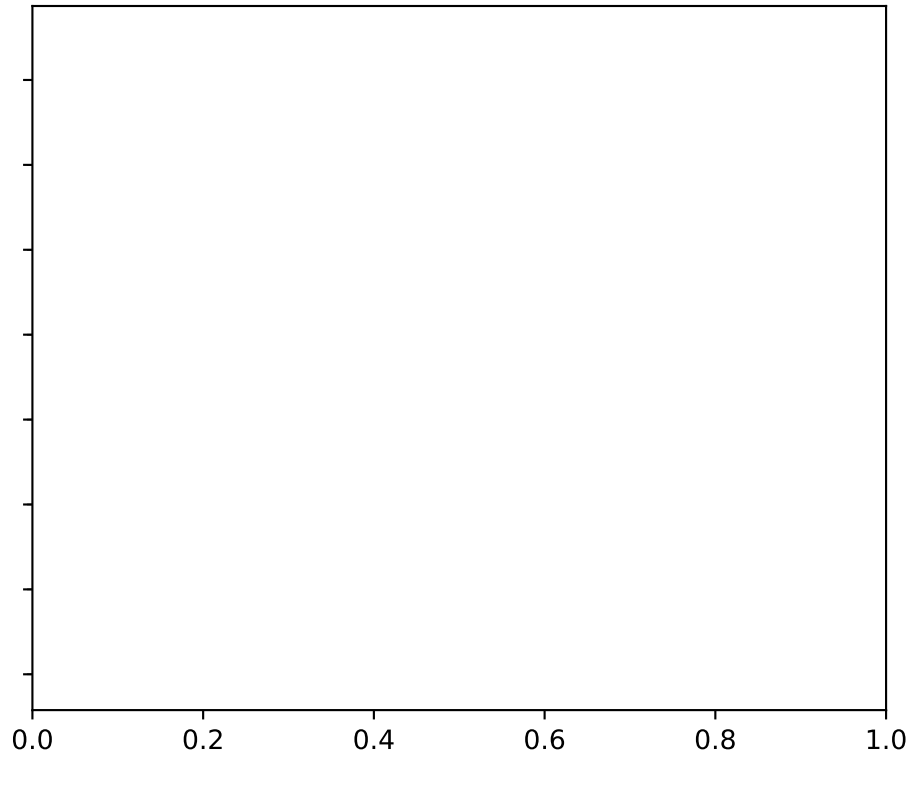
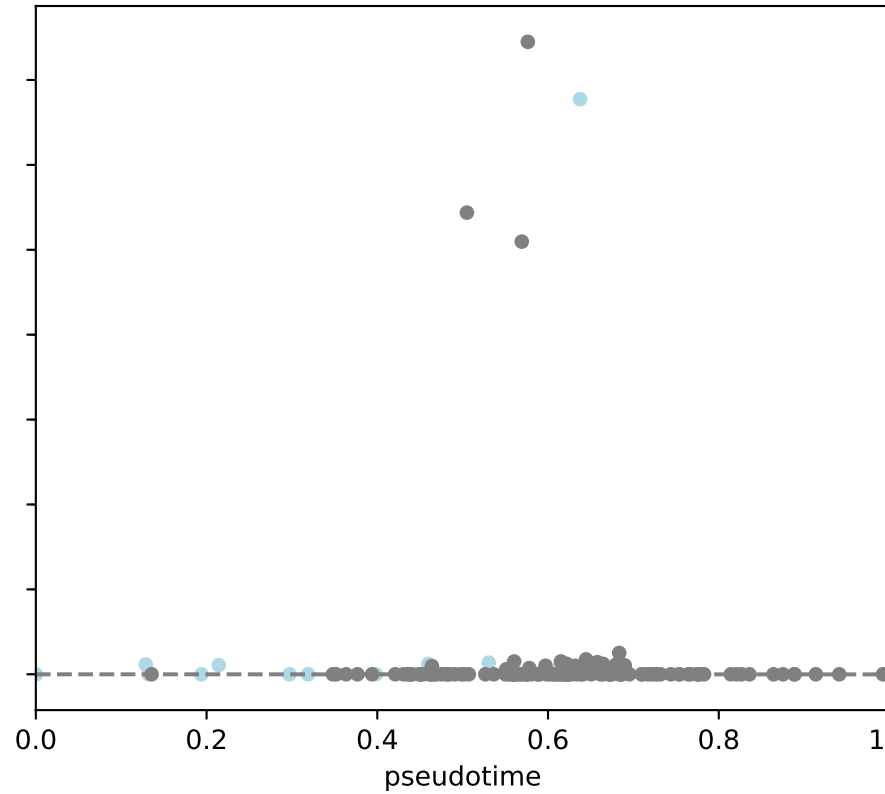
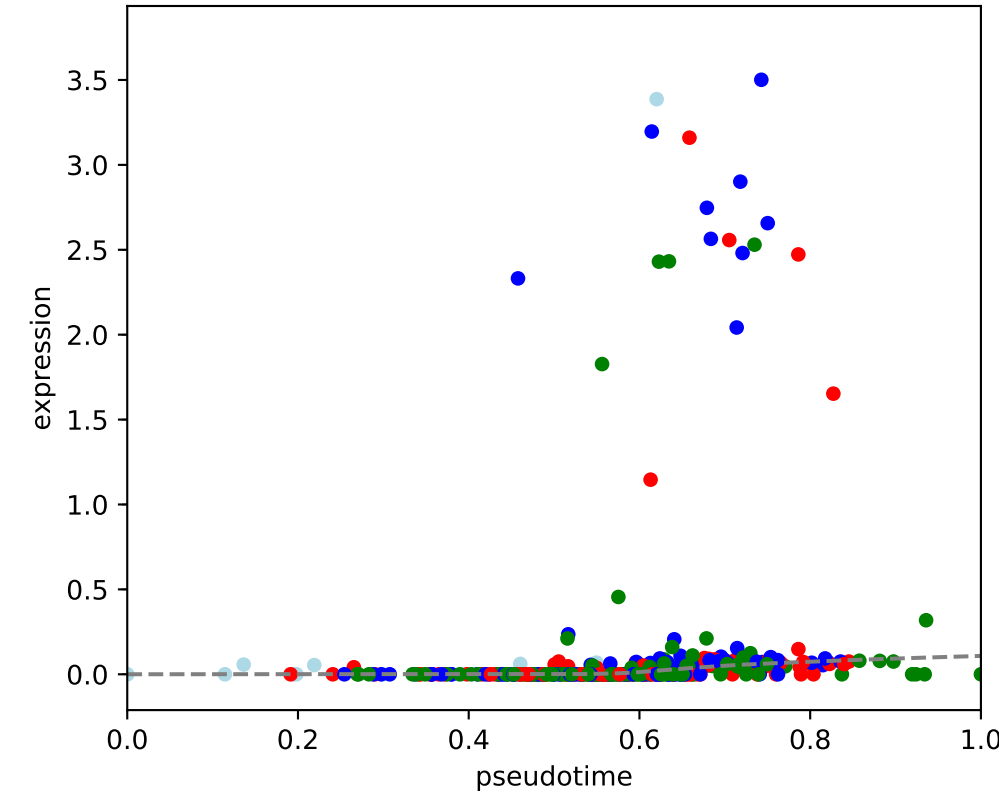
733 1spearman=-0.20



ENST00000467787 GNAT1 (G protein subunit alpha transducin 1)

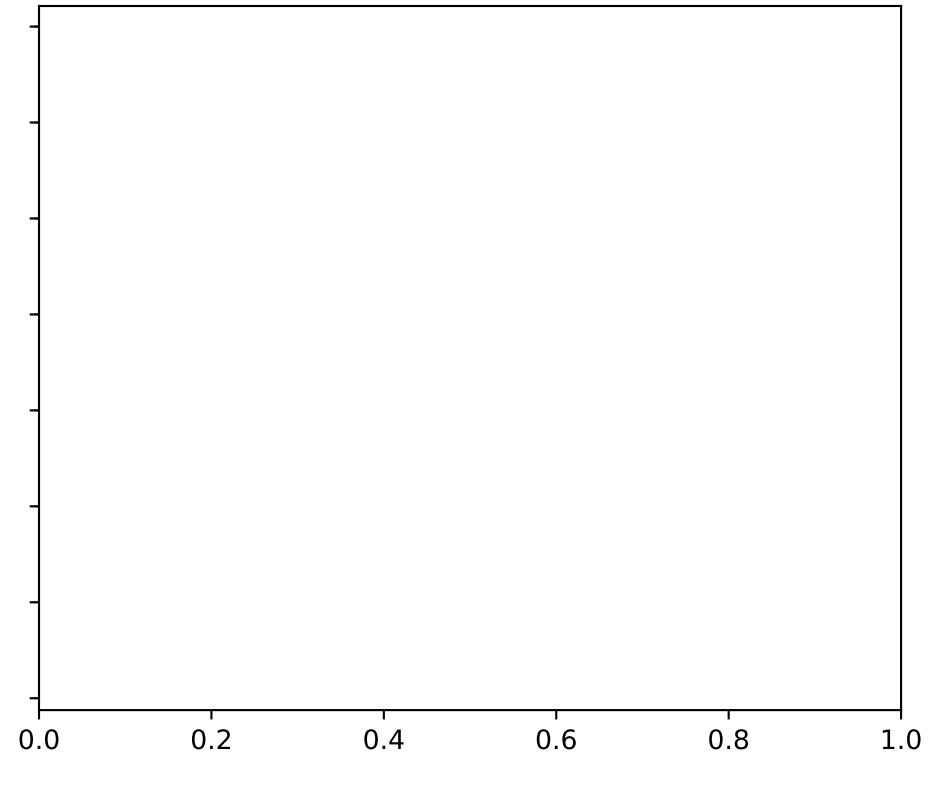
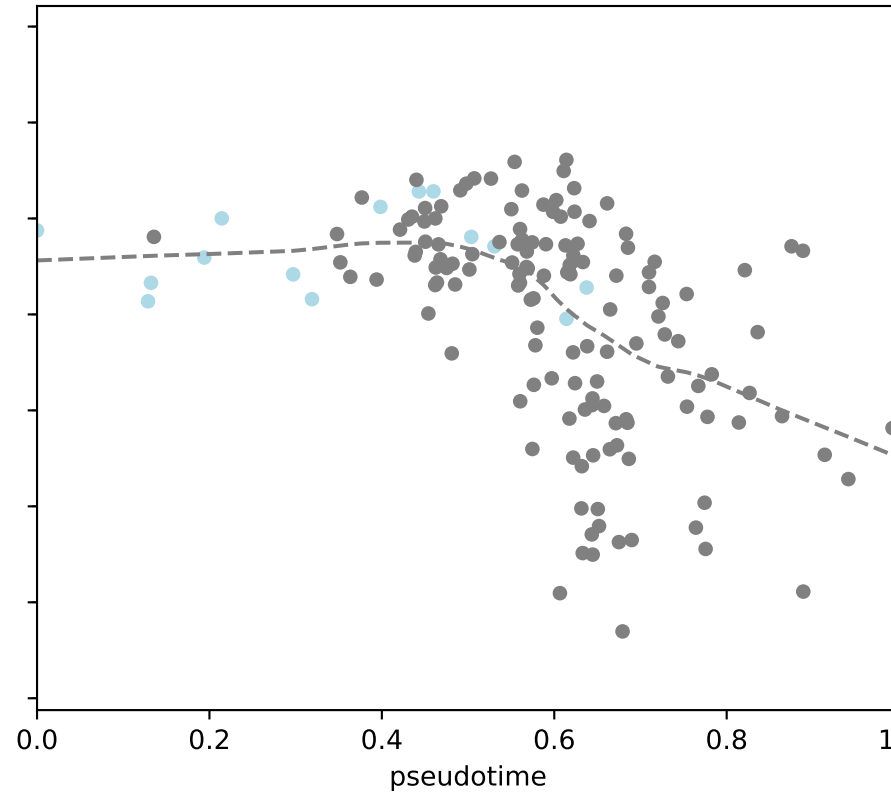
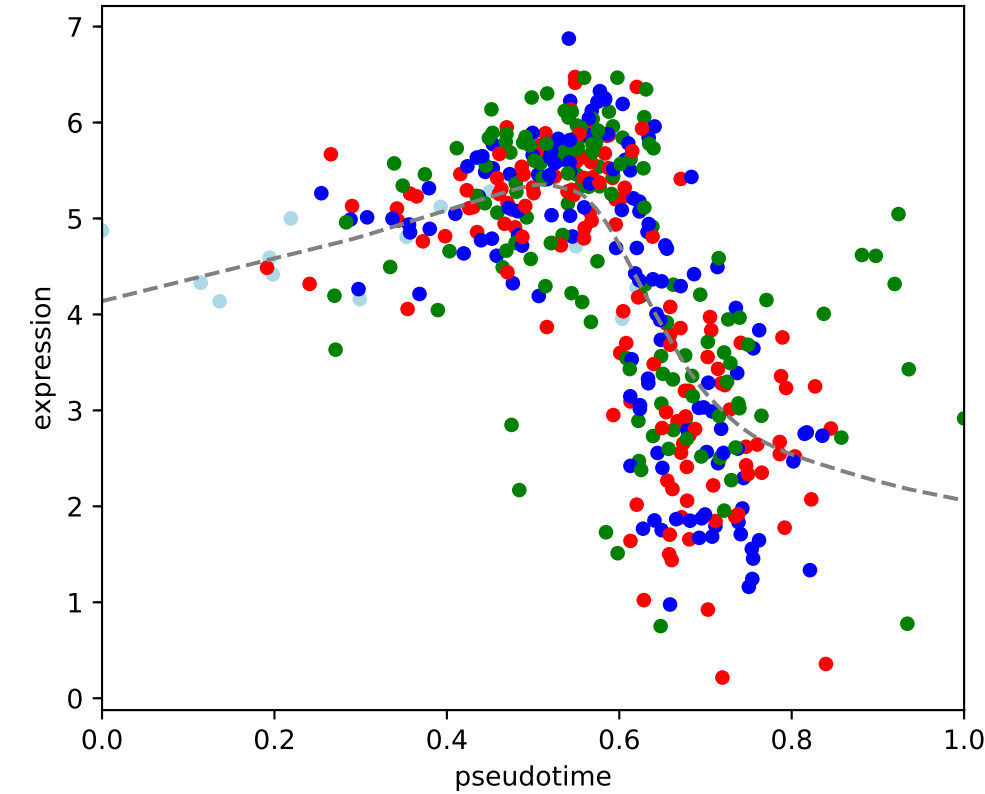
733 0spearman=0.61

733 1spearman=0.04



733 0spearman=-0.61

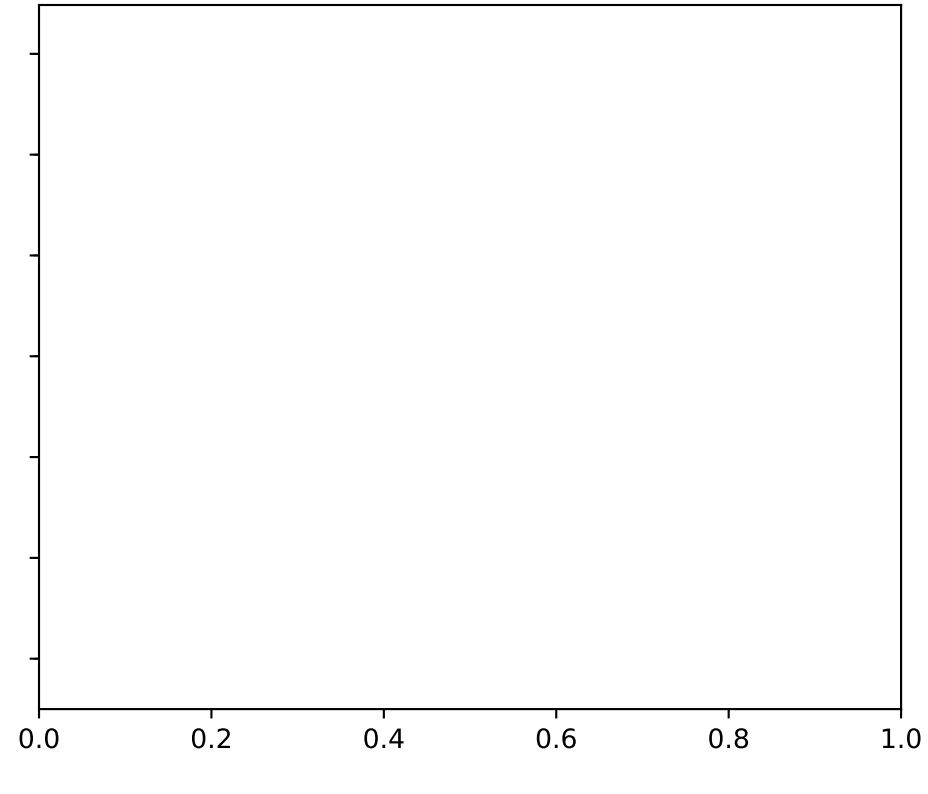
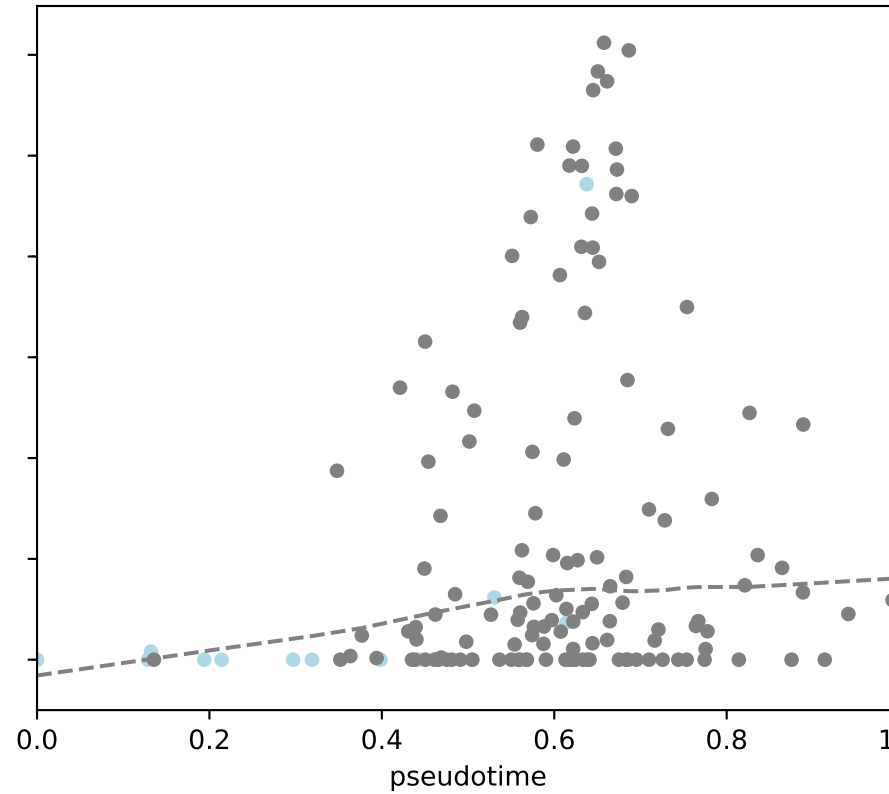
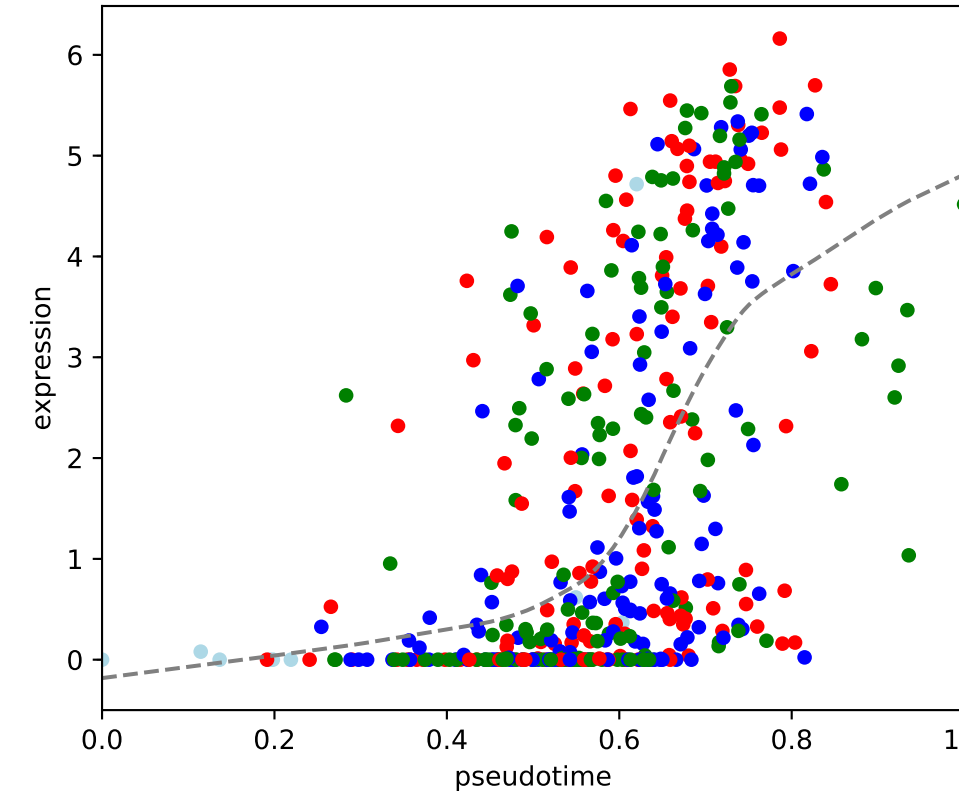
733 1spearman=-0.58



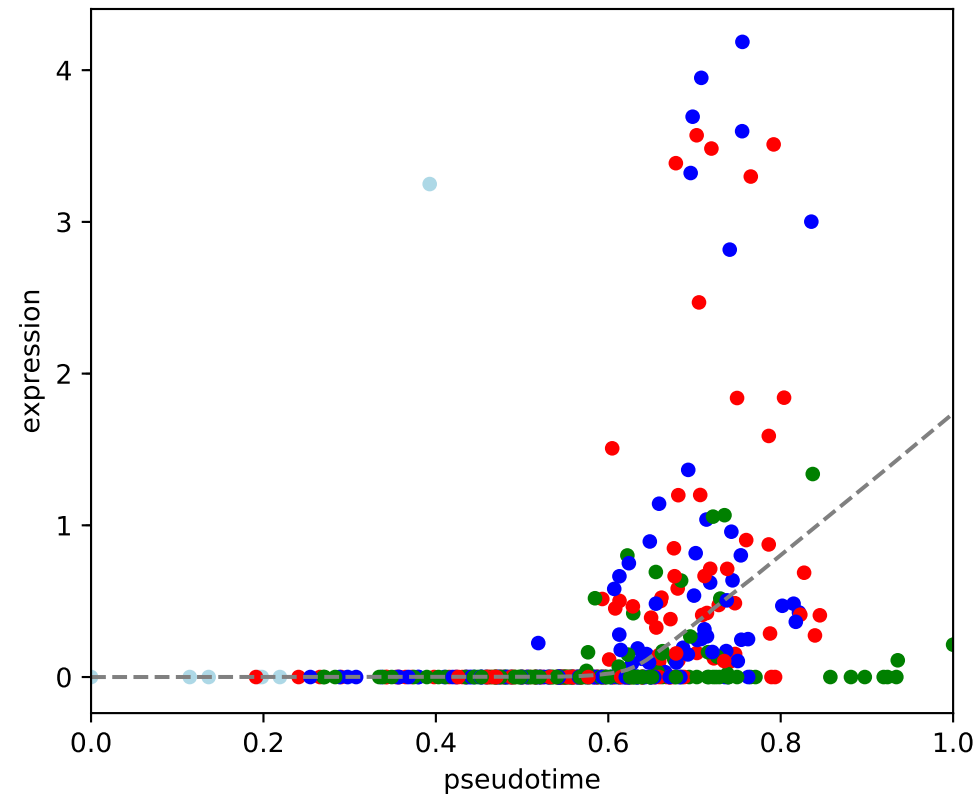
ENST00000455979 SAMD11 (sterile alpha motif domain containing 11)

733 0spearman=0.61

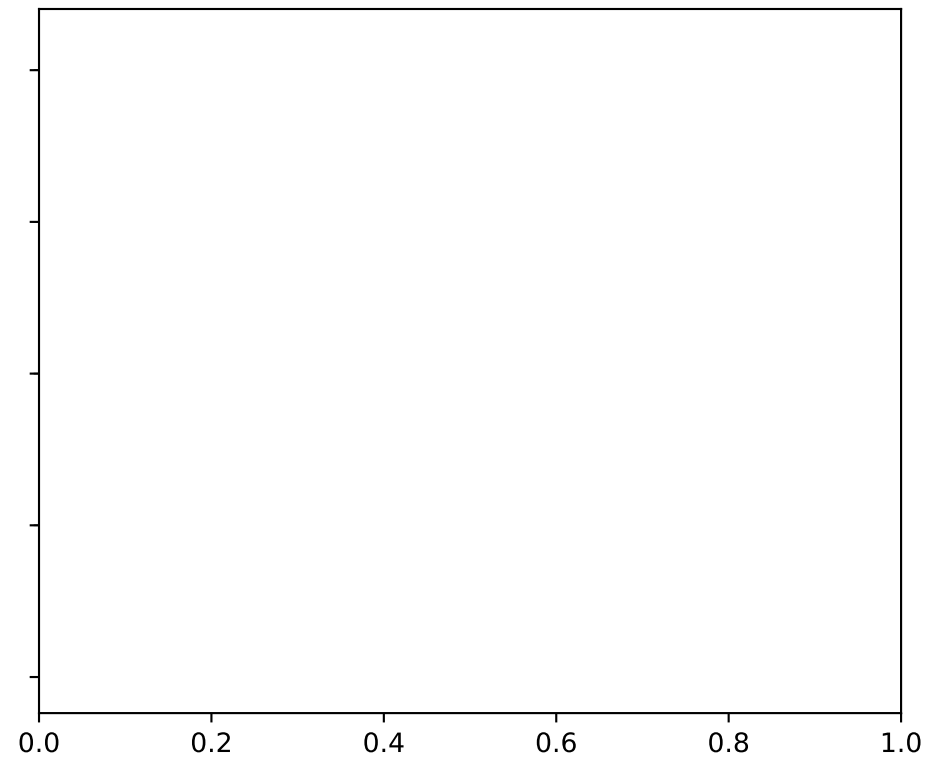
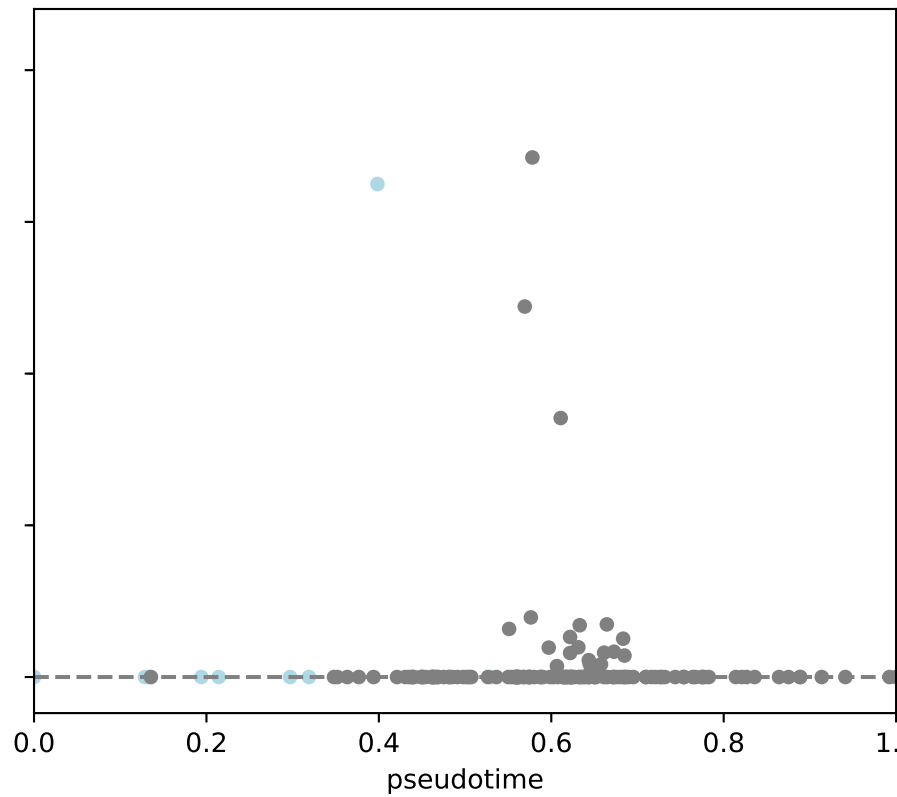
733 1spearman=0.28



733 0spearman=0.60



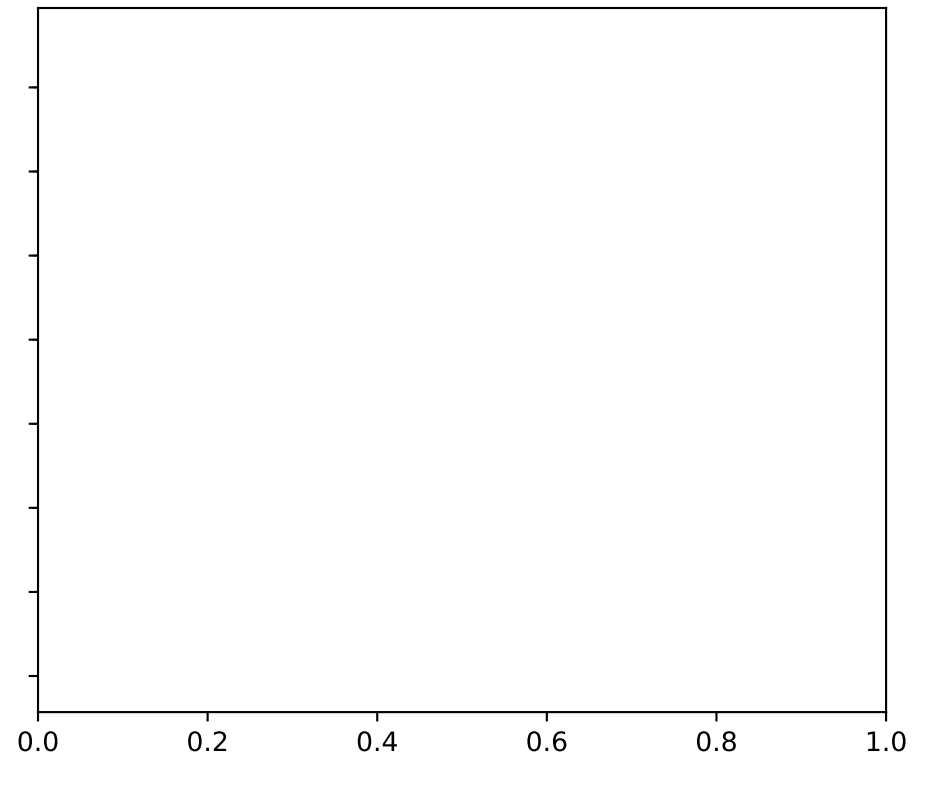
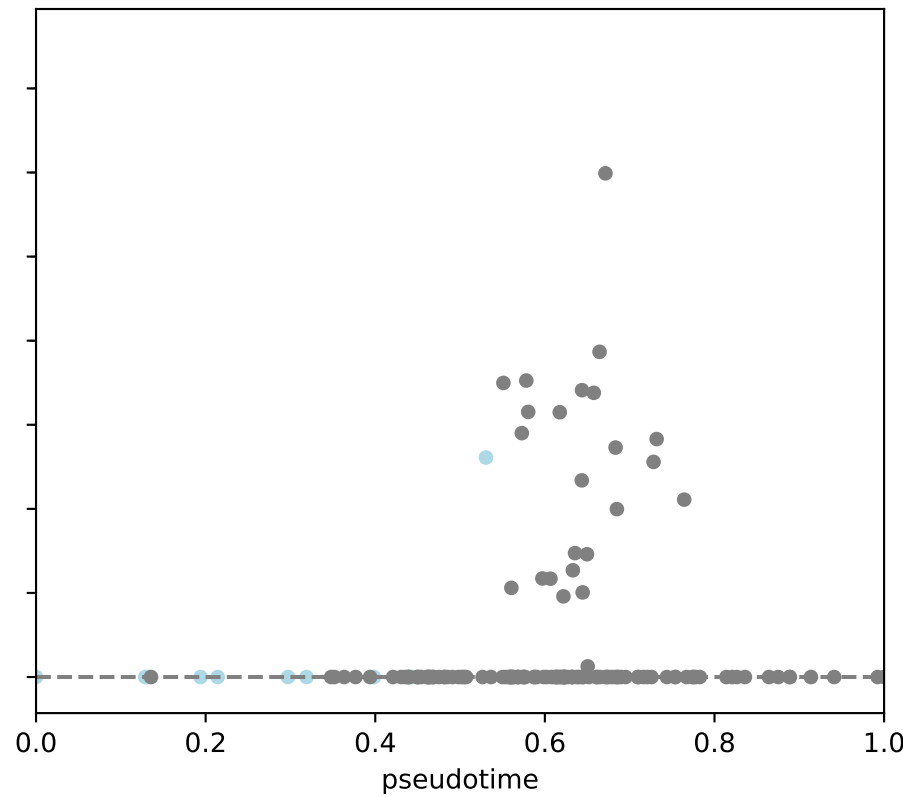
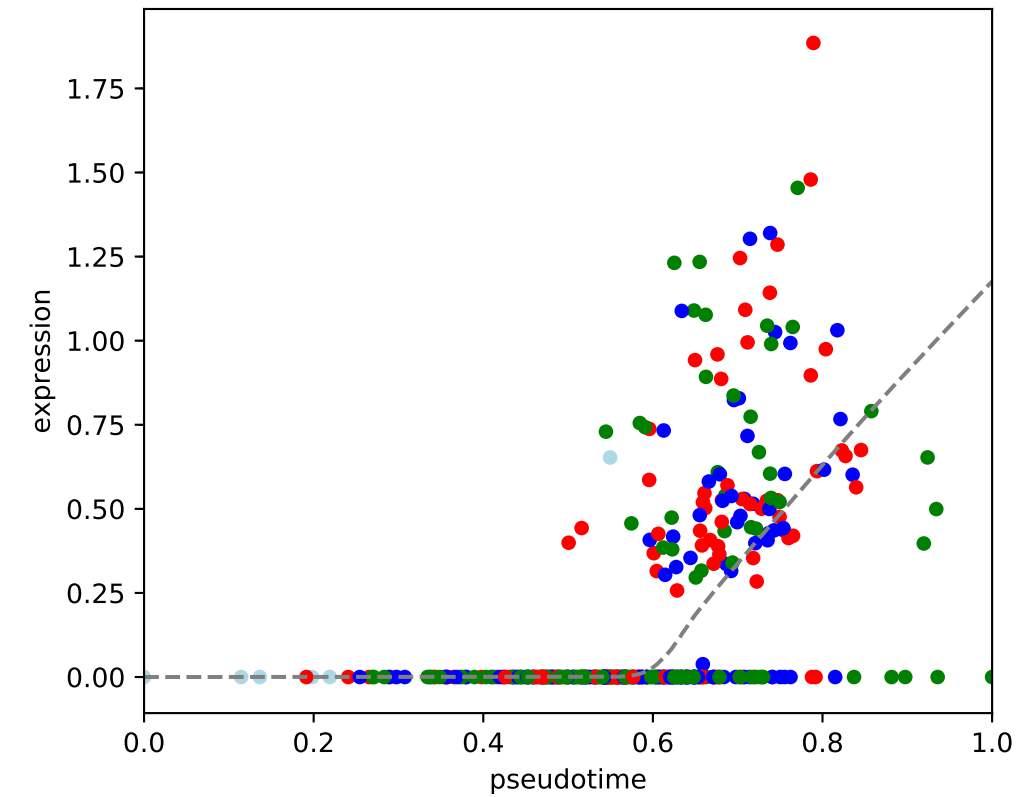
733 1spearman=0.08



ENST00000602051 MRLN (myoregulin)

733 0spearman=0.60

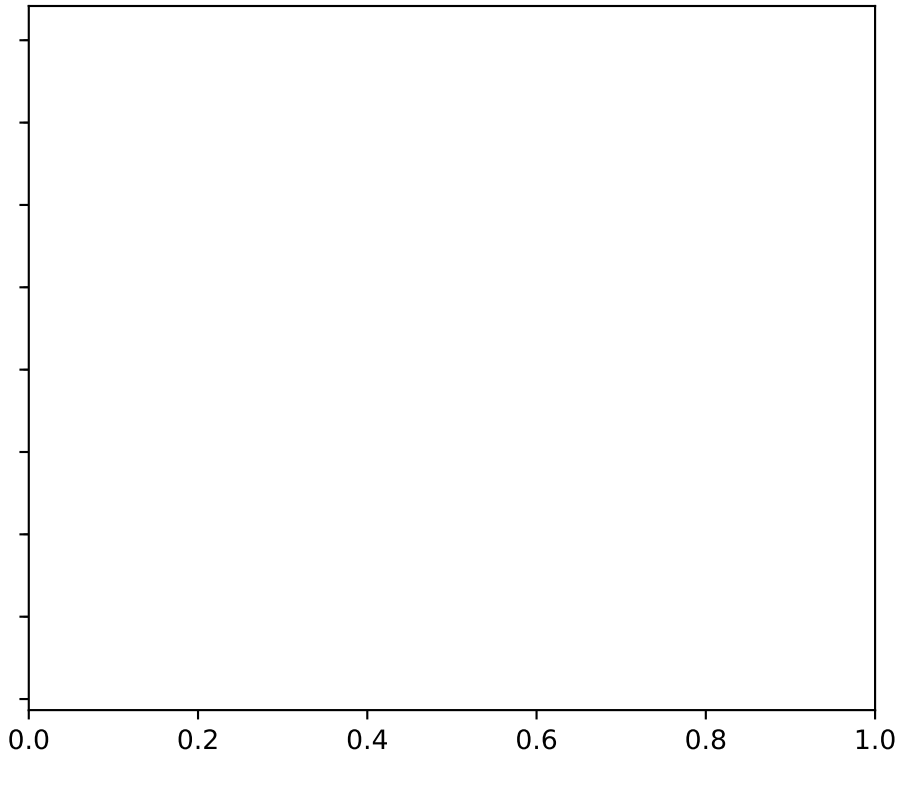
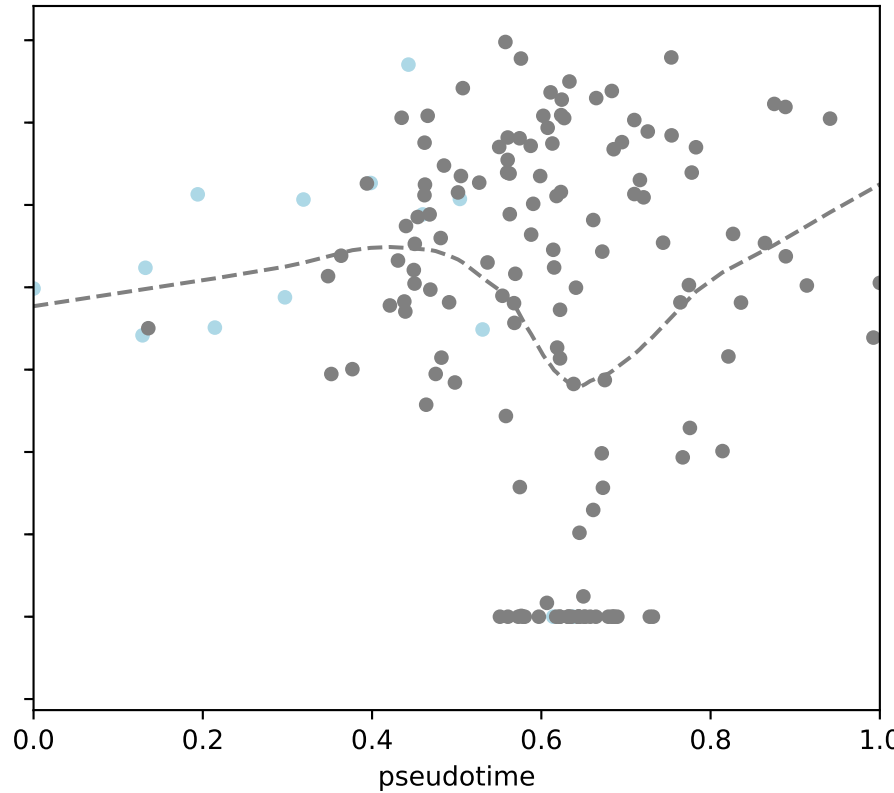
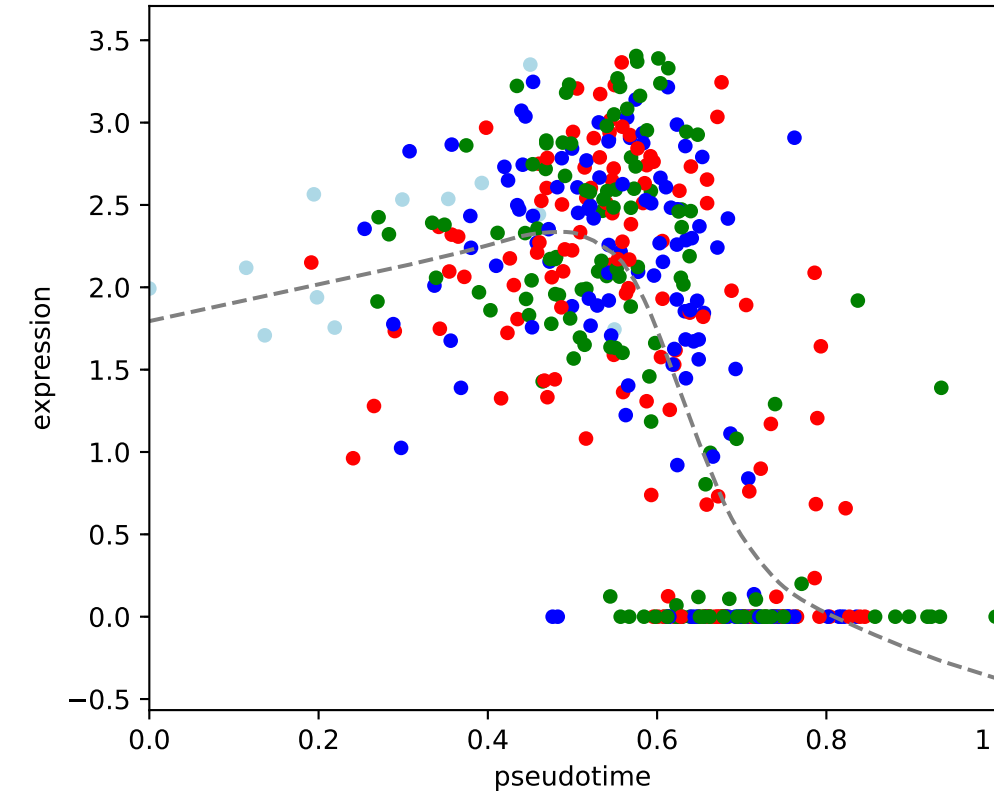
733 1spearman=0.17



ENST00000622865 GNAT2 (G protein subunit alpha transducin 2)

733 0spearman=-0.60

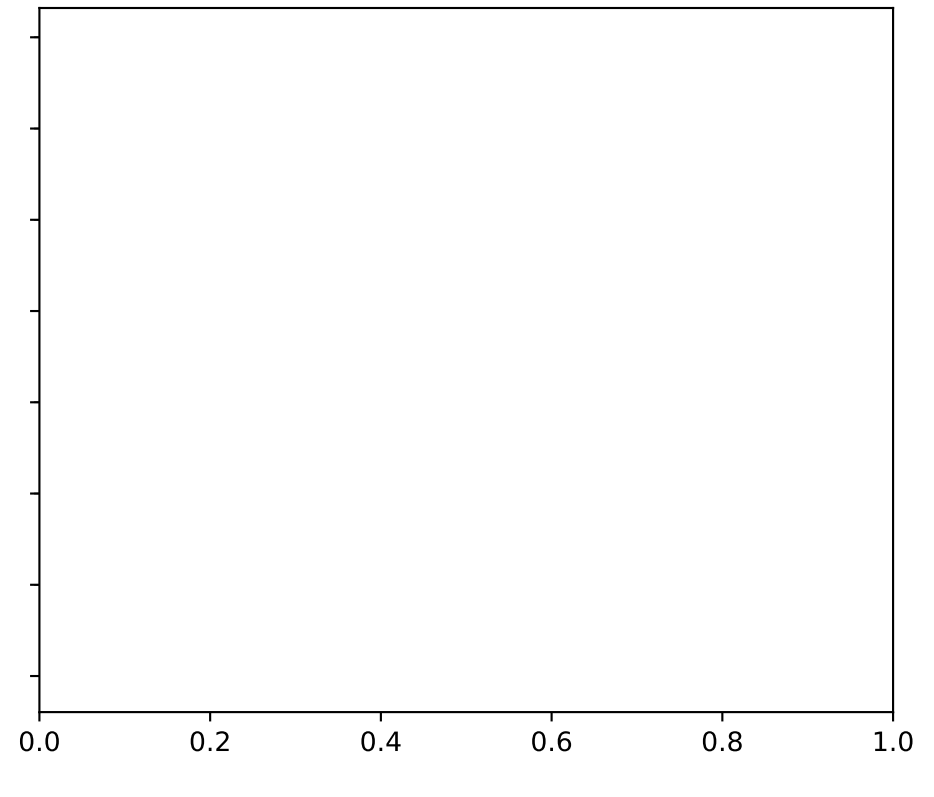
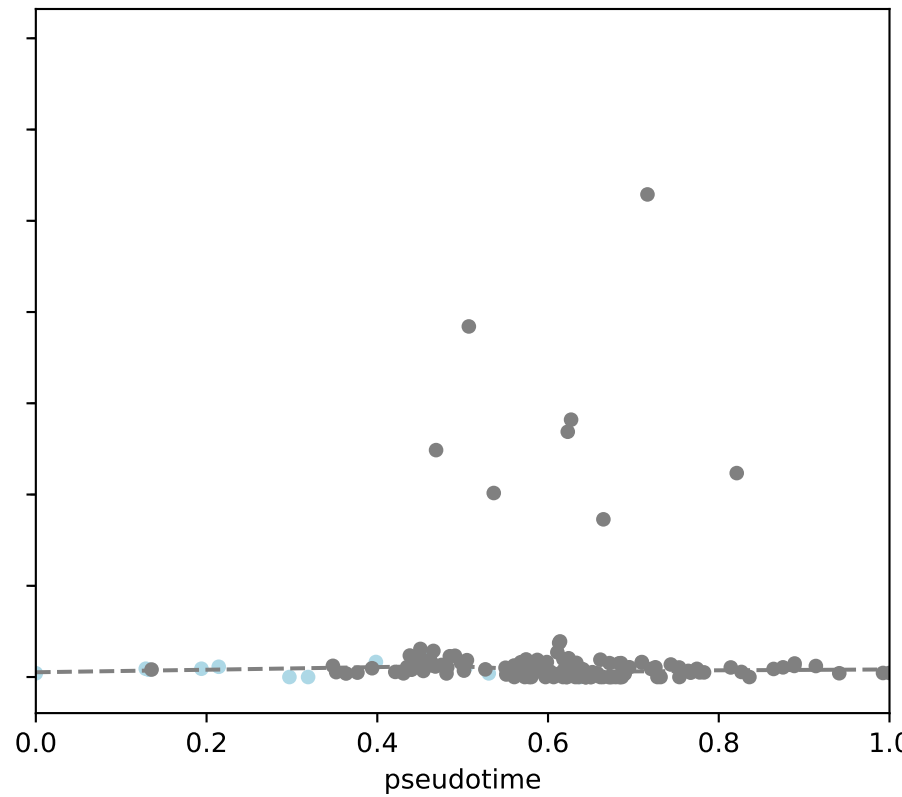
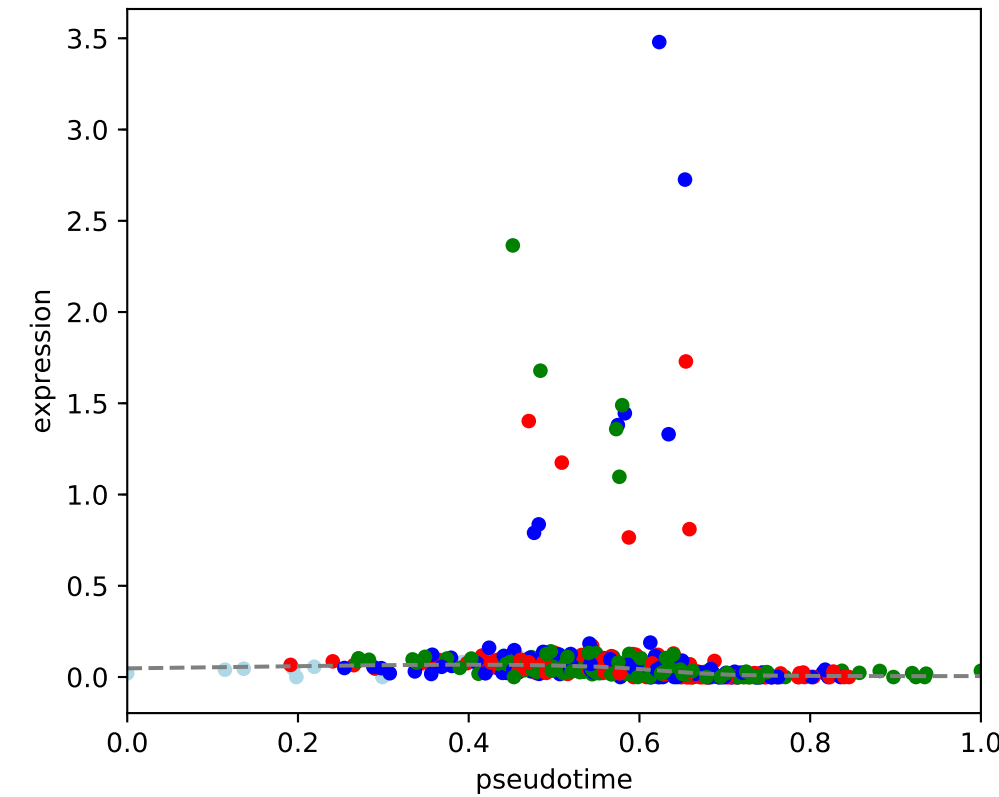
733 1spearman=-0.11



ENST00000373230 PPA1 (pyrophosphatase (inorganic) 1)

733 0spearman=-0.60

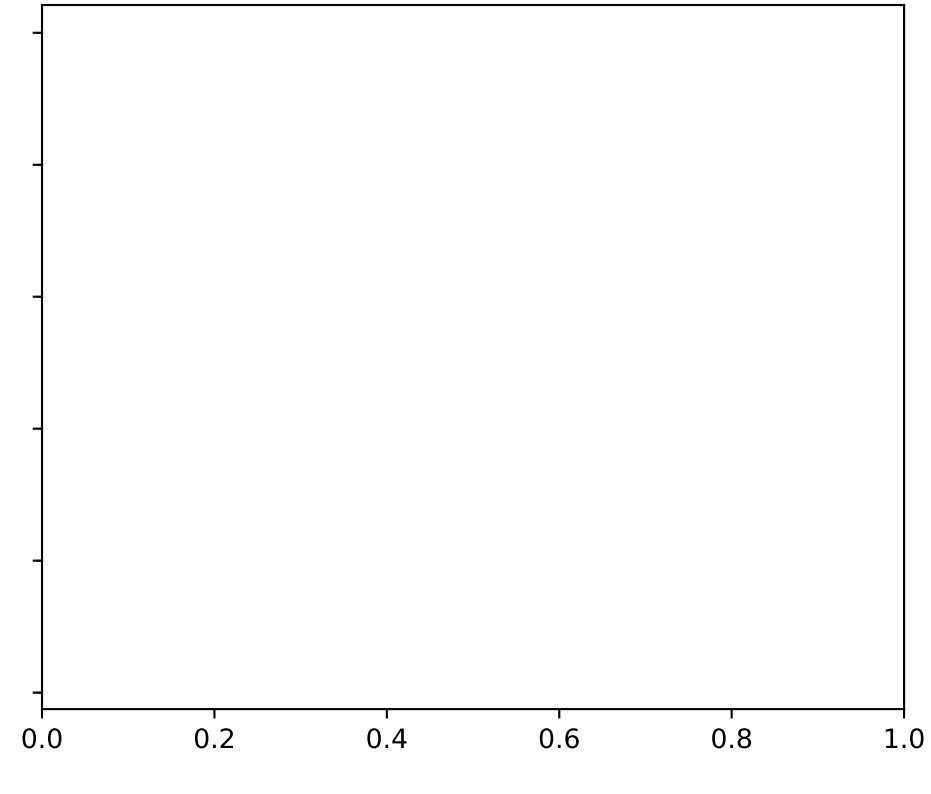
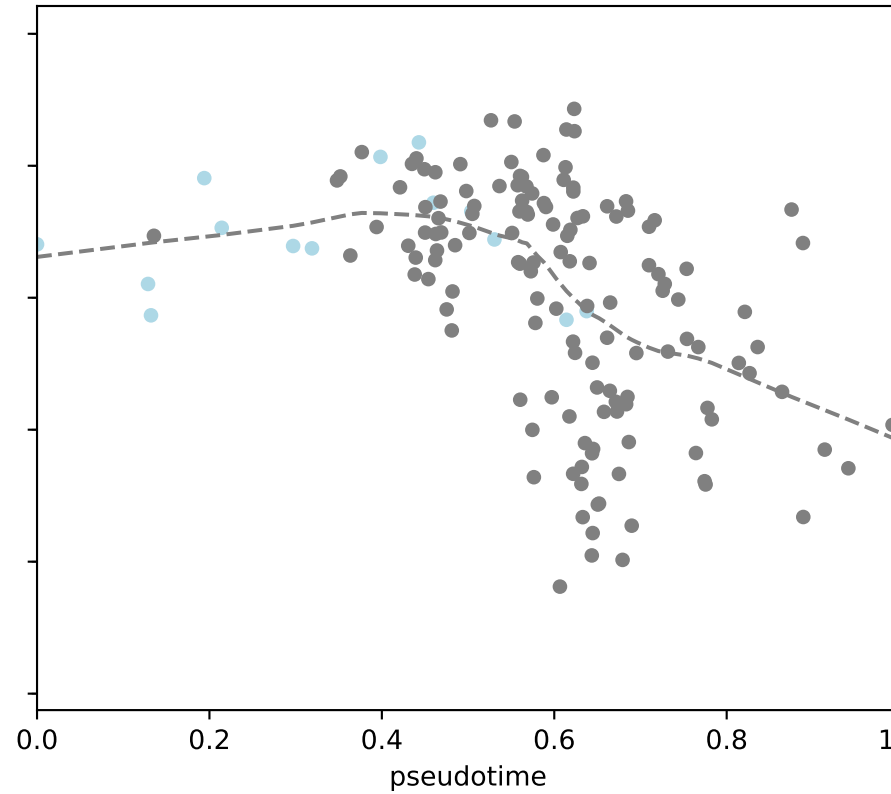
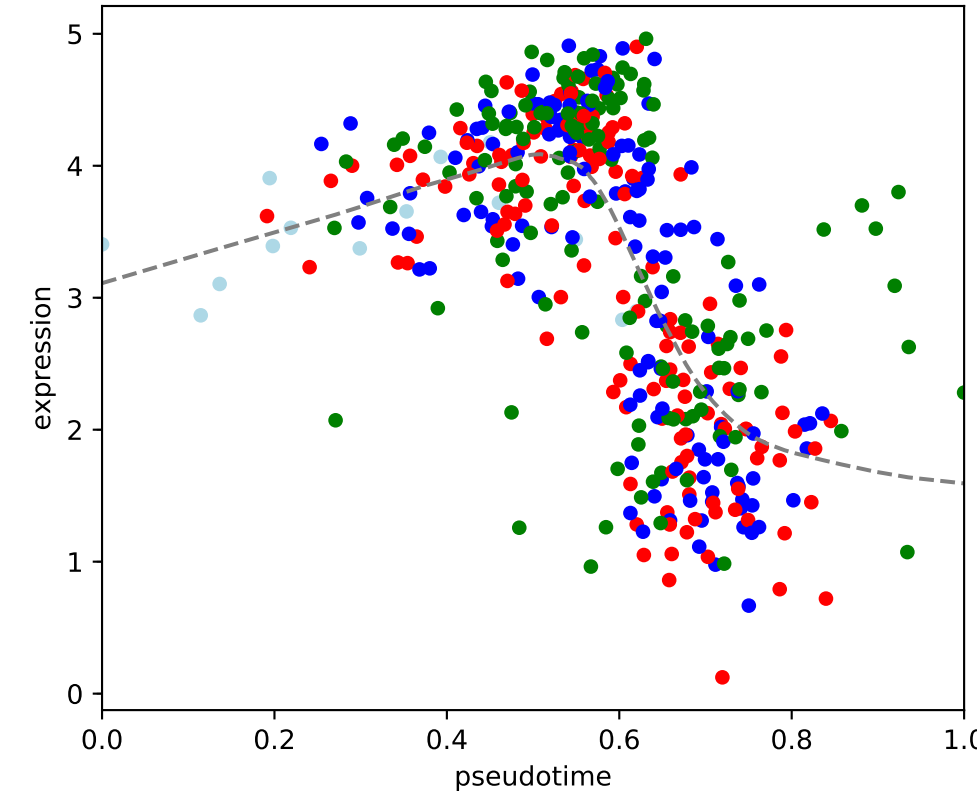
733 1spearman=-0.20



ENST00000548149 AC011603.2

733 0spearman=-0.60

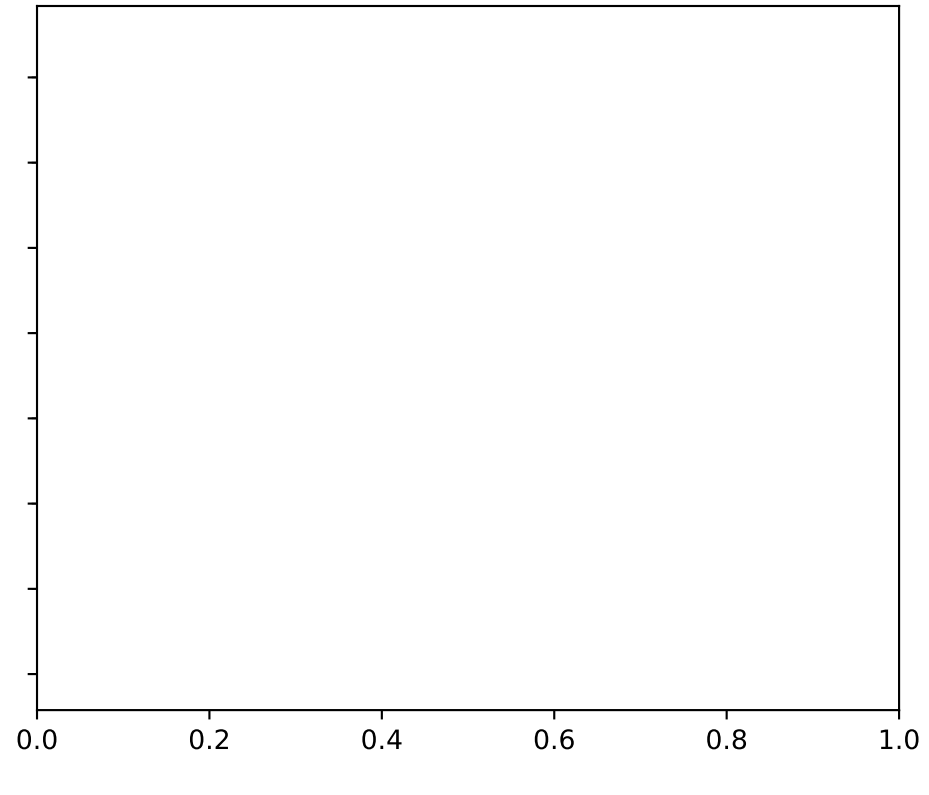
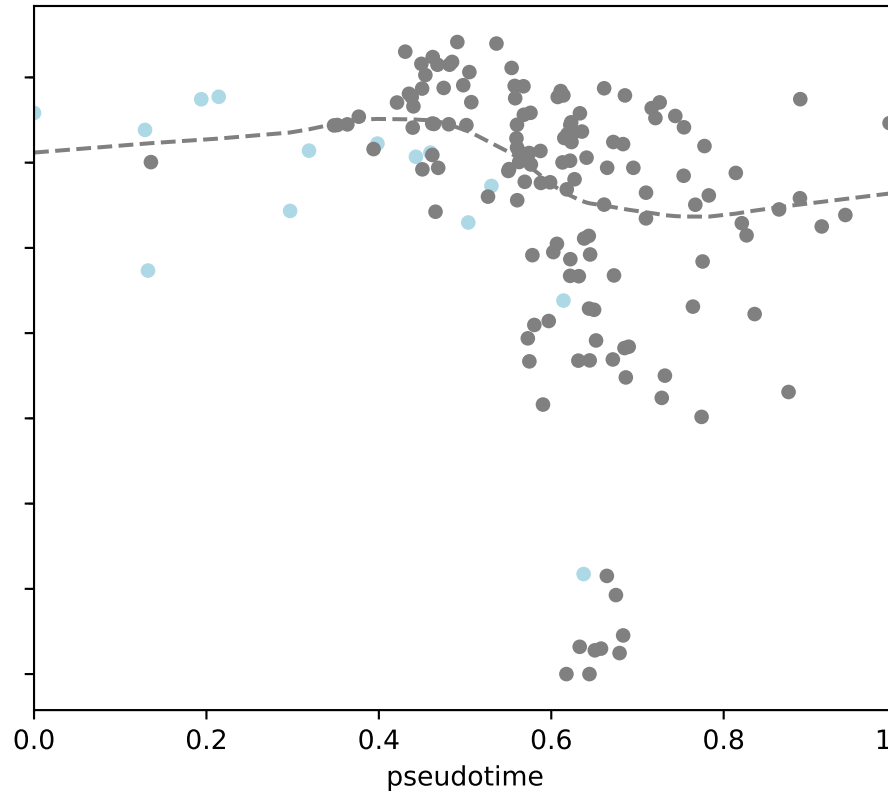
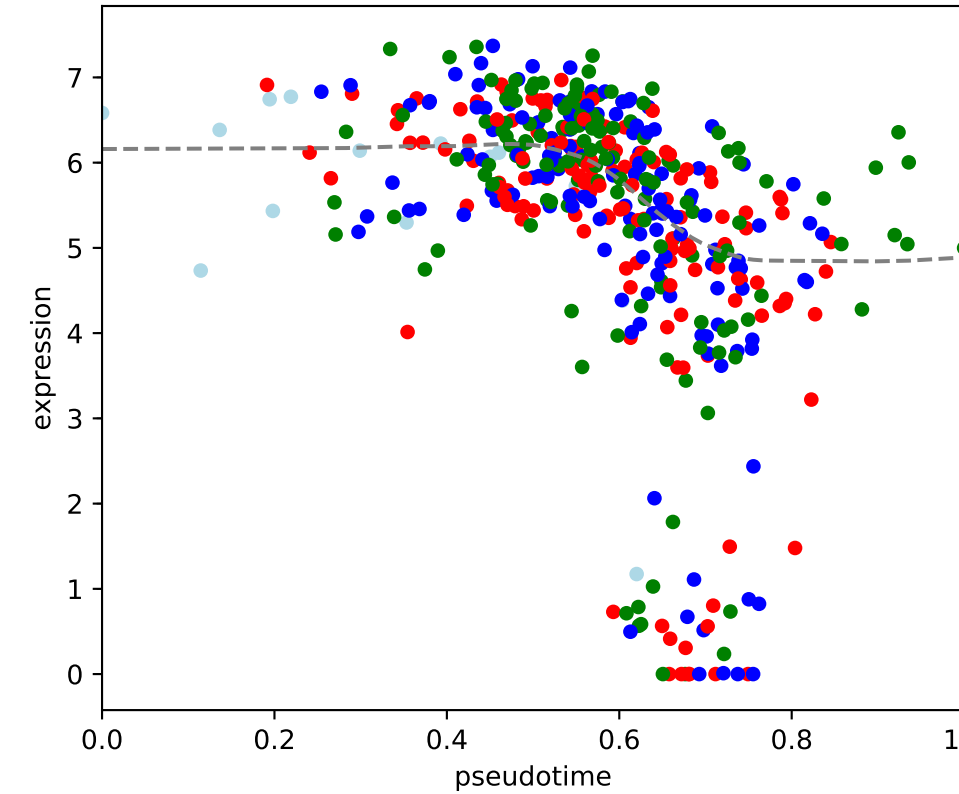
733 1spearman=-0.55



ENST00000274606 NHP2 (NHP2 ribonucleoprotein)

733 0spearman=-0.60

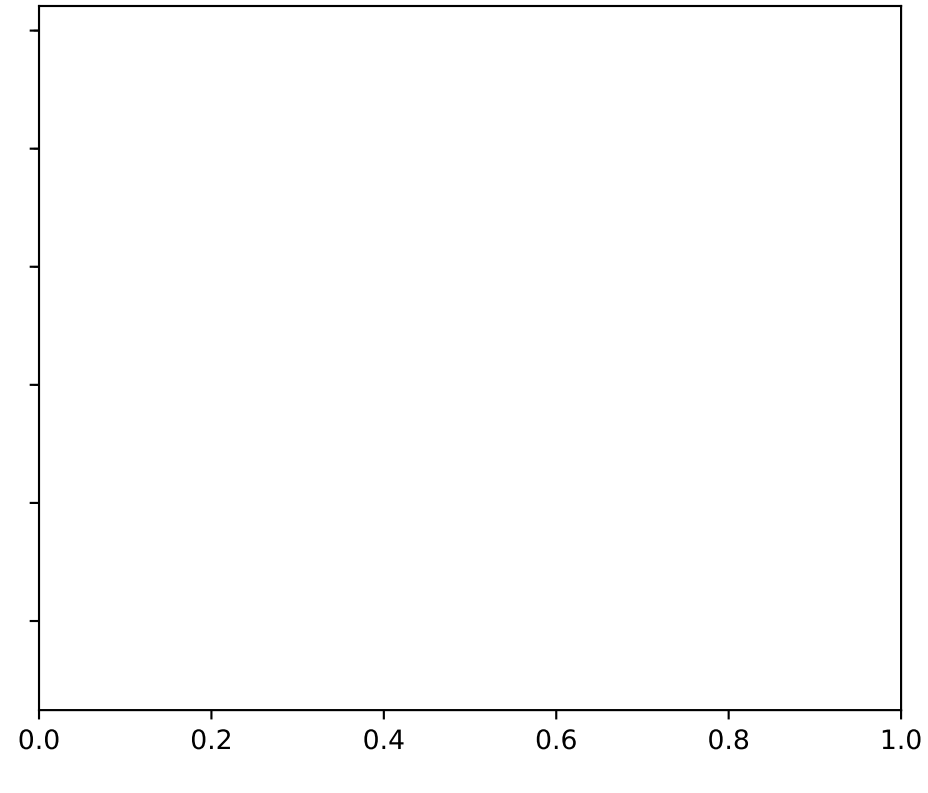
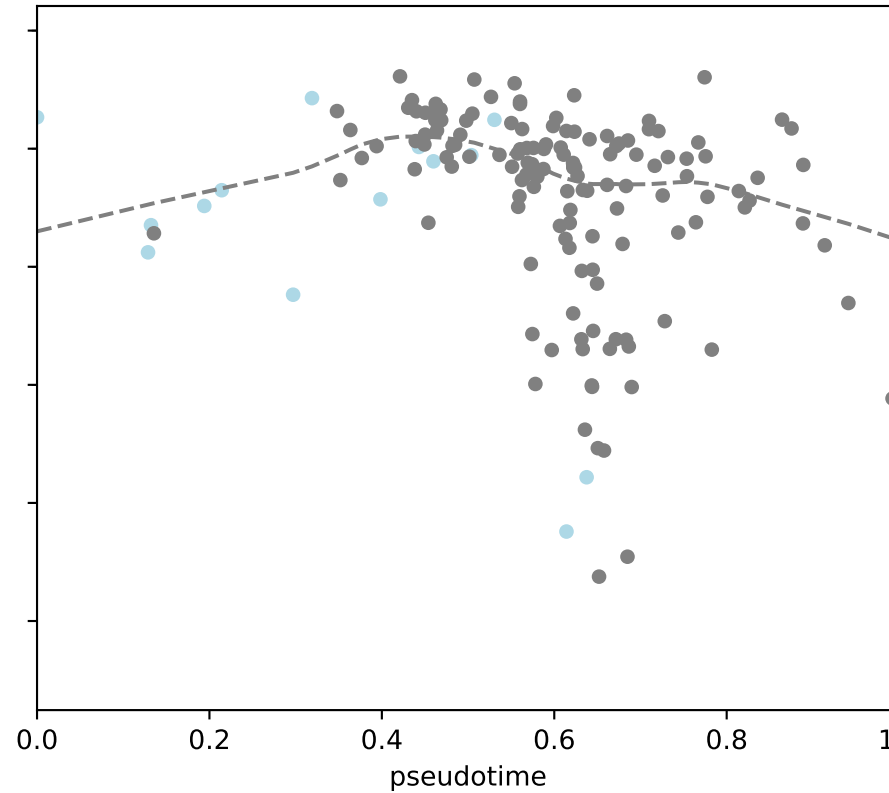
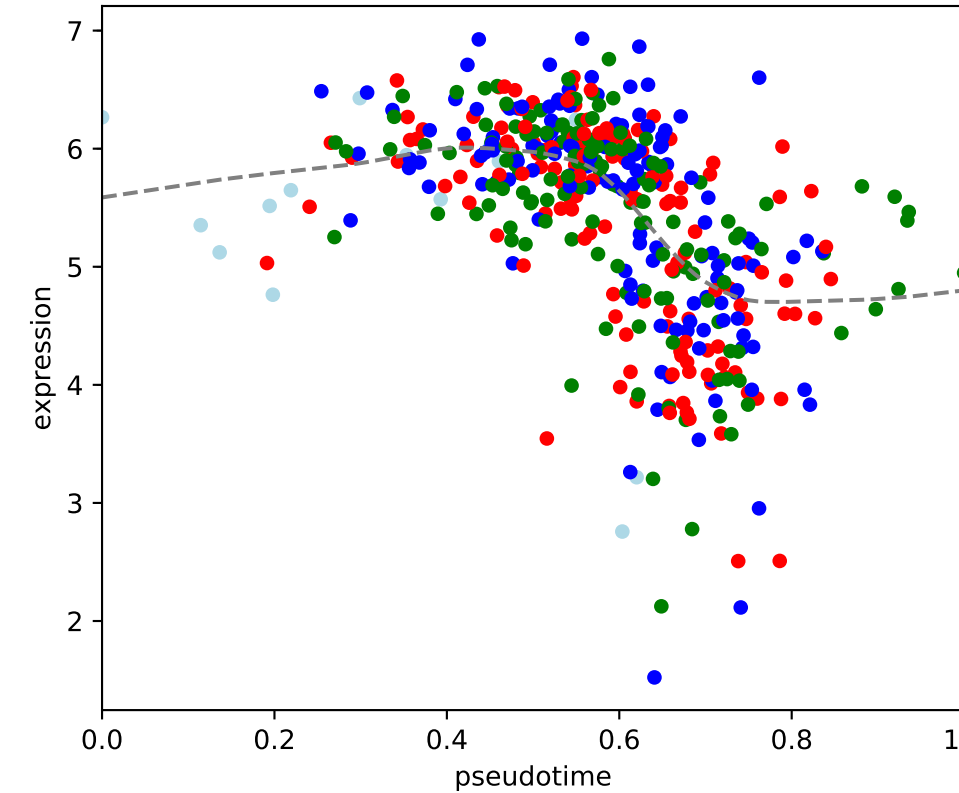
733 1spearman=-0.48



ENST00000520468 COX6C (cytochrome c oxidase subunit 6C)

733 0spearman=-0.60

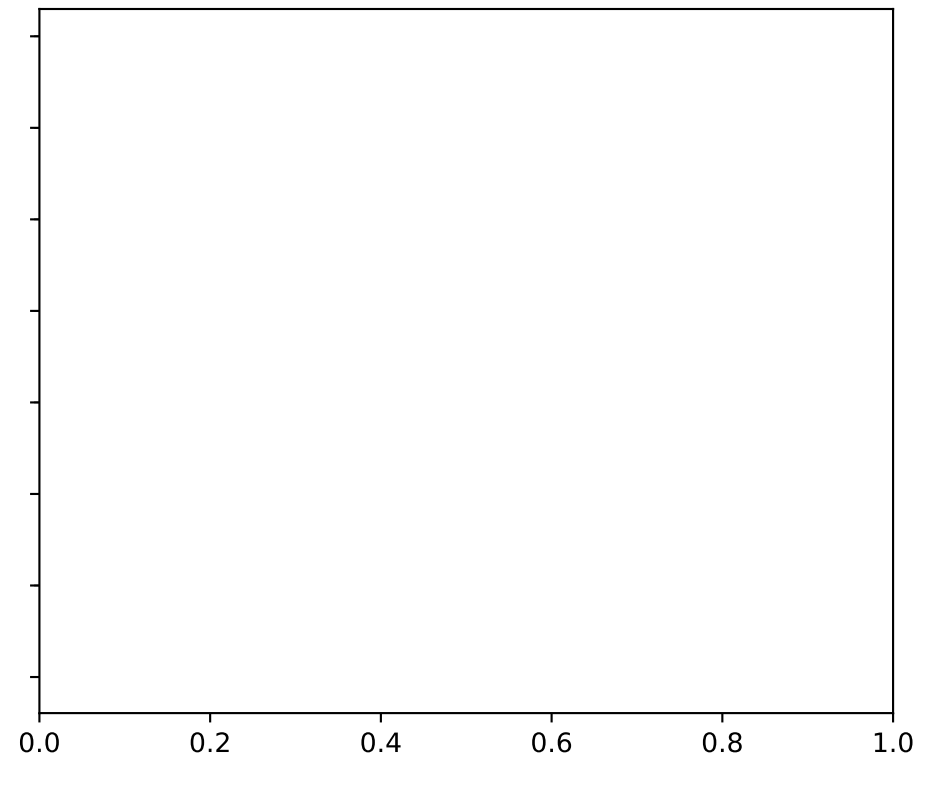
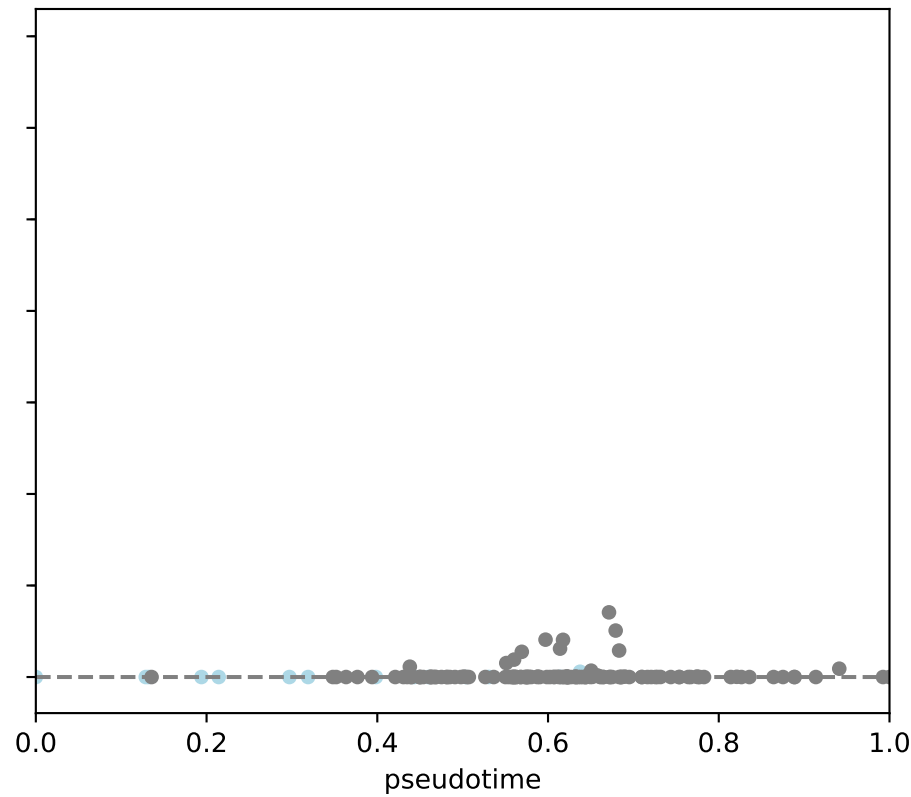
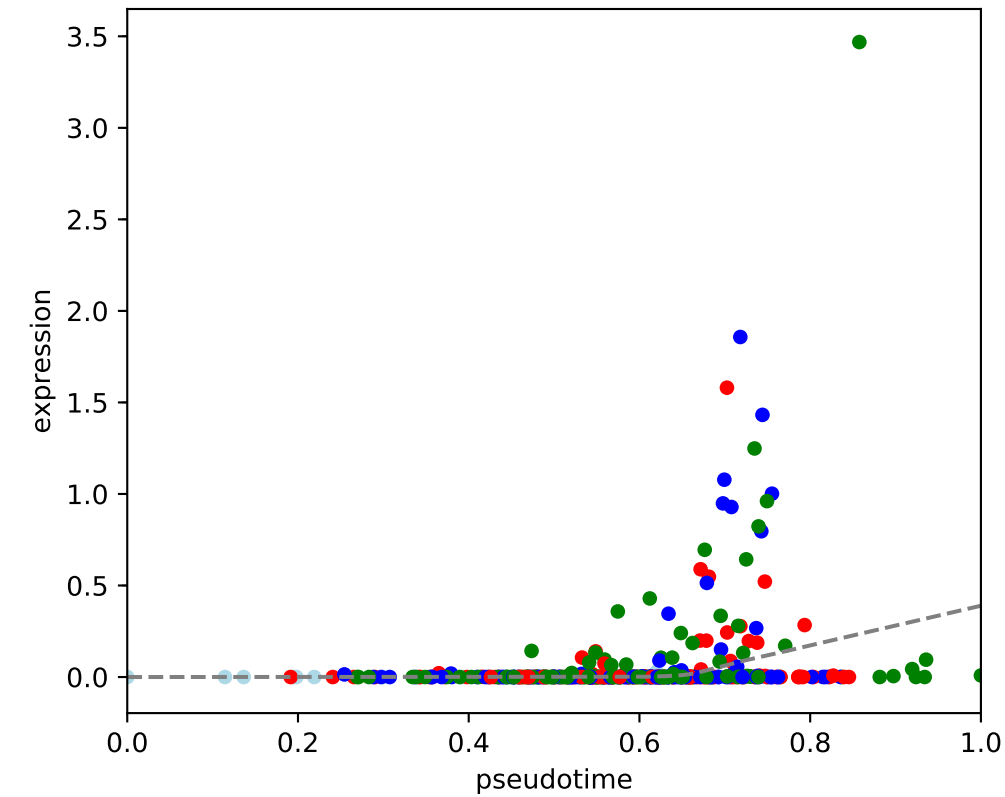
733 1spearman=-0.39



ENST00000600486 MRLN (myoregulin)

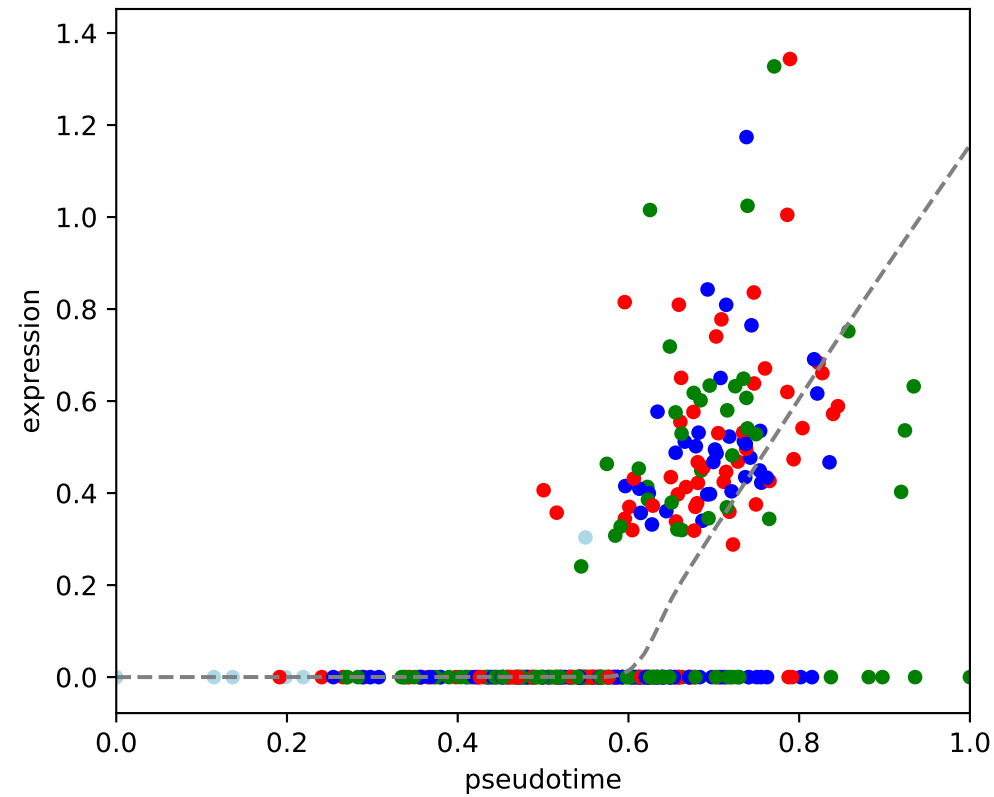
733 0spearman=0.60

733 1spearman=0.22

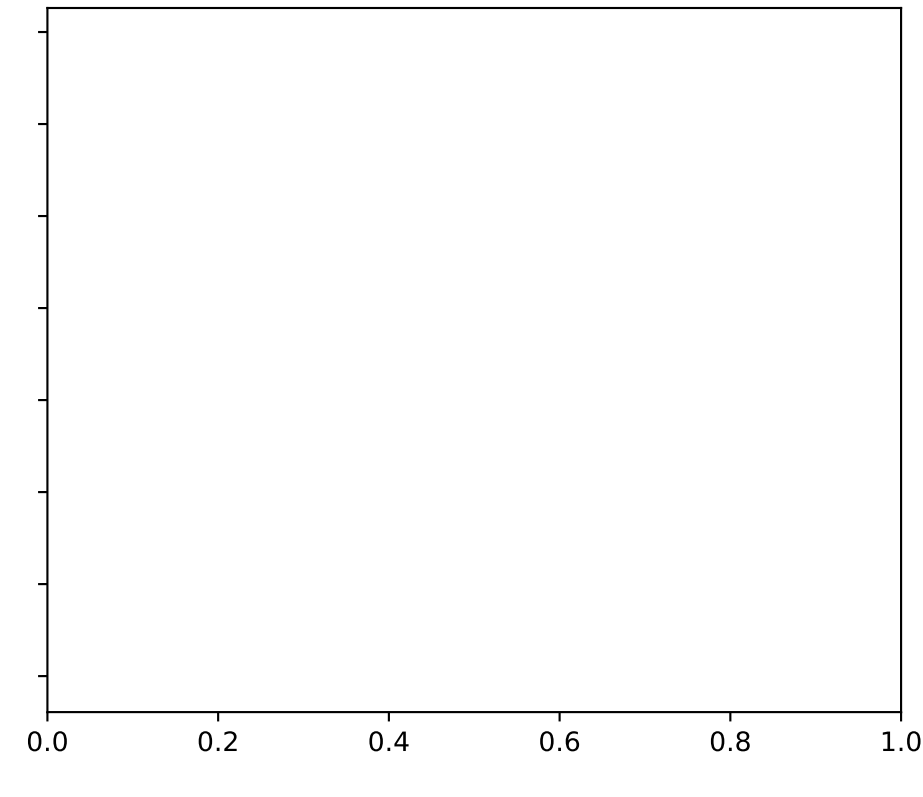
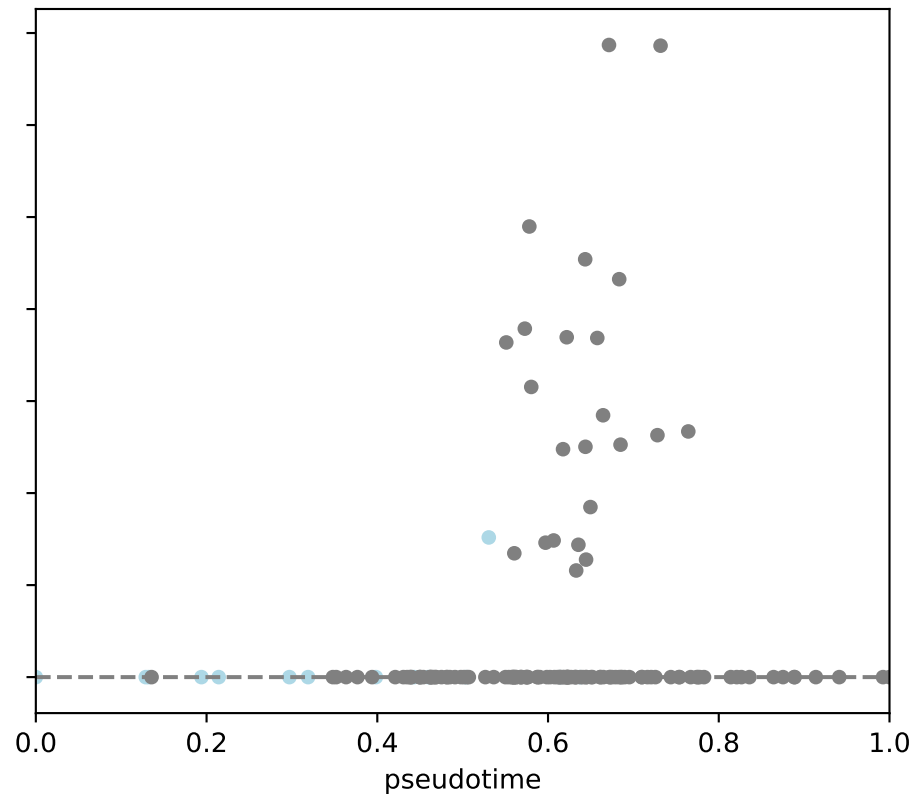


ENST00000599605 MRLN (myoregulin)

733 0spearman=0.60

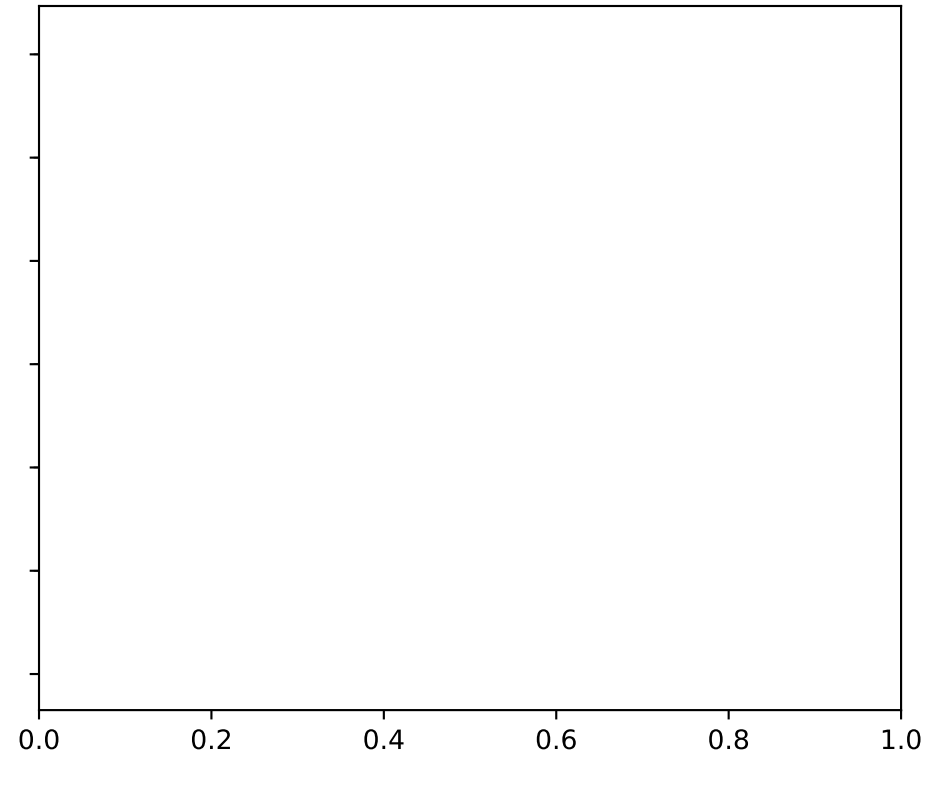
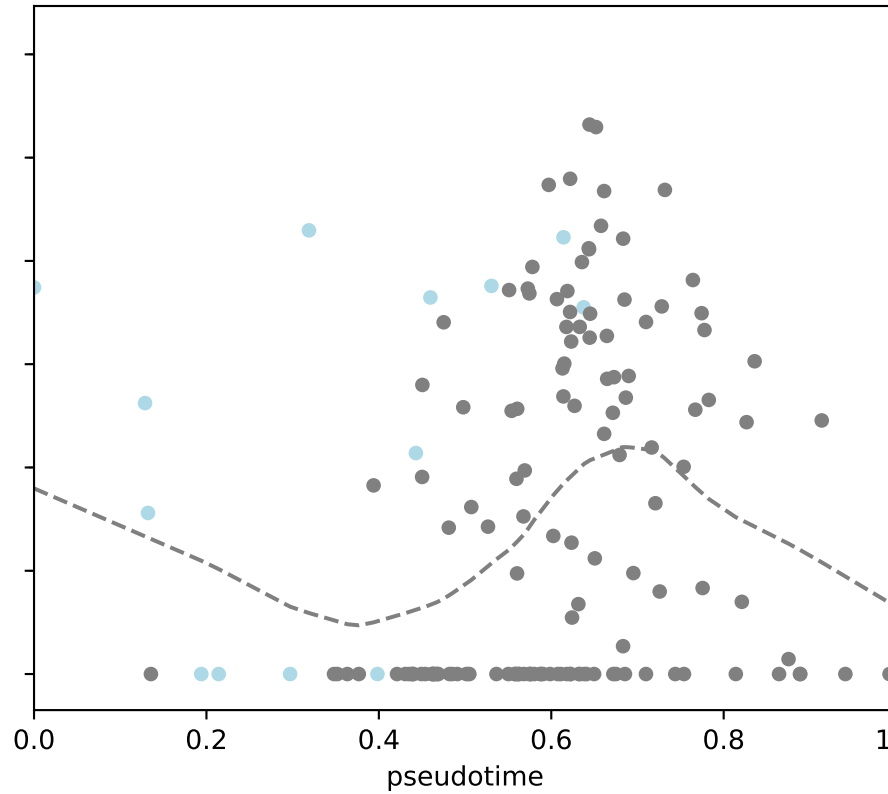
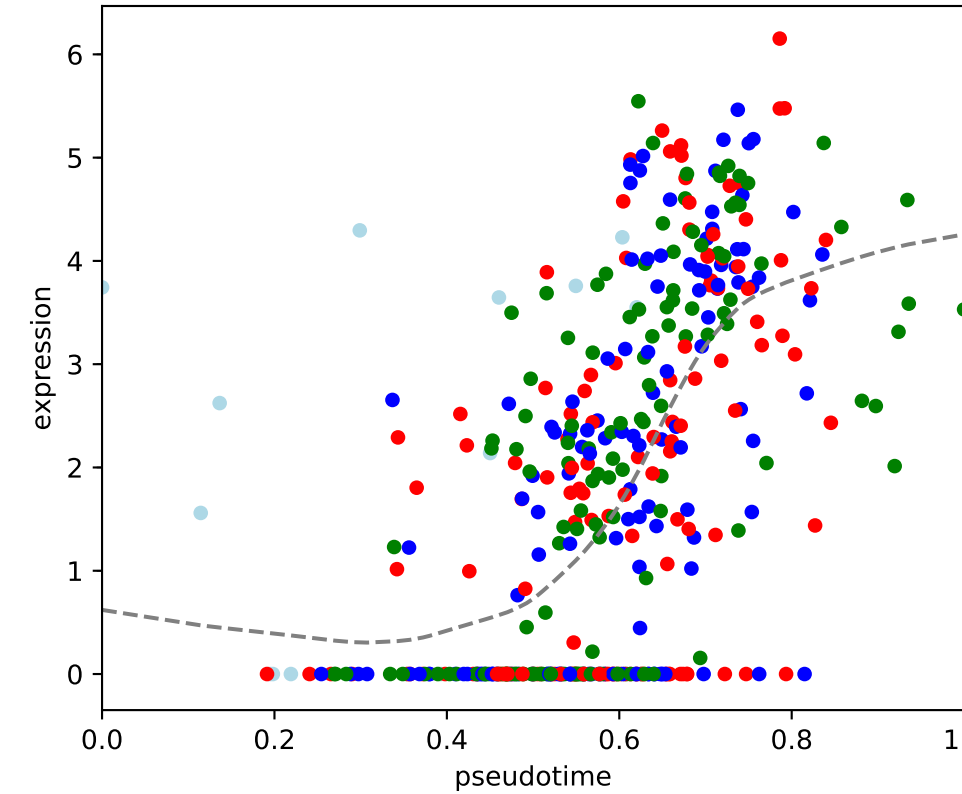


733 1spearman=0.16



733 0spearman=0.60

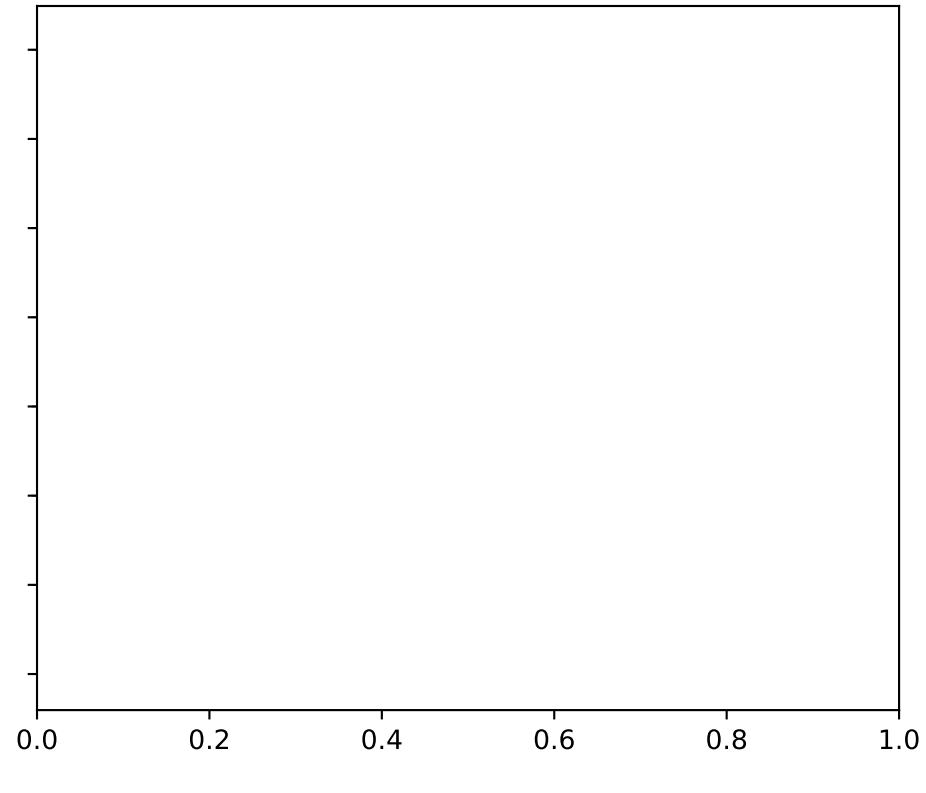
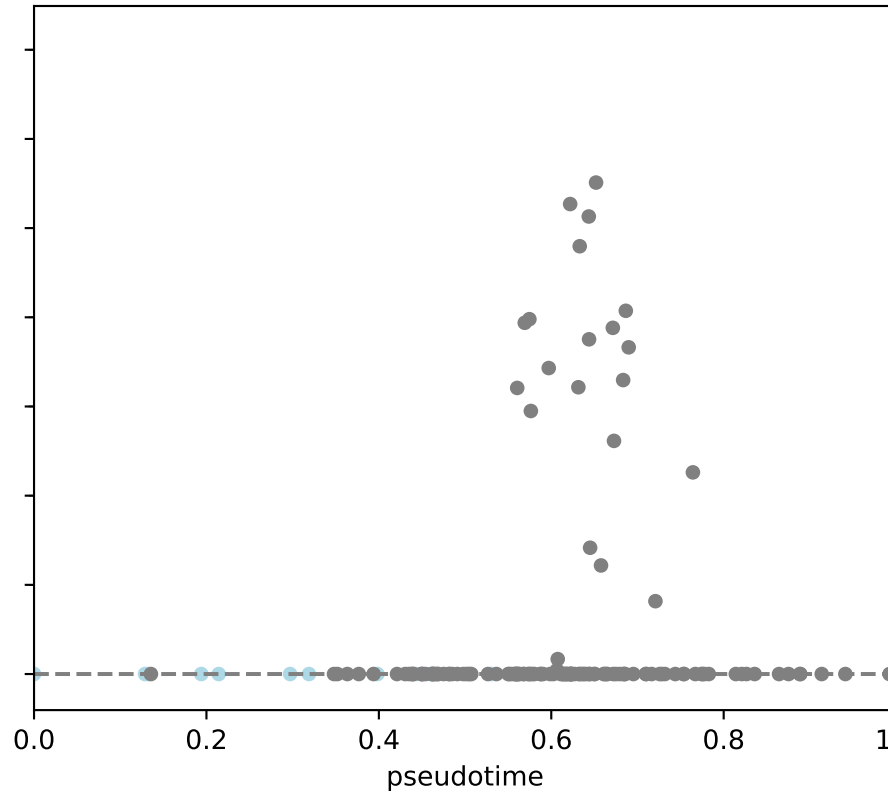
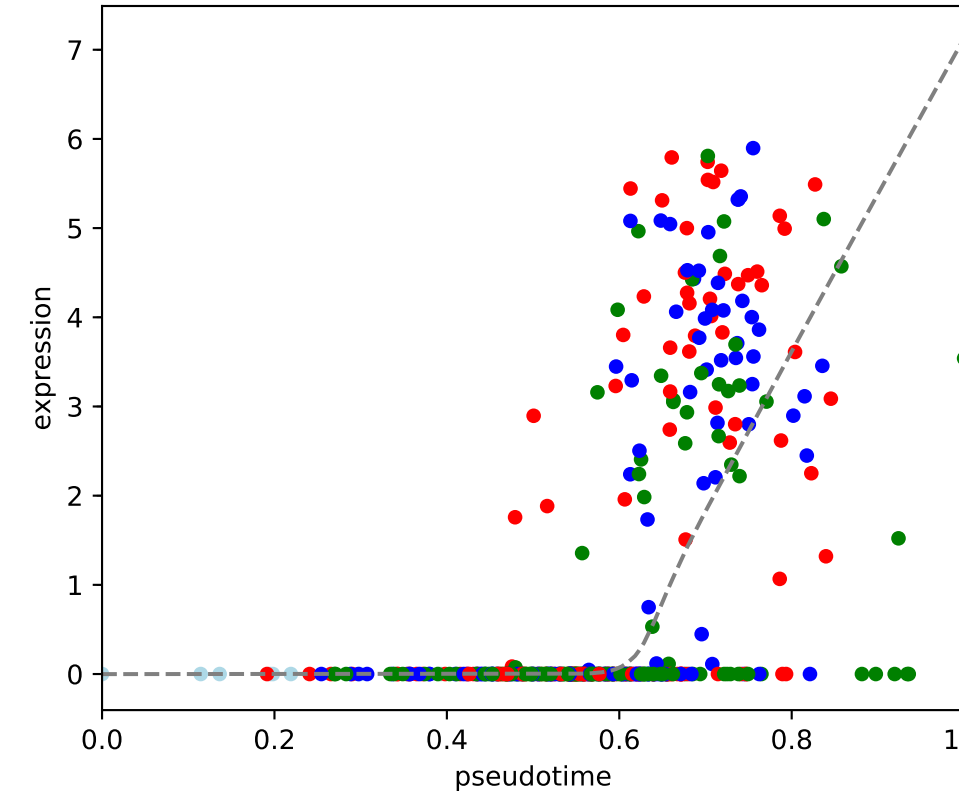
733 1spearman=0.28



ENST00000621172 GABRR3 (gamma-aminobutyric acid type A receptor rho3 subunit (gene/pseudogene))

733 0spearman=0.60

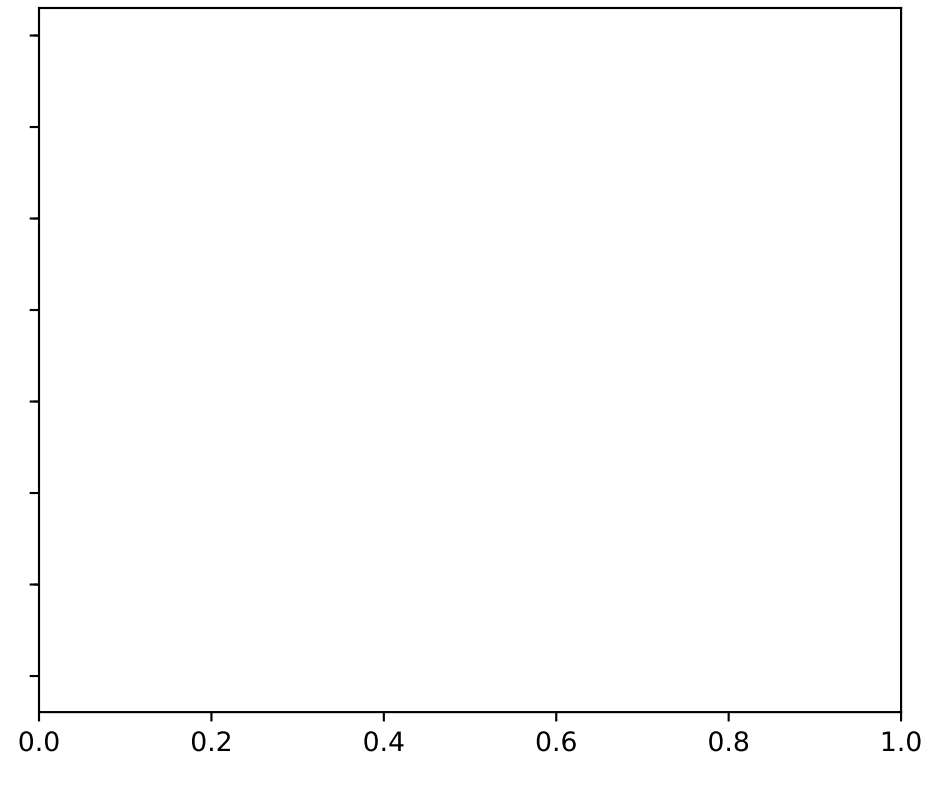
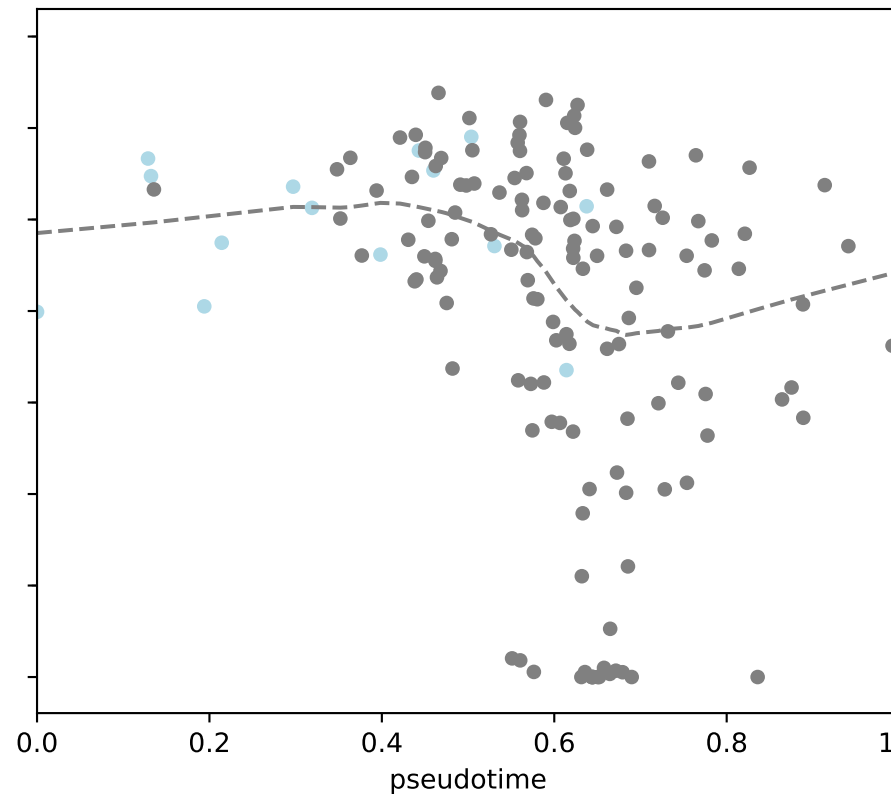
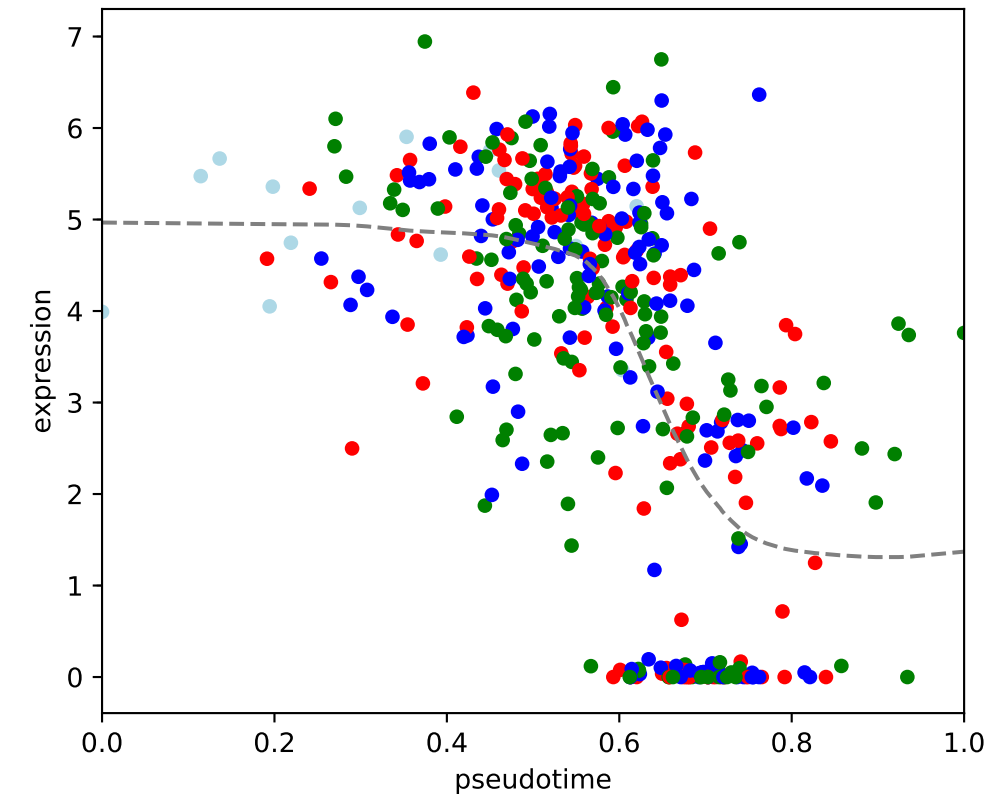
733 1spearman=0.18



ENST00000327720 SIX6 (SIX homeobox 6)

733 0spearman=-0.60

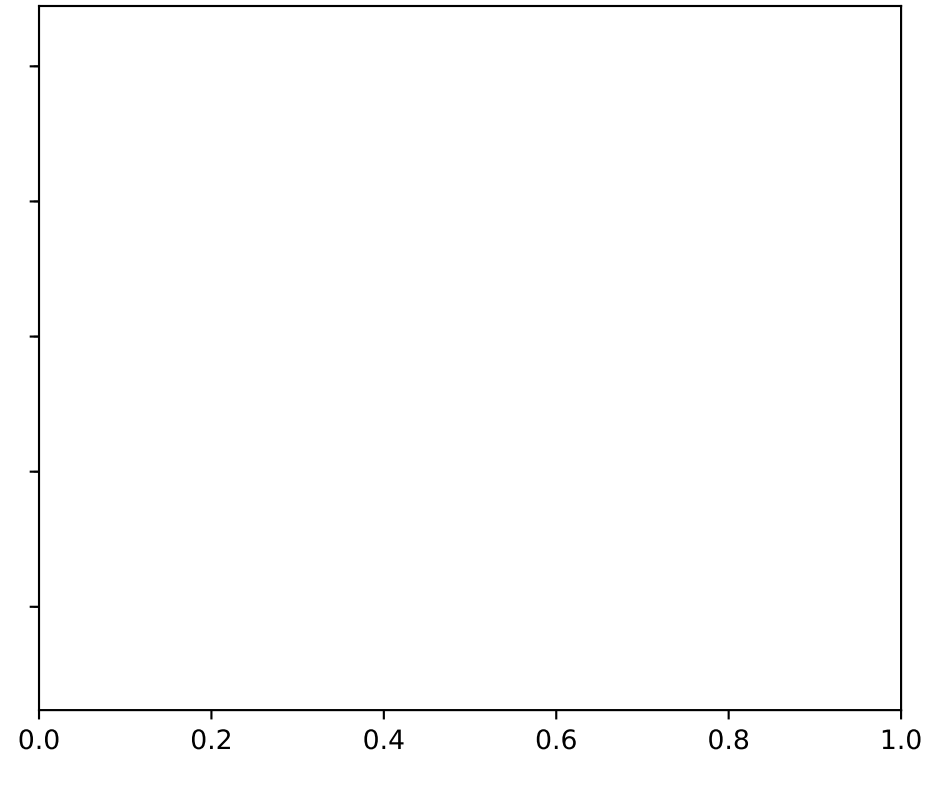
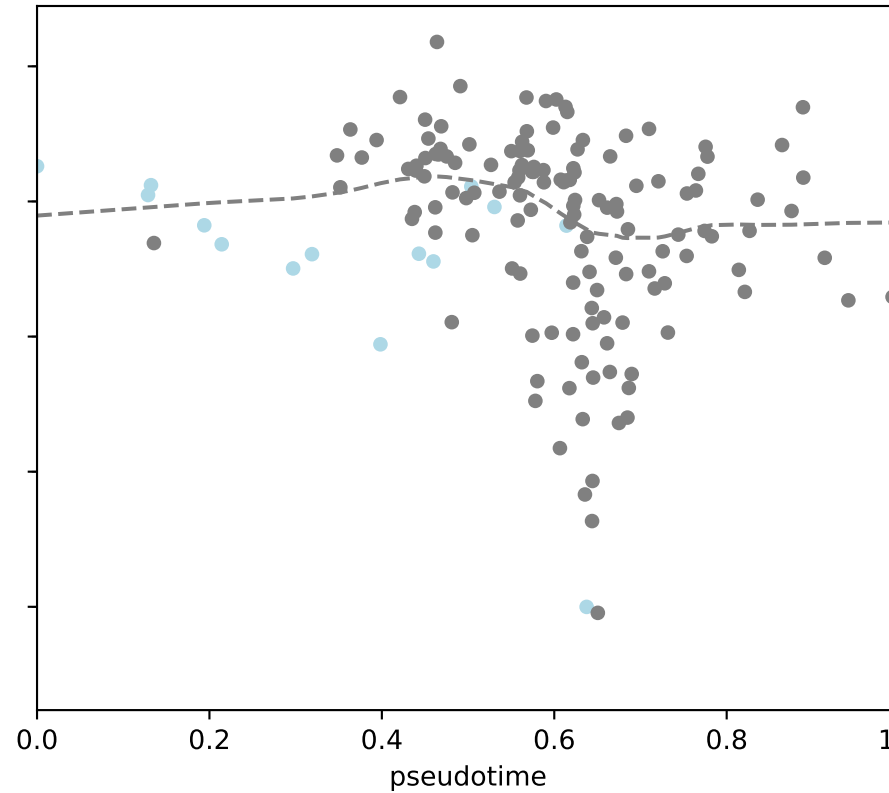
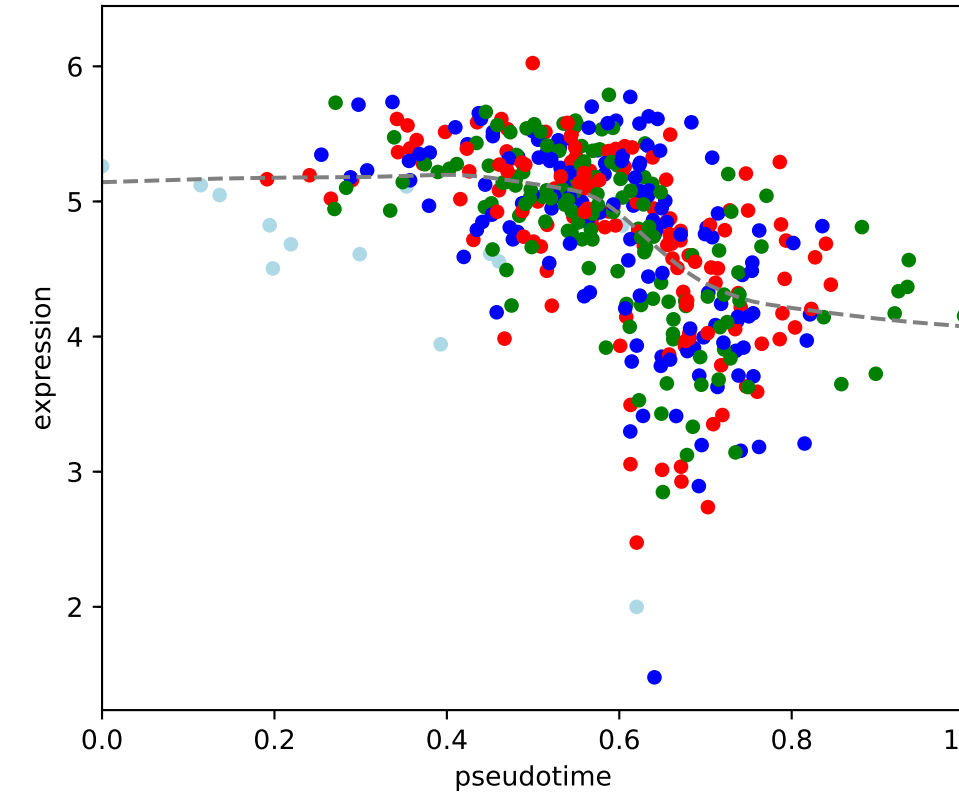
733 1spearman=-0.41



ENST00000472311 COX7A2 (cytochrome c oxidase subunit 7A2)

733 0spearman=-0.59

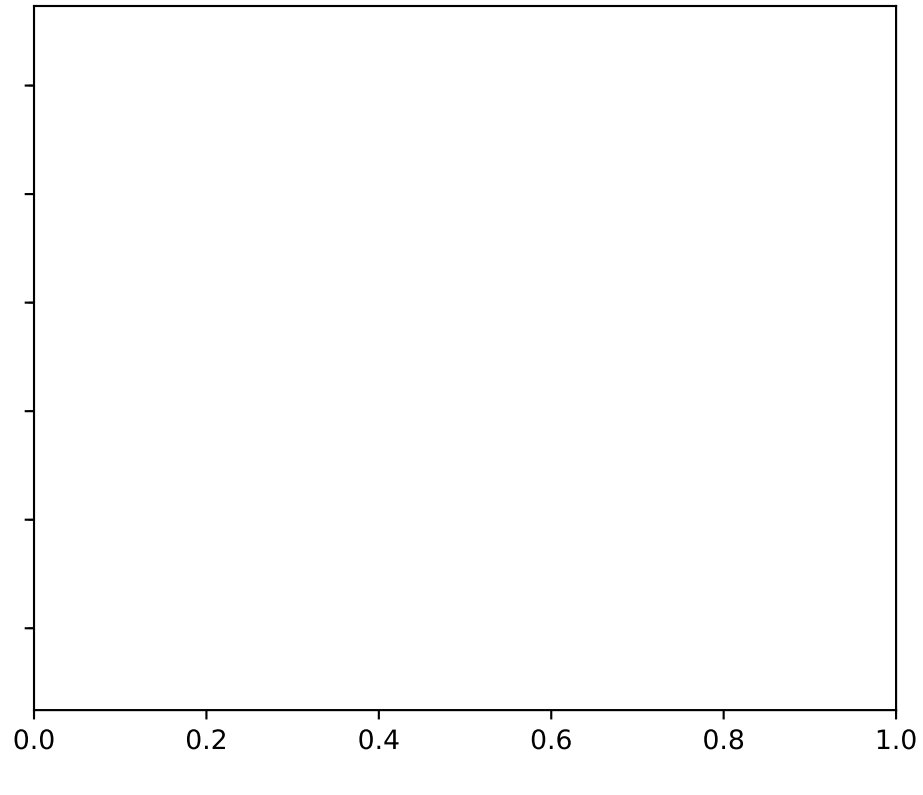
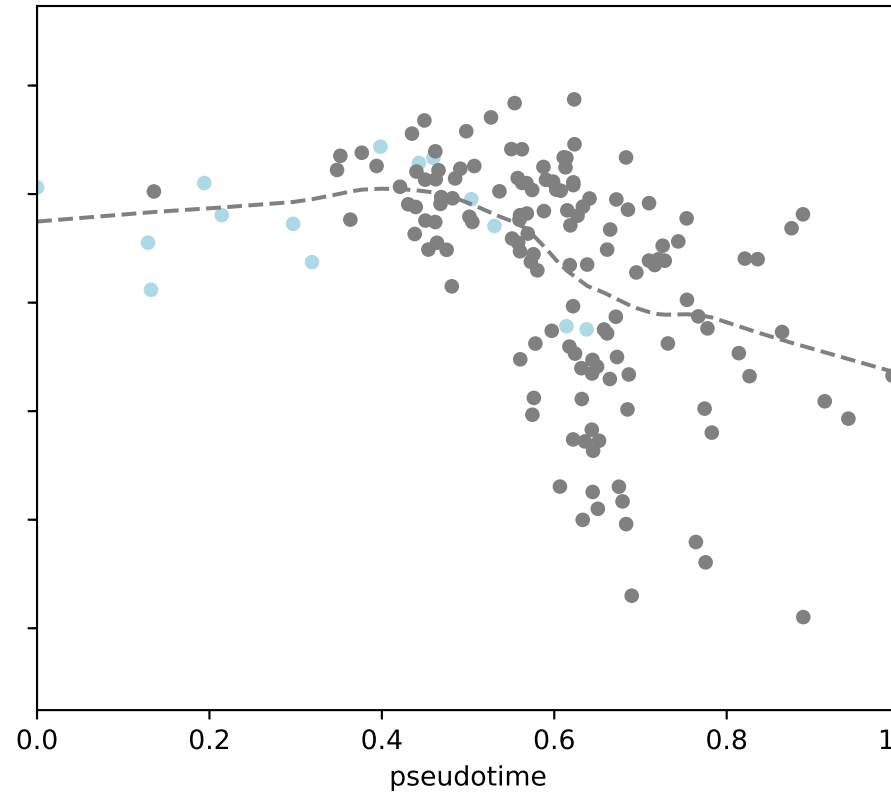
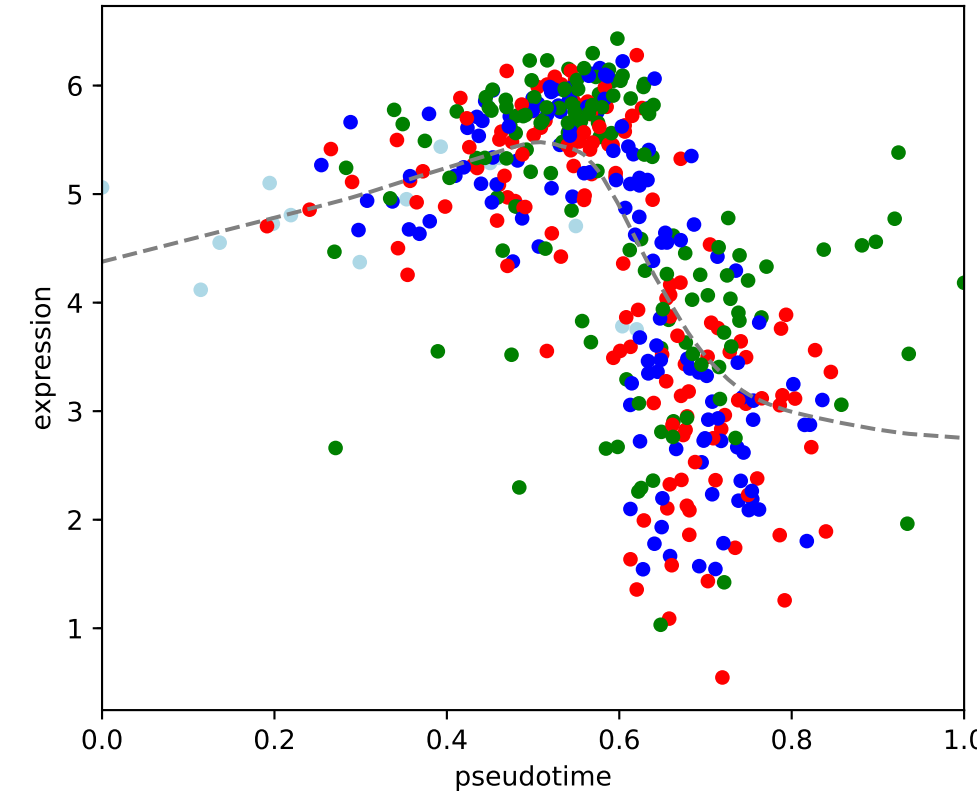
733 1spearman=-0.32



ENST00000301072 TUBA1C (tubulin alpha 1c)

733 0spearman=-0.59

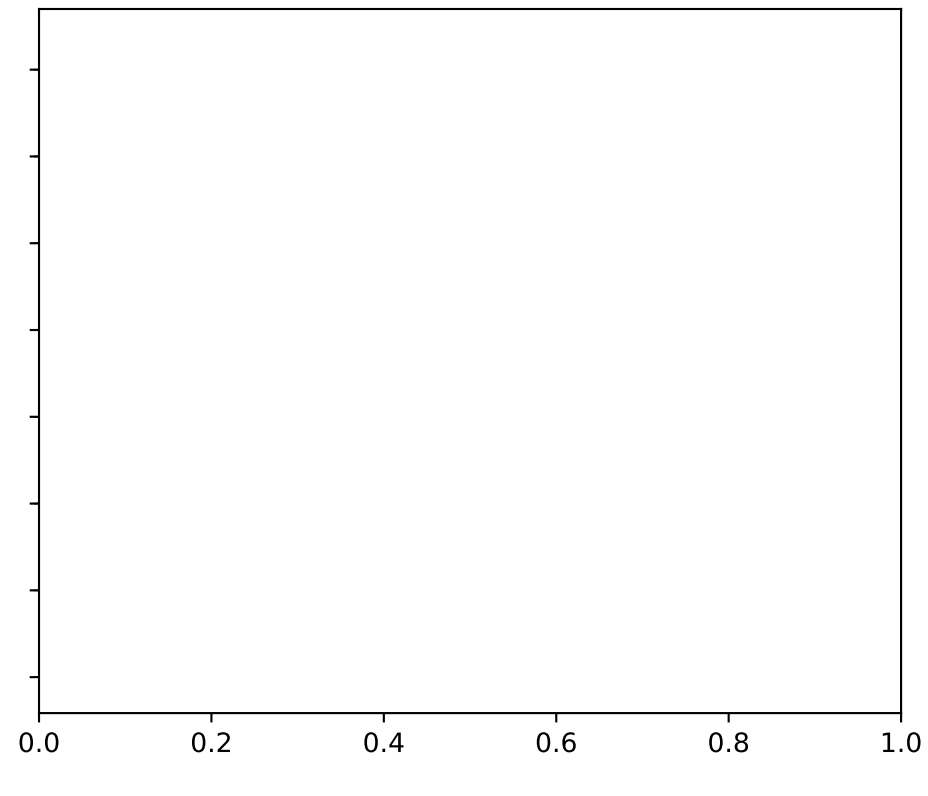
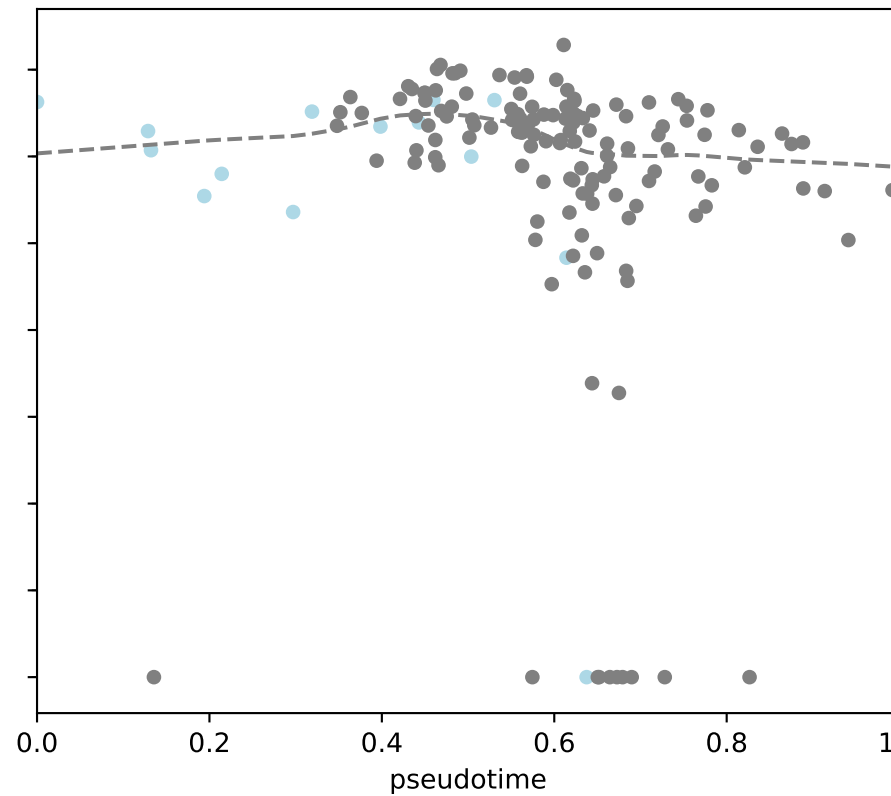
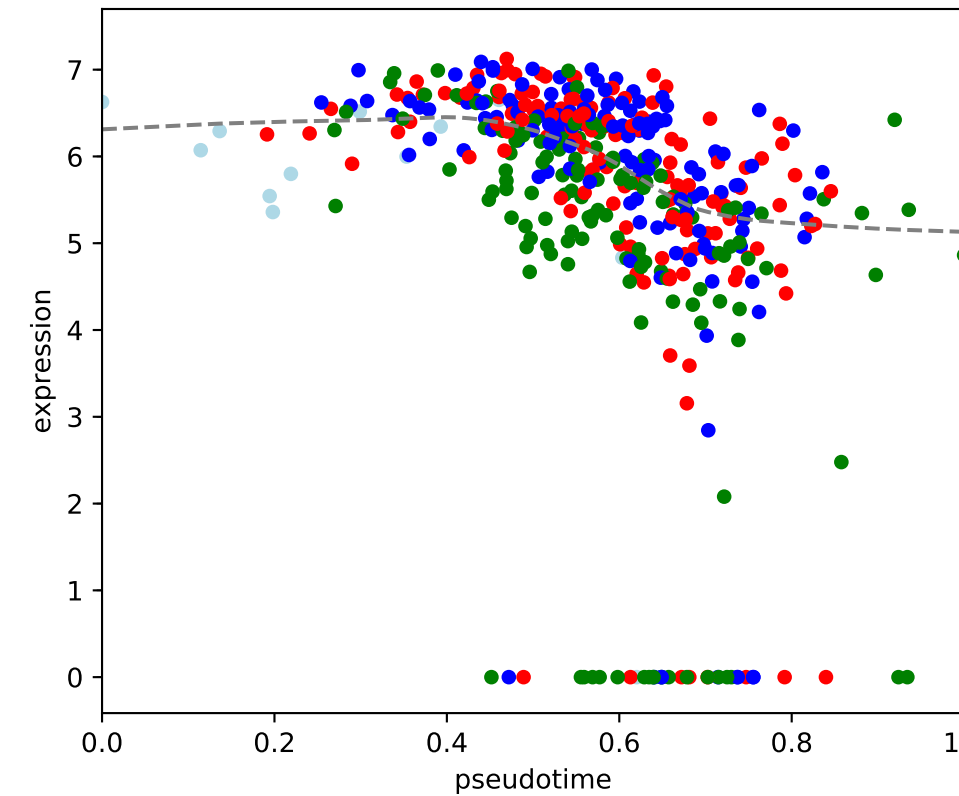
733 1spearman=-0.59



ENST00000606722 NDUFA13 (NADH:ubiquinone oxidoreductase subunit A13)

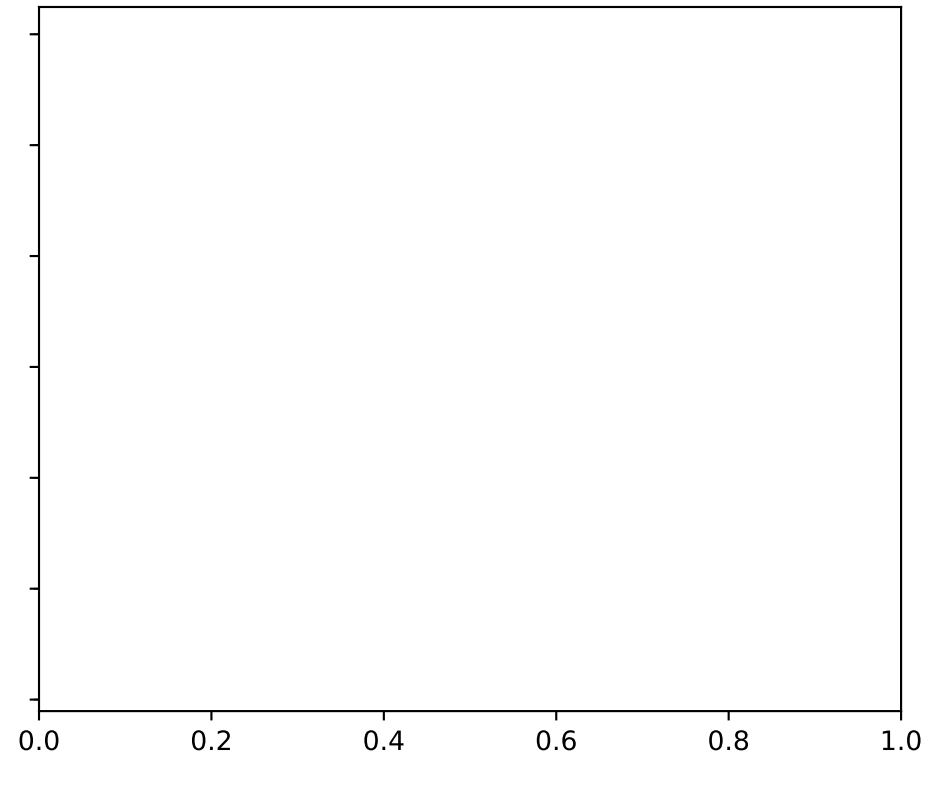
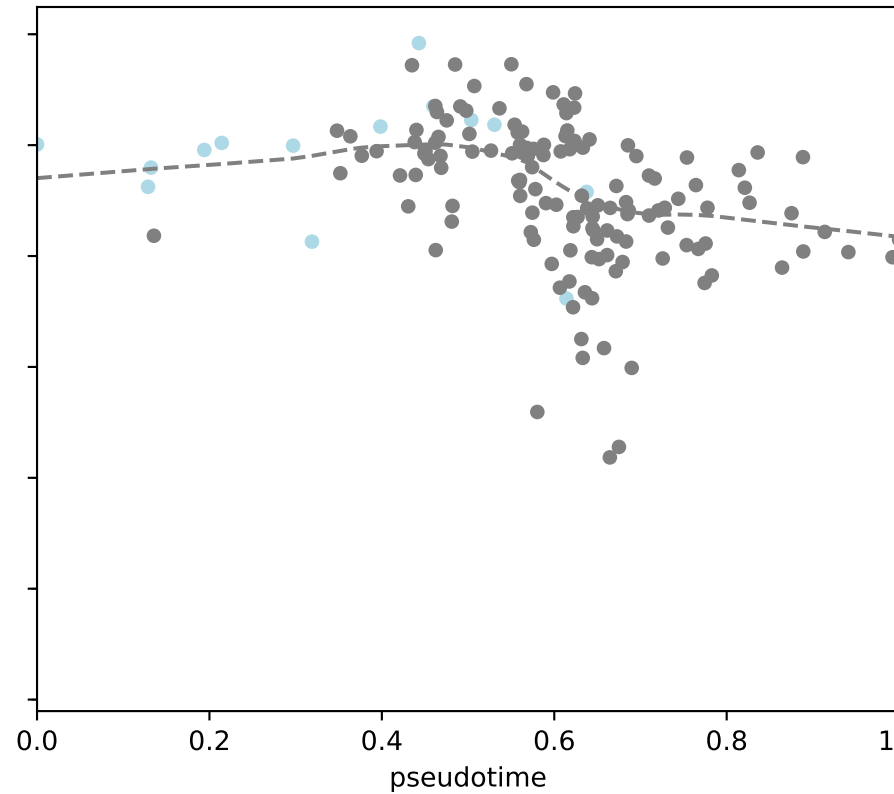
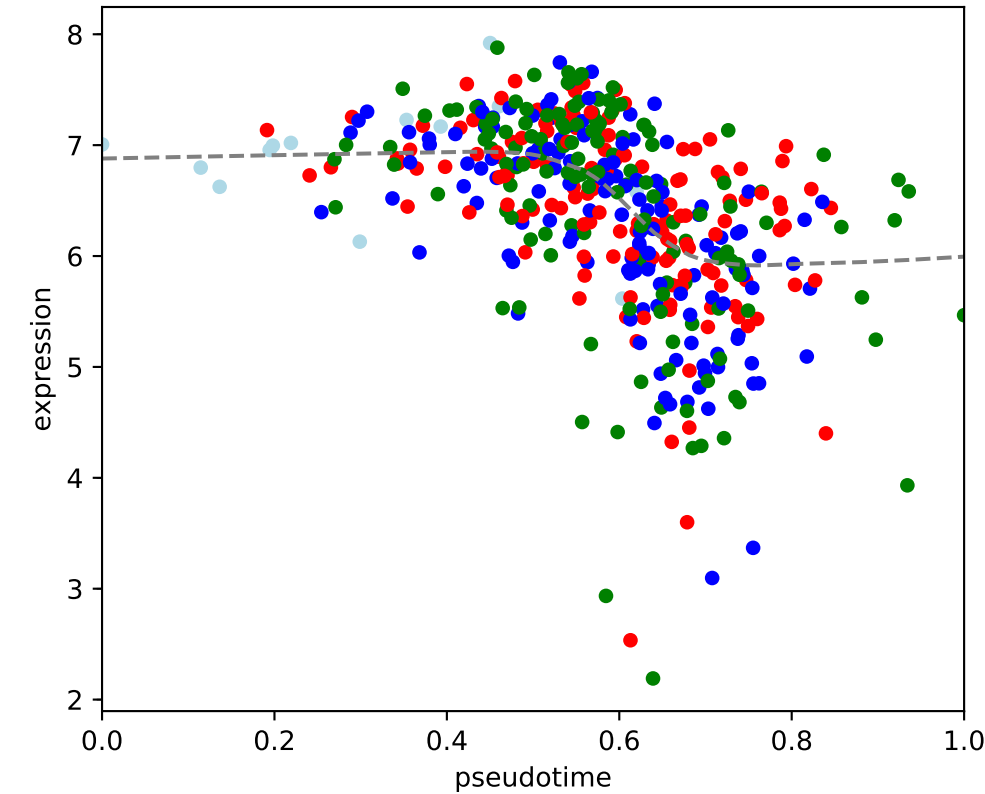
733 0spearman=-0.59

733 1spearman=-0.42



733 0spearman=-0.59

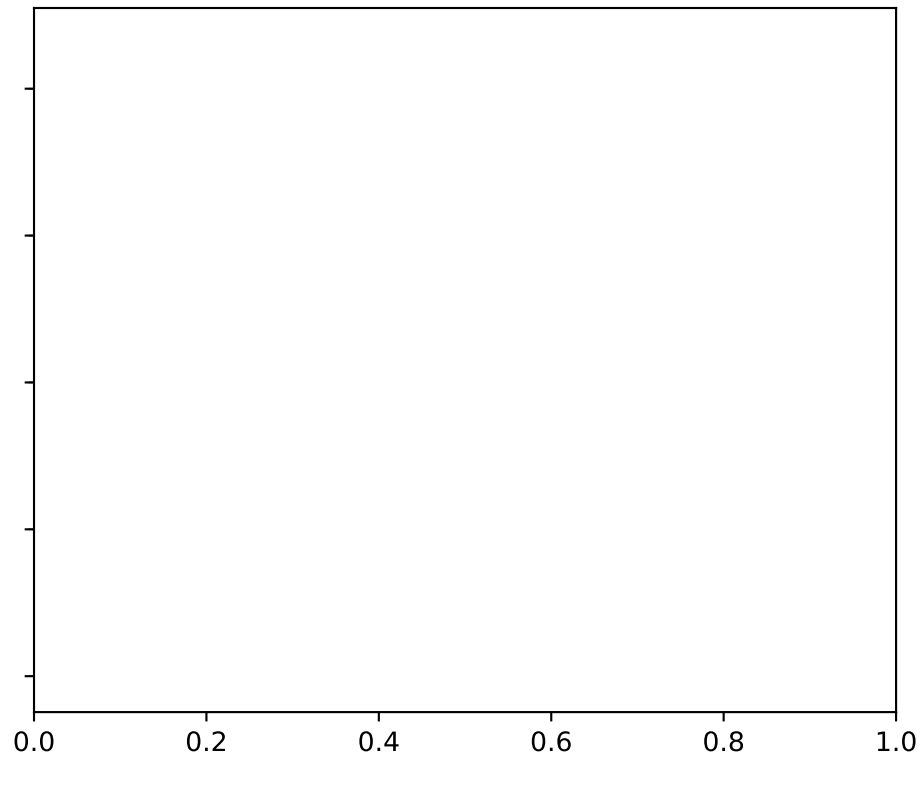
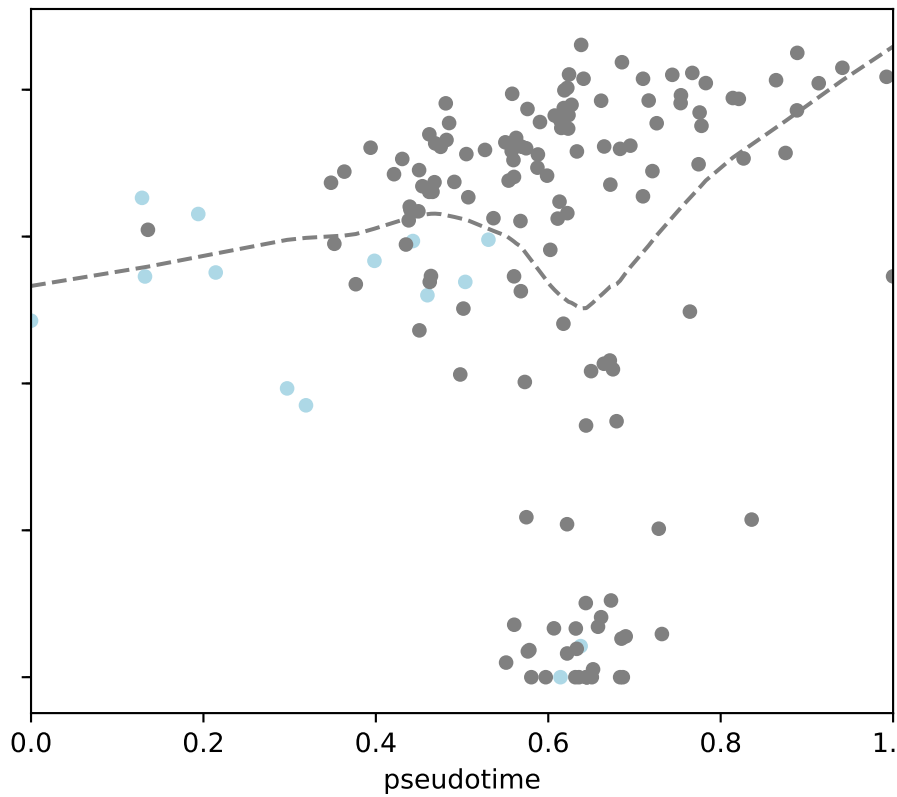
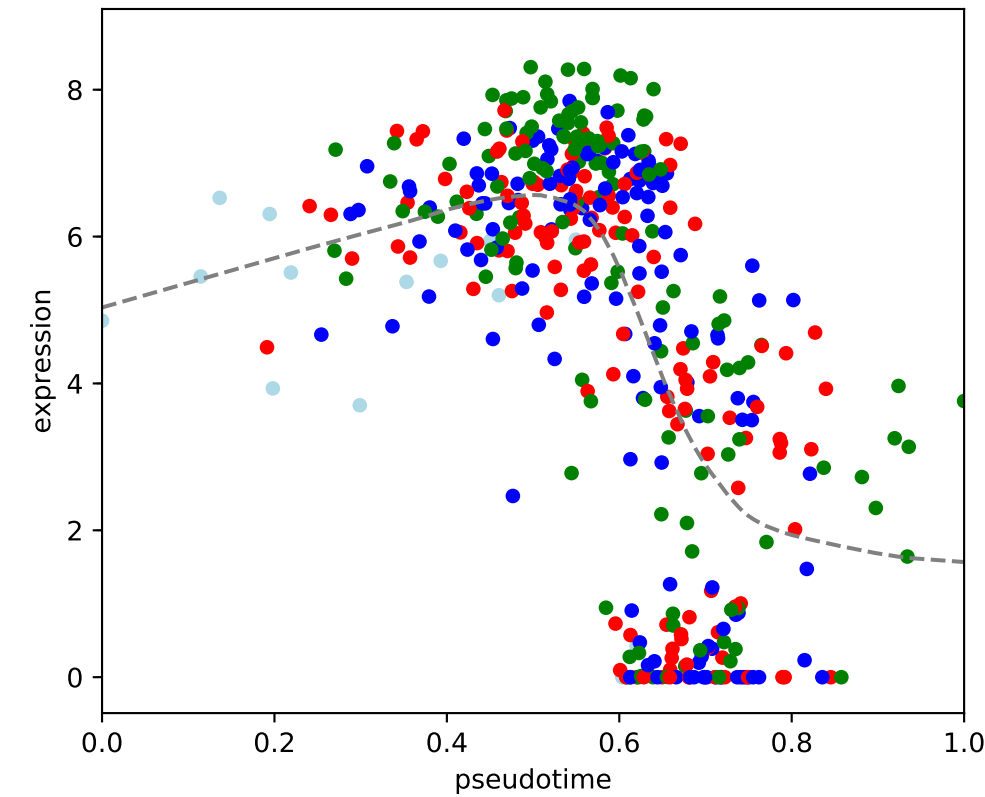
733 1spearman=-0.53



ENST00000277415 OLFM1 (olfactomedin 1)

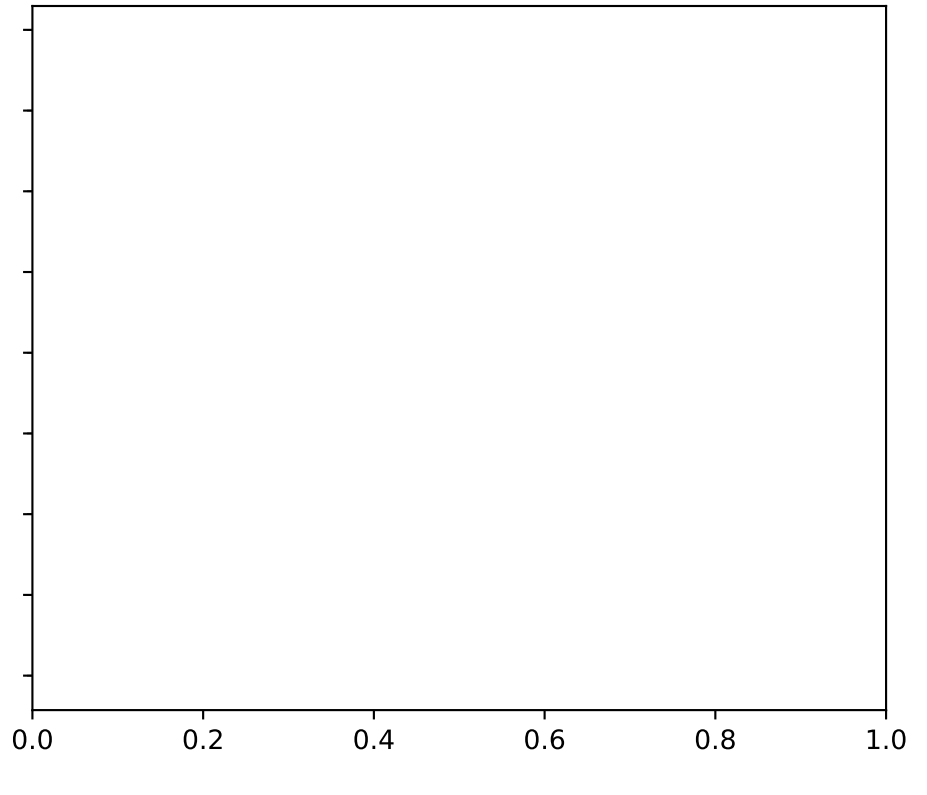
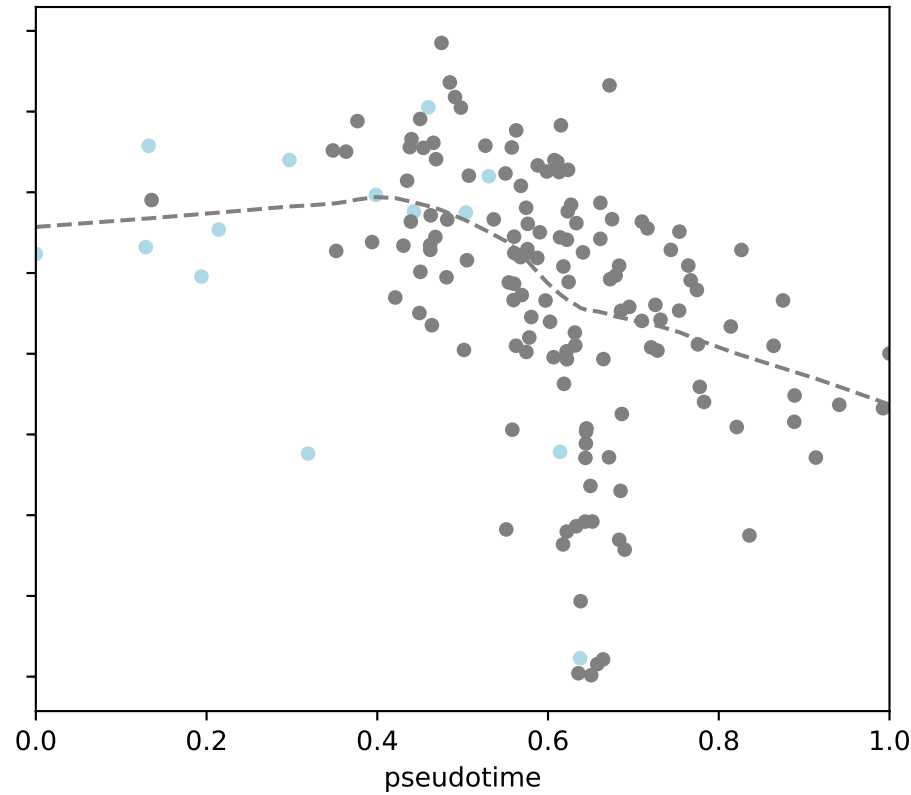
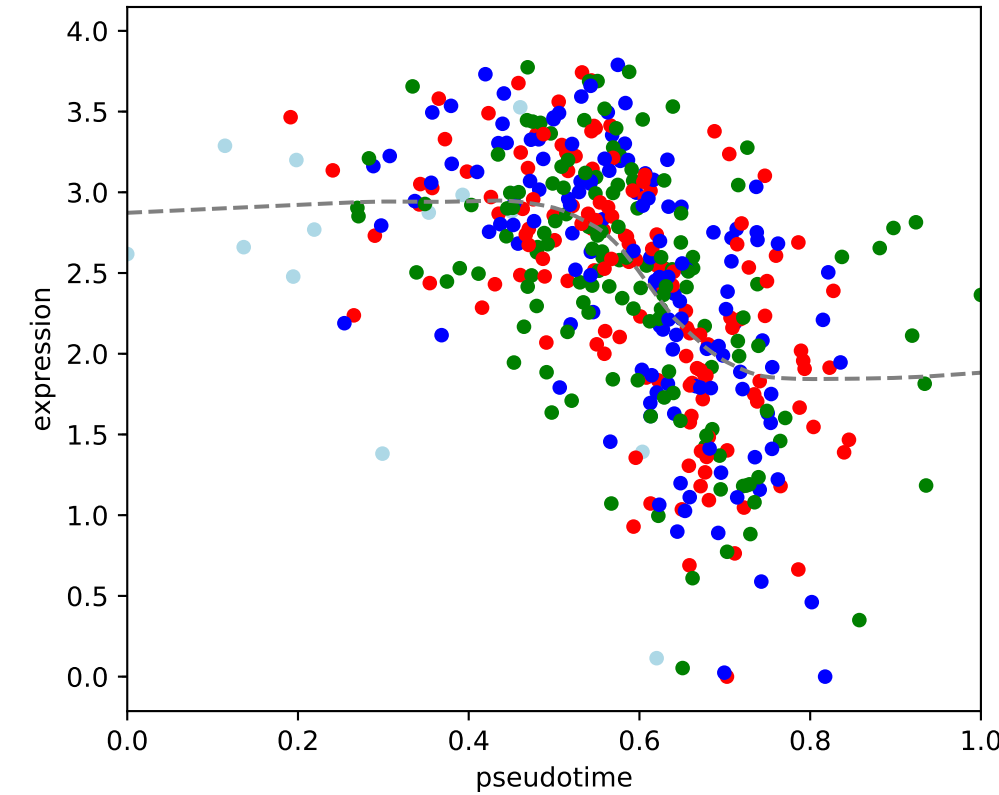
733 0spearman=-0.59

733 1spearman=0.16



733 0spearman=-0.59

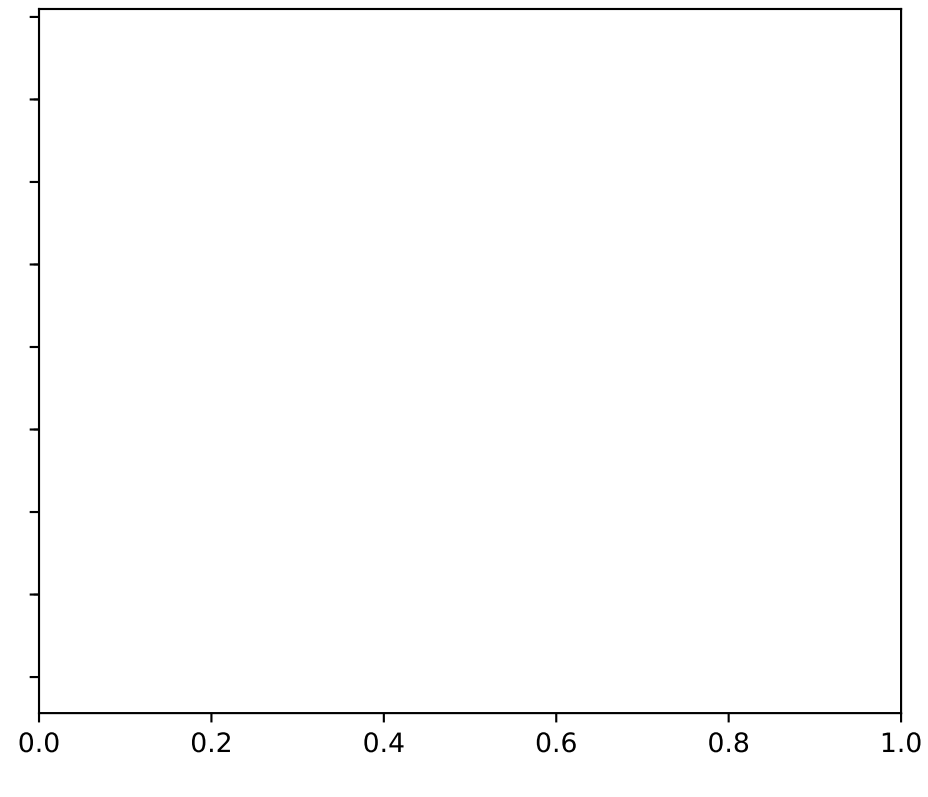
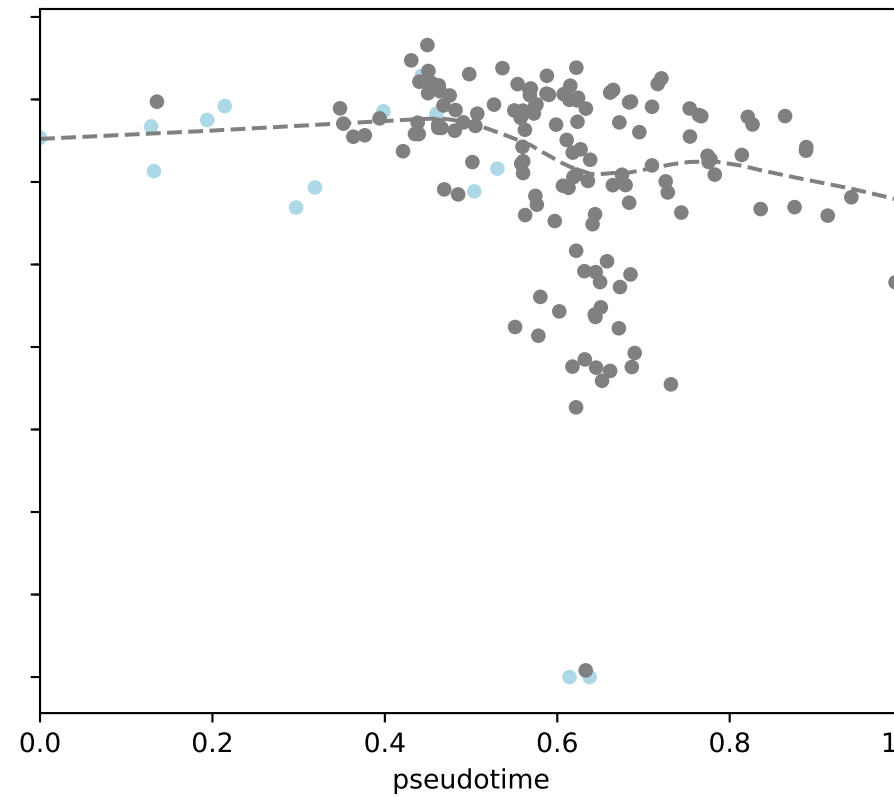
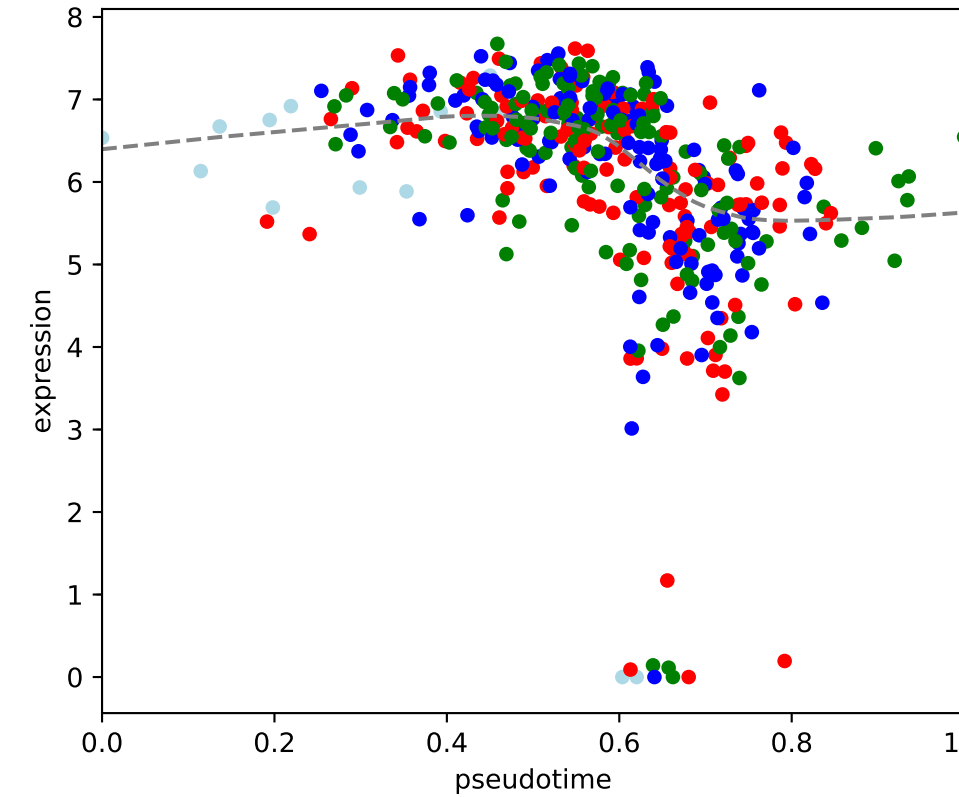
733 1spearman=-0.54



ENST00000225698 C1QBP (complement C1q binding protein)

733 0spearman=-0.59

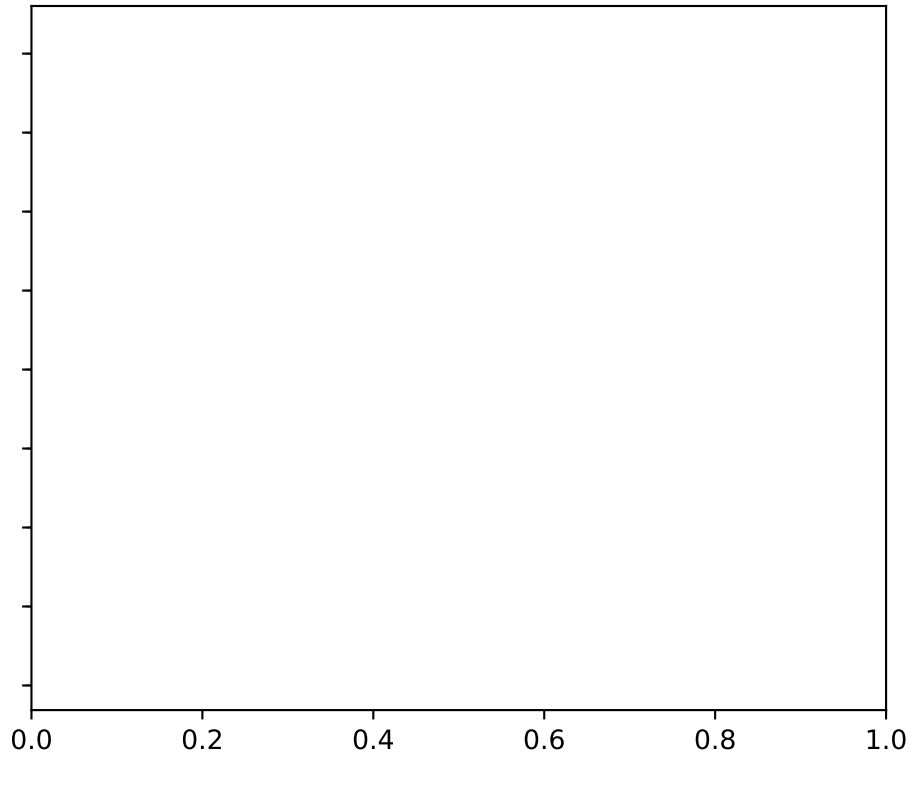
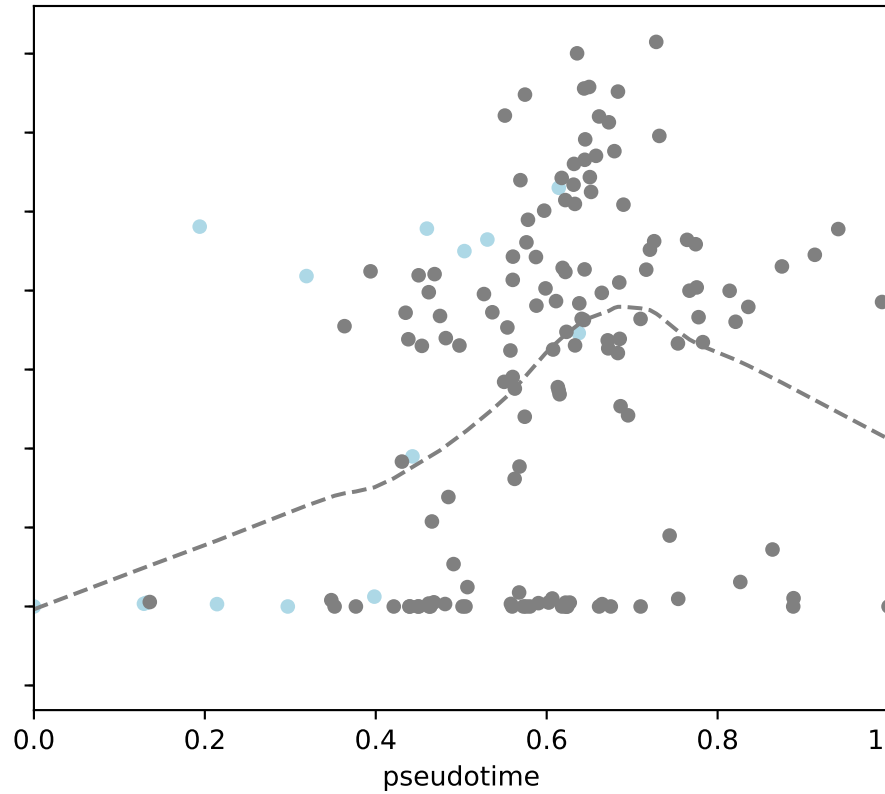
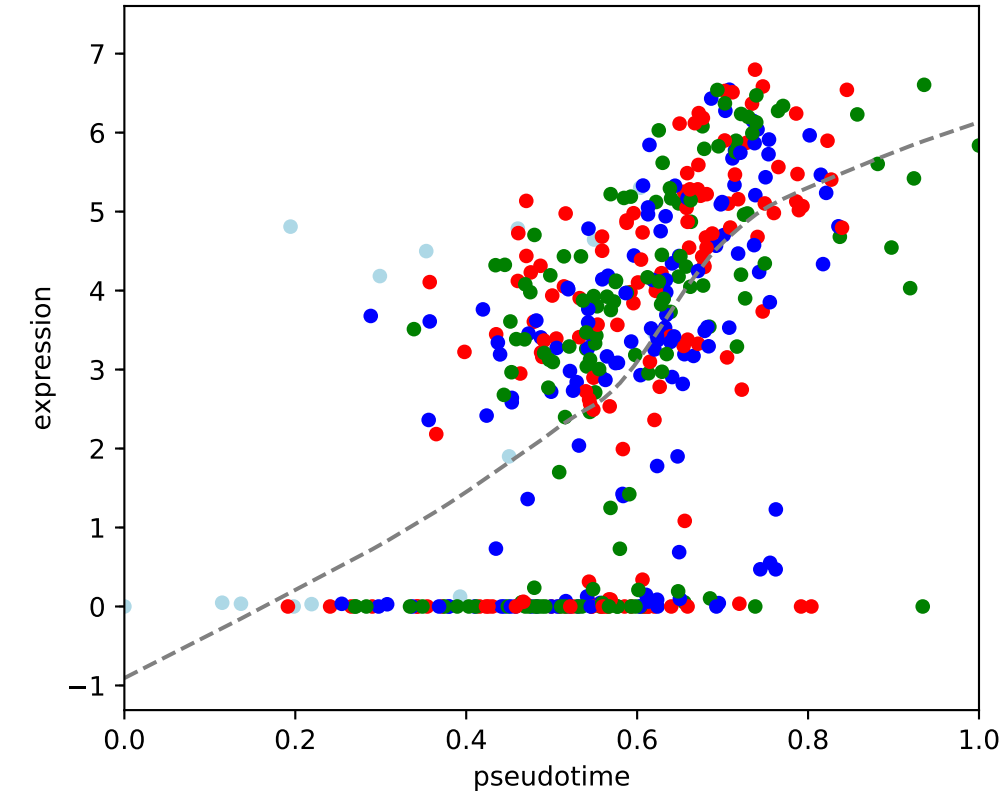
733 1spearman=-0.36



ENST00000355295 TDRD7 (tudor domain containing 7)

733 0spearman=0.59

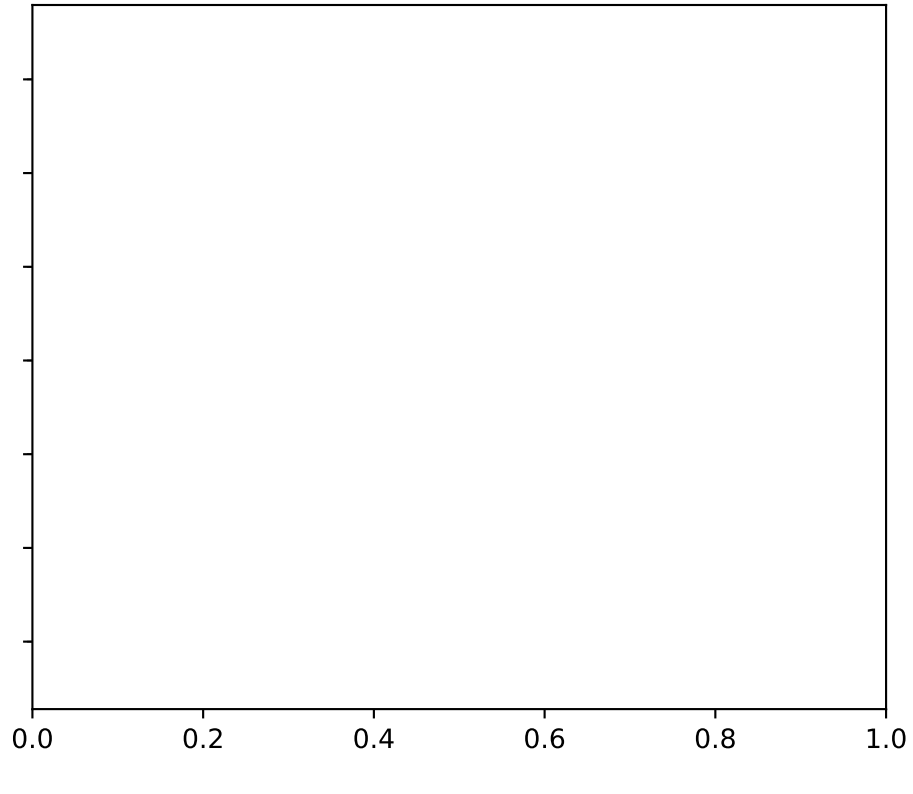
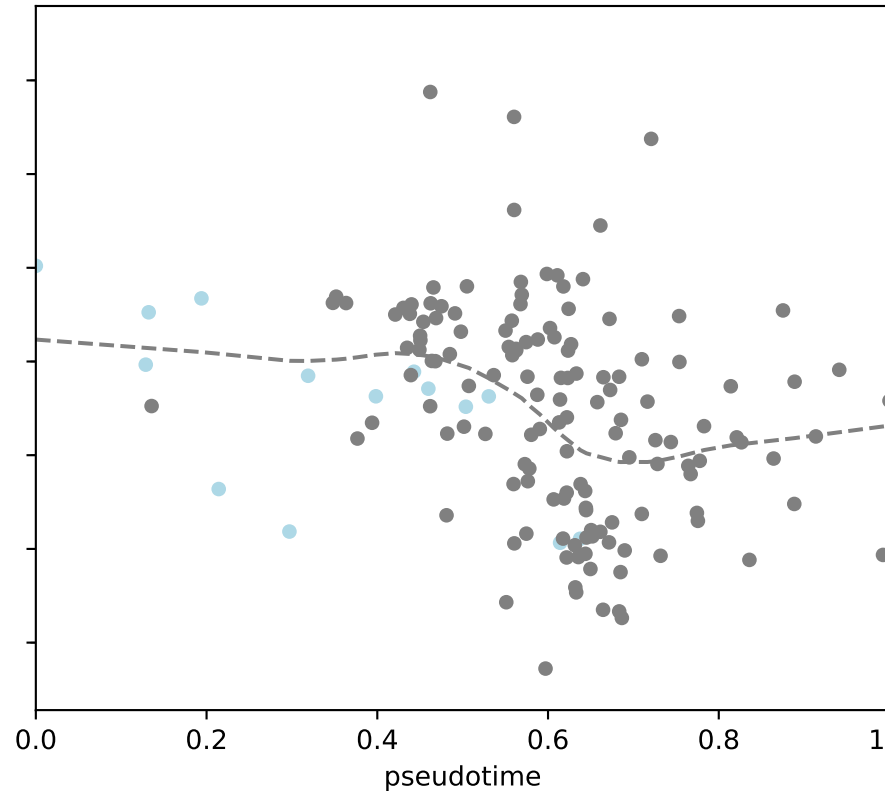
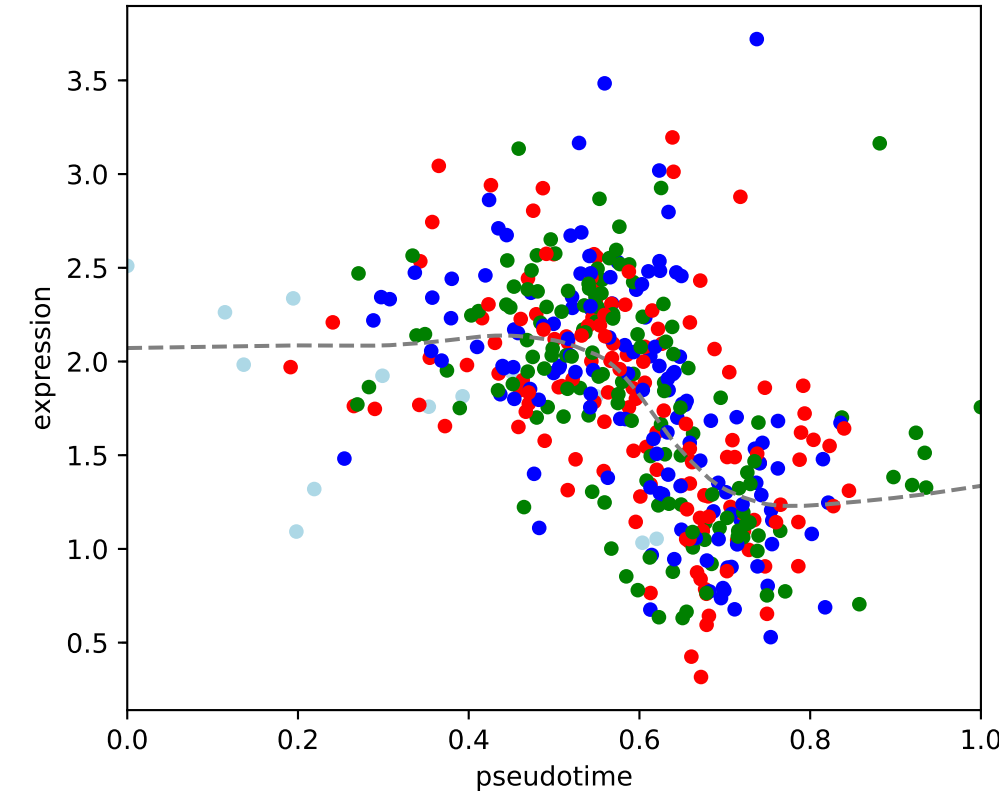
733 1spearman=0.32



ENST00000551570 ATP5F1B (ATP synthase F1 subunit beta)

733 0spearman=-0.59

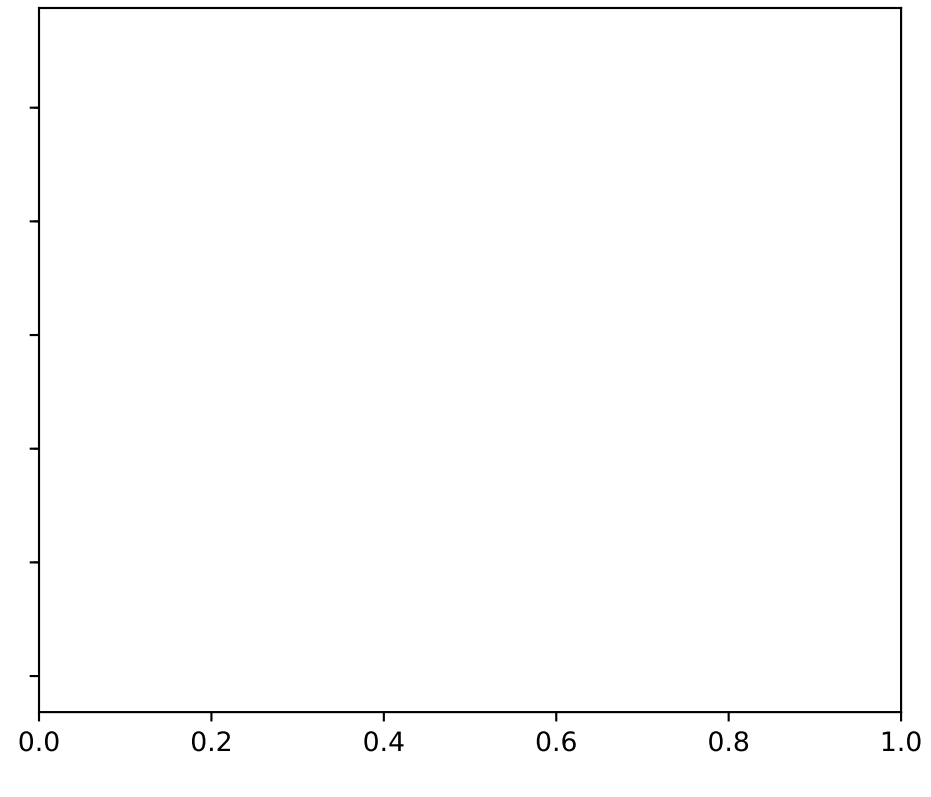
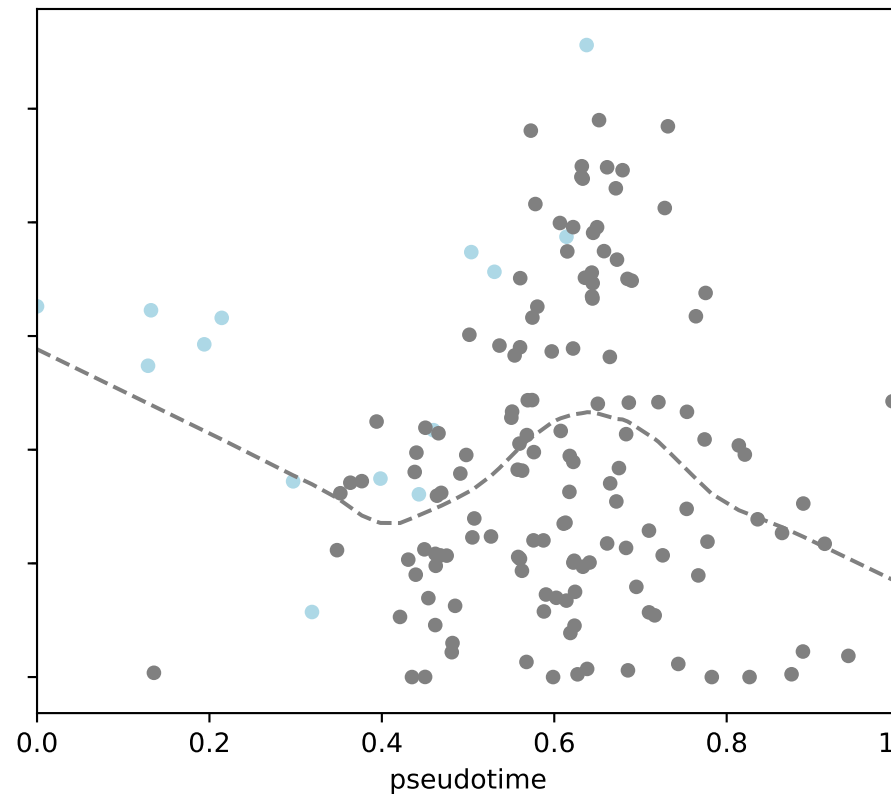
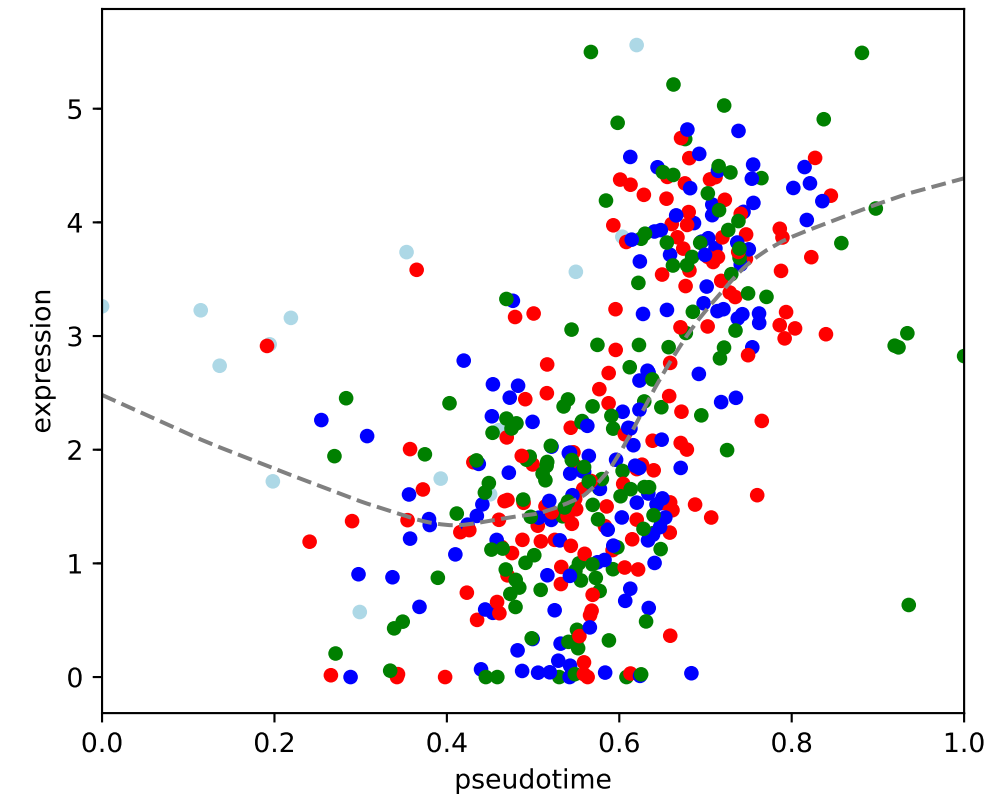
733 1spearman=-0.43



ENST00000244745 SOX4 (SRY-box 4)

733 0spearman=0.59

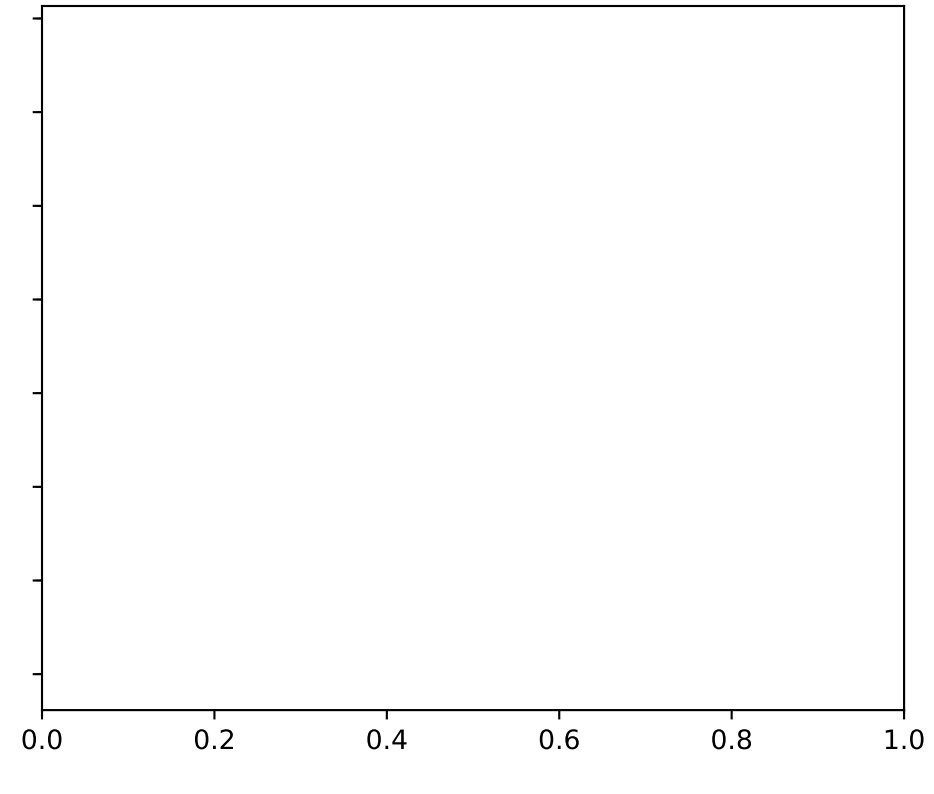
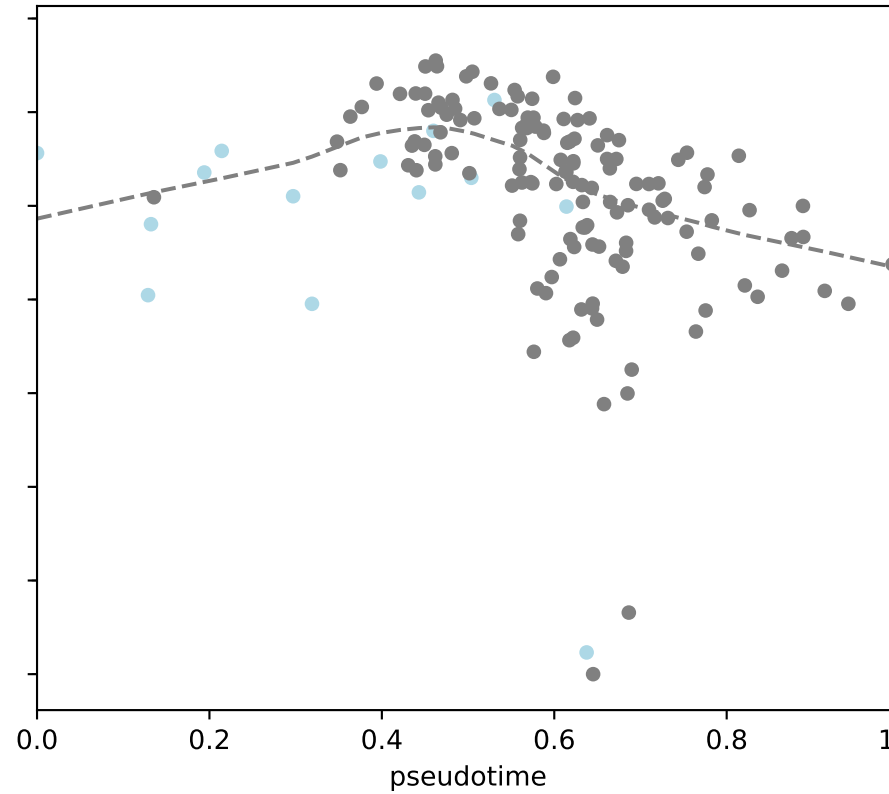
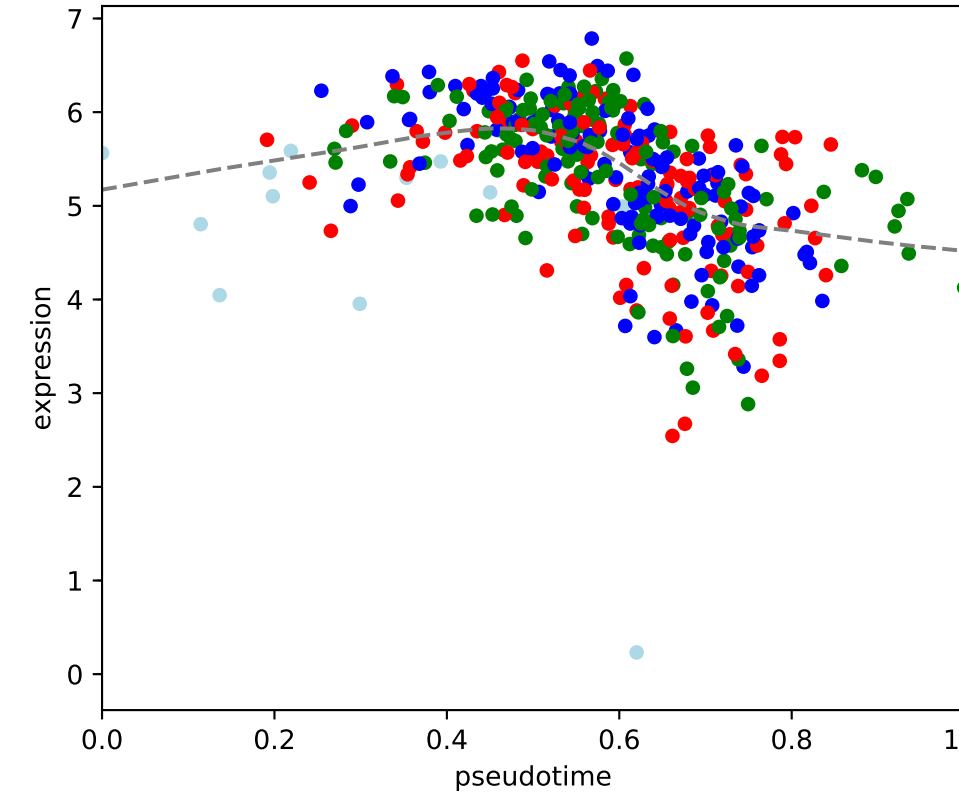
733 1spearman=0.07



ENST00000260379 RPLP1 (ribosomal protein lateral stalk subunit P1)

733 0spearman=-0.59

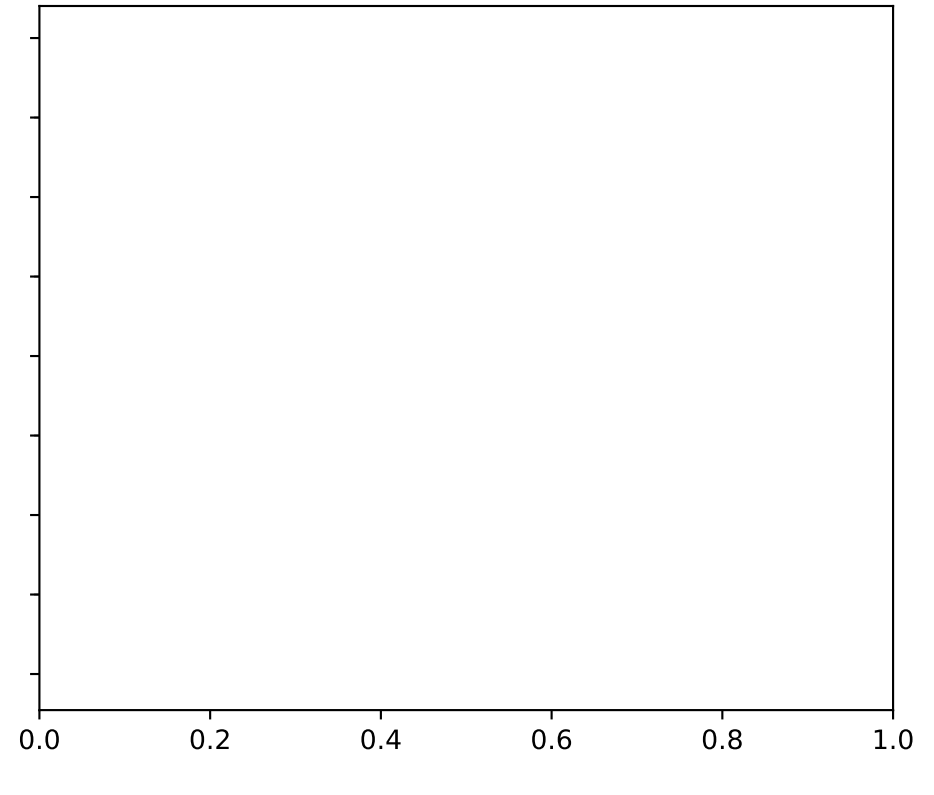
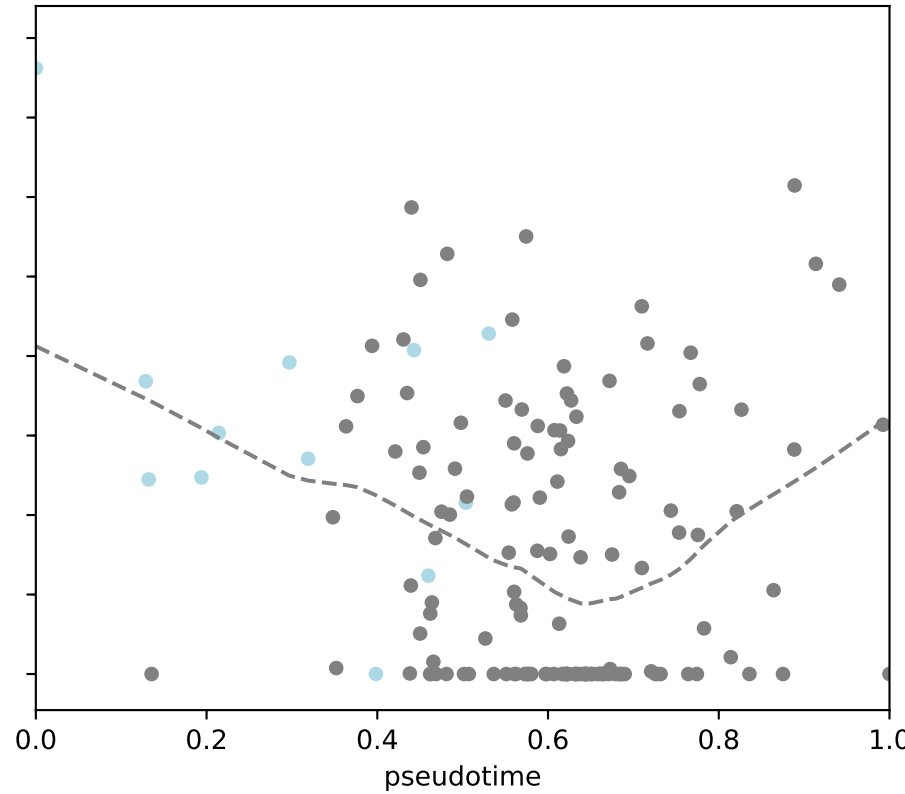
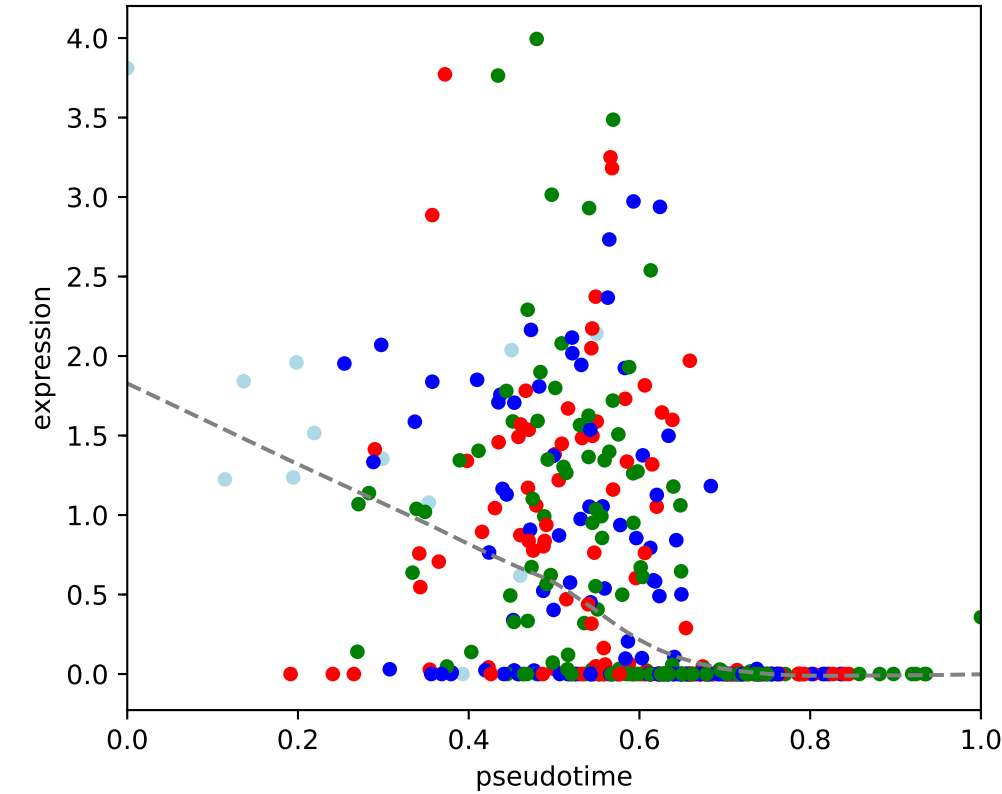
733 1spearman=-0.54



ENST00000447459 GSG1L (GSG1 like)

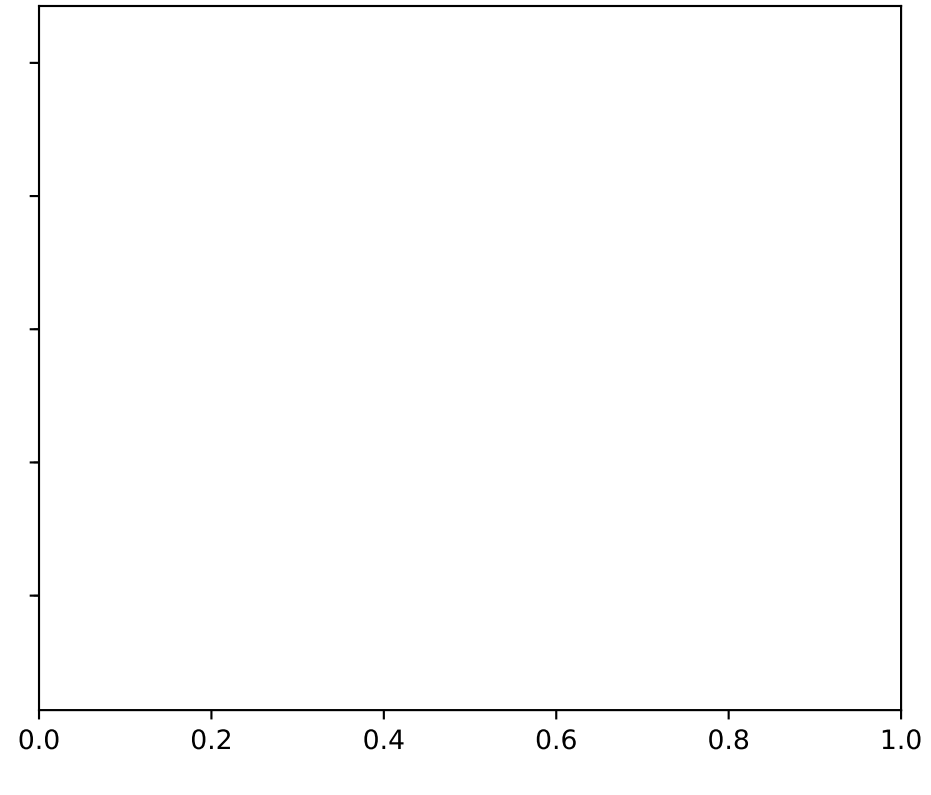
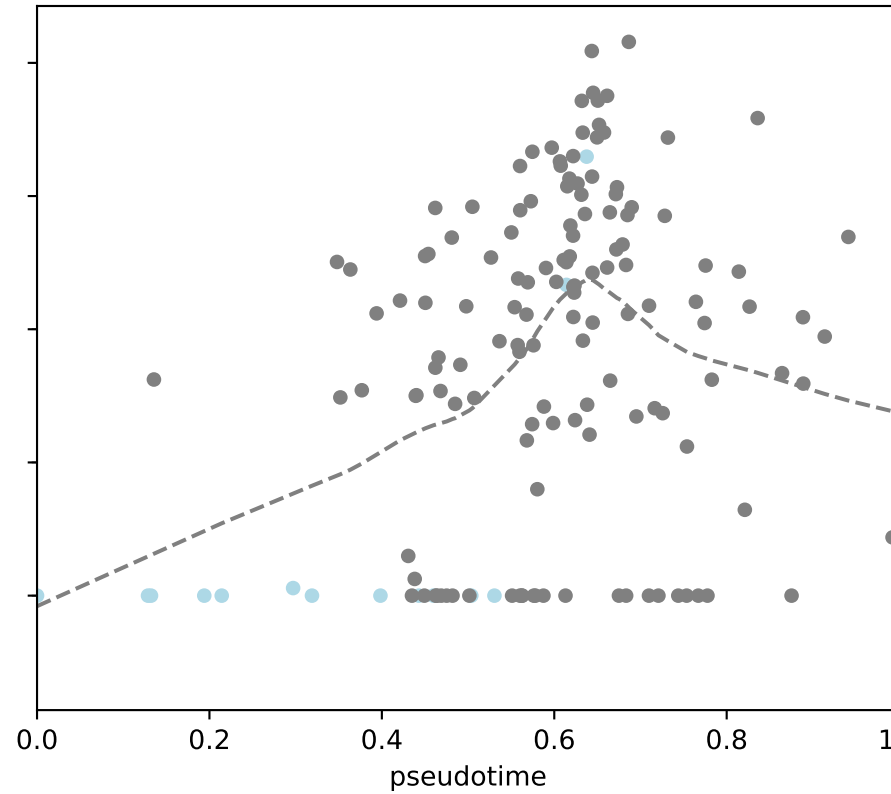
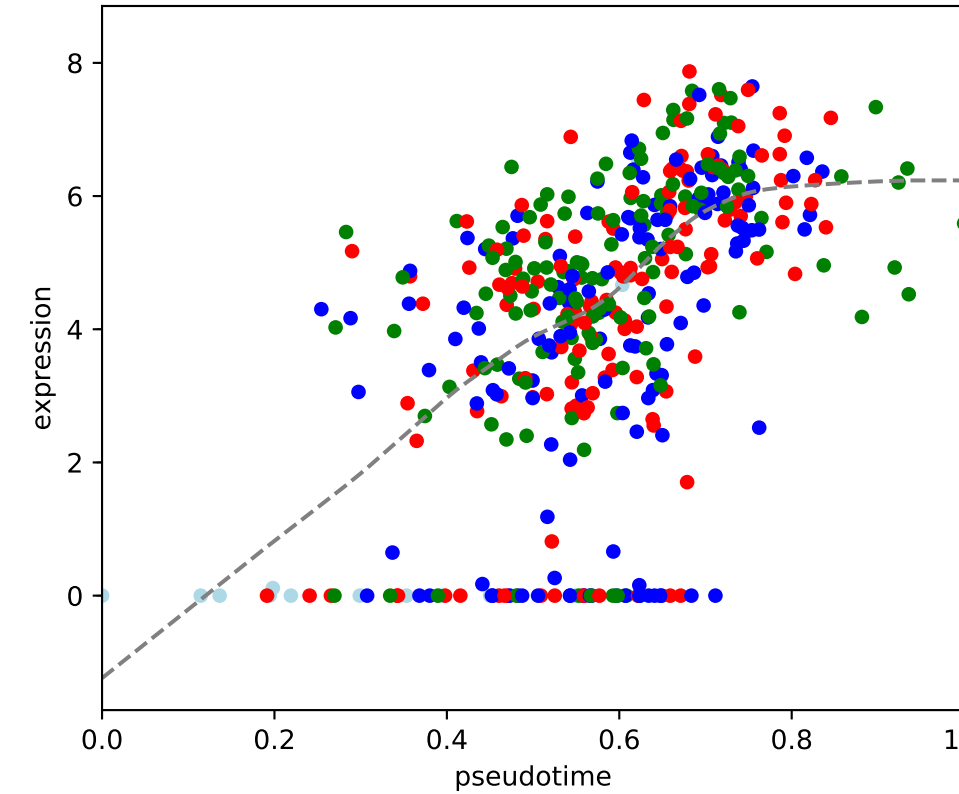
733 0spearman=-0.59

733 1spearman=-0.19



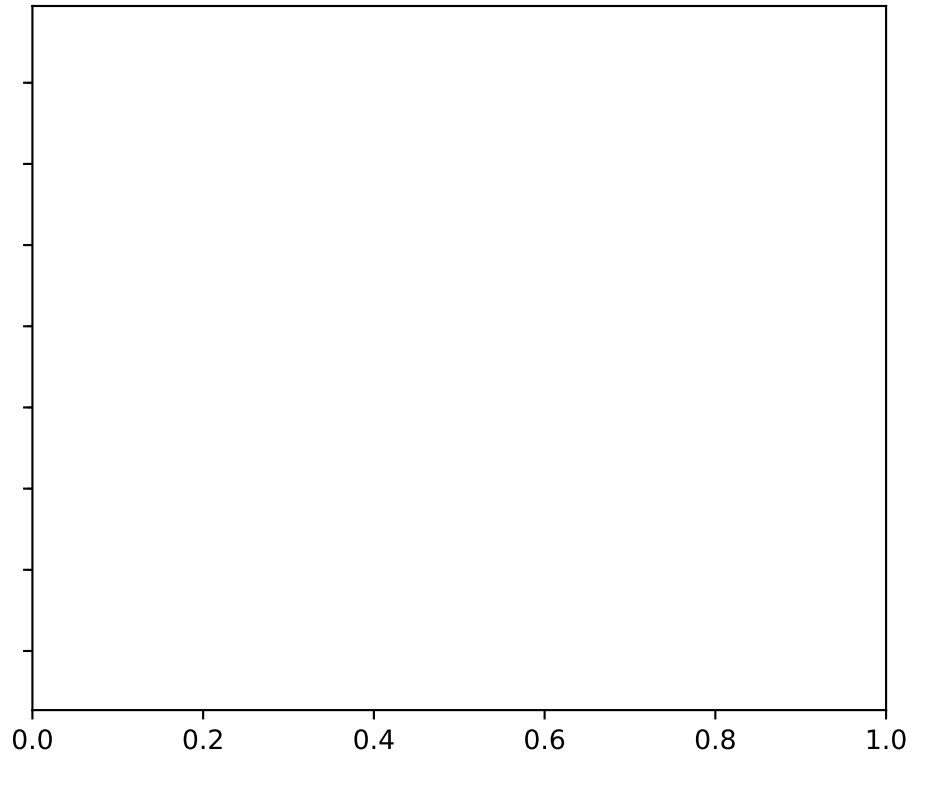
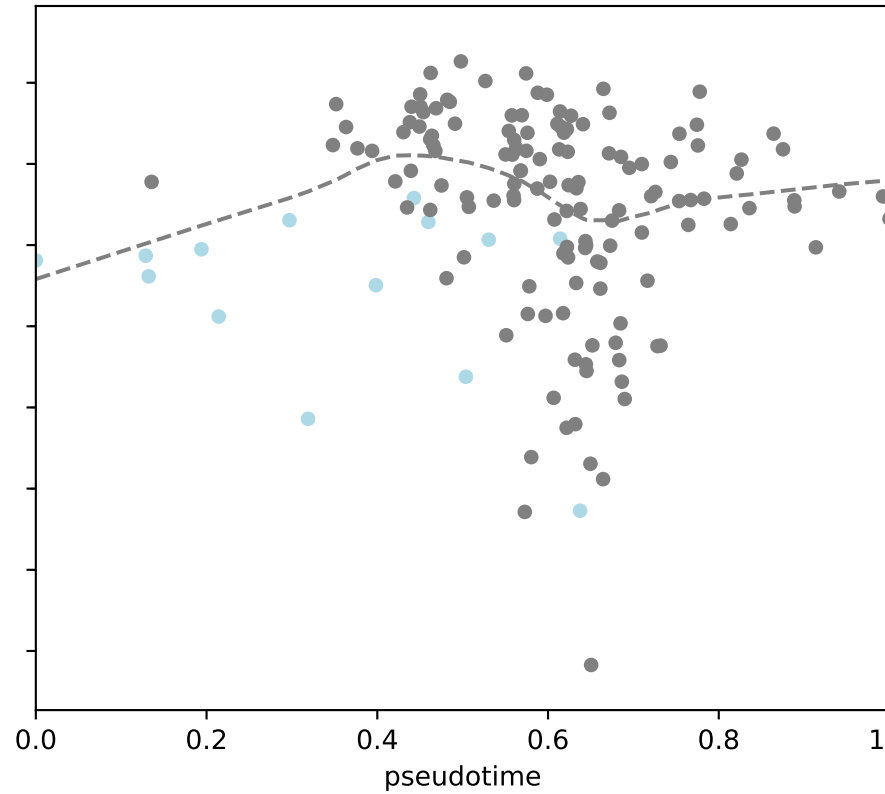
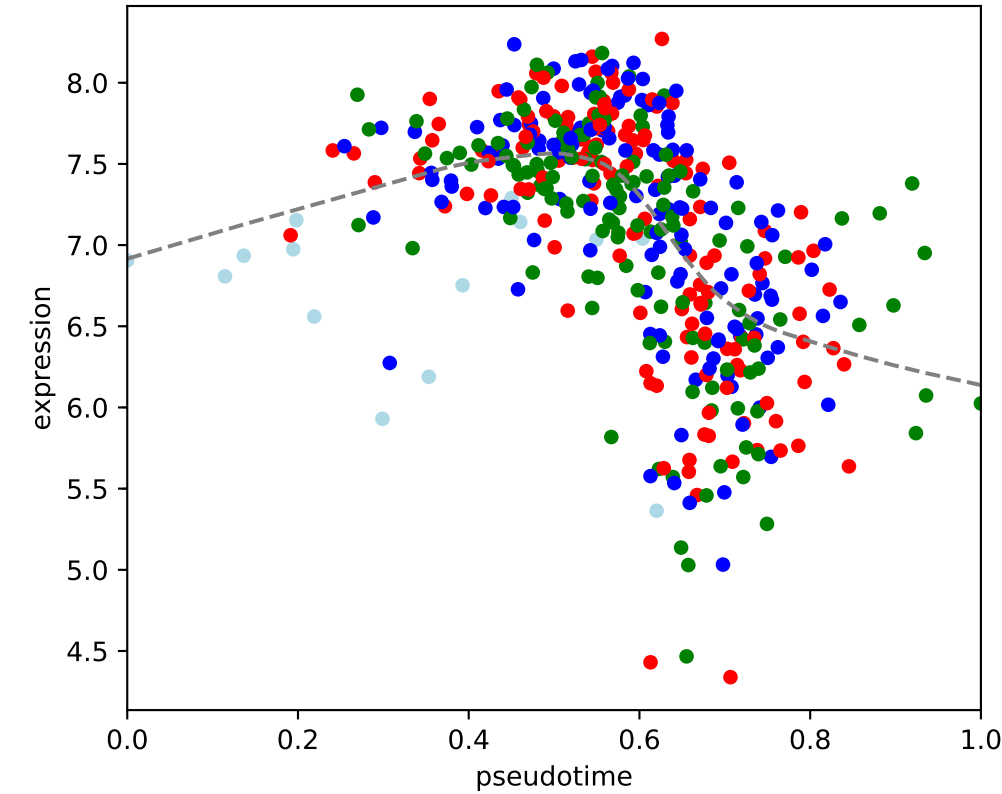
733 0spearman=0.58

733 1spearman=0.28



733 0spearman=-0.58

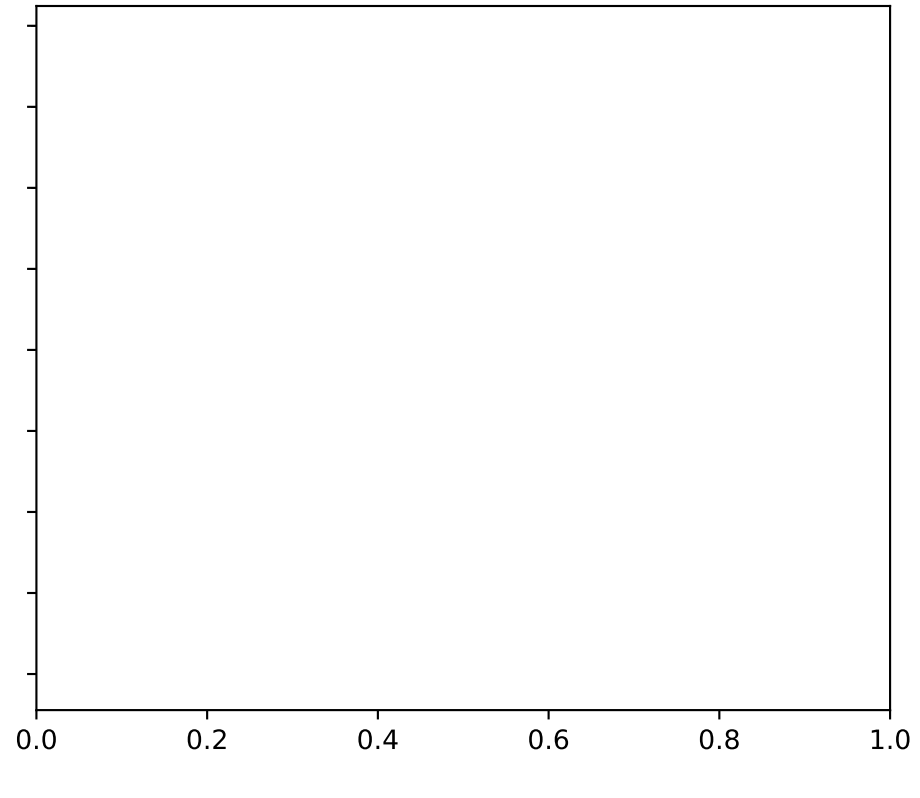
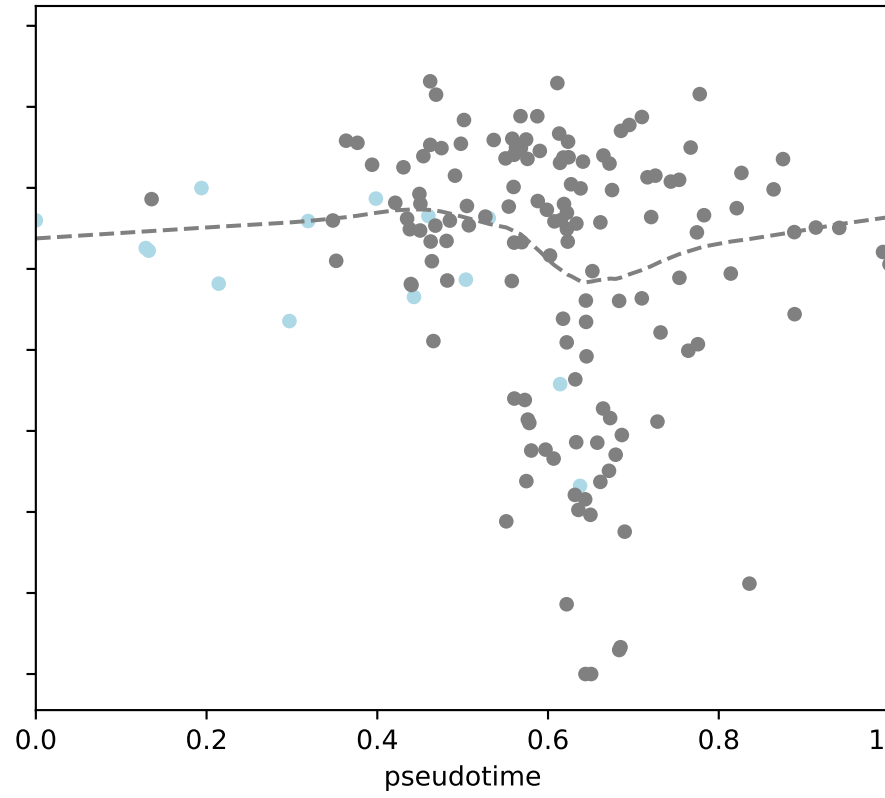
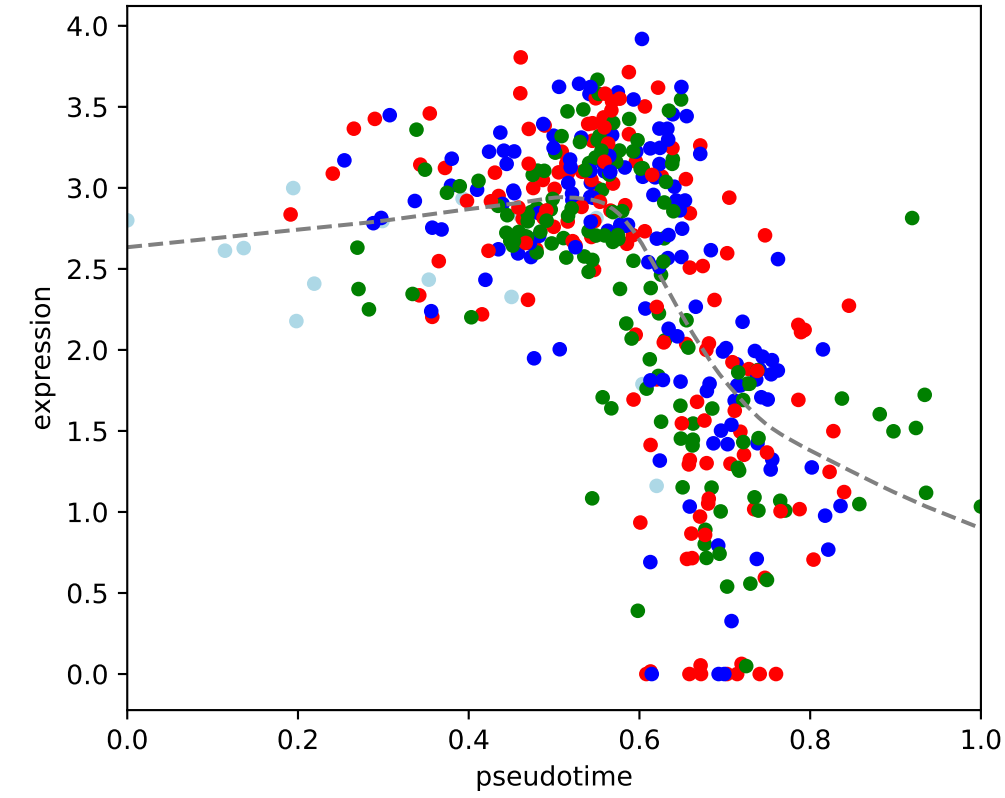
733 1spearman=-0.22



ENST00000596415 NDUFV2P1 (NADH:ubiquinone oxidoreductase core subunit V2 pseudogene 1)

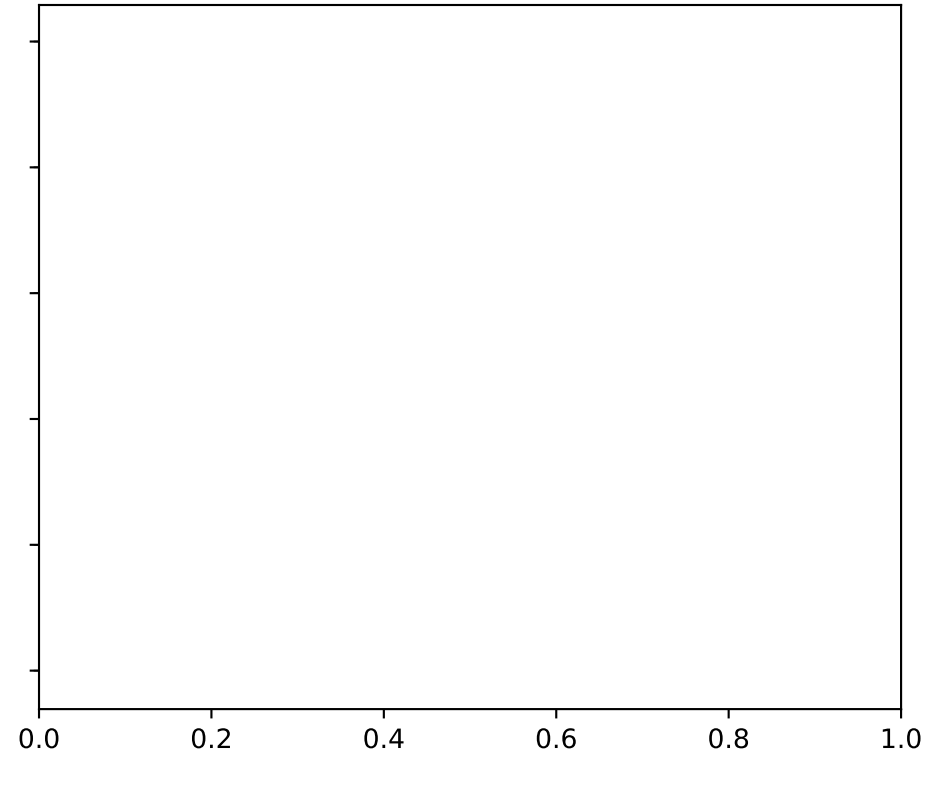
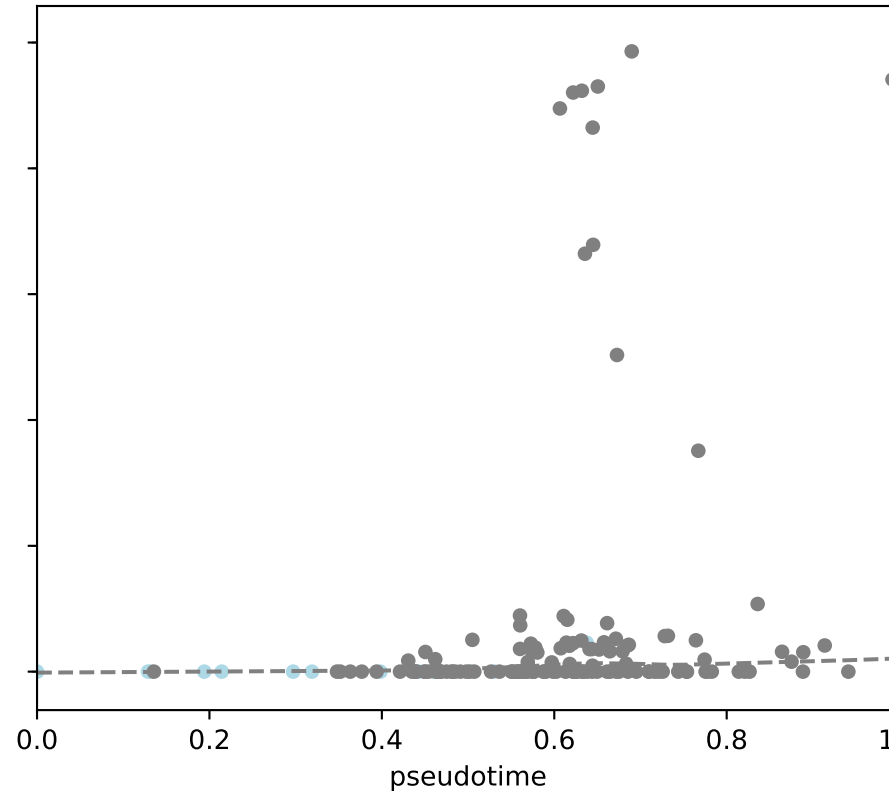
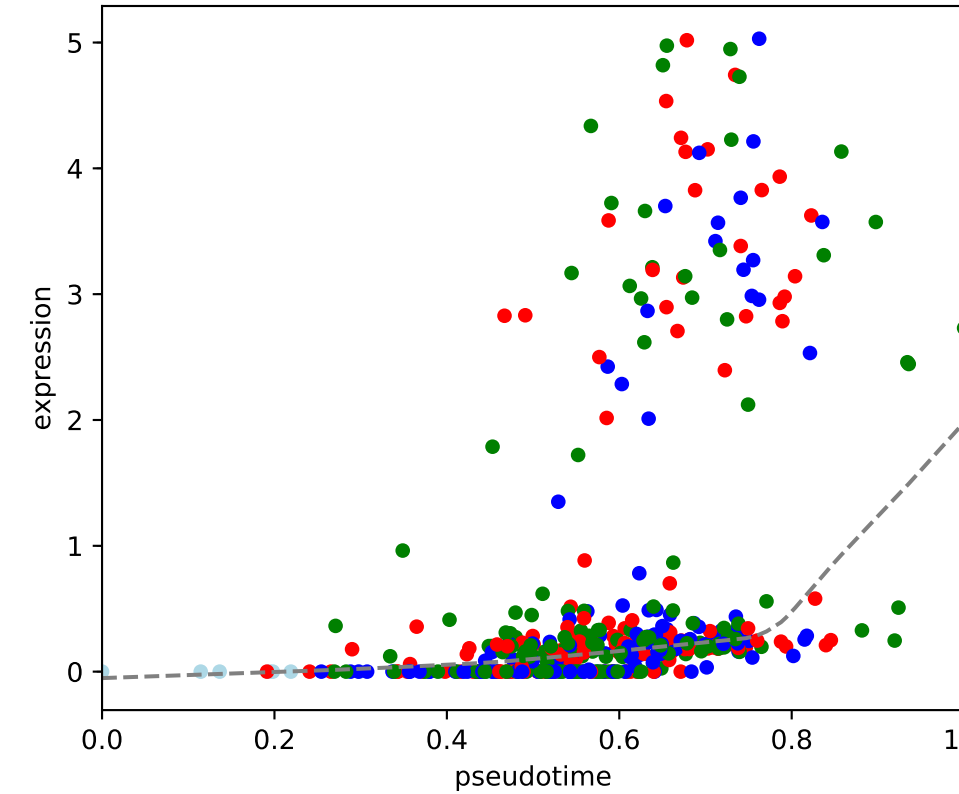
733 0spearman=-0.58

733 1spearman=-0.20



733 0spearman=0.58

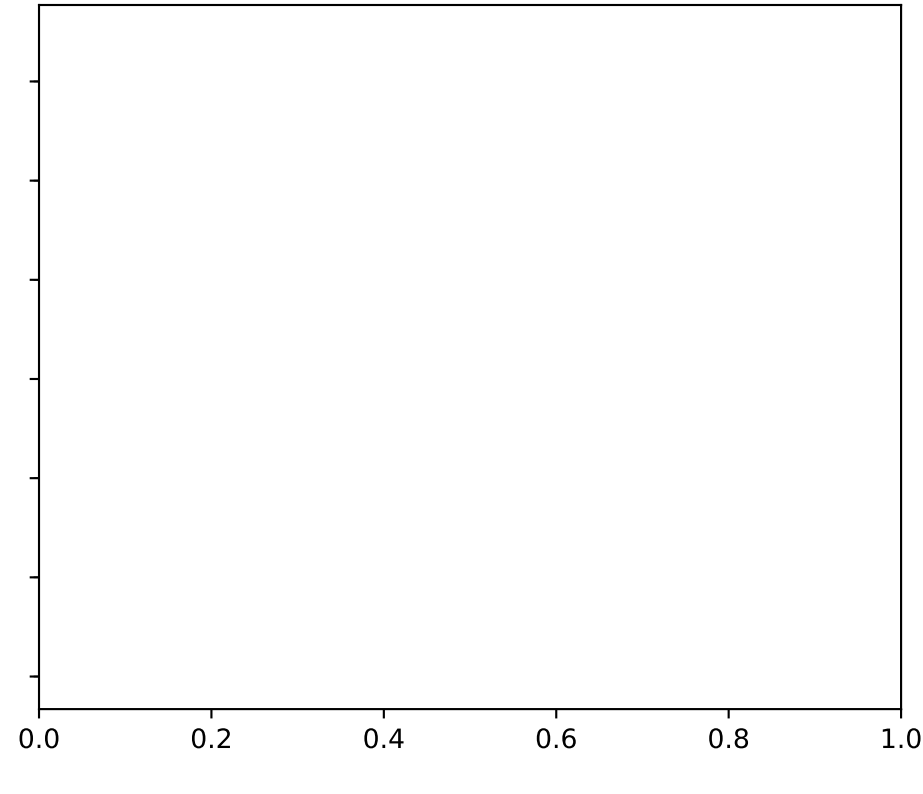
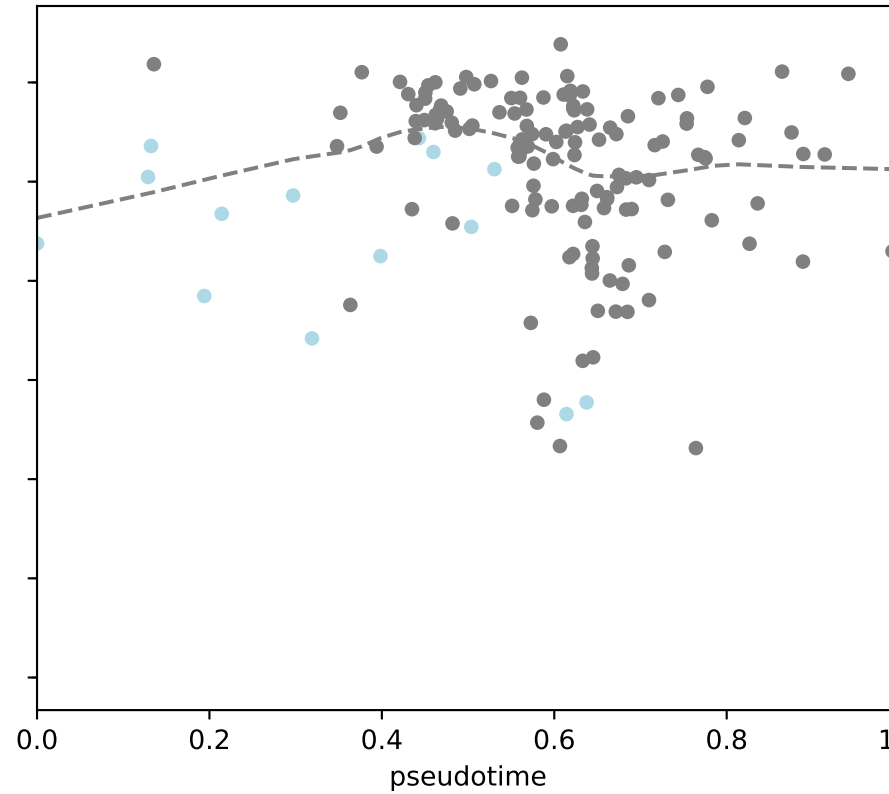
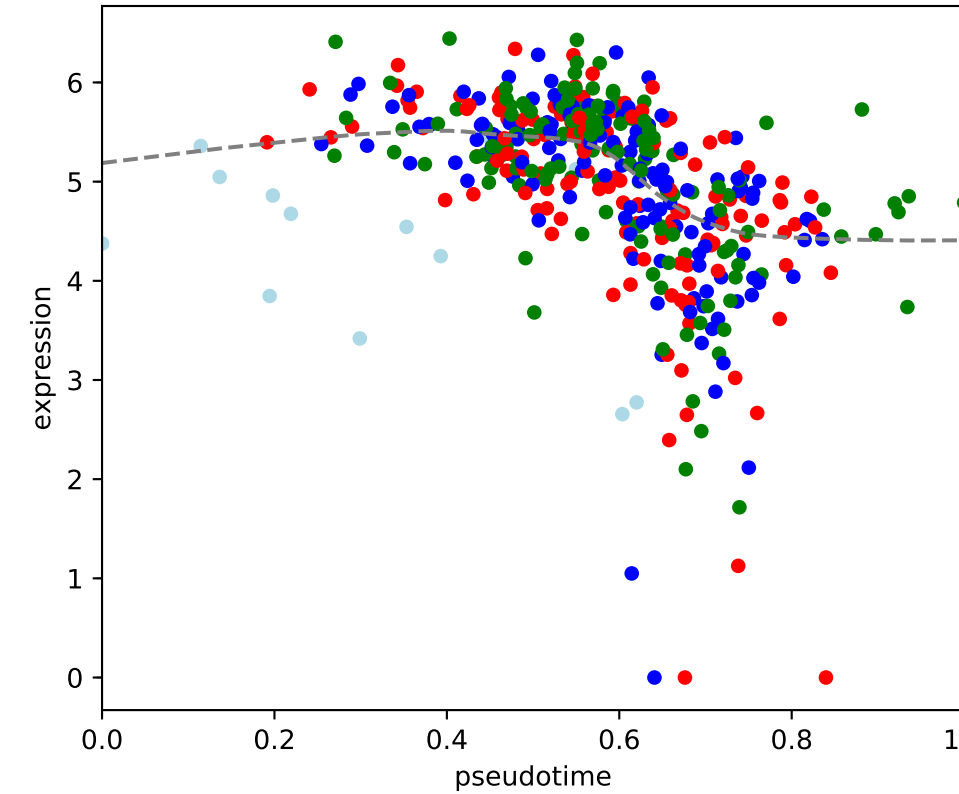
733 1spearman=0.36



ENST00000525085 NDUFC2 (NADH:ubiquinone oxidoreductase subunit C2)

733 0spearman=-0.58

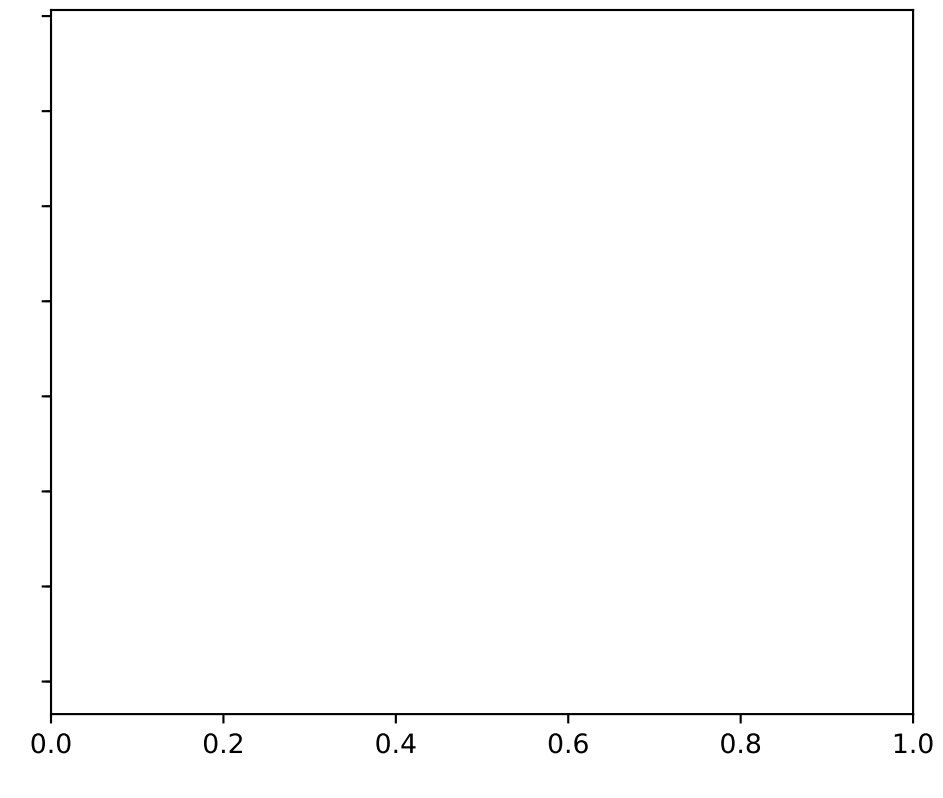
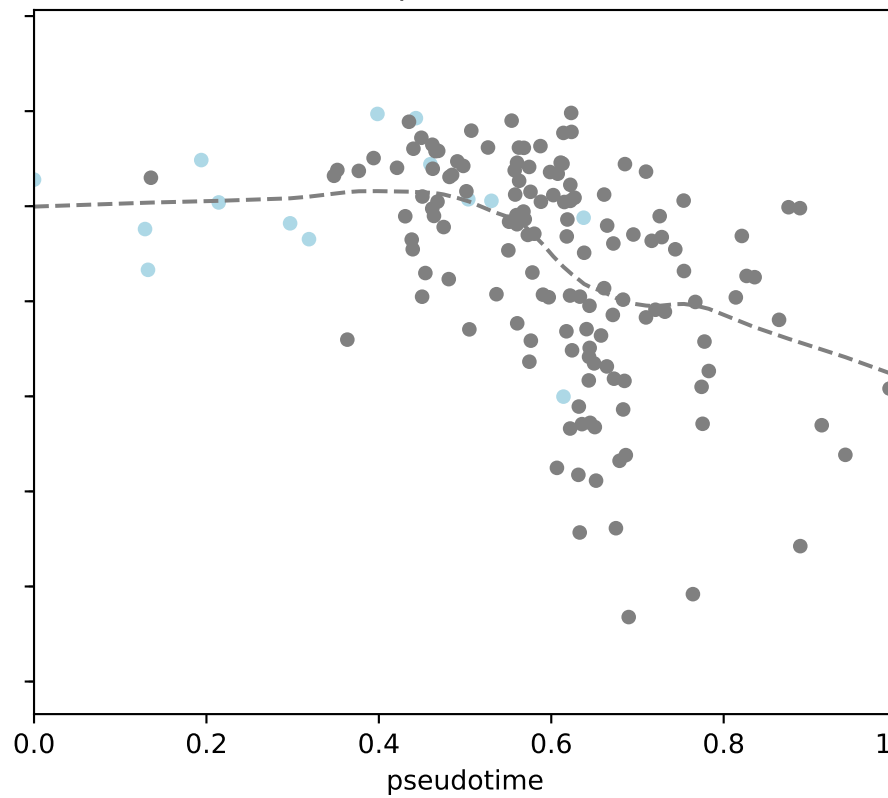
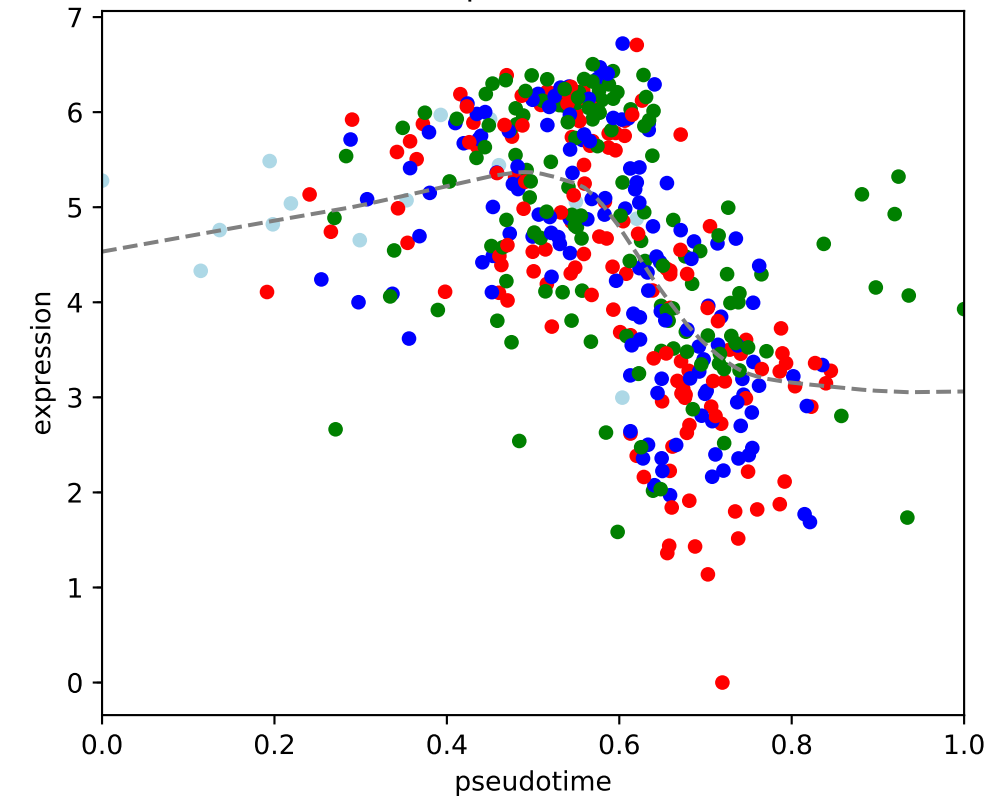
733 1spearman=-0.24



ENST00000550367 TUBA1B (tubulin alpha 1b)

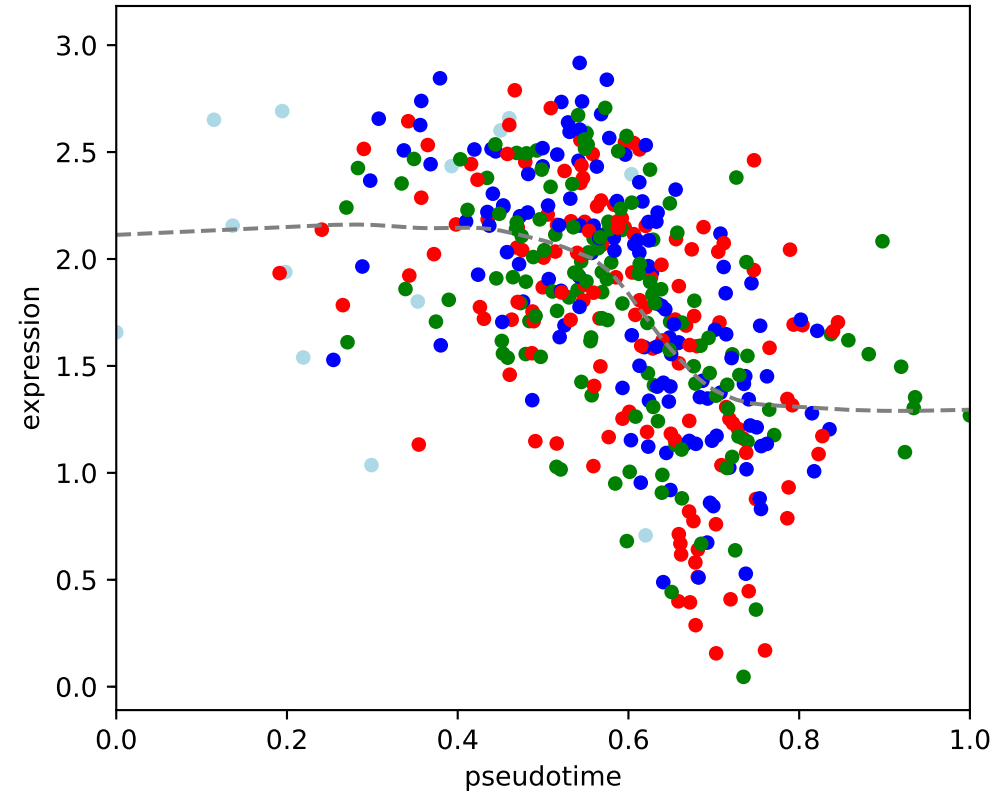
733 0spearman=-0.58

733 1spearman=-0.56

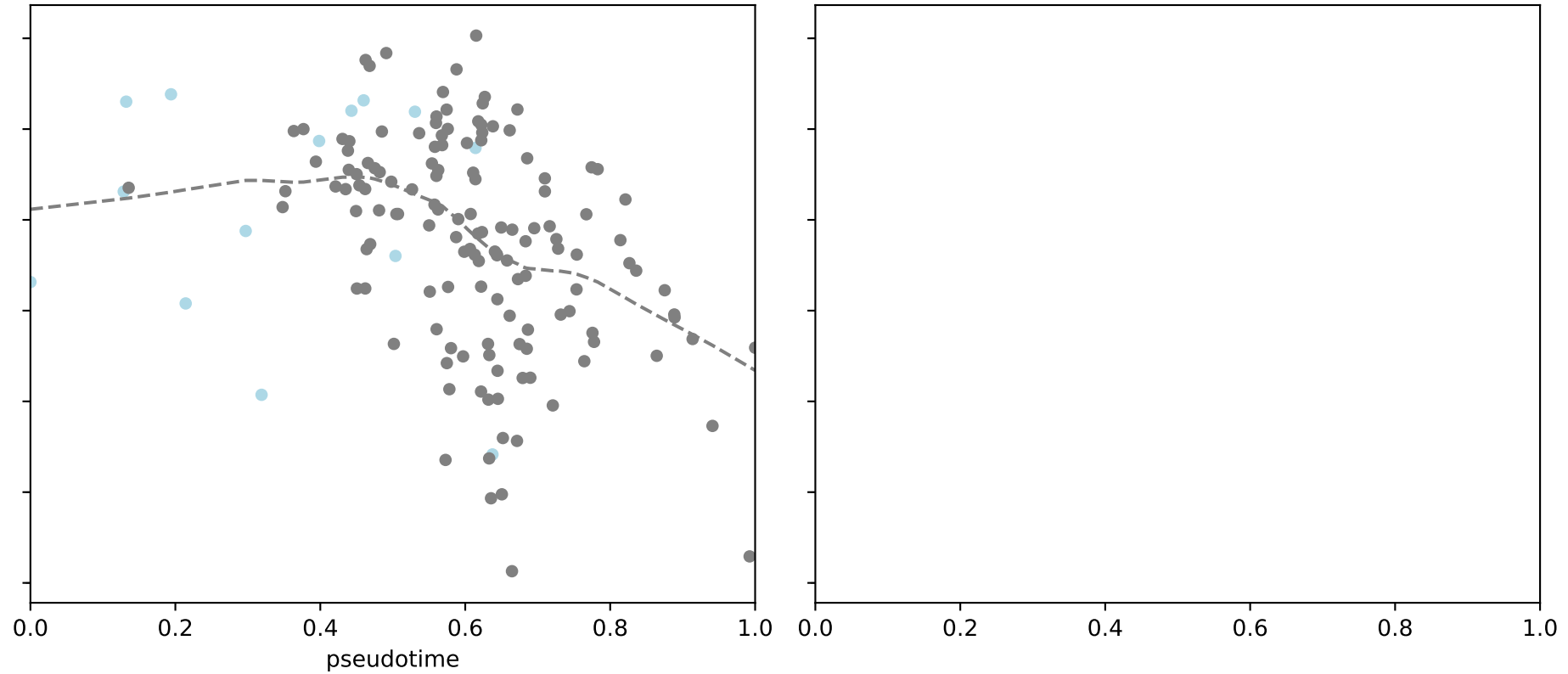


ENST00000456046 AC079250.1

733 0spearman=-0.58



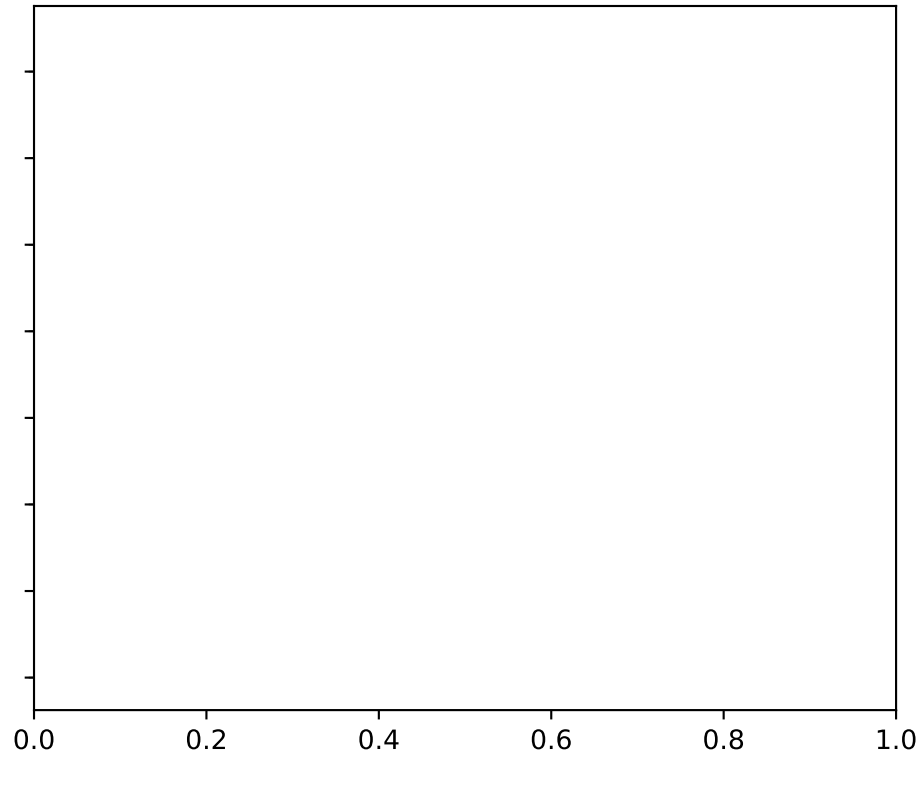
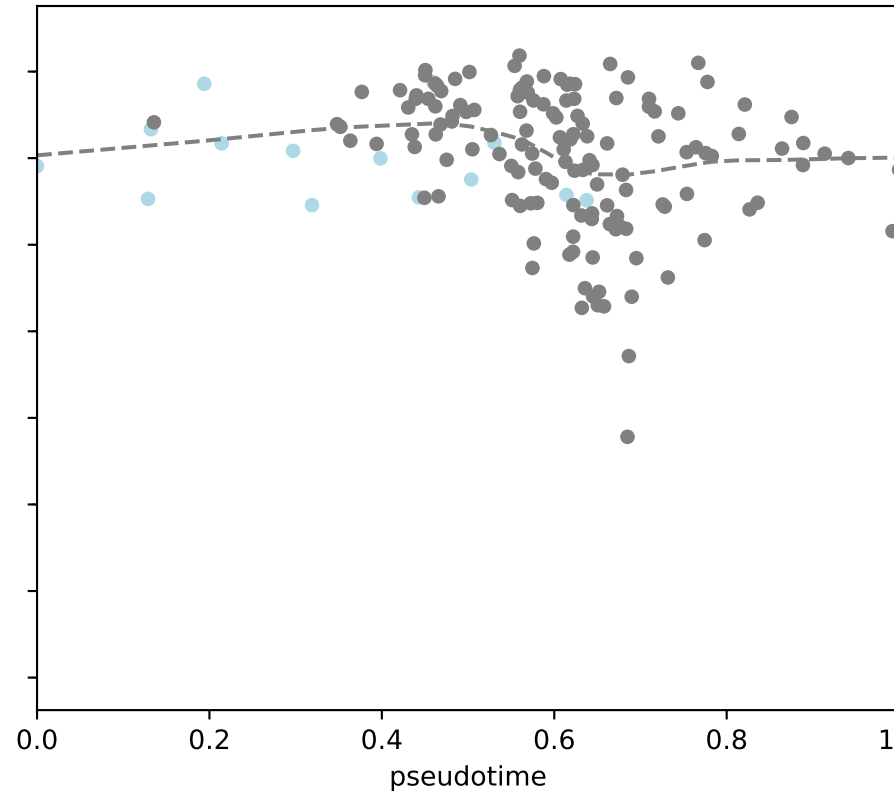
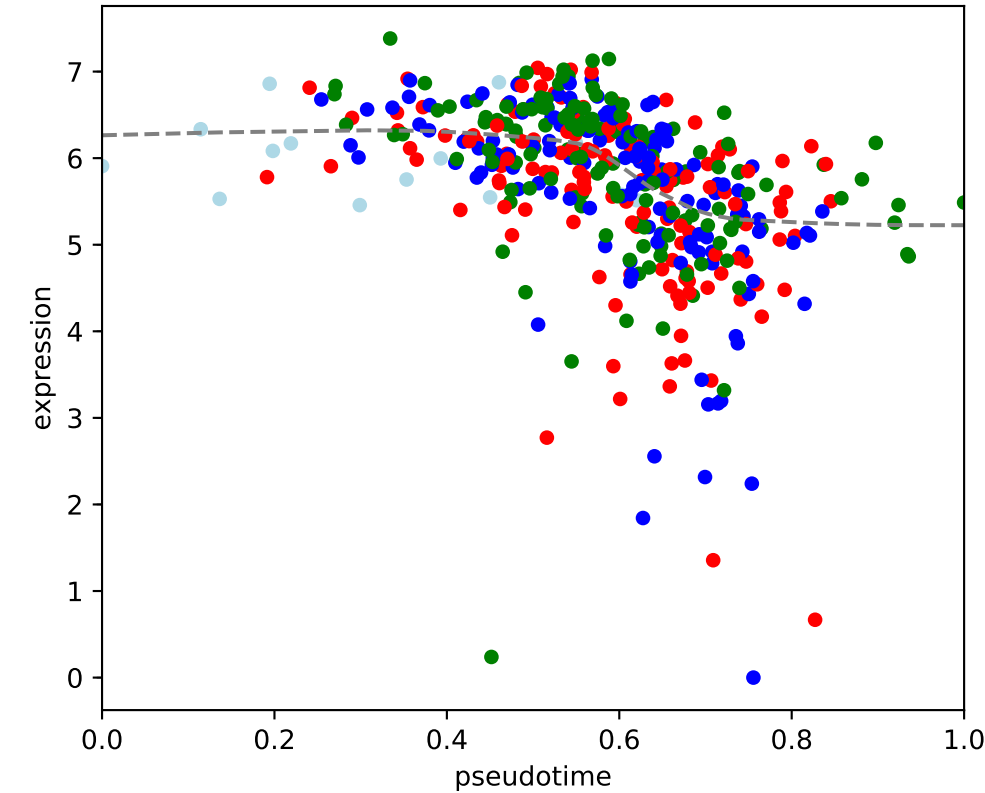
733 1spearman=-0.46



ENST00000343262 RPS2 (ribosomal protein S2)

733 0spearman=-0.58

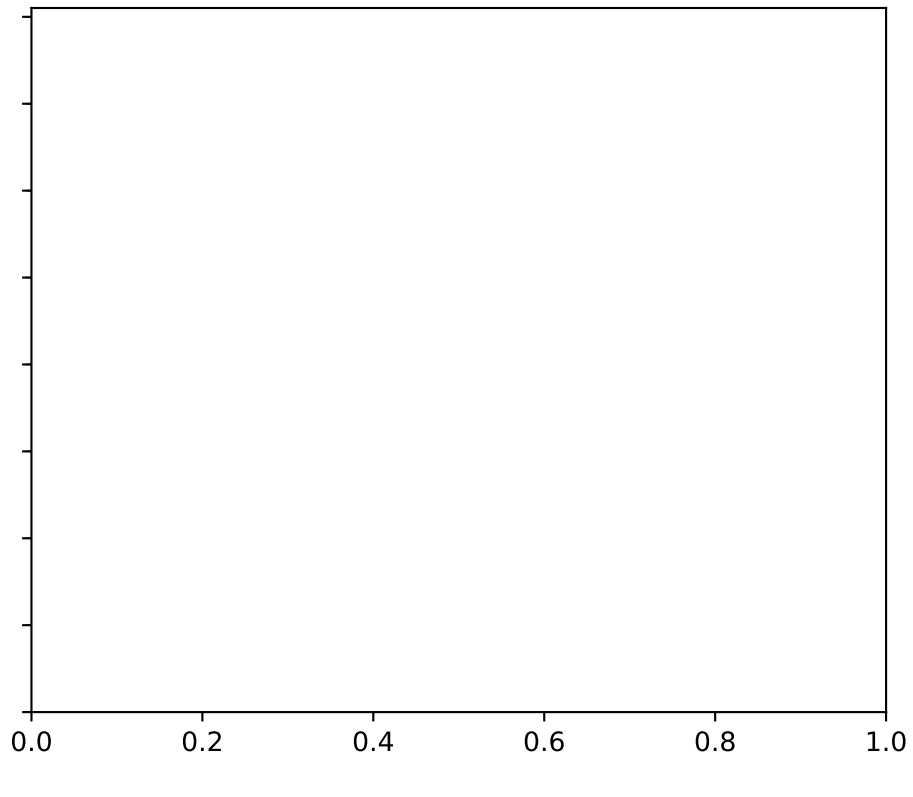
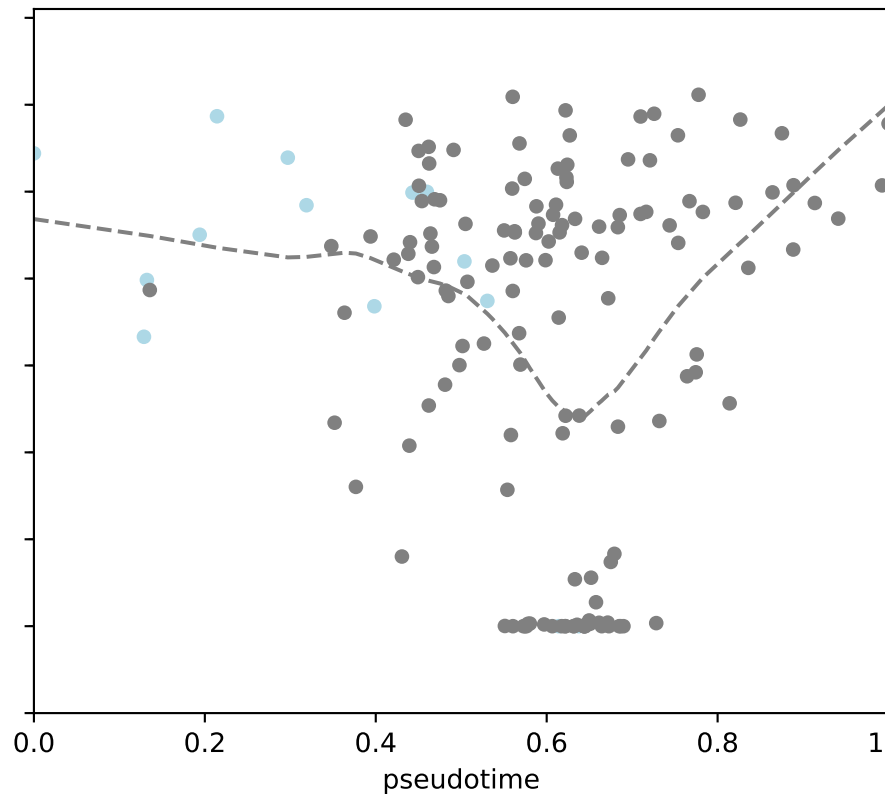
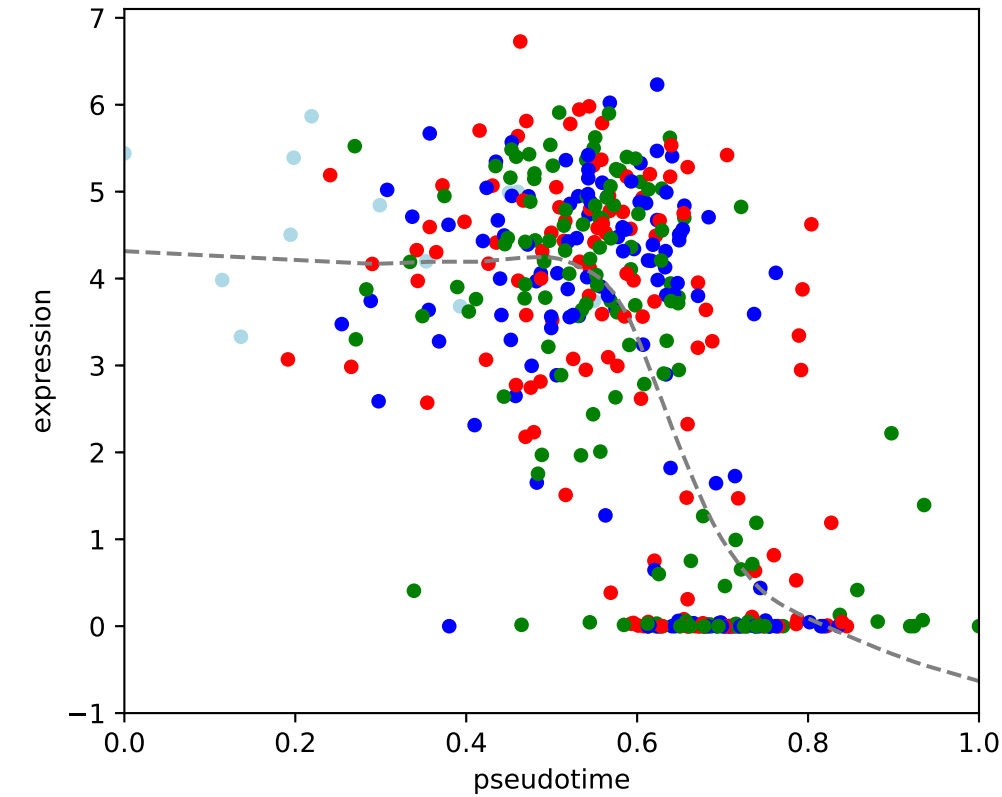
733 1spearman=-0.33



ENST00000521104 DST (dystonin)

733 0spearman=-0.58

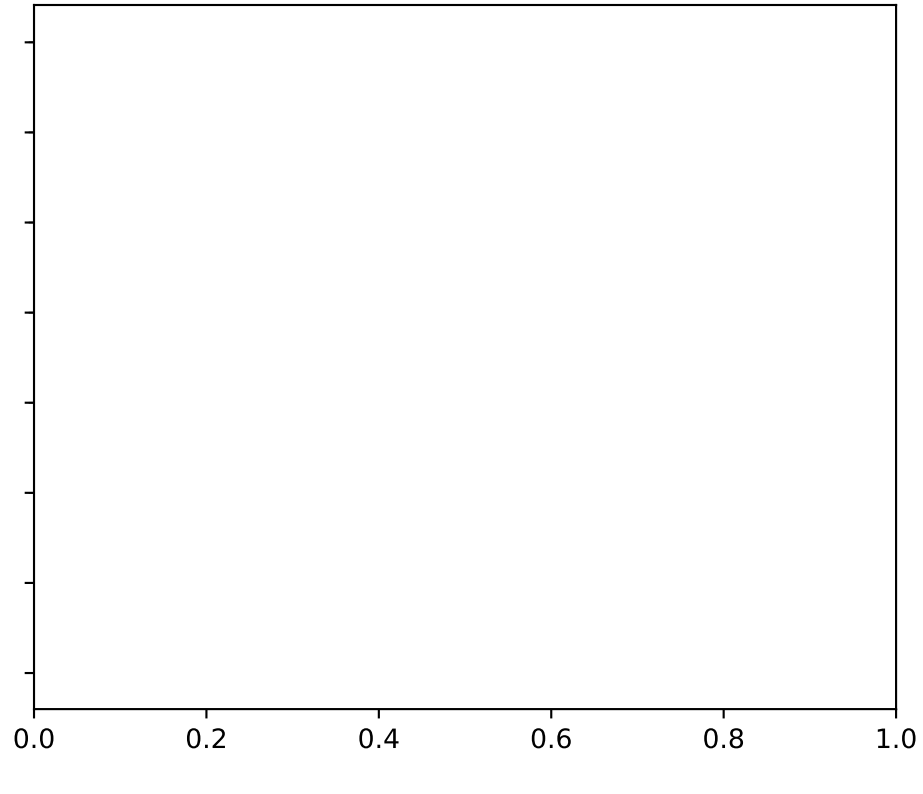
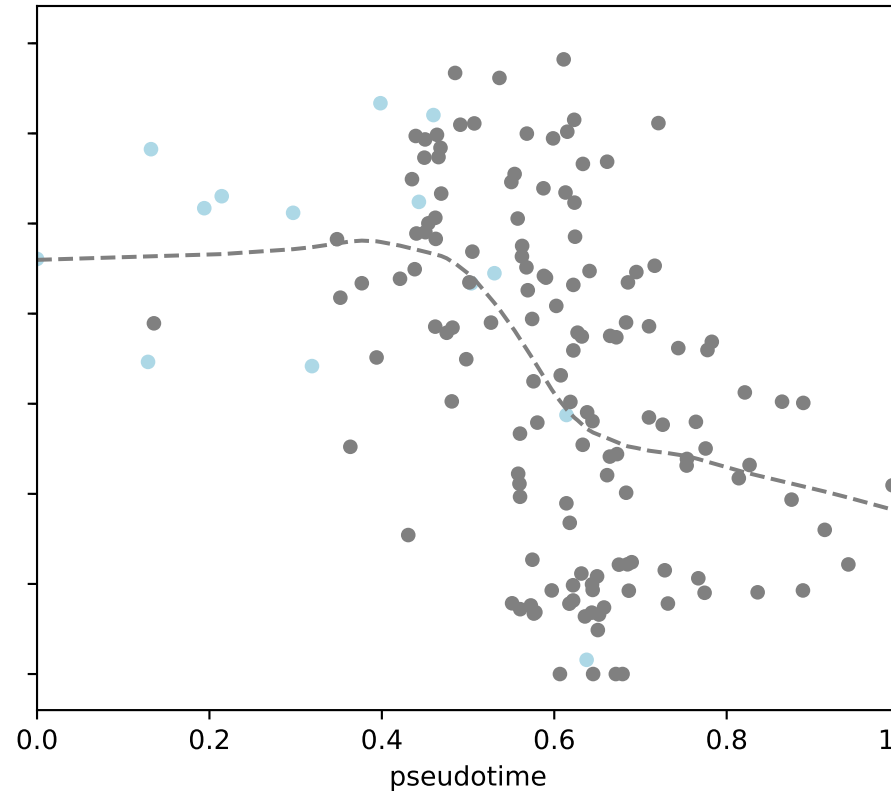
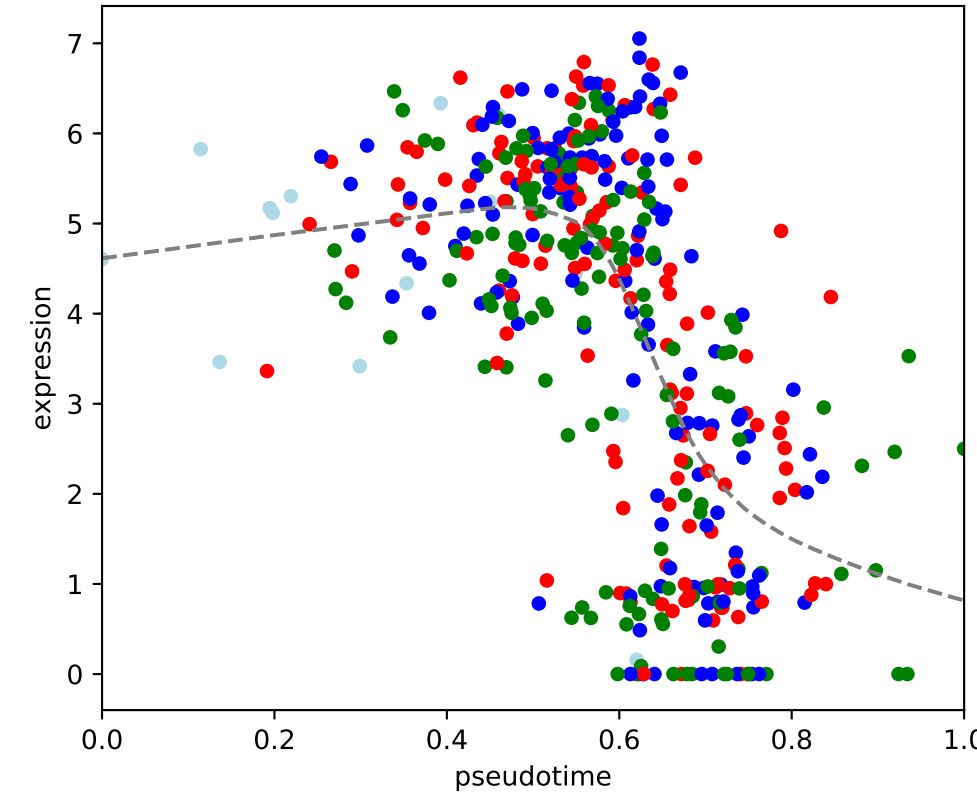
733 1spearman=-0.04



ENST00000366718 GUK1 (guanylate kinase 1)

733 0spearman=-0.58

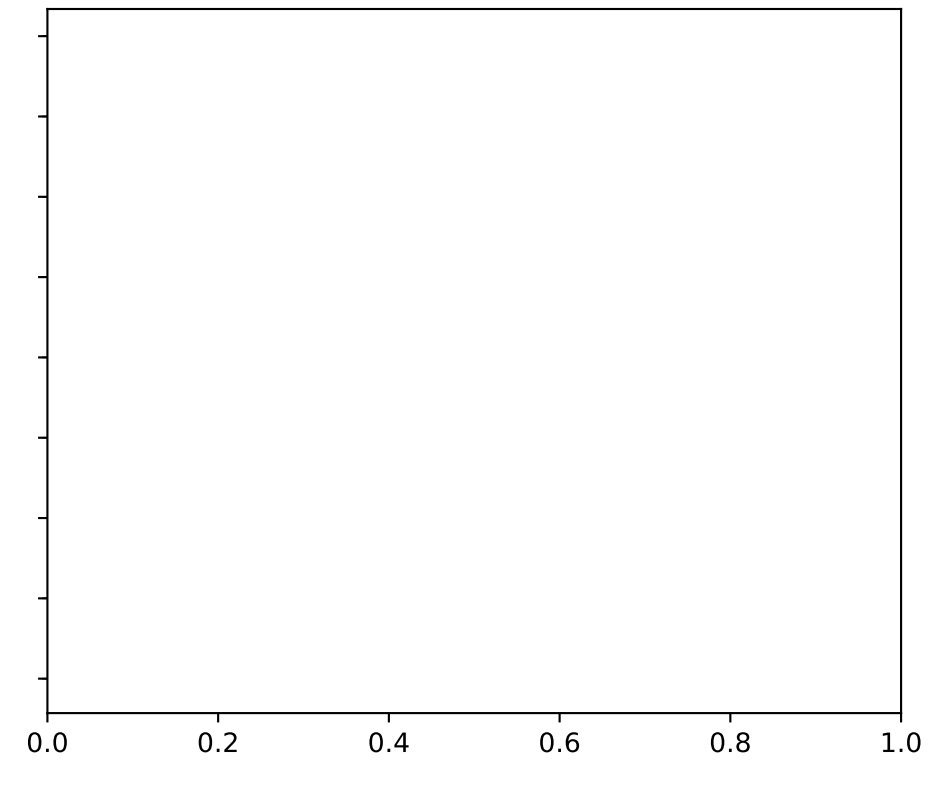
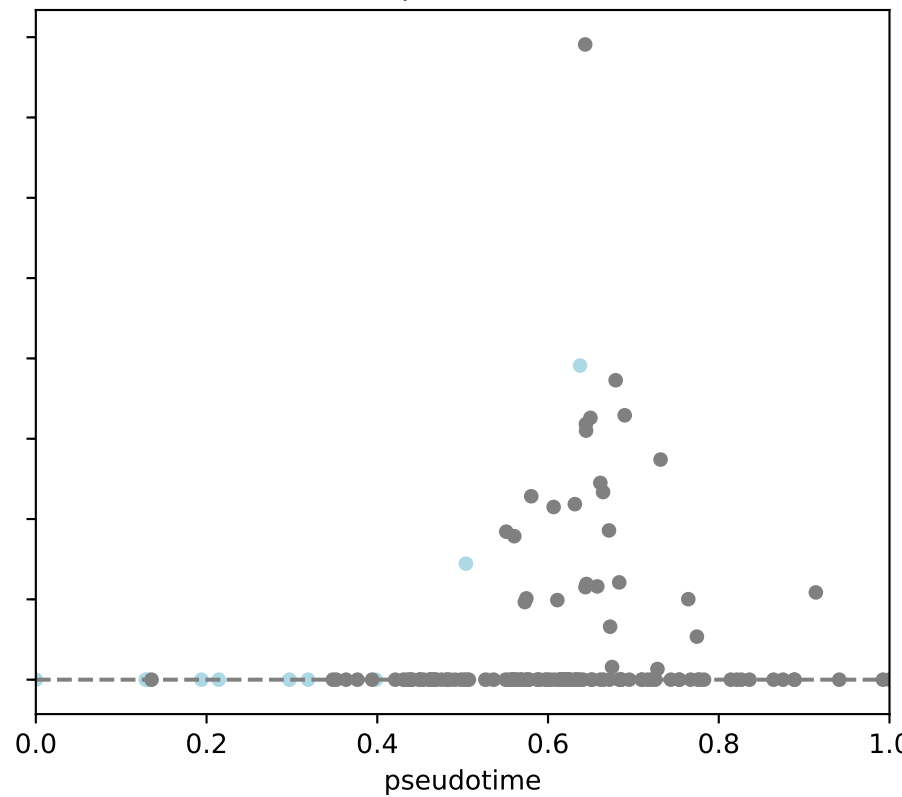
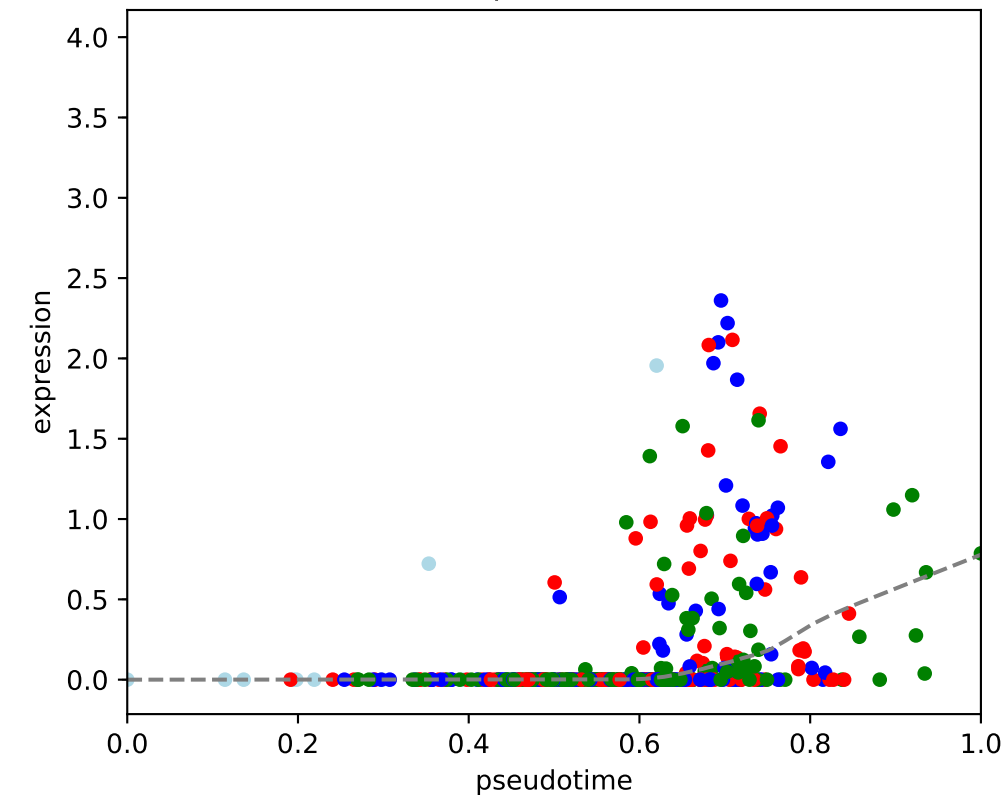
733 1spearman=-0.51



ENST00000559194 CGNL1 (cingulin like 1)

733 0spearman=0.58

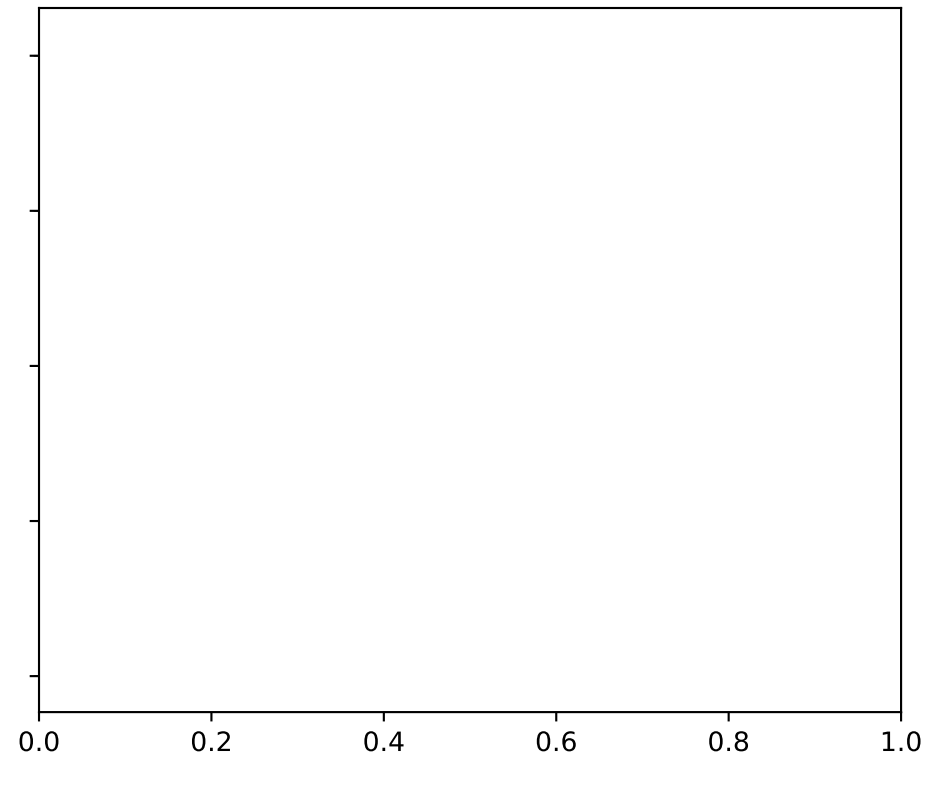
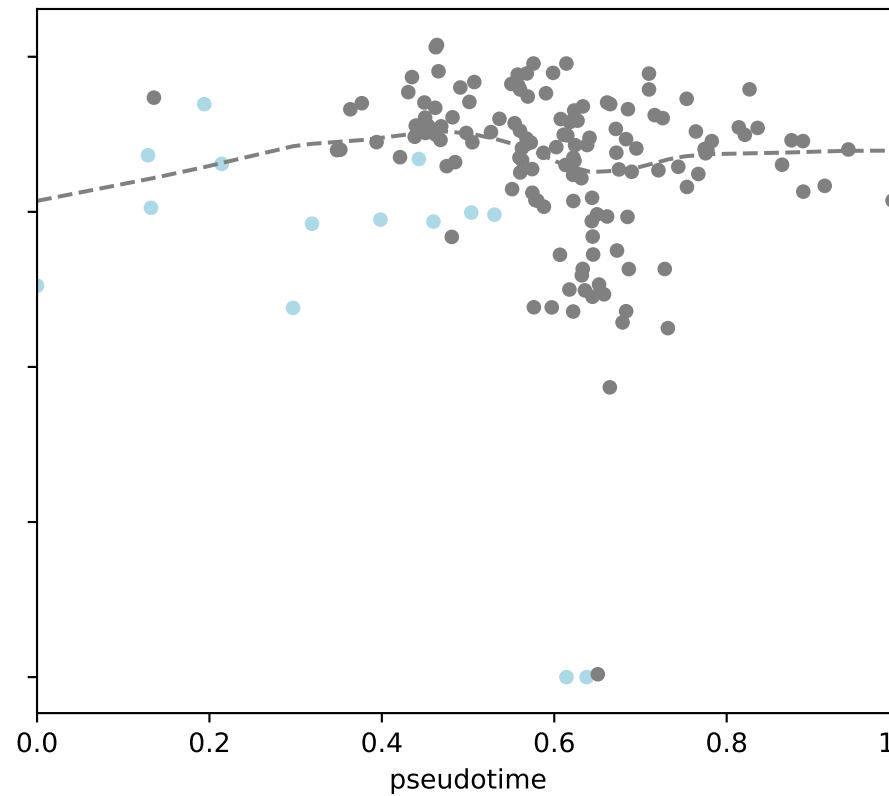
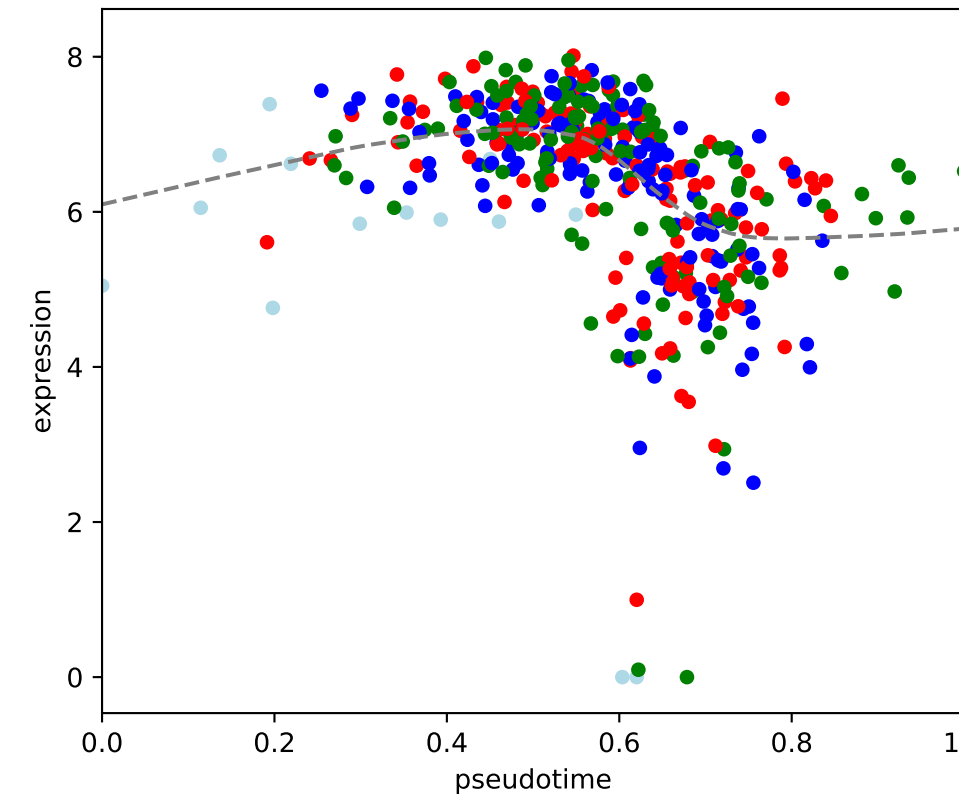
733 1spearman=0.26



ENST00000304312 ATP5ME (ATP synthase membrane subunit e)

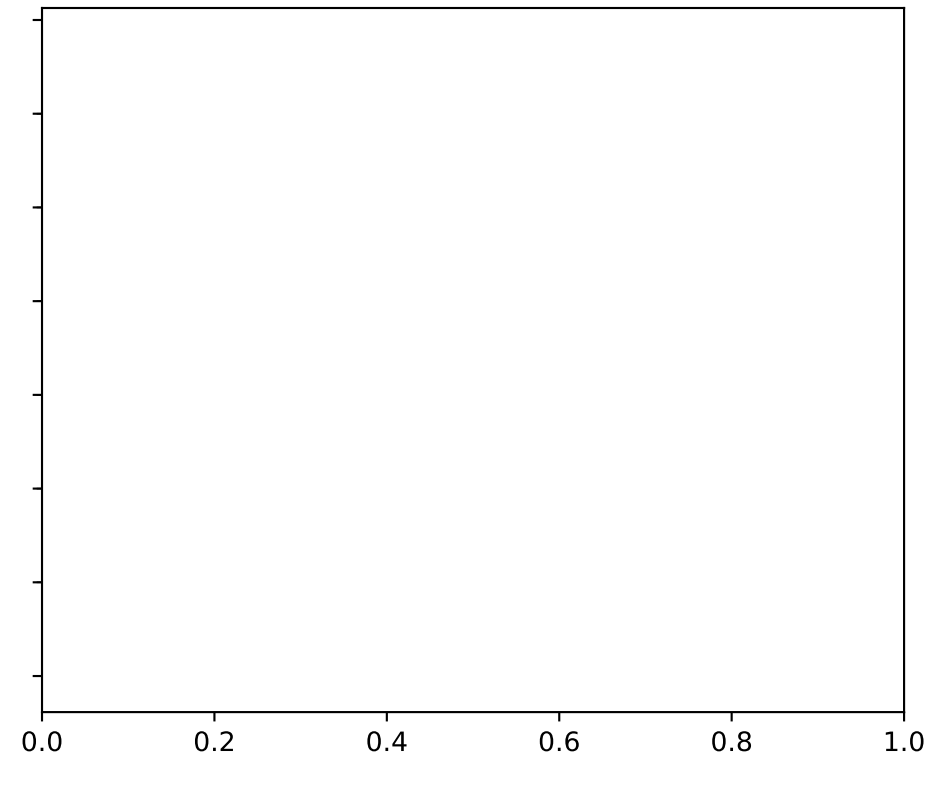
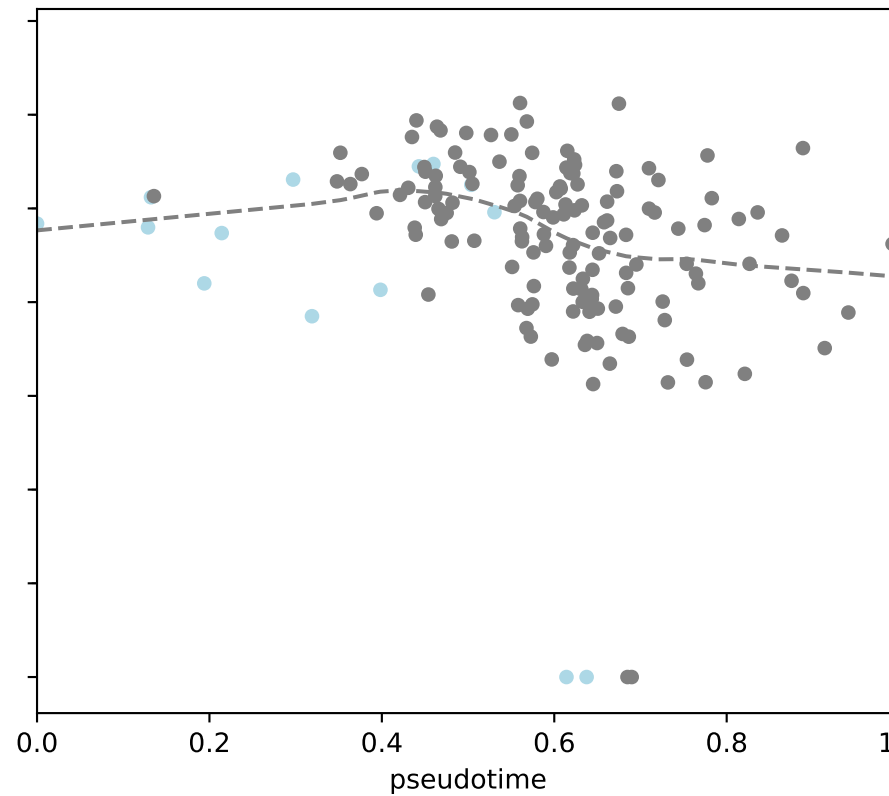
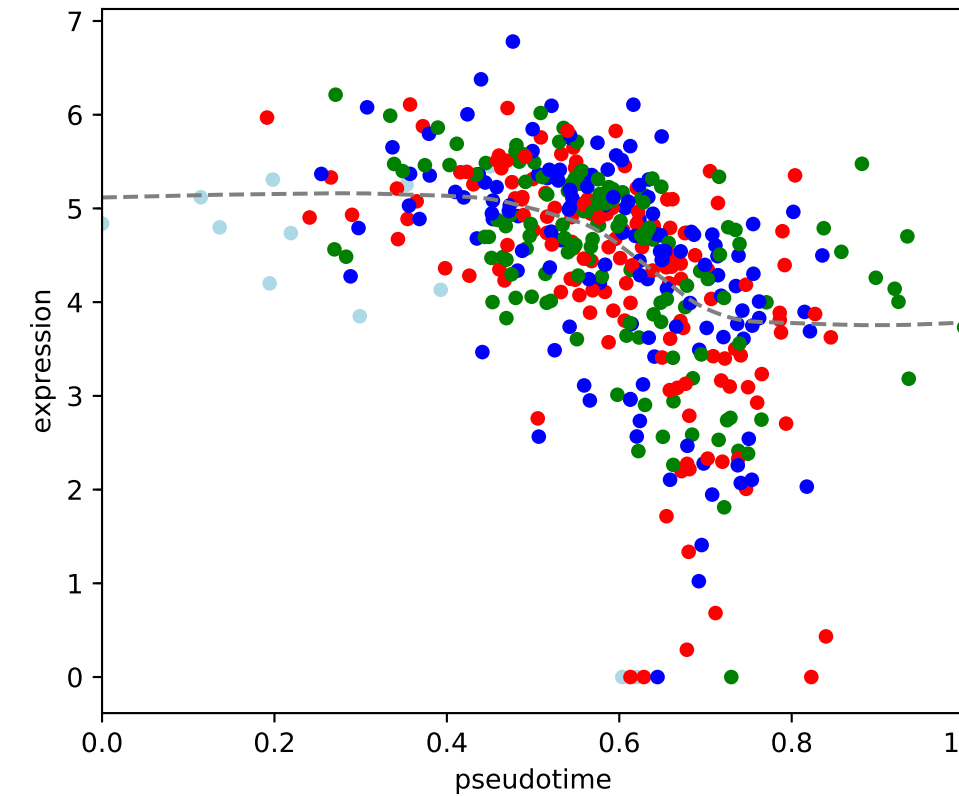
733 0spearman=-0.58

733 1spearman=-0.21



733 0spearman=-0.58

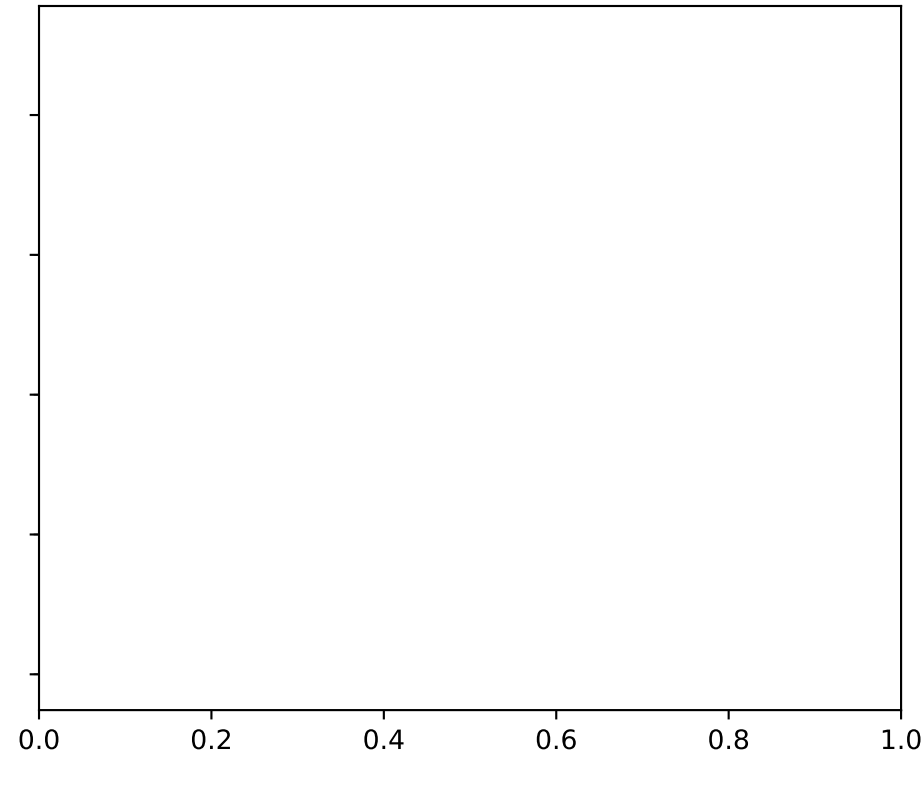
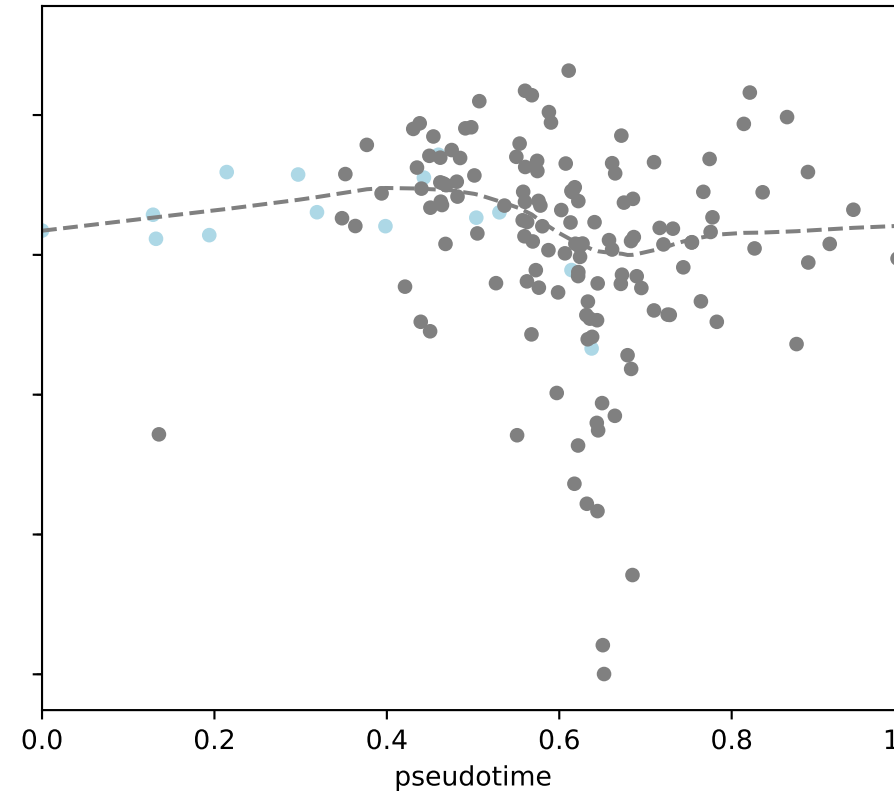
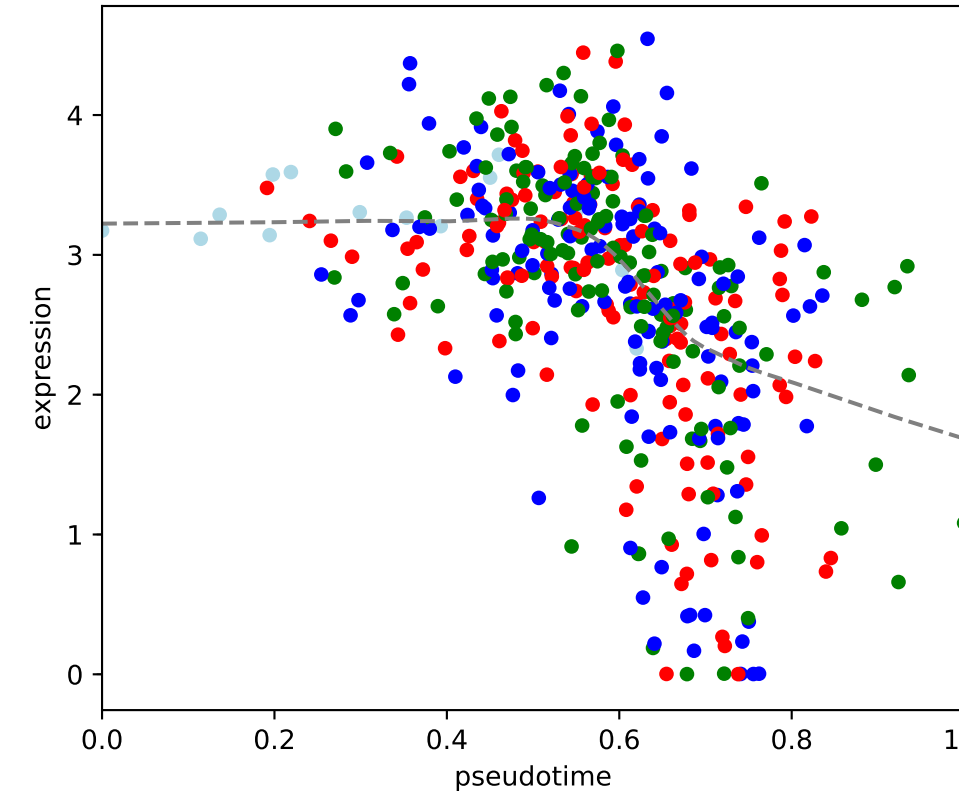
733 1spearman=-0.41



ENST00000544151 LDHB (lactate dehydrogenase B)

733 0spearman=-0.57

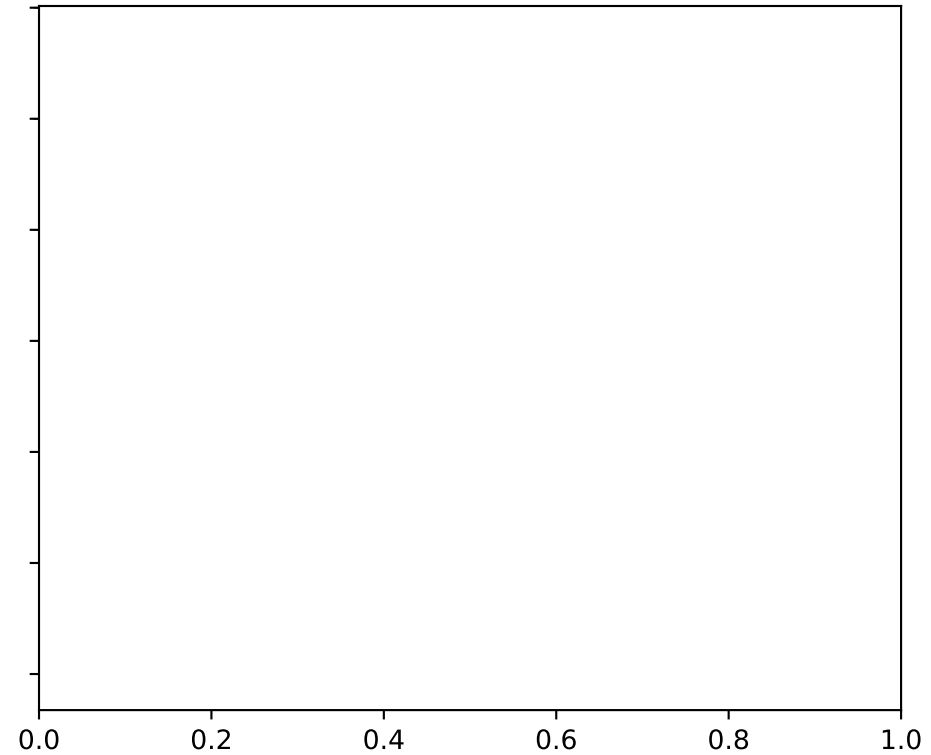
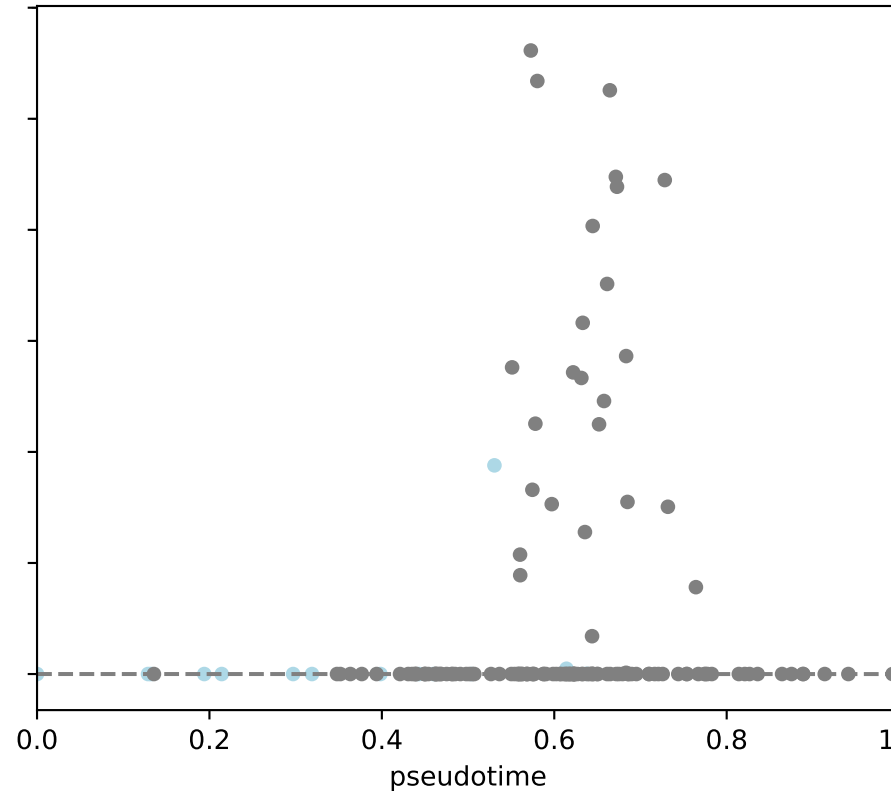
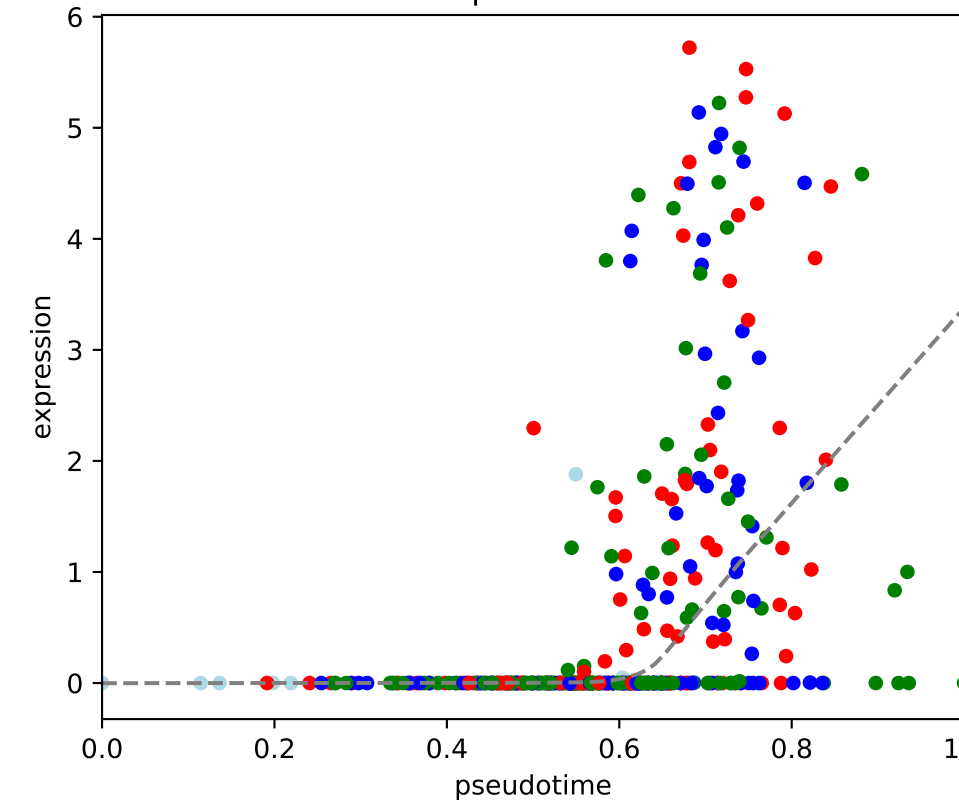
733 1spearman=-0.32



ENST00000594536 MRLN (myoregulin)

733 0spearman=0.57

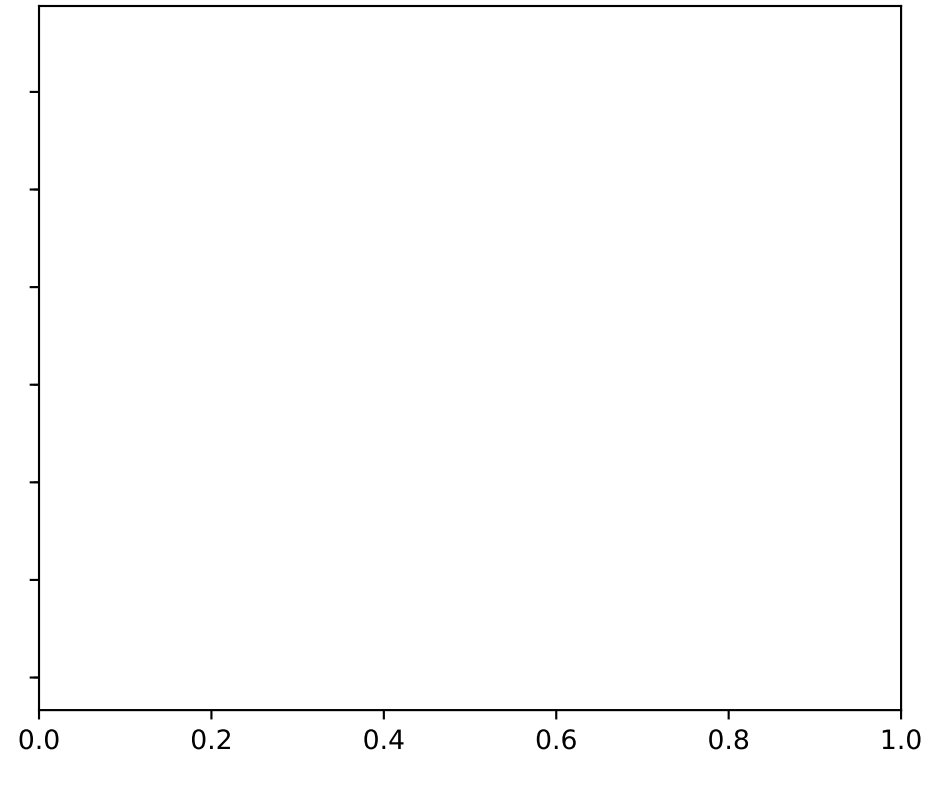
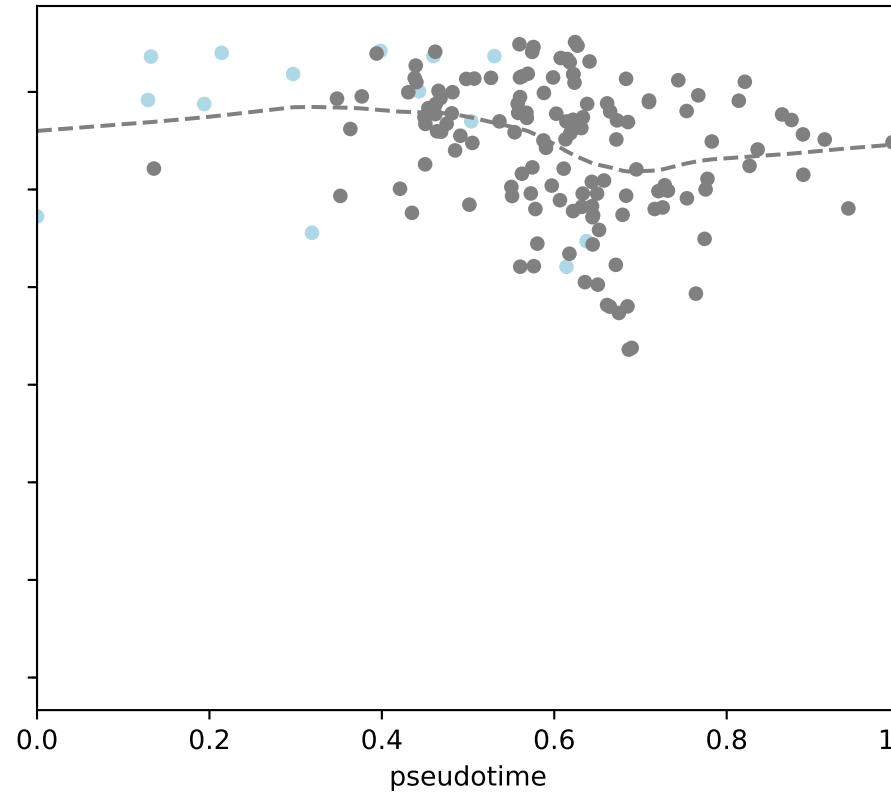
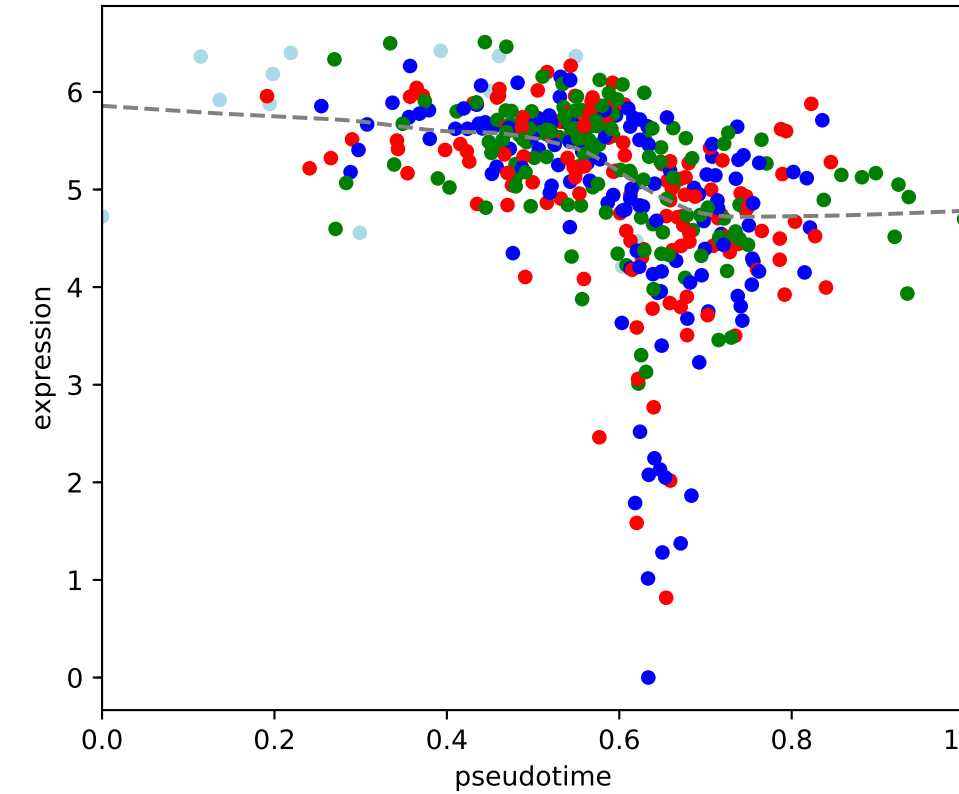
733 1spearman=0.19



ENST00000434005 EEF1A1P6 (eukaryotic translation elongation factor 1 alpha 1 pseudogene 6)

733 0spearman=-0.57

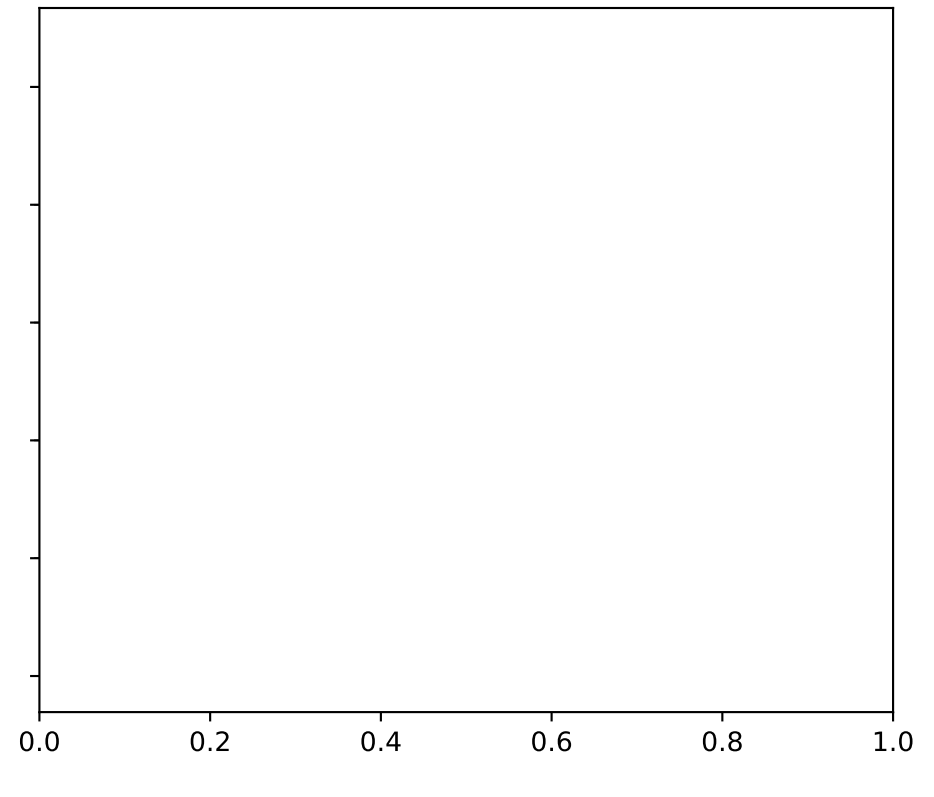
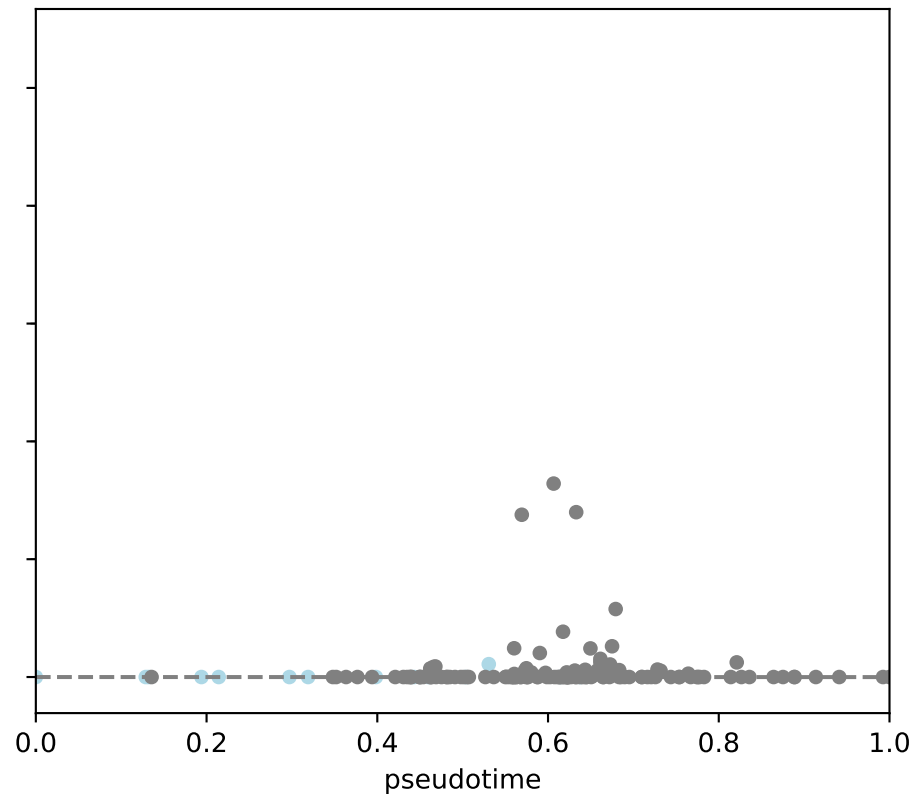
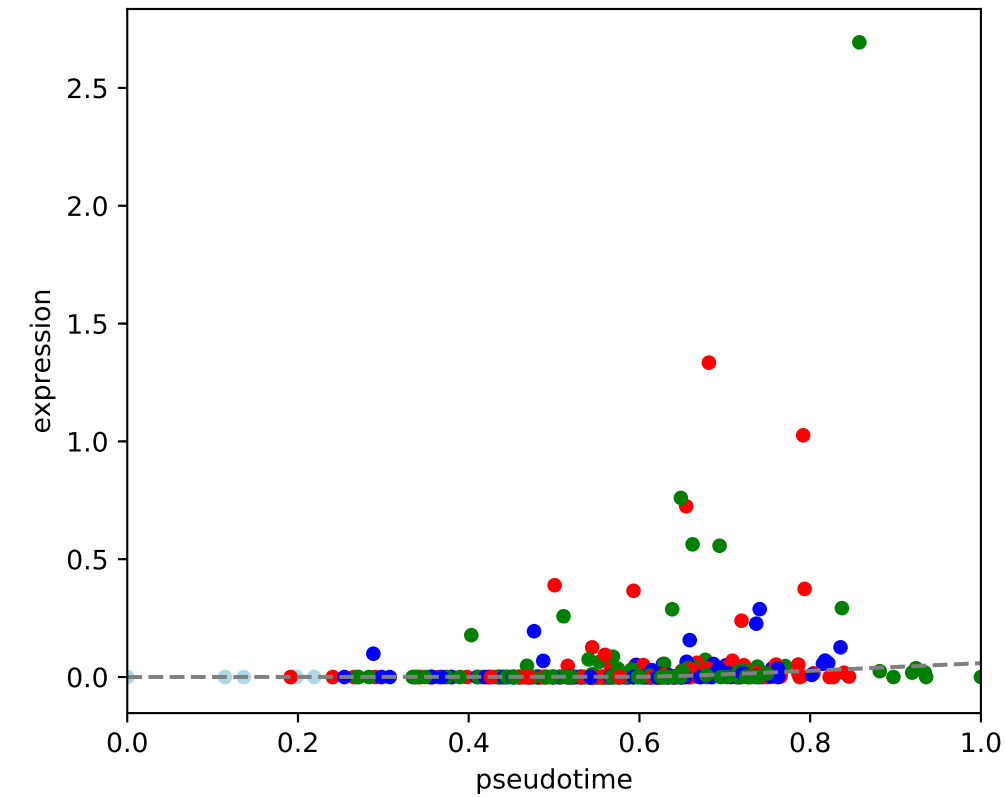
733 1spearman=-0.35



ENST00000621566 MRLN (myoregulin)

733 0spearman=0.57

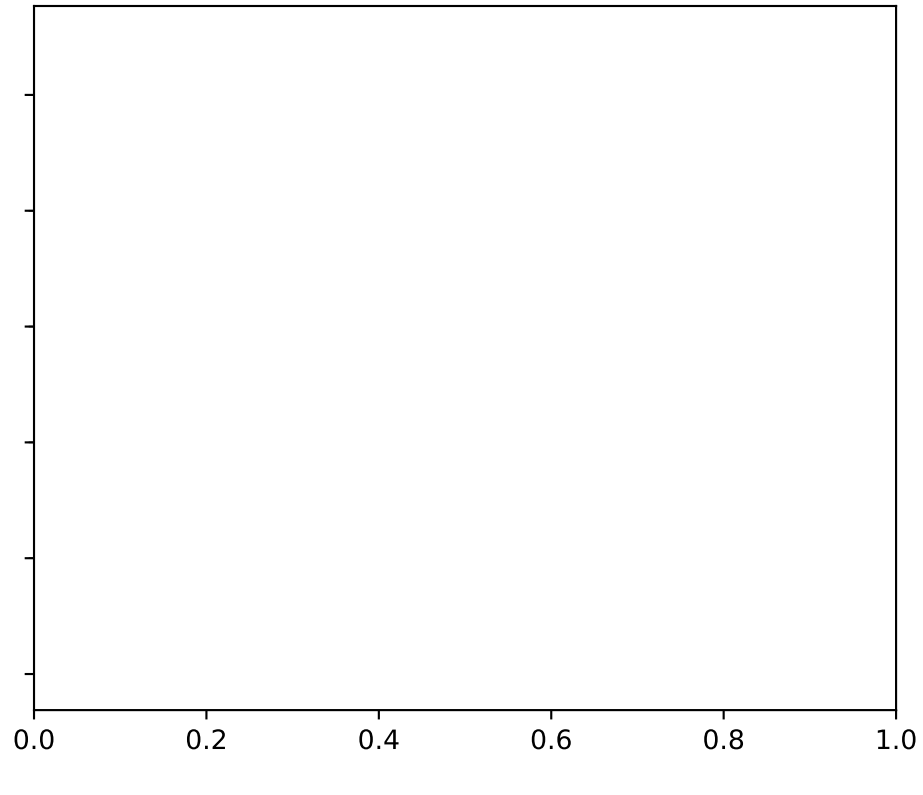
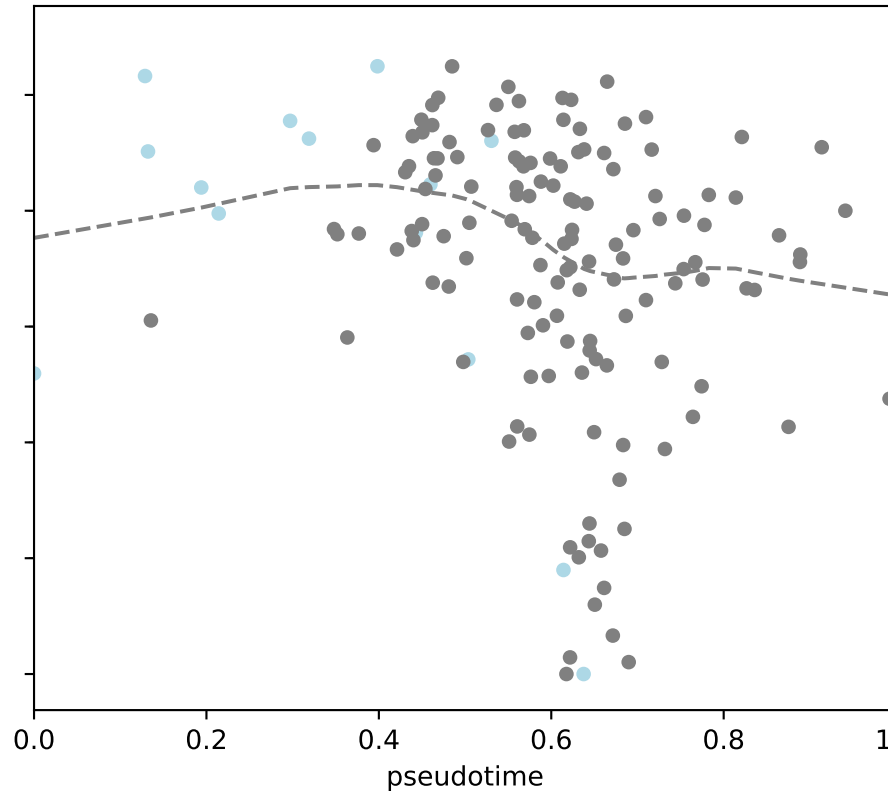
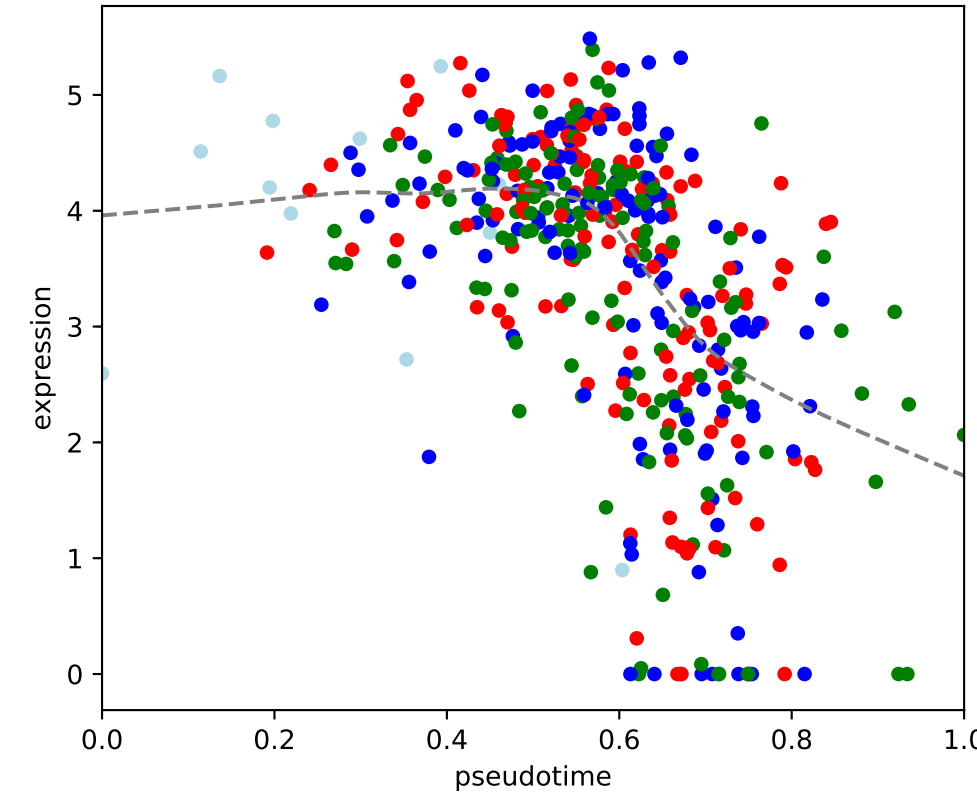
733 1spearman=0.17



ENST00000312726 GUK1 (guanylate kinase 1)

733 0spearman=-0.57

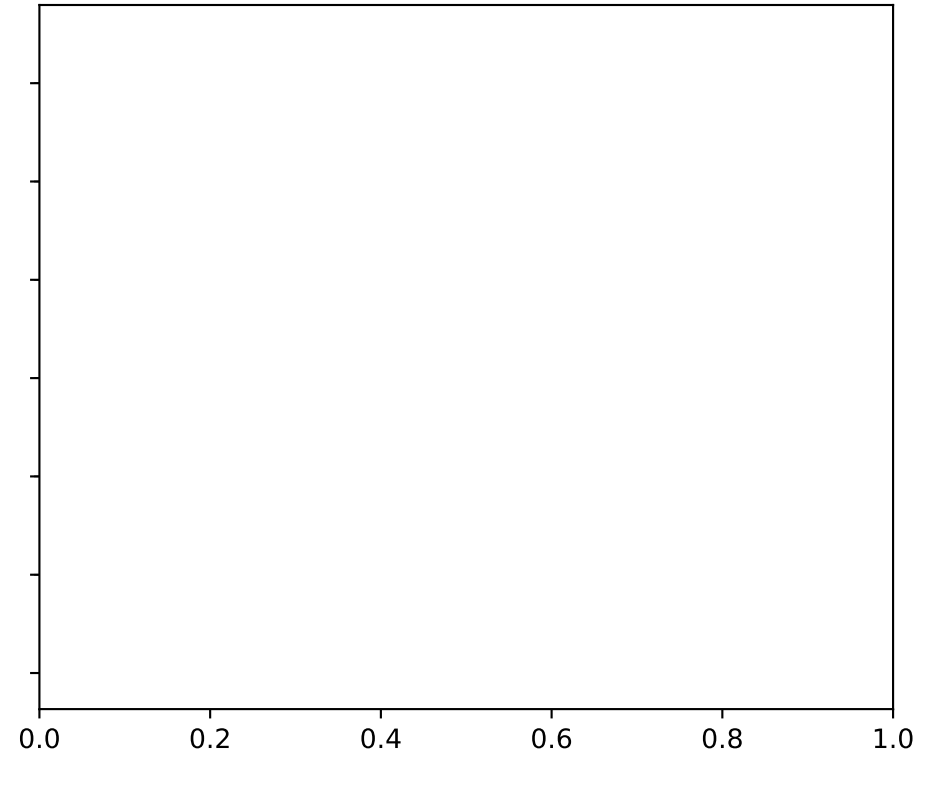
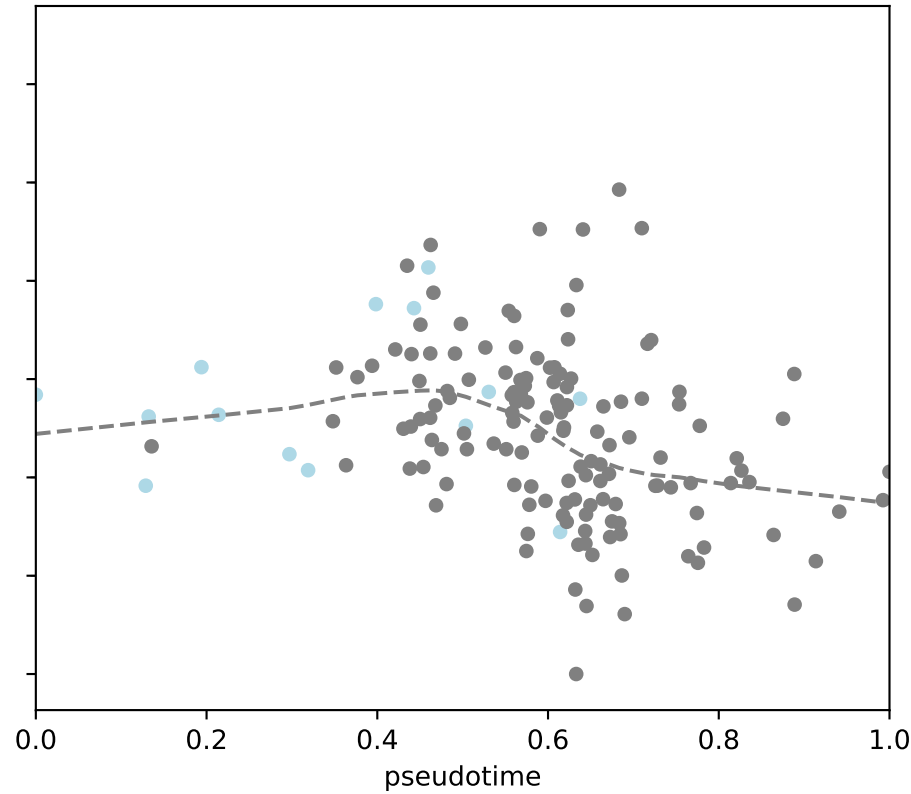
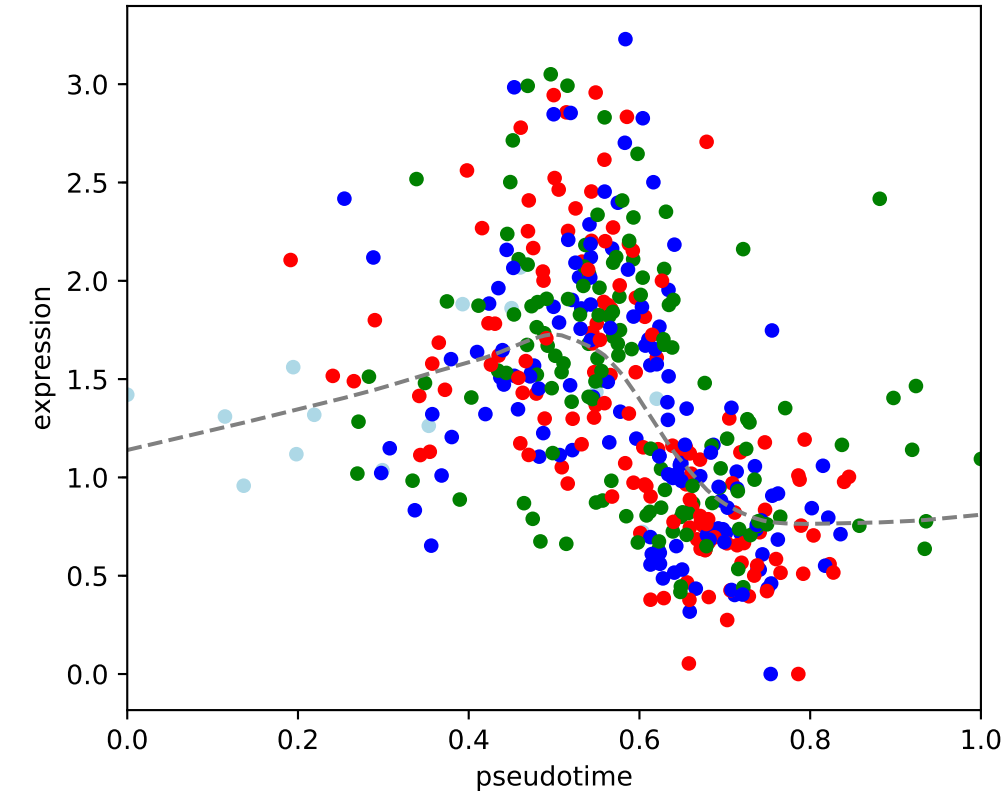
733 1spearman=-0.36



ENST00000549870 TUBA1B (tubulin alpha 1b)

733 0spearman=-0.57

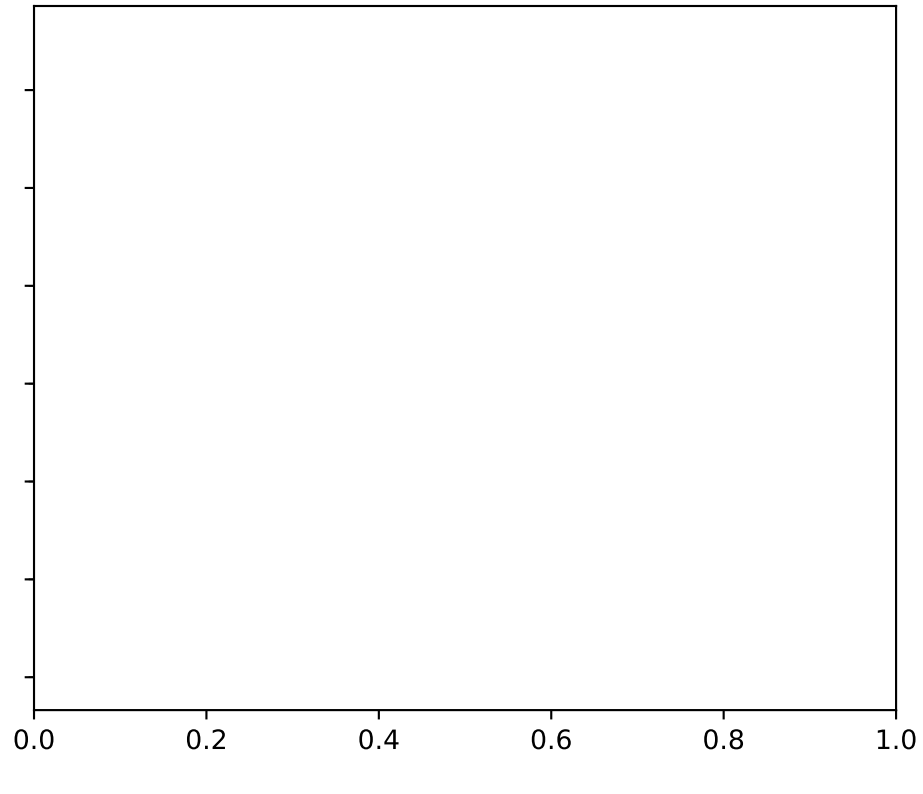
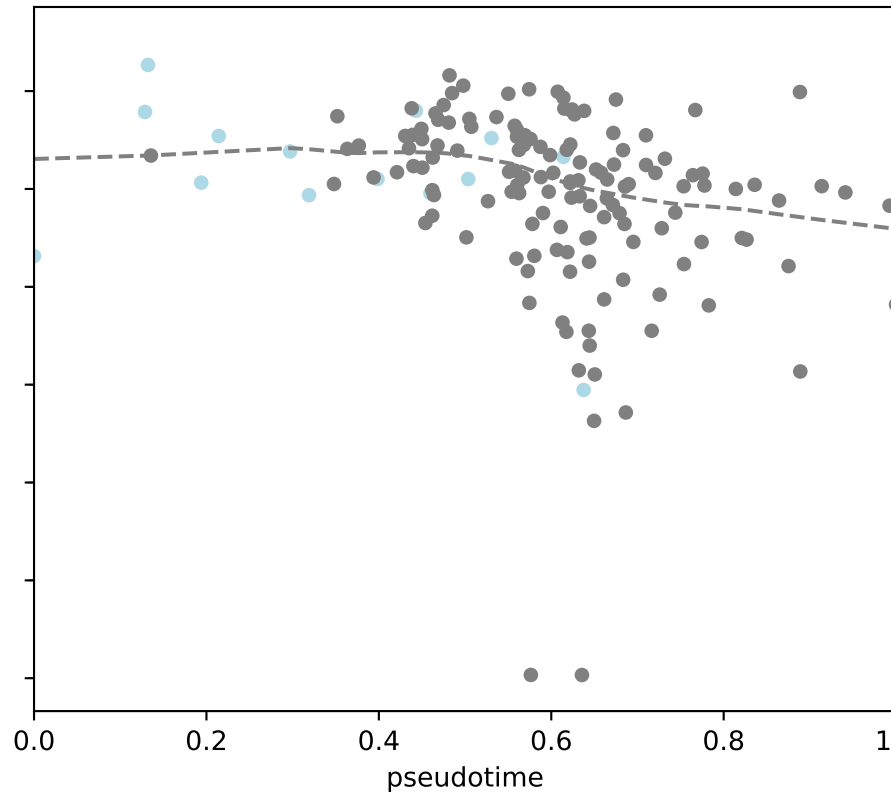
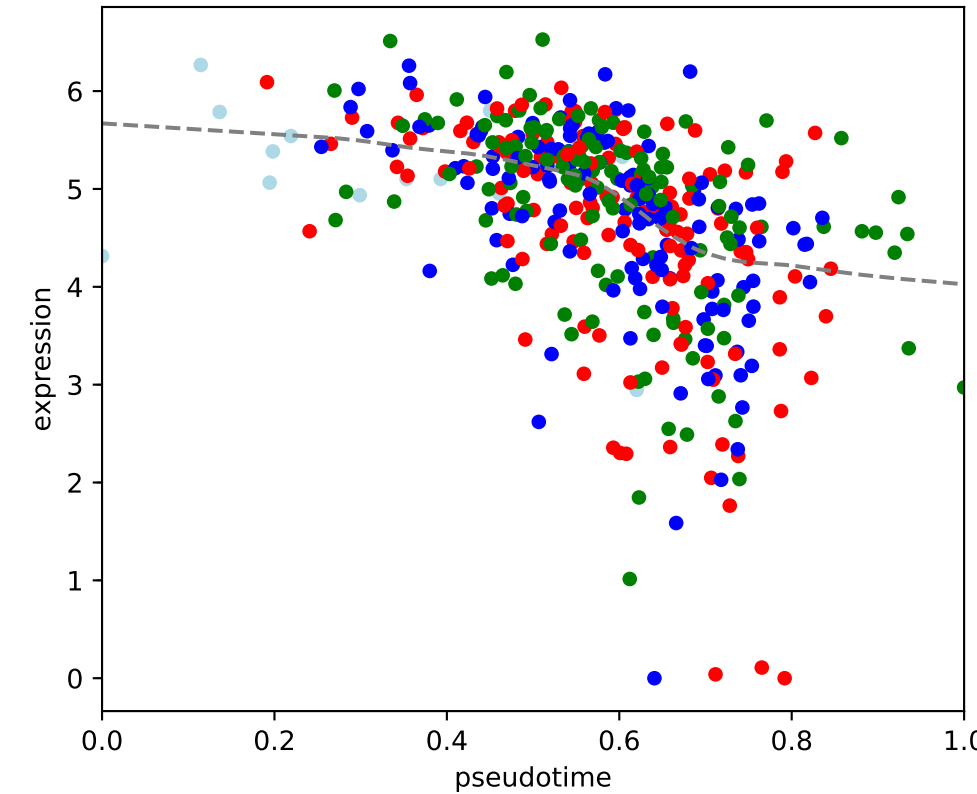
733 1spearman=-0.42



ENST00000422514 RPL23A (ribosomal protein L23a)

733 0spearman=-0.57

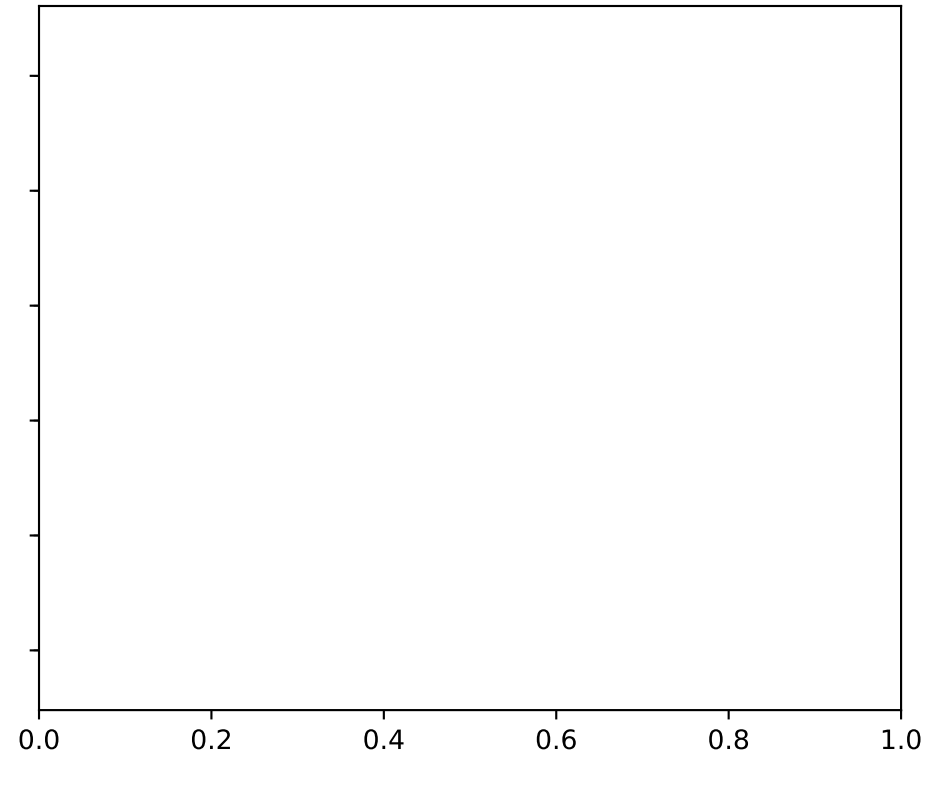
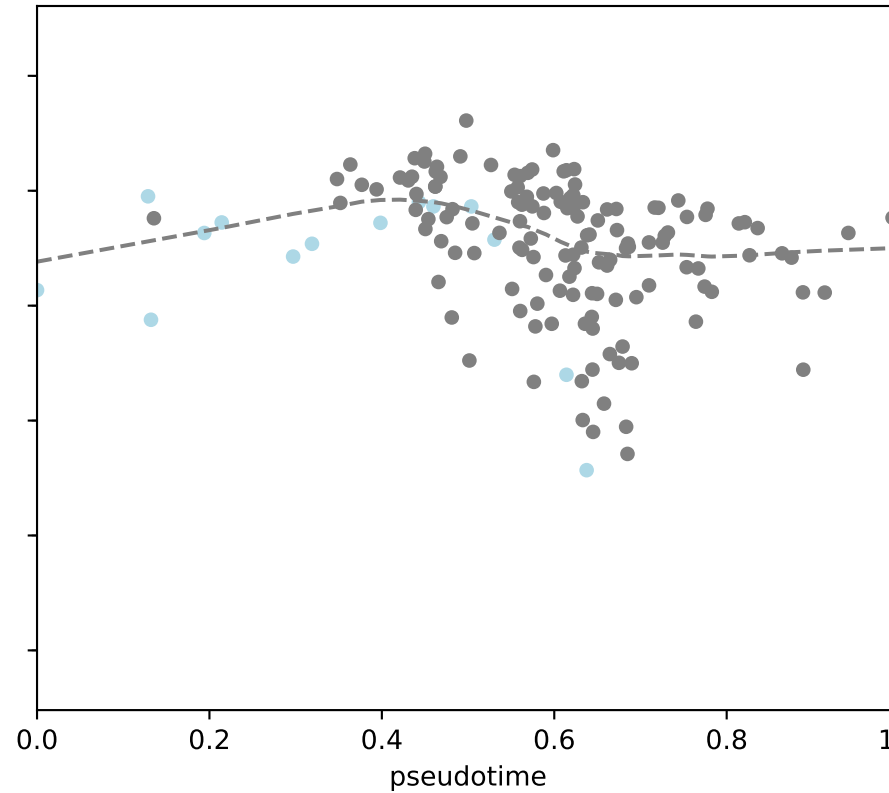
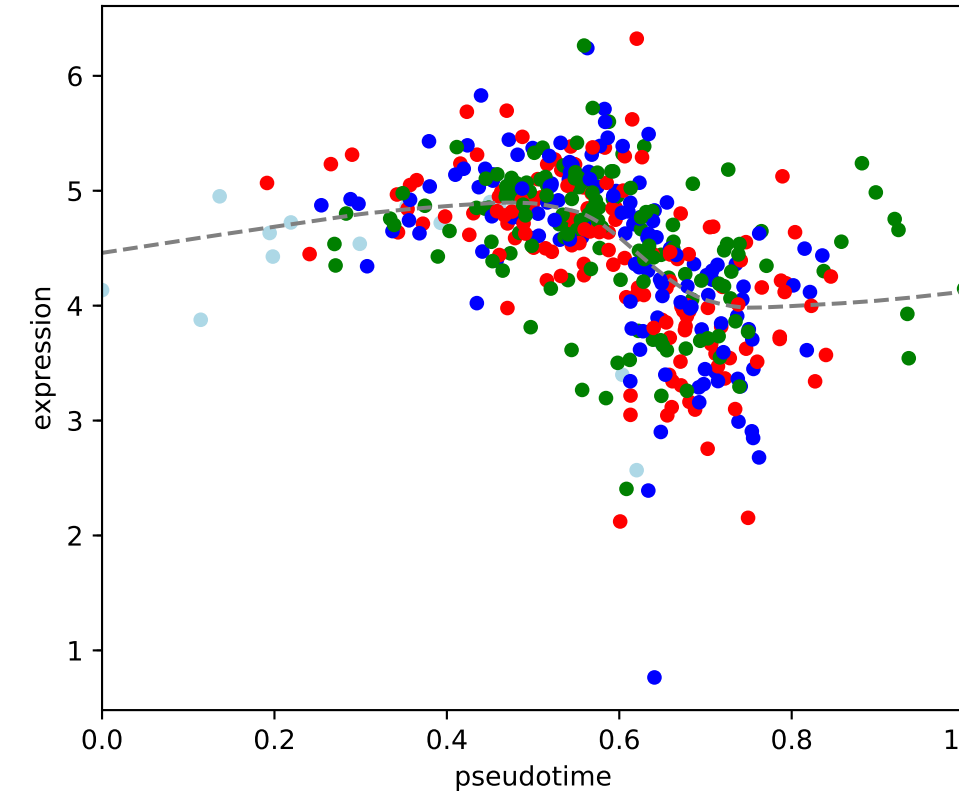
733 1spearman=-0.41



ENST00000358455 PPIAP22 (peptidylprolyl isomerase A pseudogene 22)

733 0spearman=-0.57

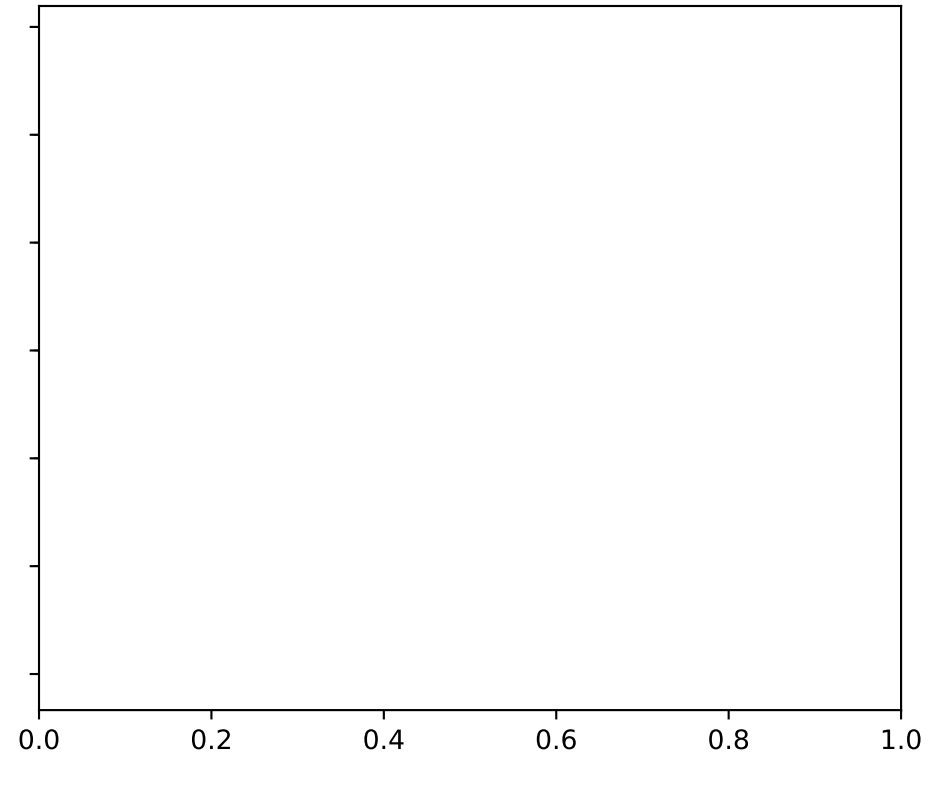
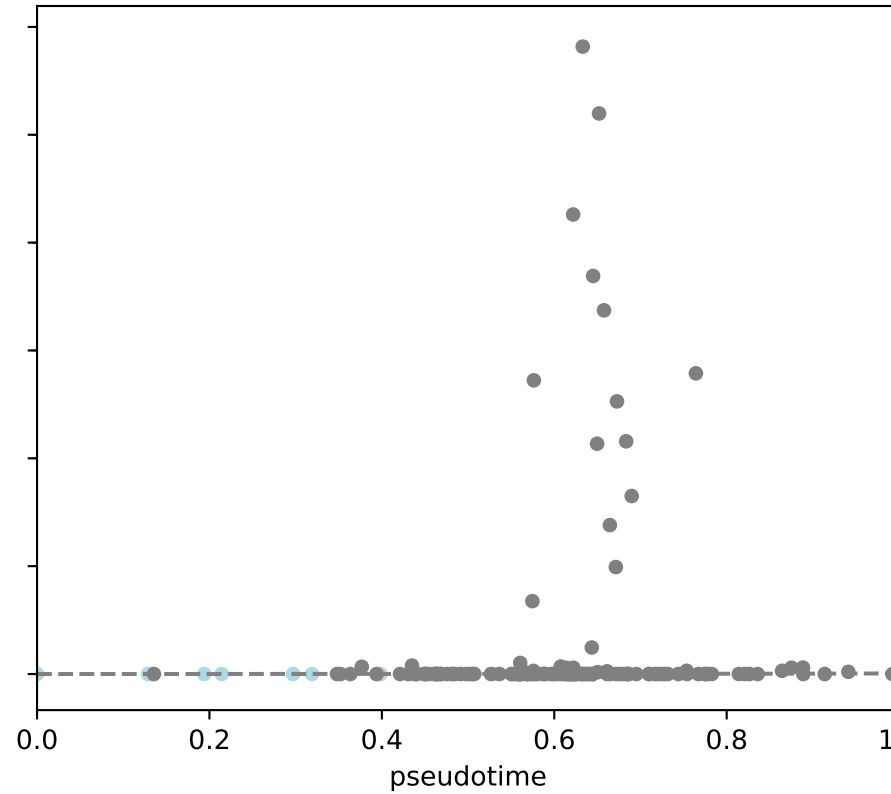
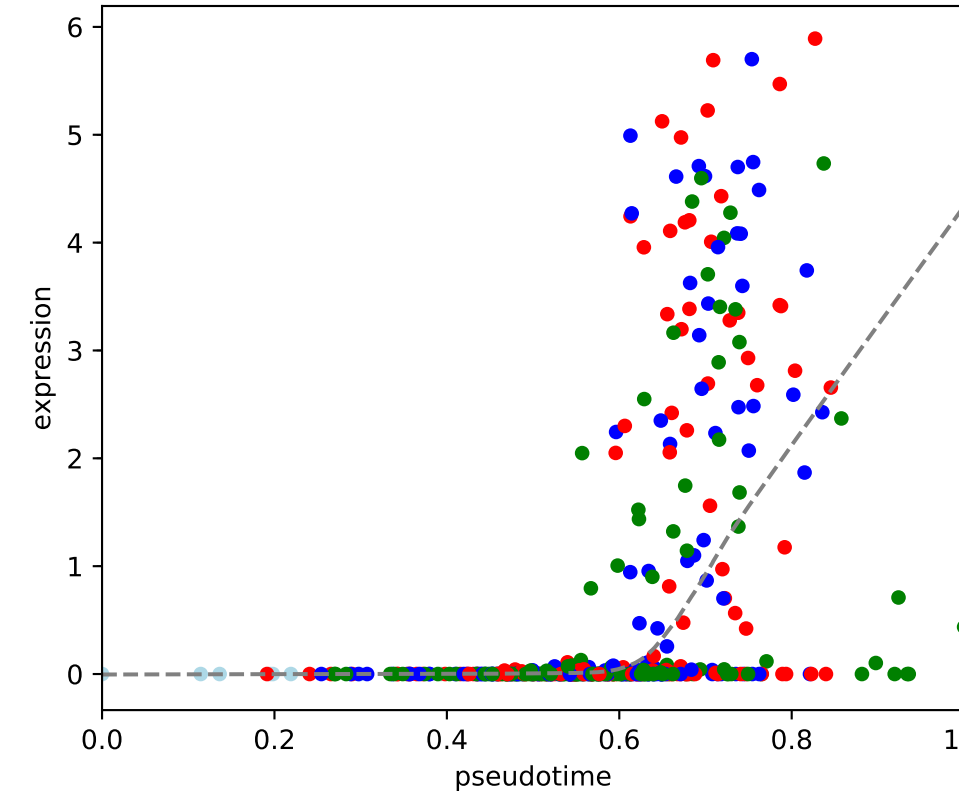
733 1spearman=-0.42



ENST00000470589 GABRR3 (gamma-aminobutyric acid type A receptor rho3 subunit (gene/pseudogene))

733 0spearman=0.57

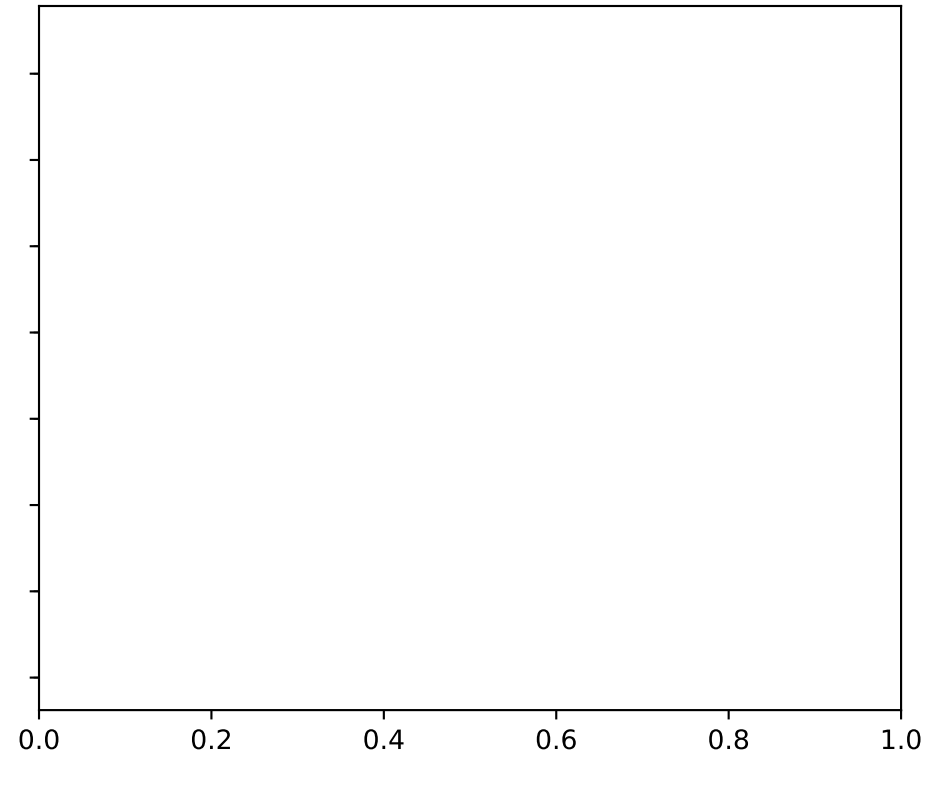
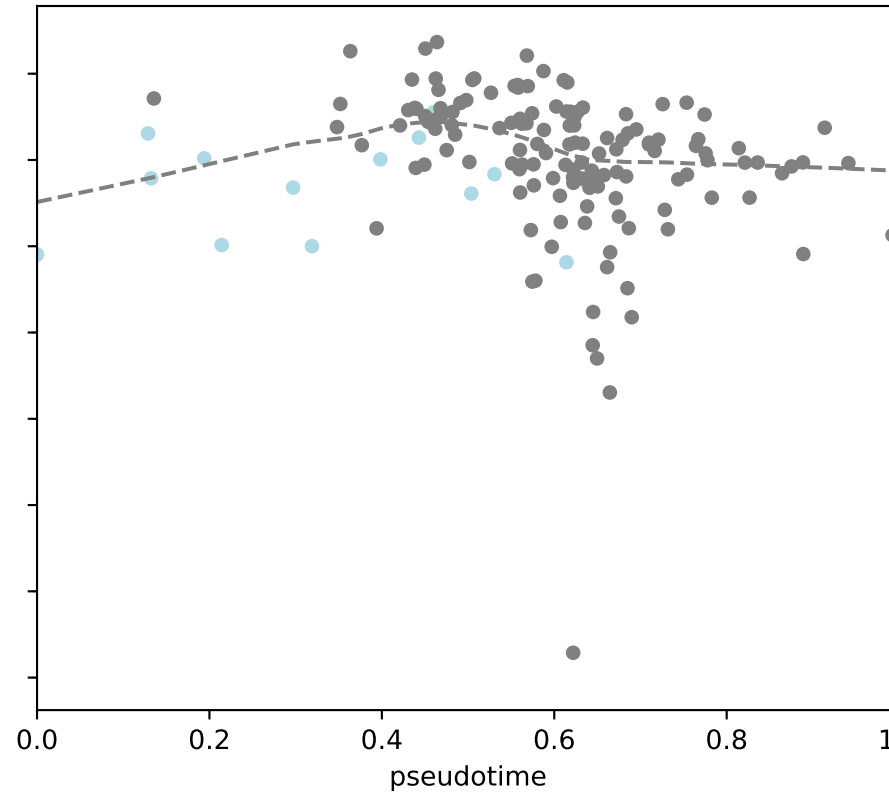
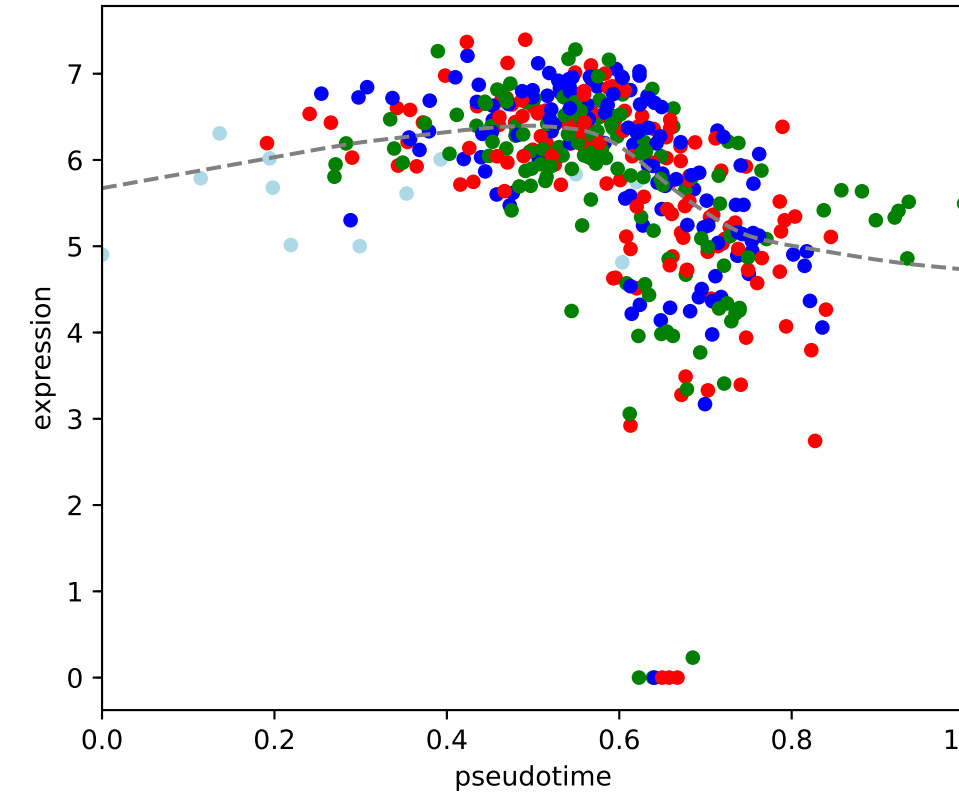
733 1spearman=0.23



ENST00000246554 COX6B1 (cytochrome c oxidase subunit 6B1)

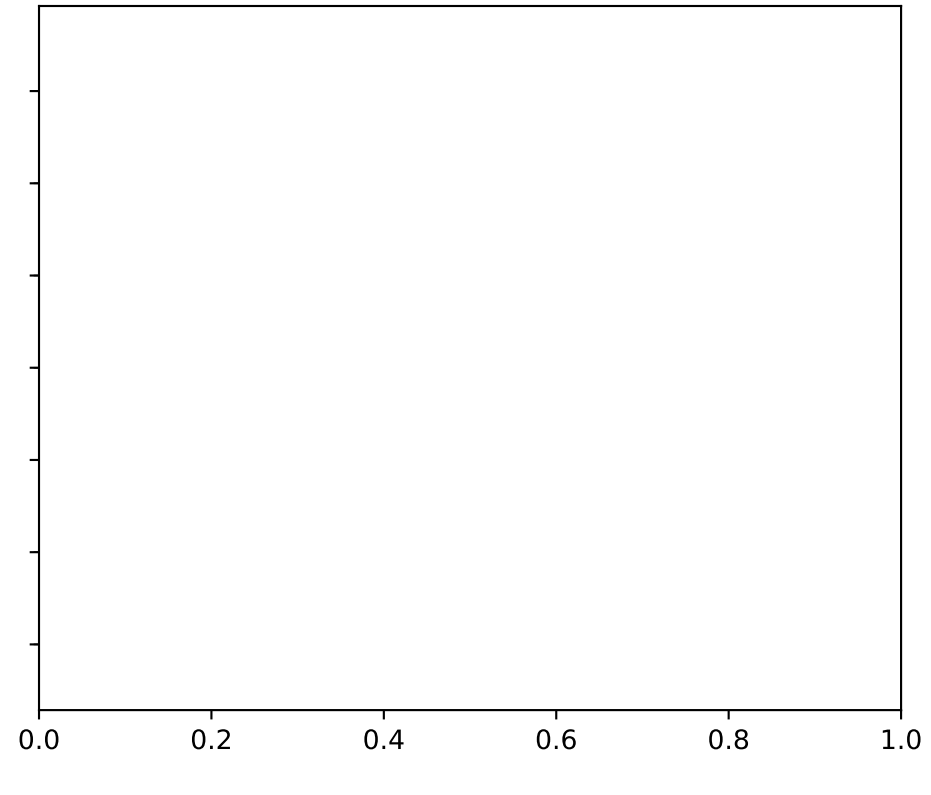
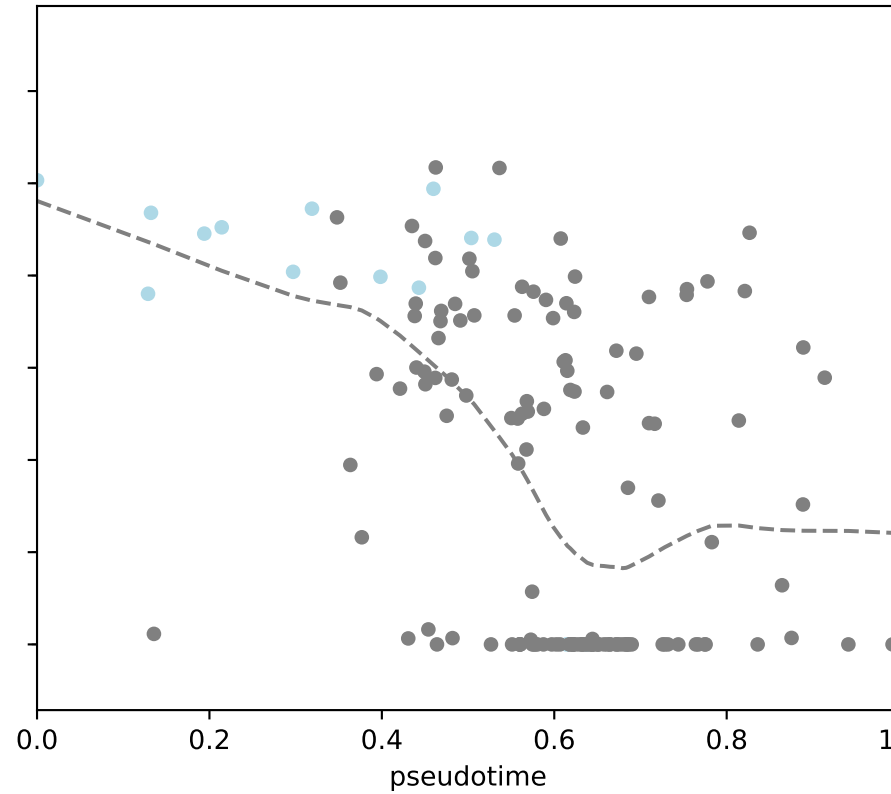
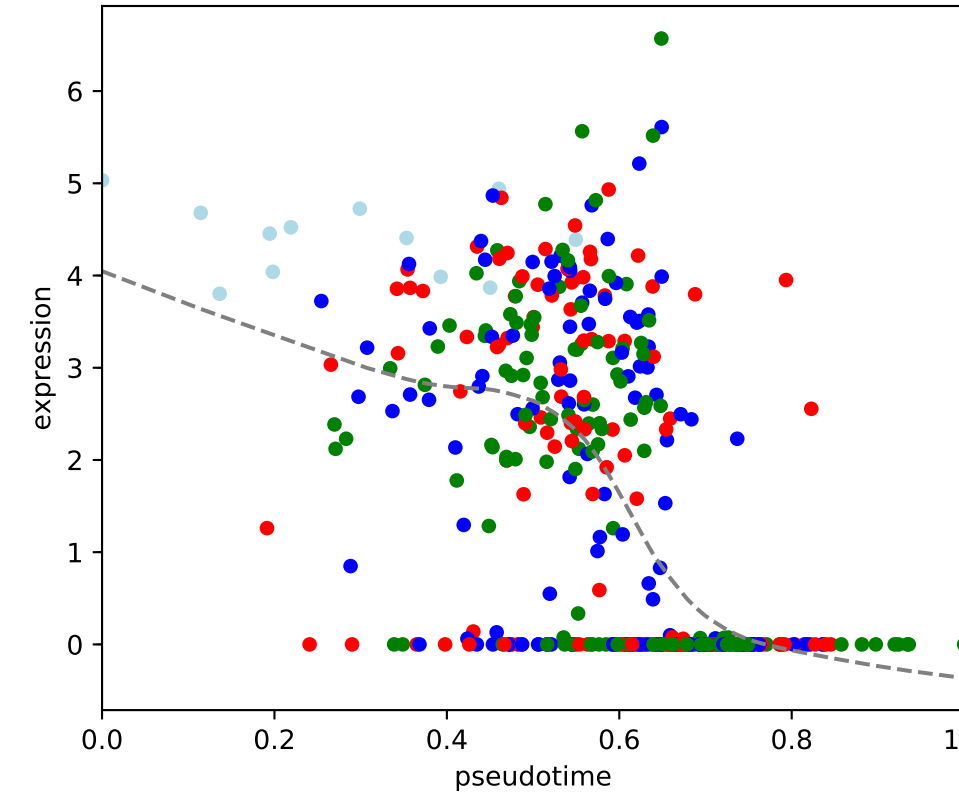
733 0spearman=-0.57

733 1spearman=-0.34



733 0spearman=-0.57

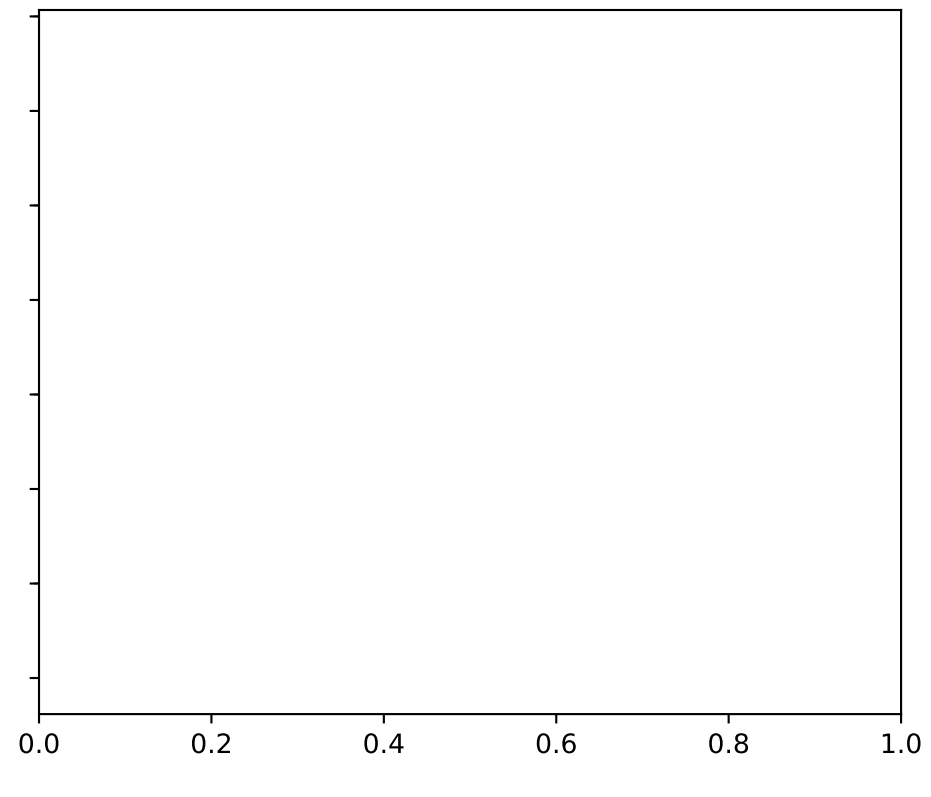
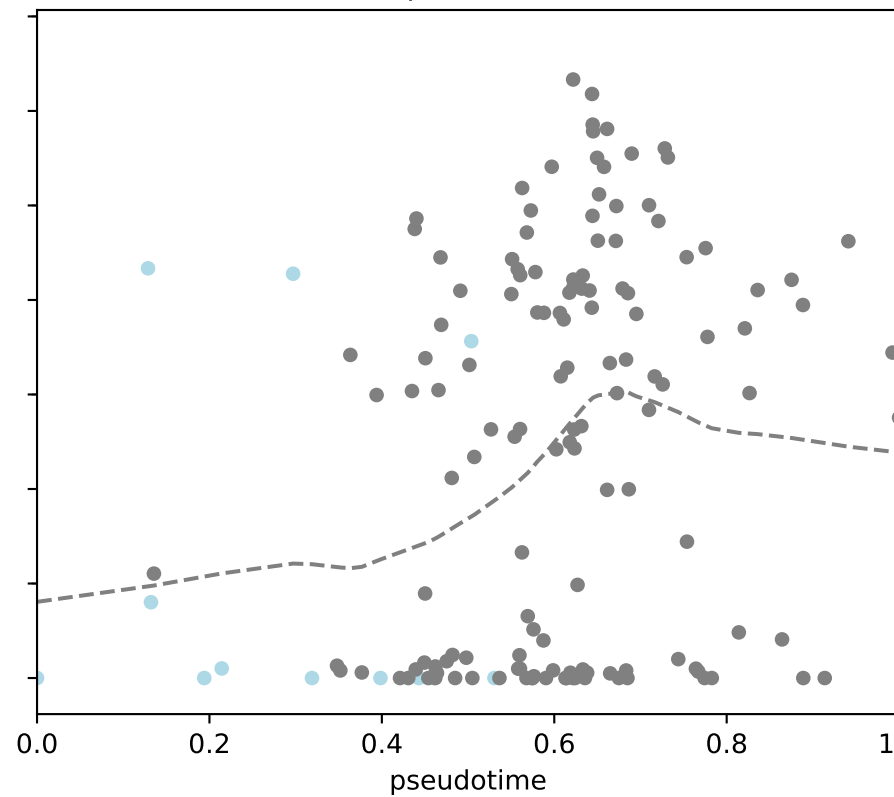
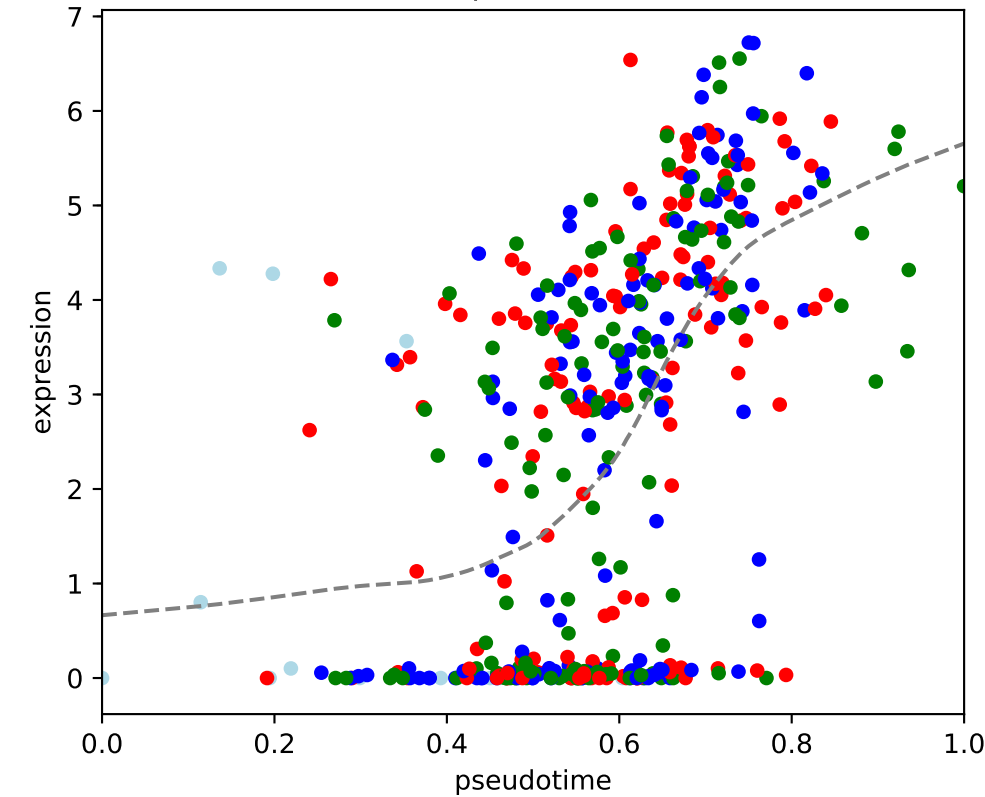
733 1spearman=-0.48



ENST00000436880 MIR181A1HG (MIR181A1 host gene)

733 0spearman=0.57

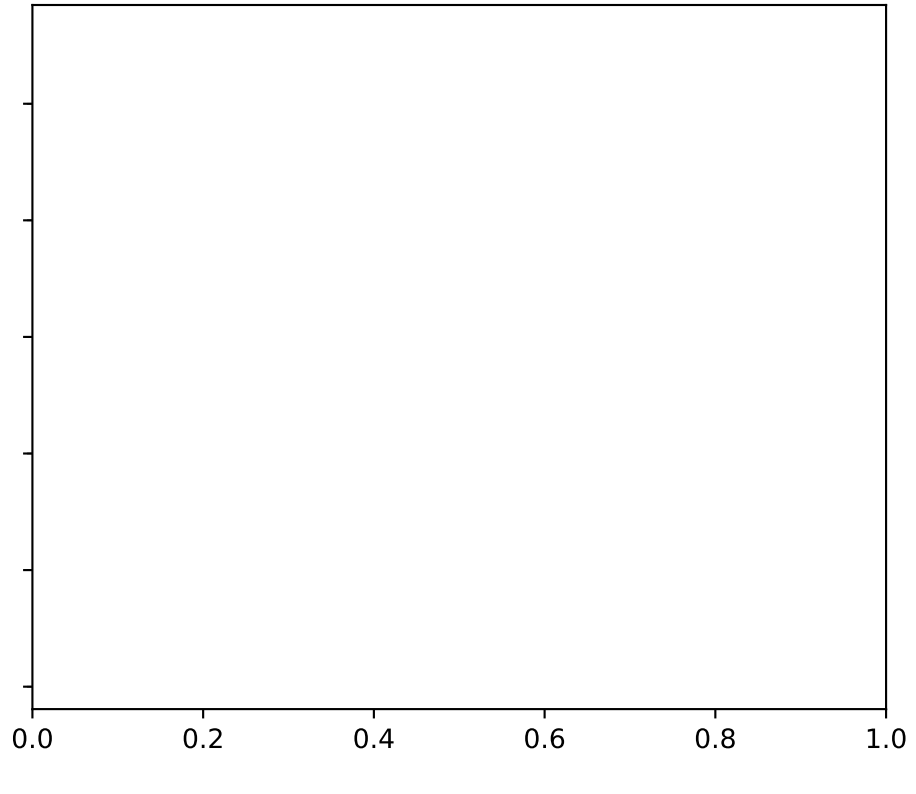
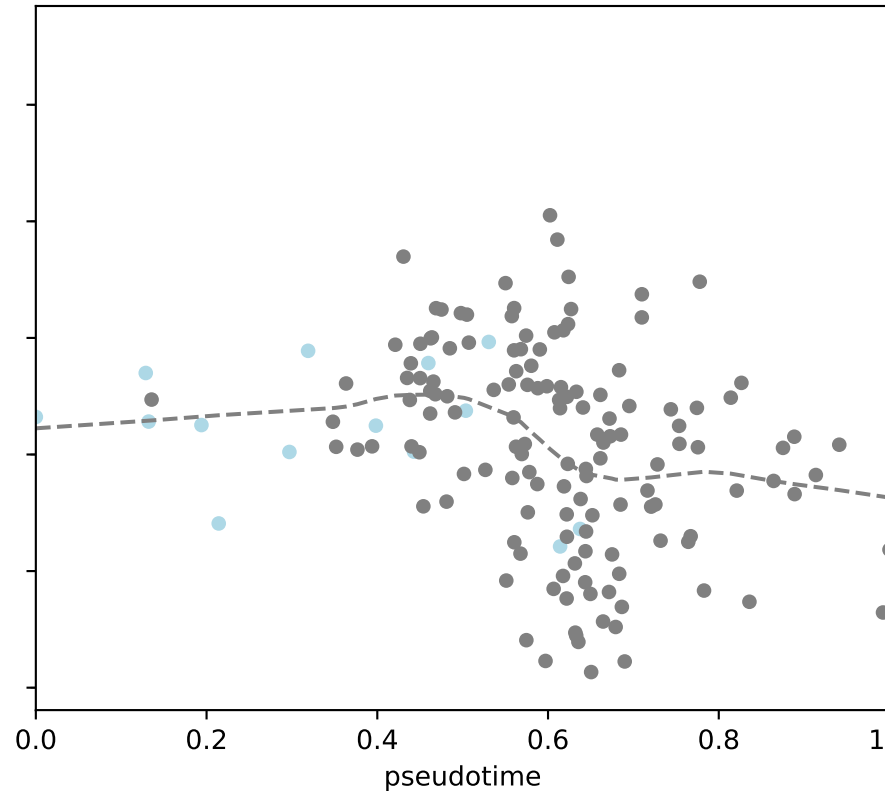
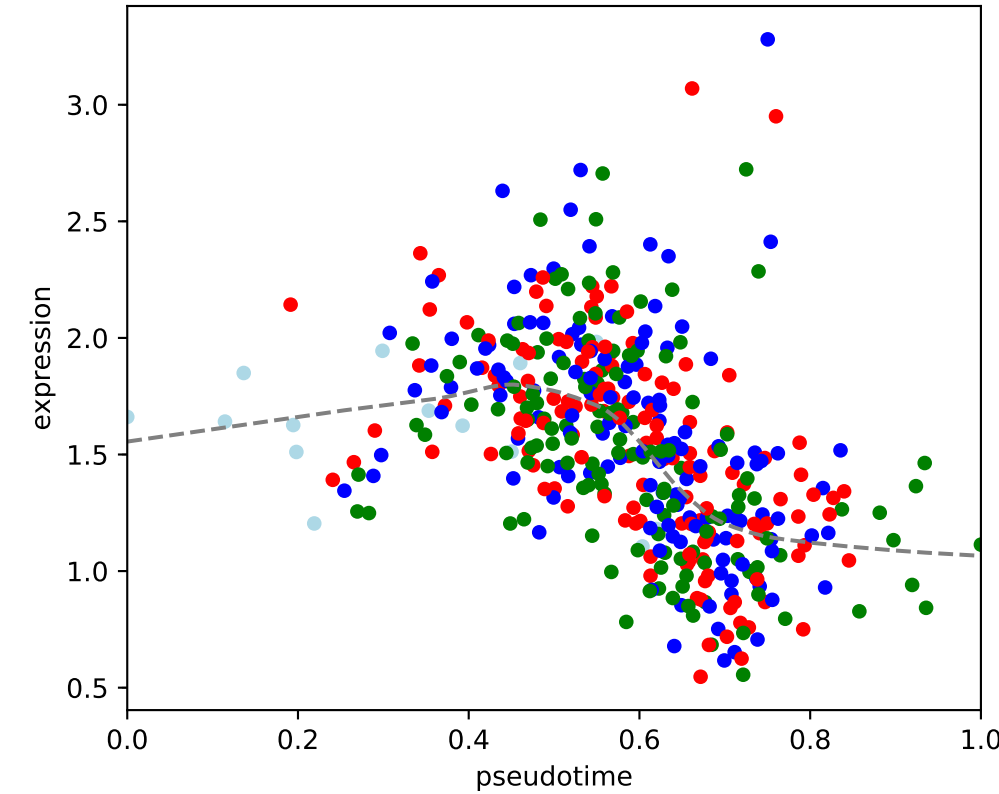
733 1spearman=0.26



ENST00000553007 ATP5F1B (ATP synthase F1 subunit beta)

733 0spearman=-0.57

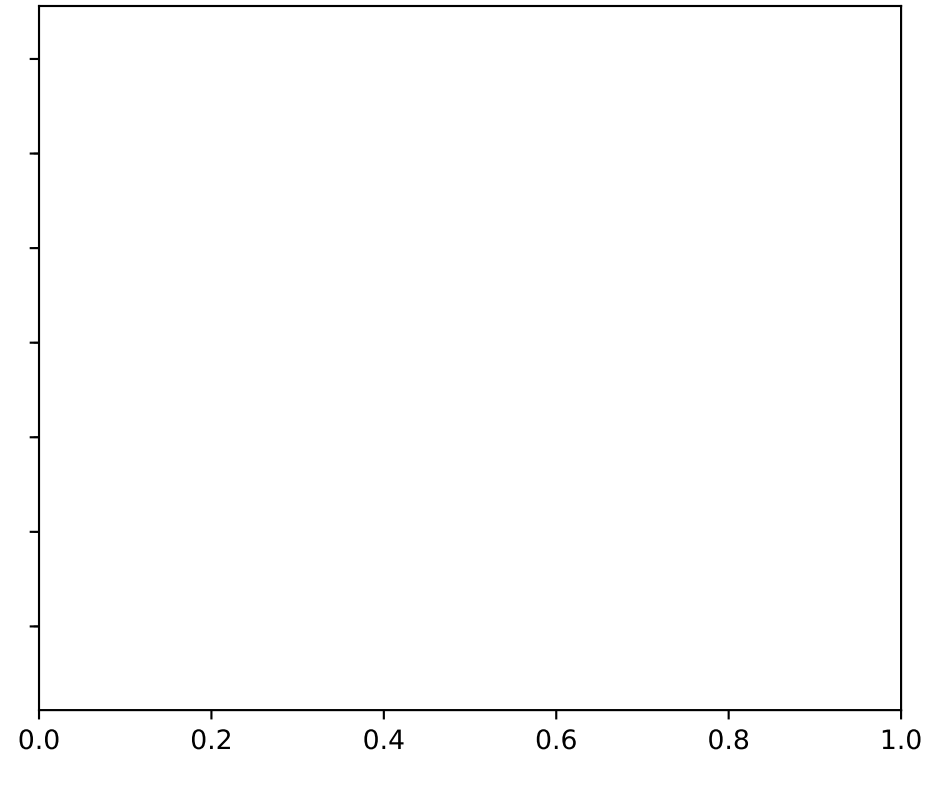
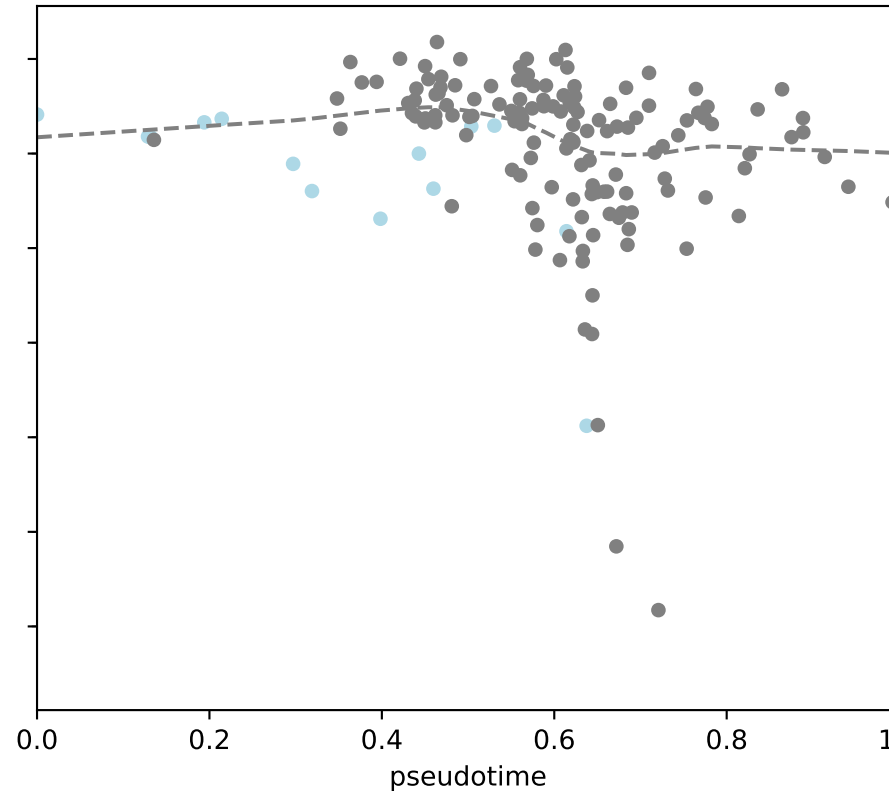
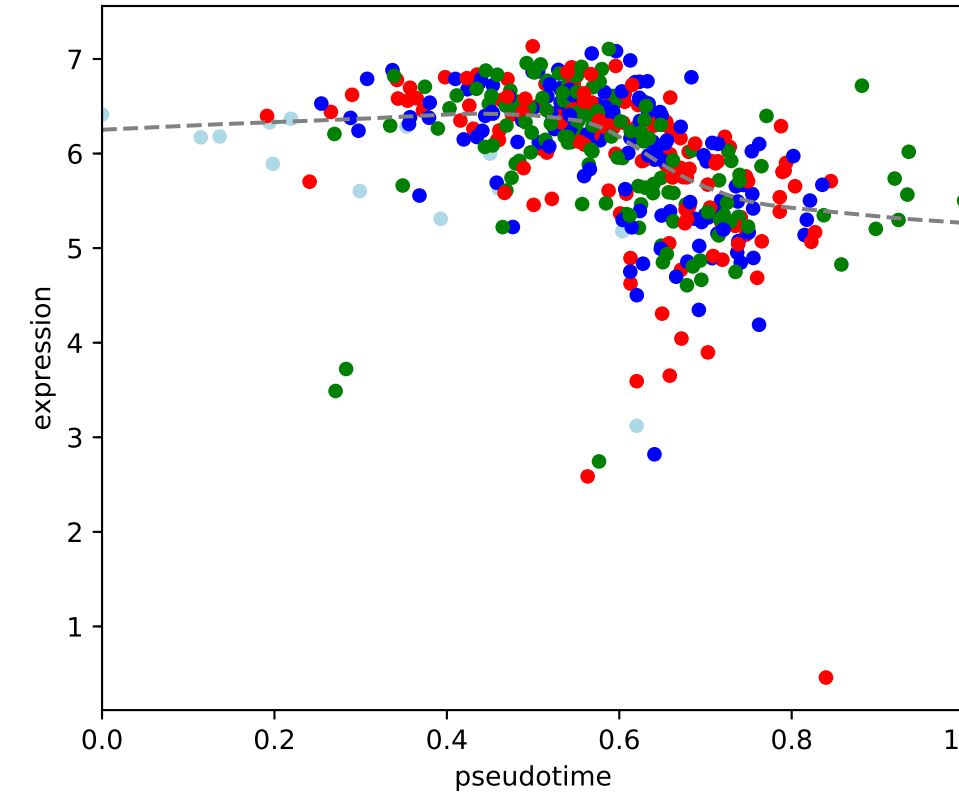
733 1spearman=-0.37



ENST00000370089 COX7A2 (cytochrome c oxidase subunit 7A2)

733 0spearman=-0.57

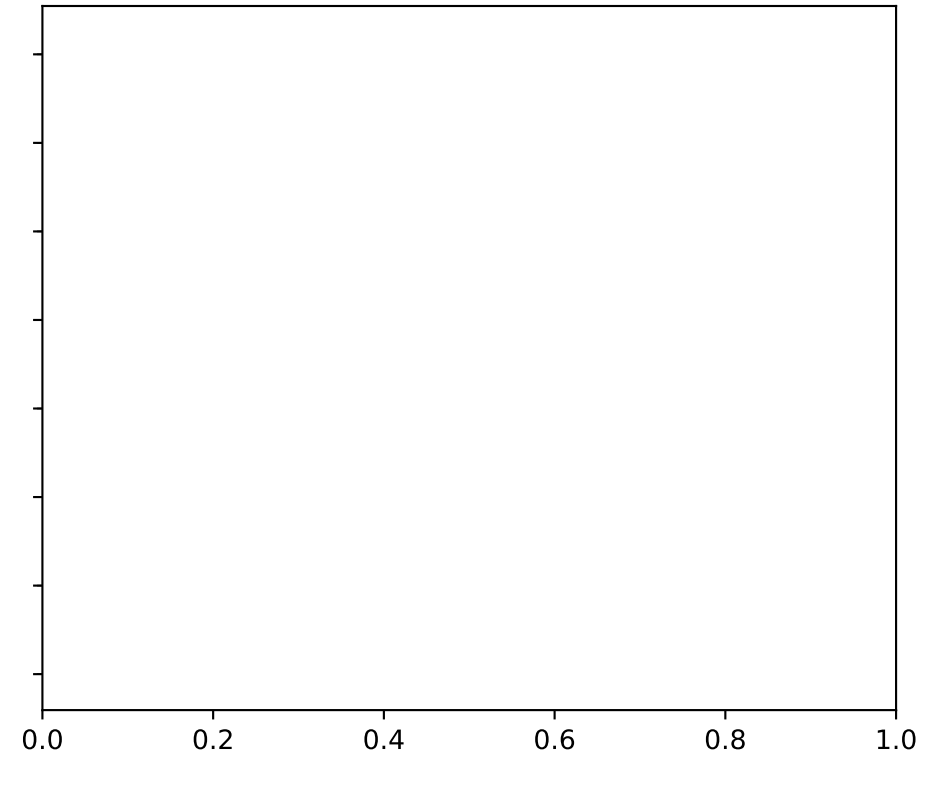
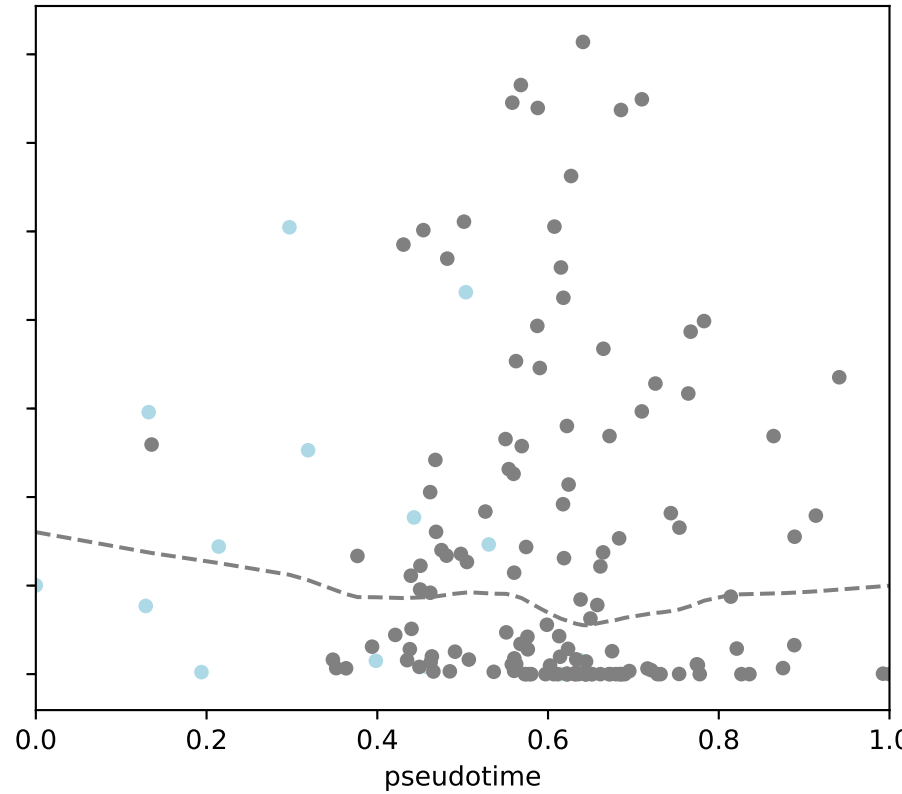
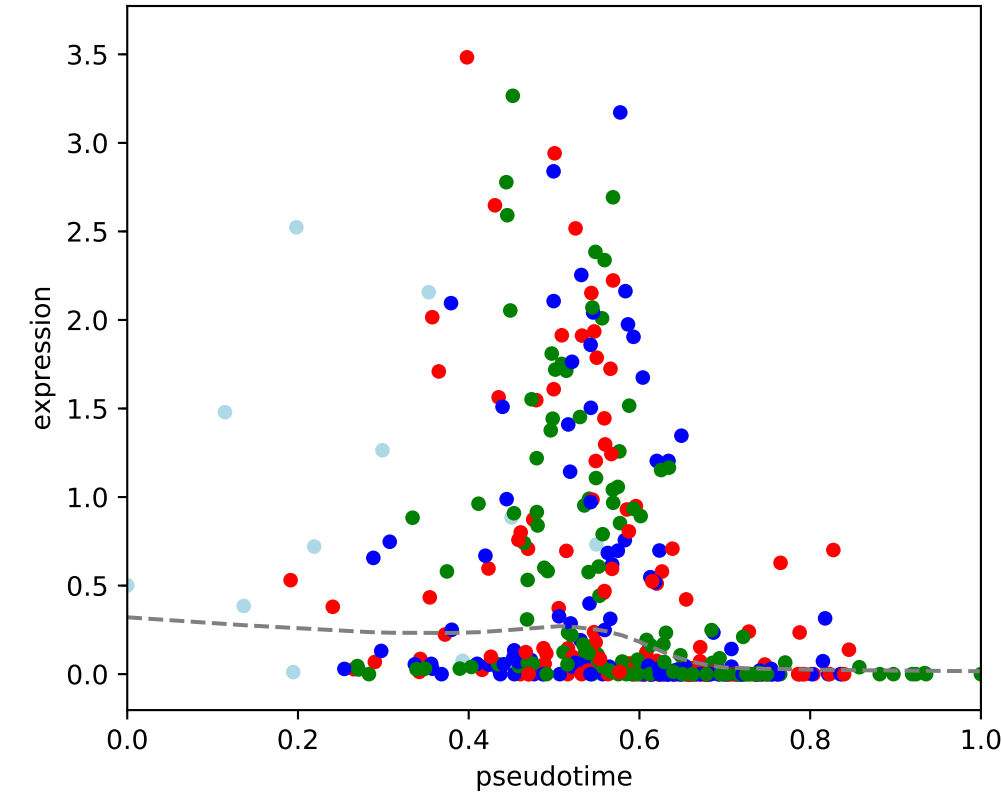
733 1spearman=-0.37



ENST00000507466 SPON2 (spondin 2)

733 0spearman=-0.57

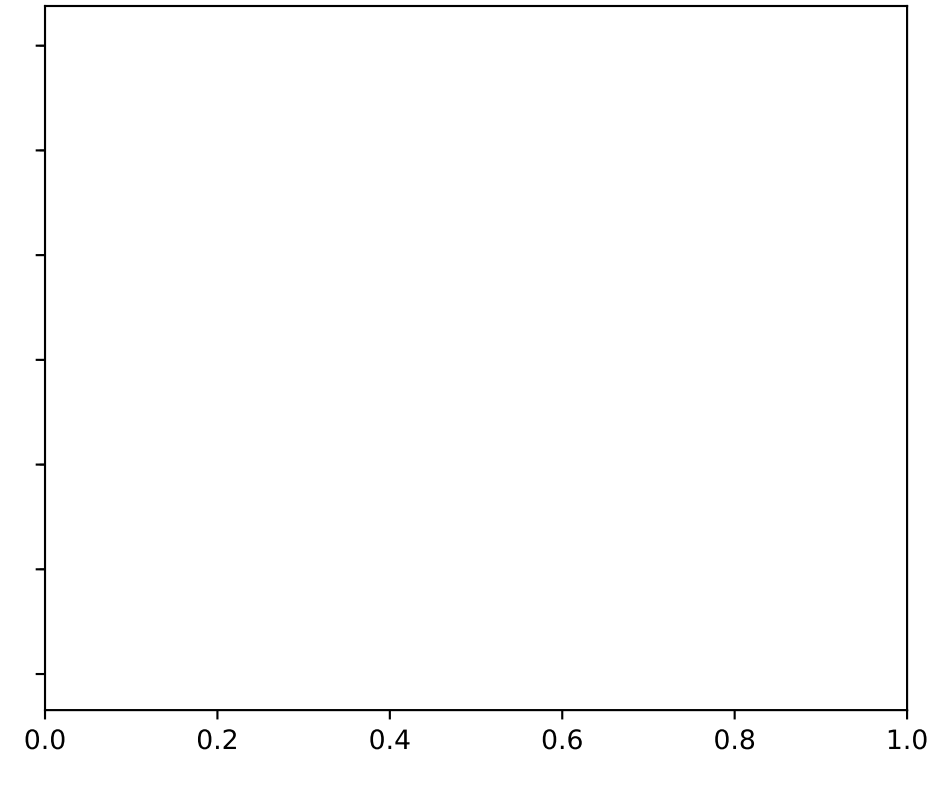
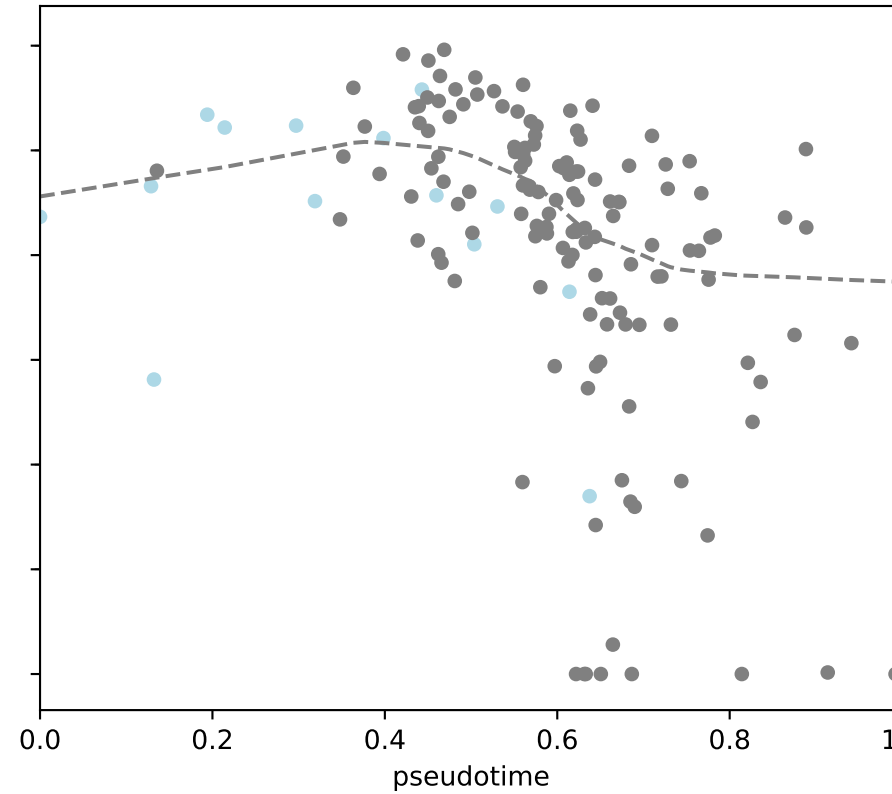
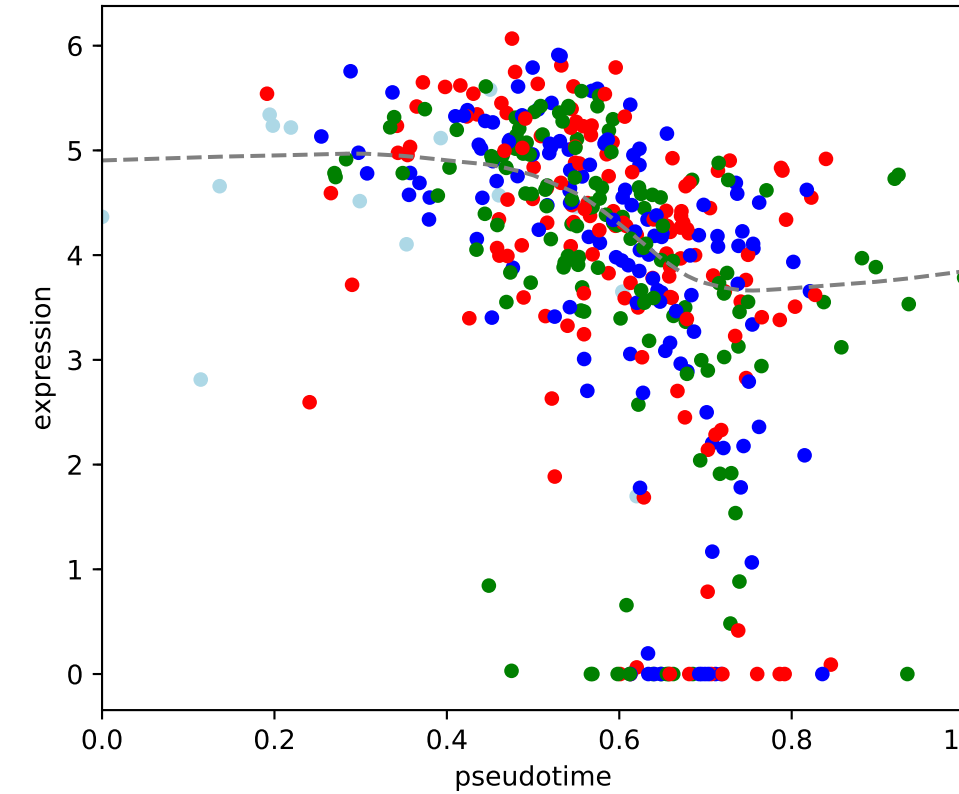
733 1spearman=-0.20



ENST00000419113 NDUFA3 (NADH:ubiquinone oxidoreductase subunit A3)

733 0spearman=-0.57

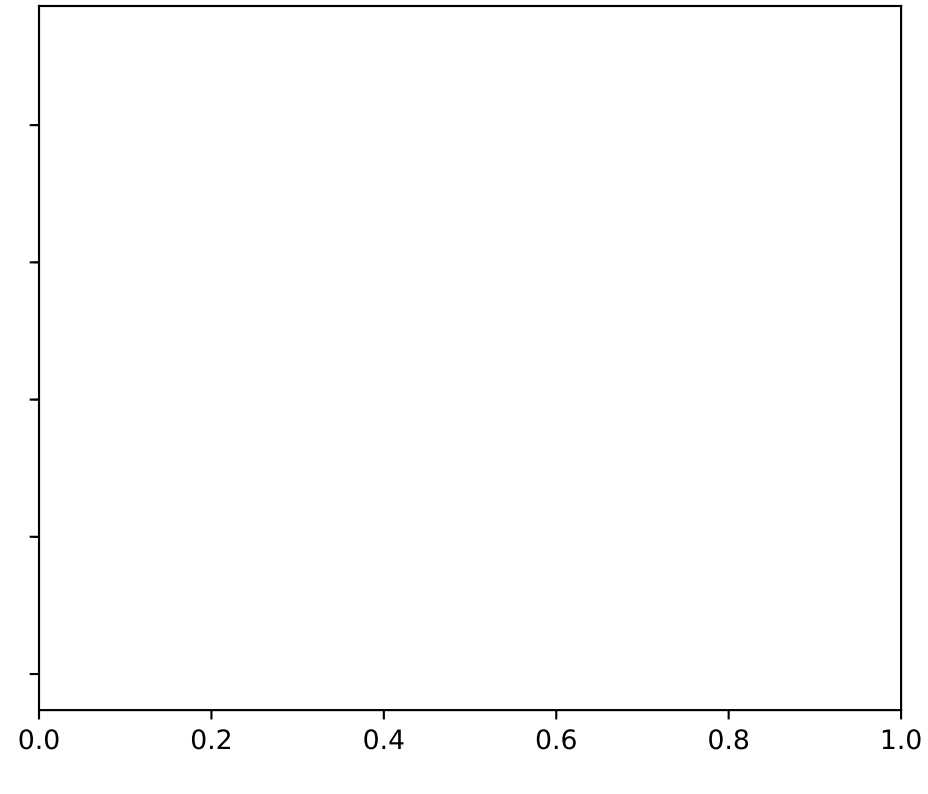
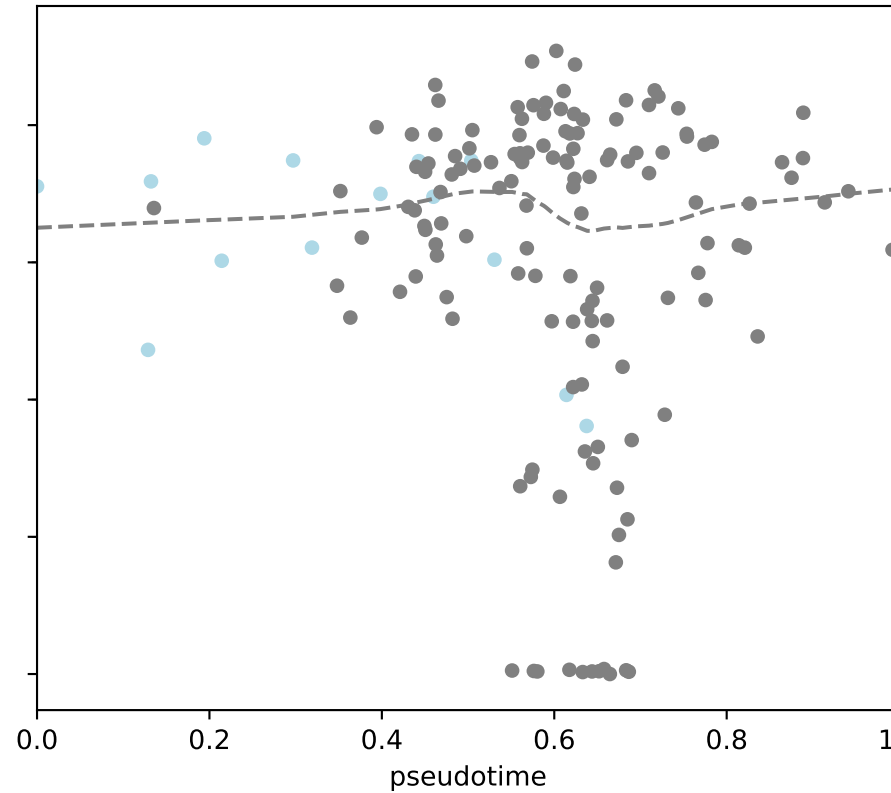
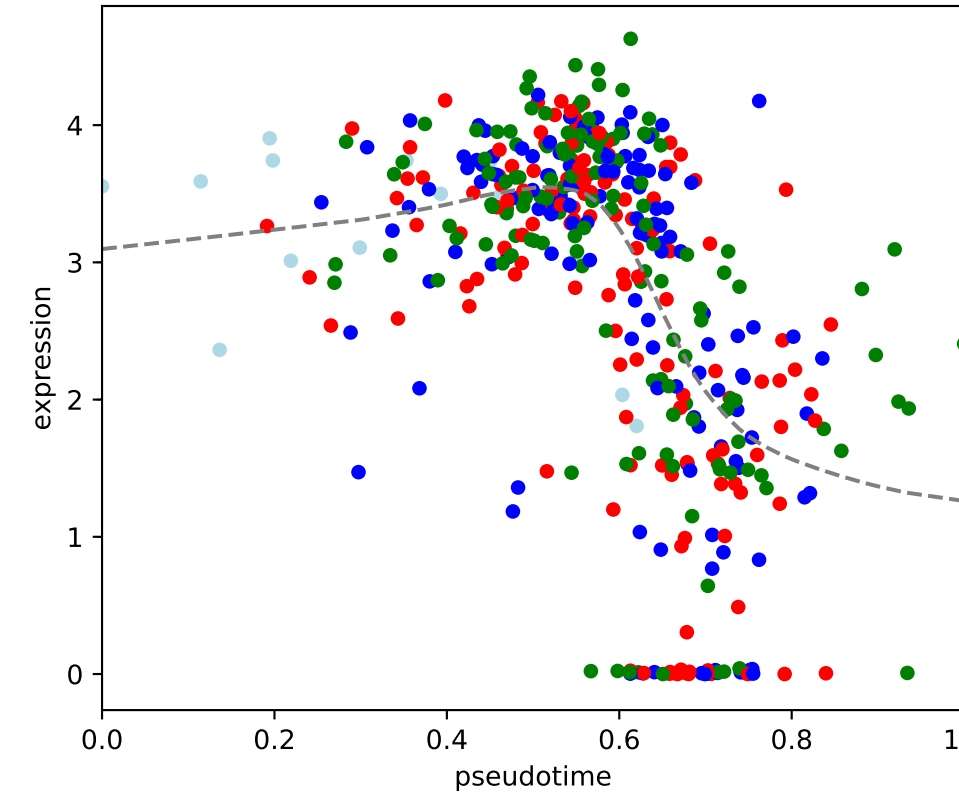
733 1spearman=-0.58



ENST00000369851 GNAI3 (G protein subunit alpha i3)

733 0spearman=-0.57

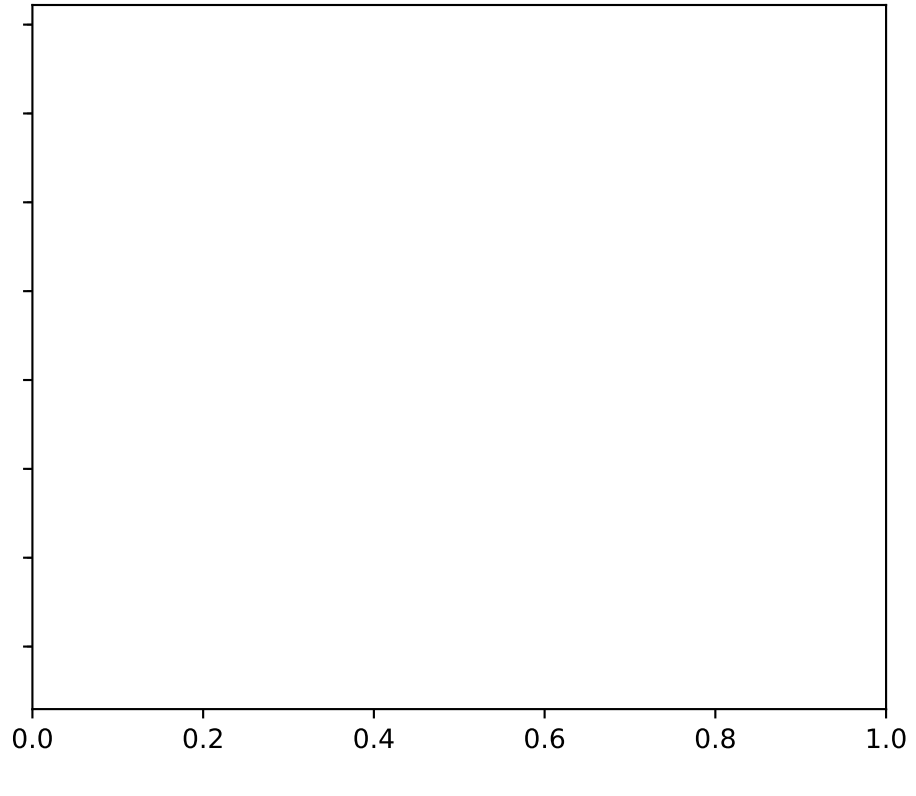
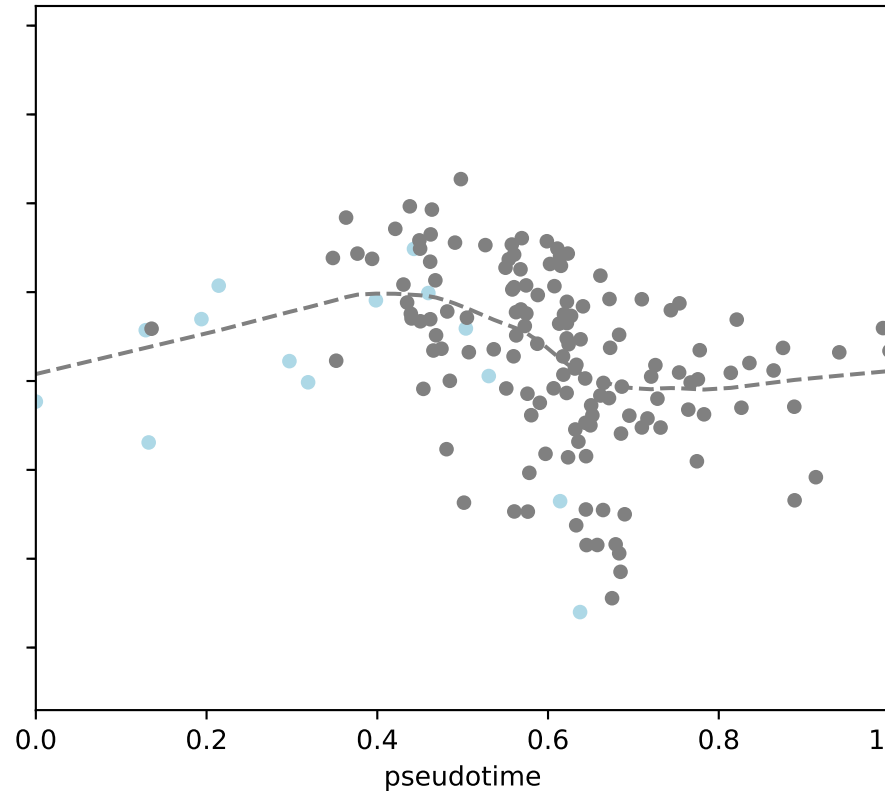
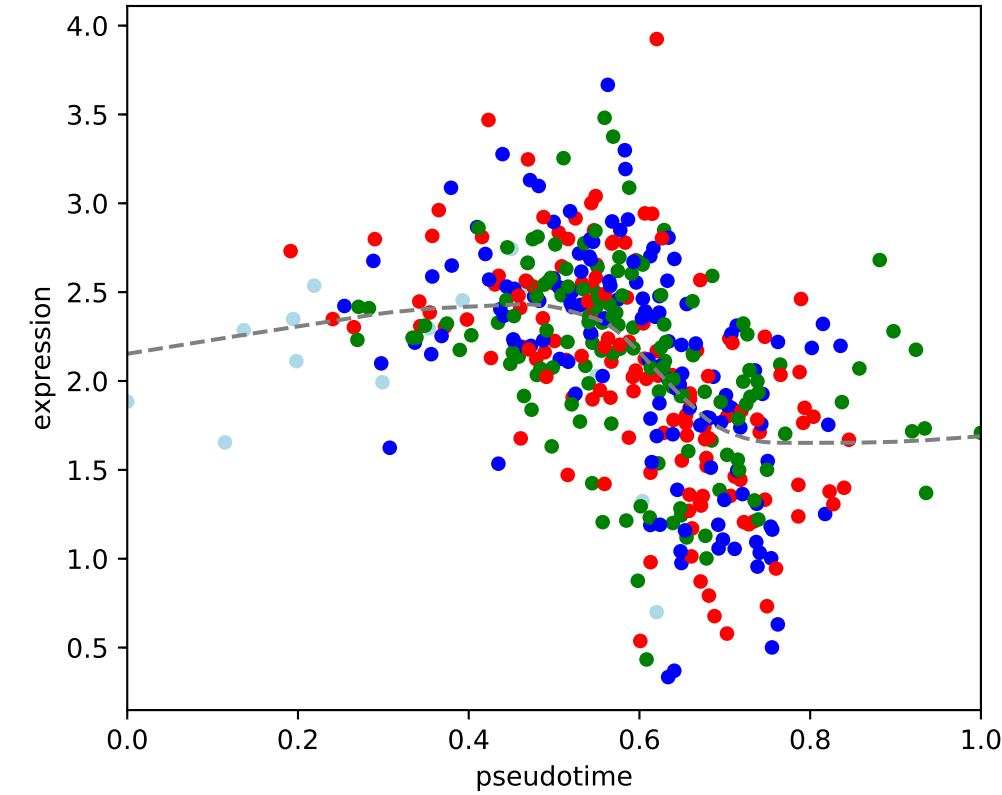
733 1spearman=-0.08



ENST00000405422 PPIAP31 (peptidylprolyl isomerase A pseudogene 31)

733 0spearman=-0.56

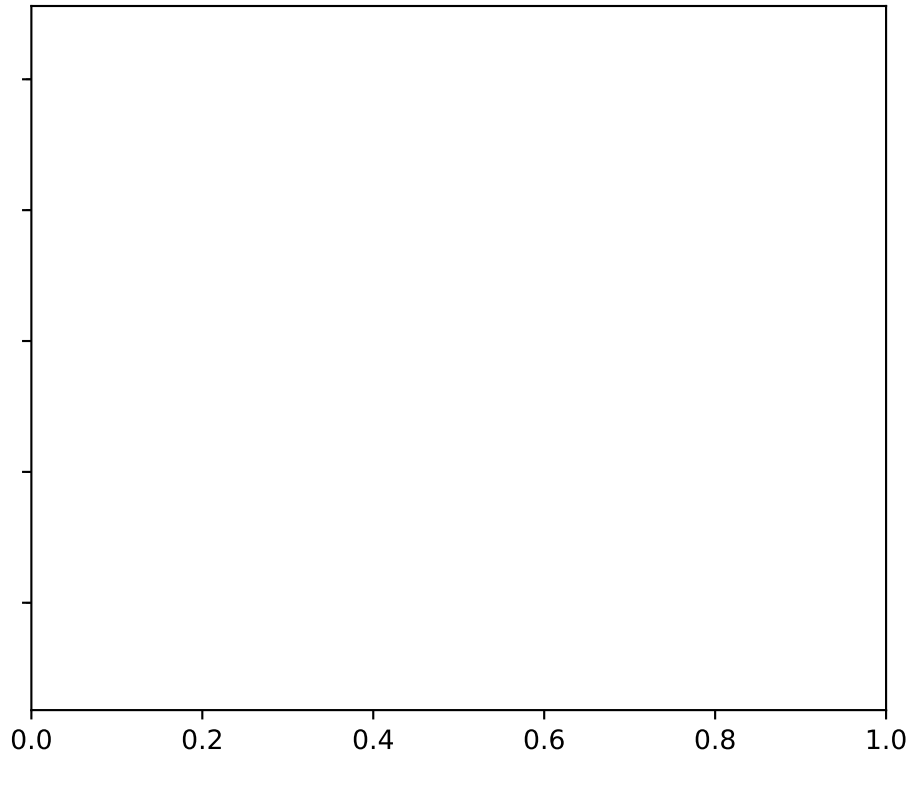
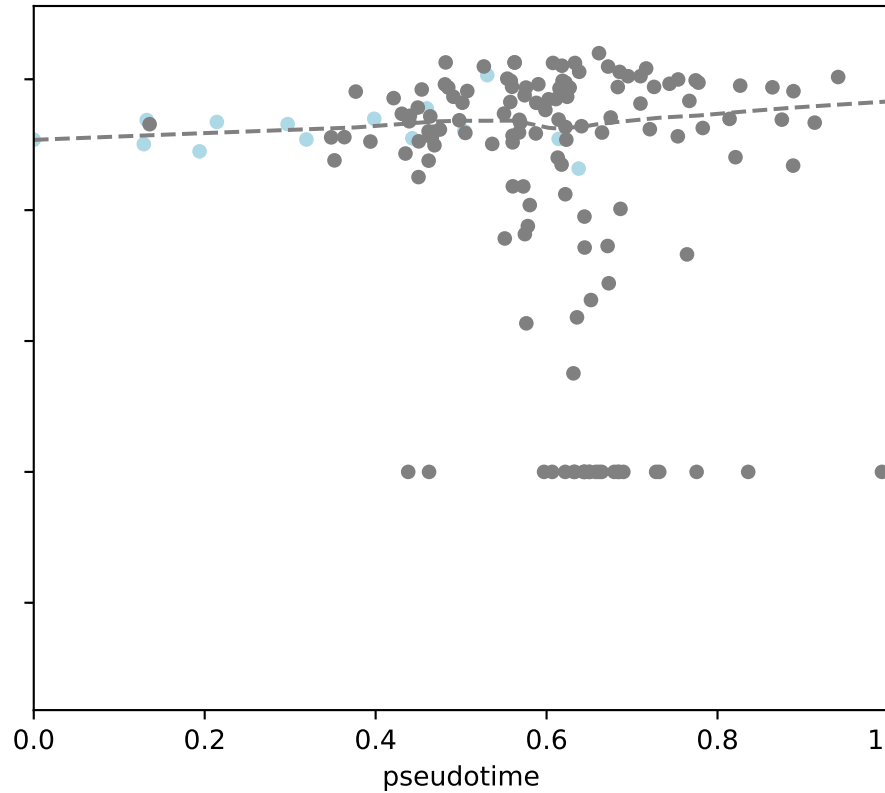
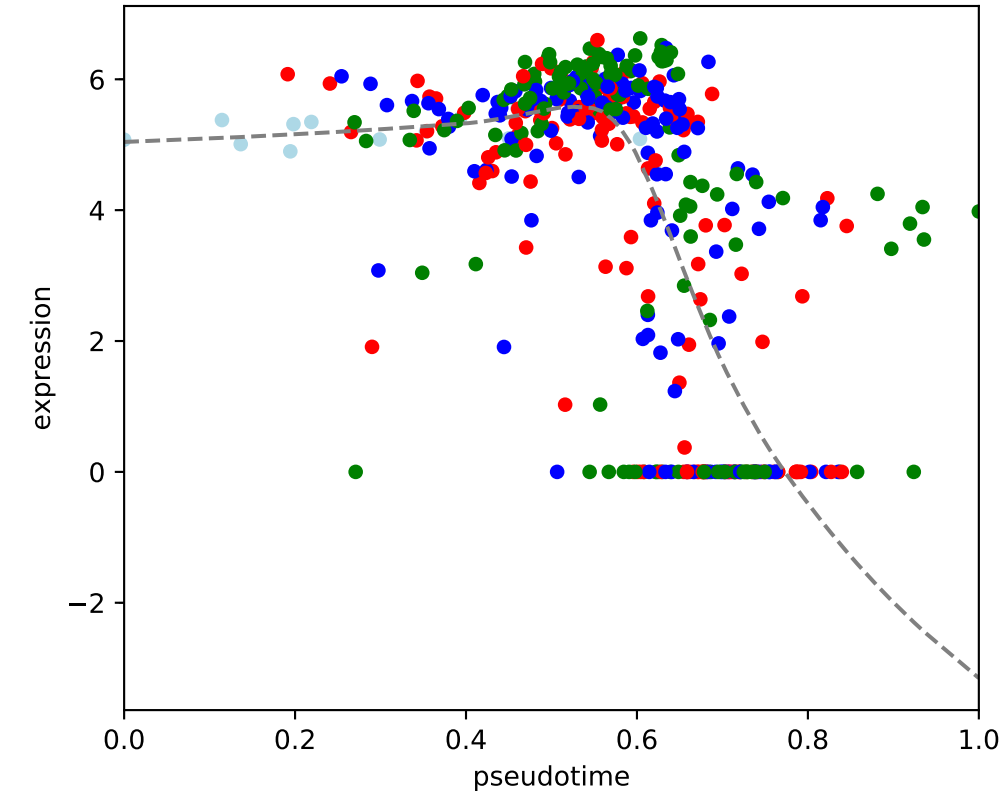
733 1spearman=-0.47



ENST00000409002 HES6 (hes family bHLH transcription factor 6)

733 0spearman=-0.56

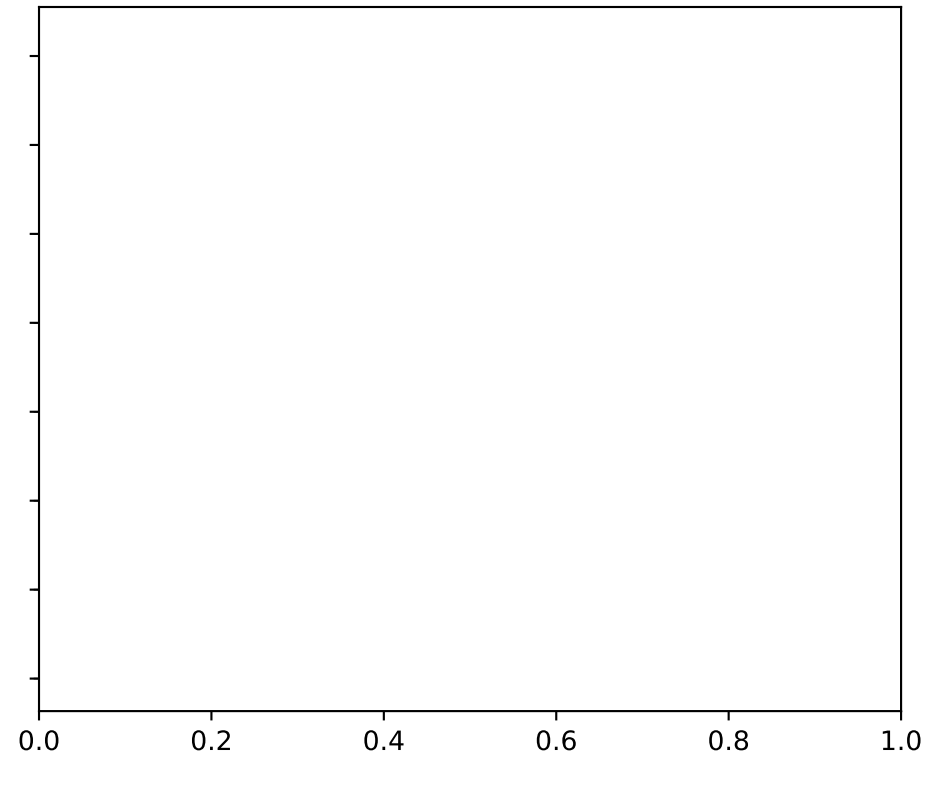
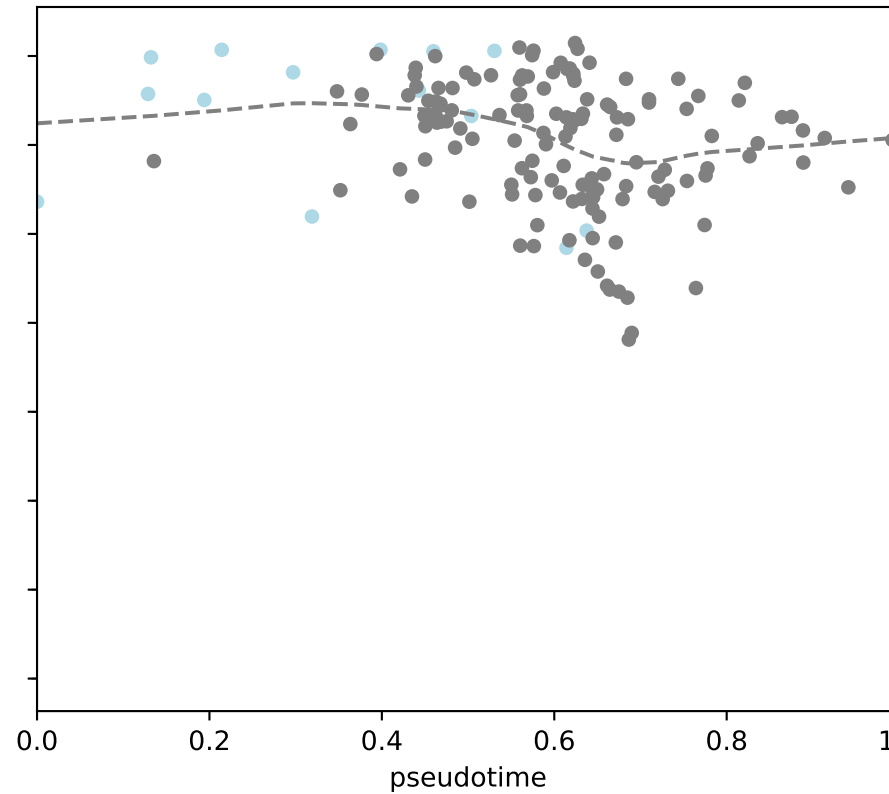
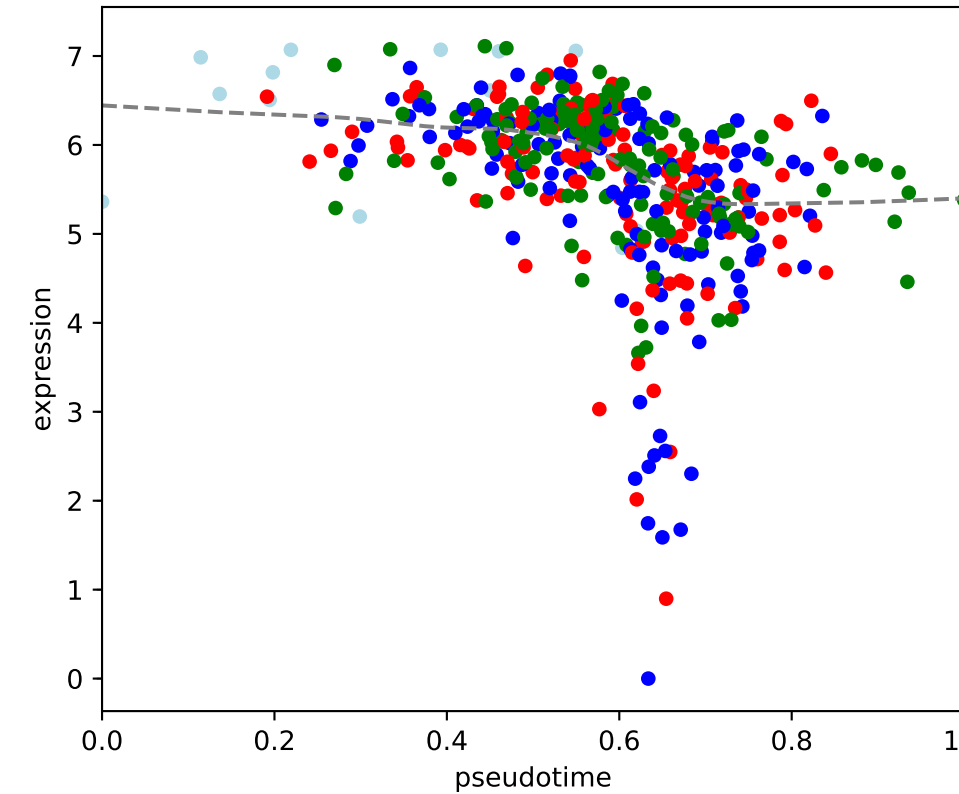
733 1spearman=-0.07



ENST00000436459 EEF1A1P5 (eukaryotic translation elongation factor 1 alpha 1 pseudogene 5)

733 0spearman=-0.56

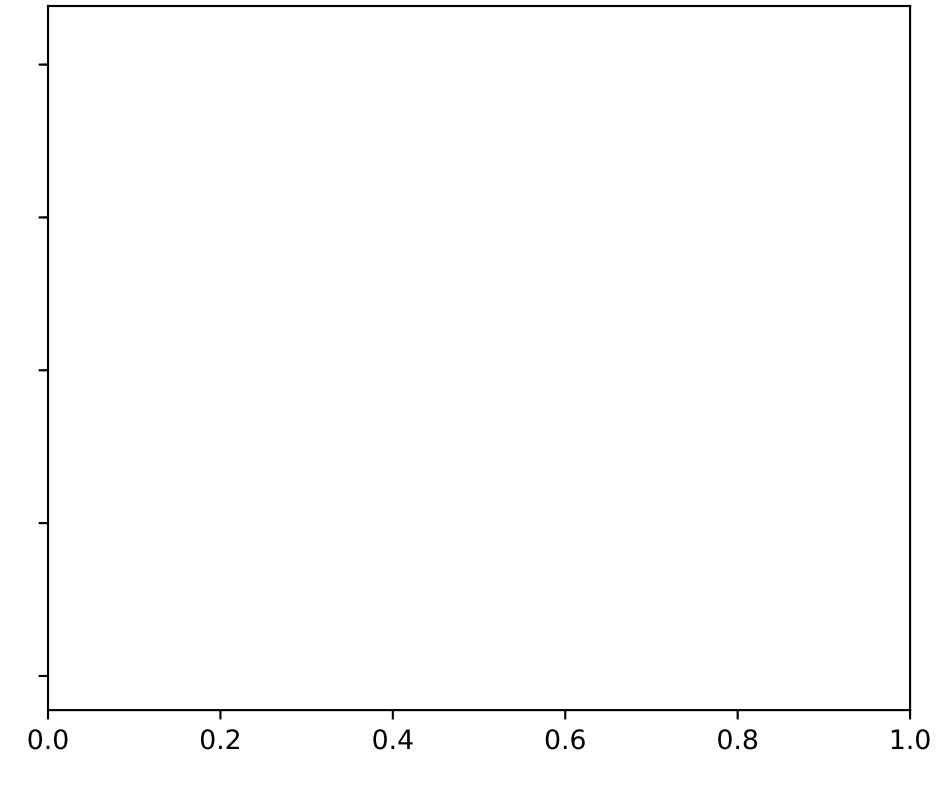
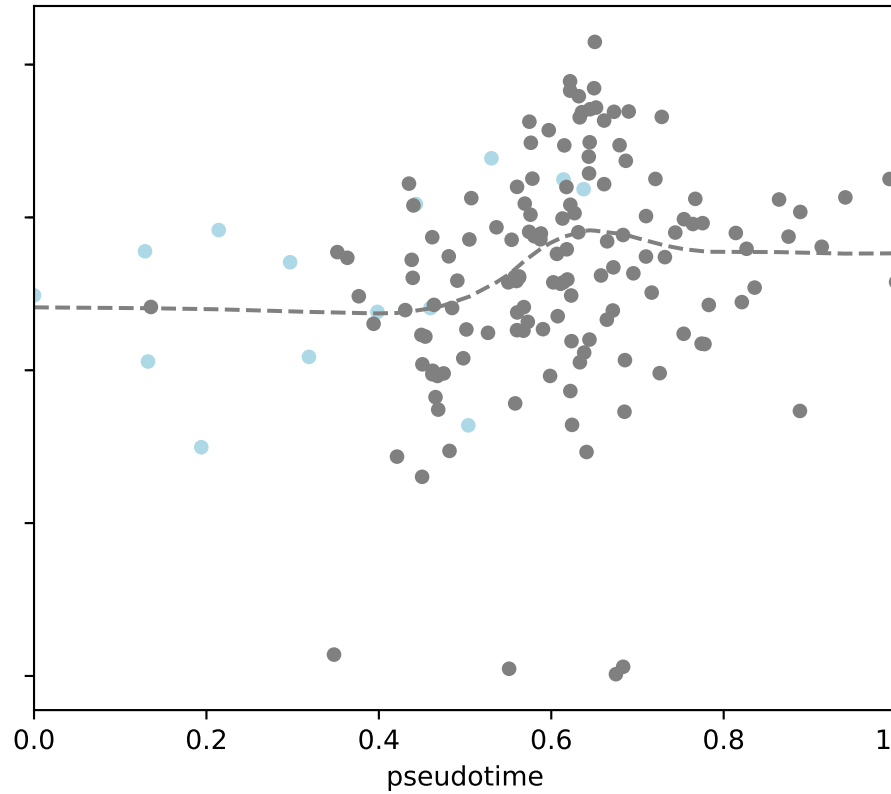
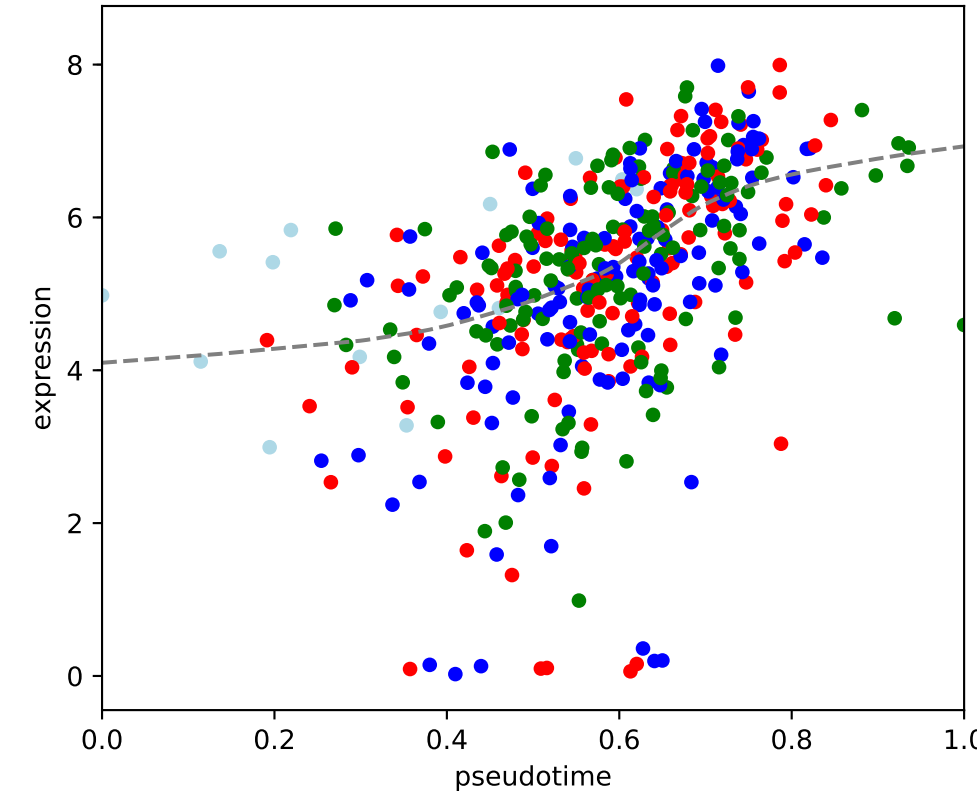
733 1spearman=-0.35



ENST00000379919 MAB21L1 (mab-21 like 1)

733 0spearman=0.56

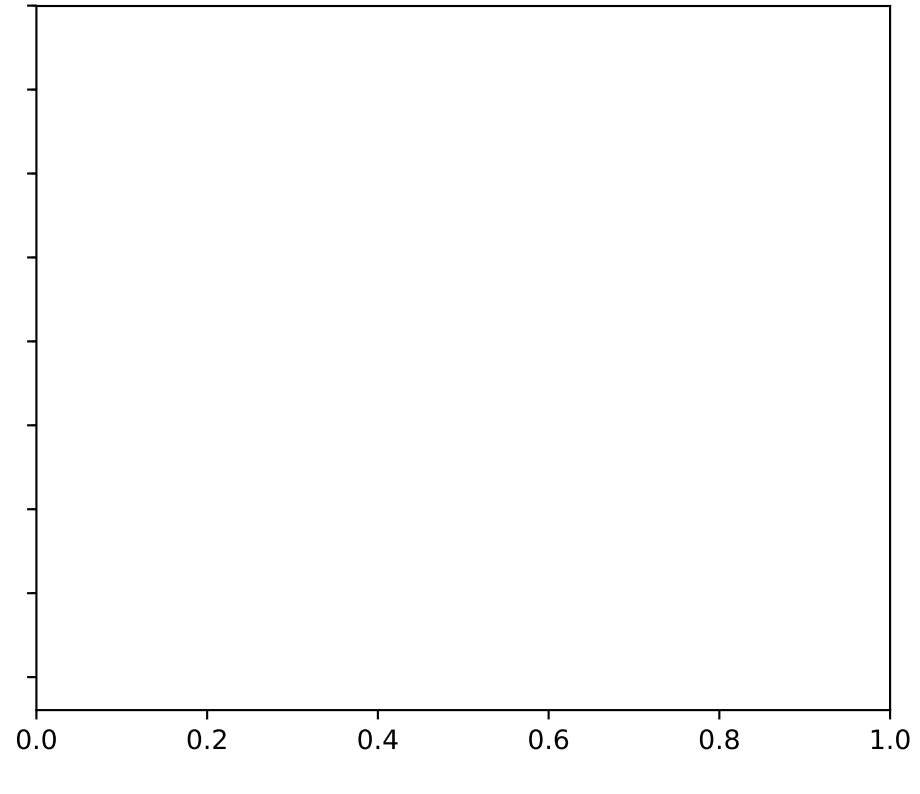
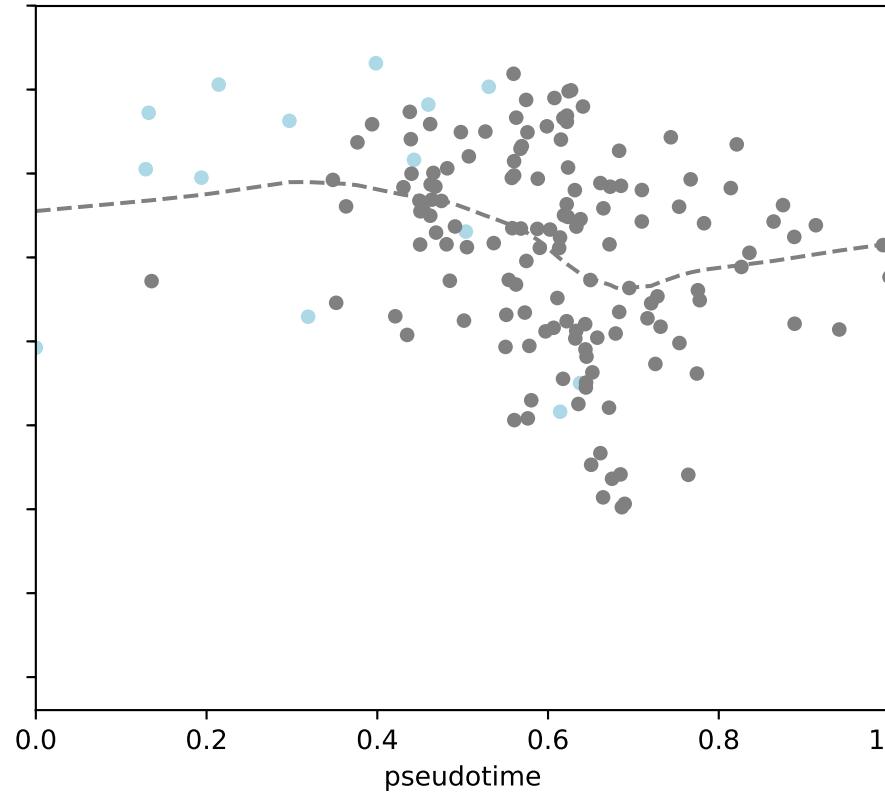
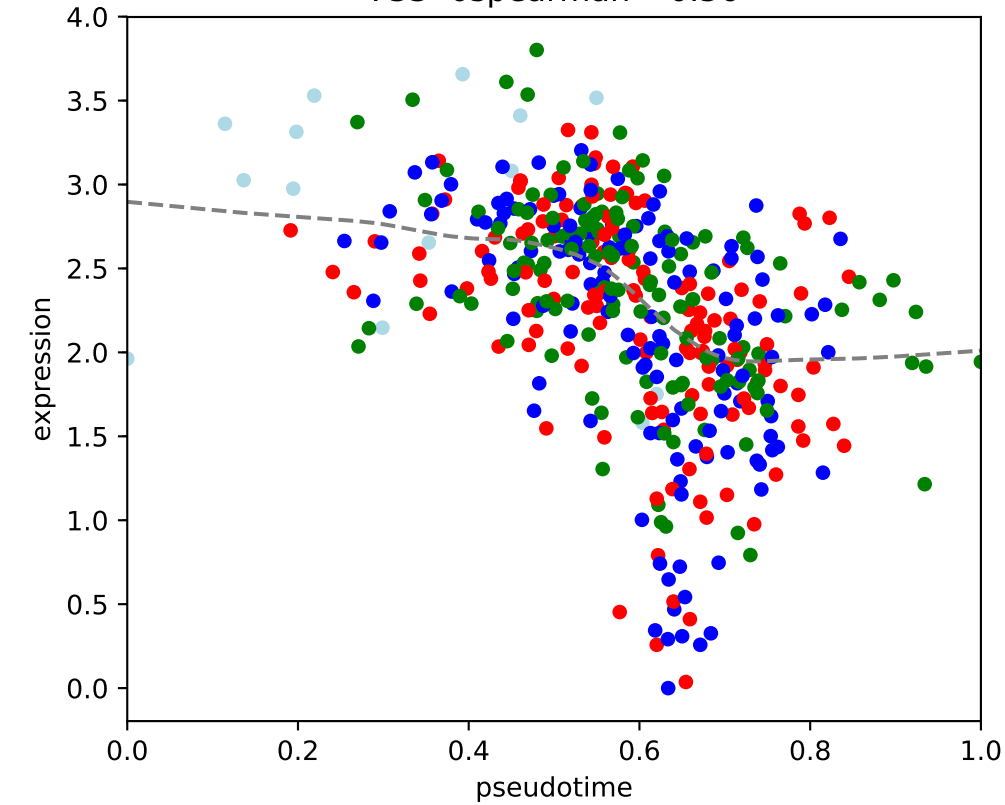
733 1spearman=0.29



ENST00000503537 EEF1A1P13 (eukaryotic translation elongation factor 1 alpha 1 pseudogene 13)

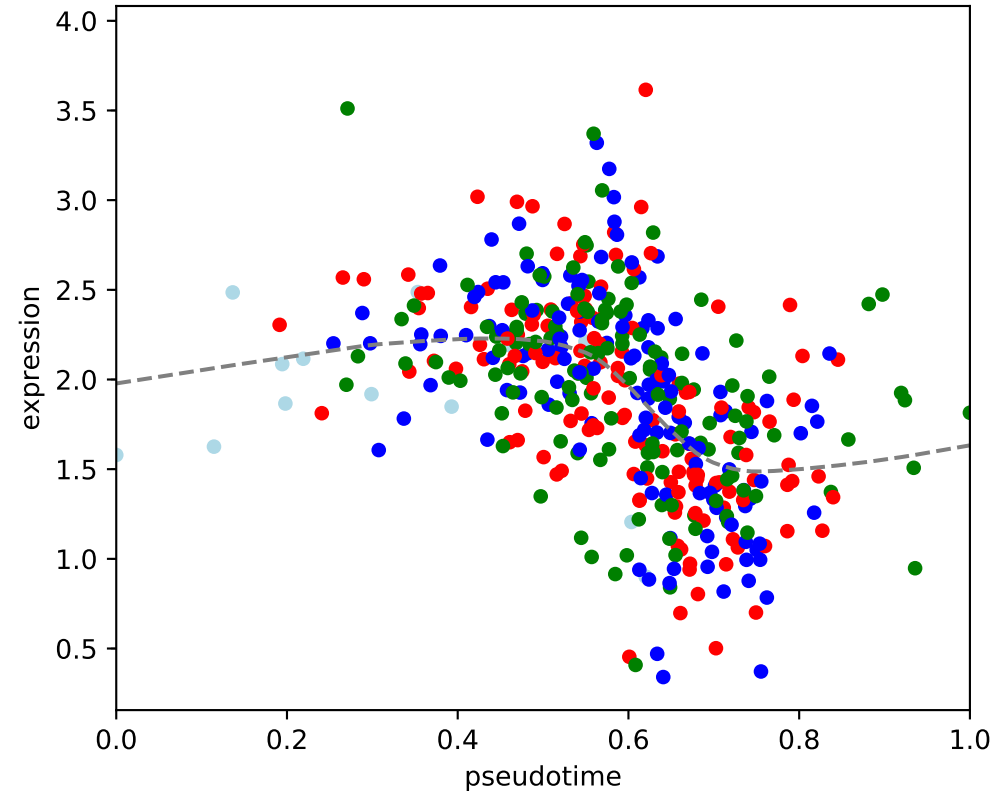
733 0spearman=-0.56

733 1spearman=-0.36

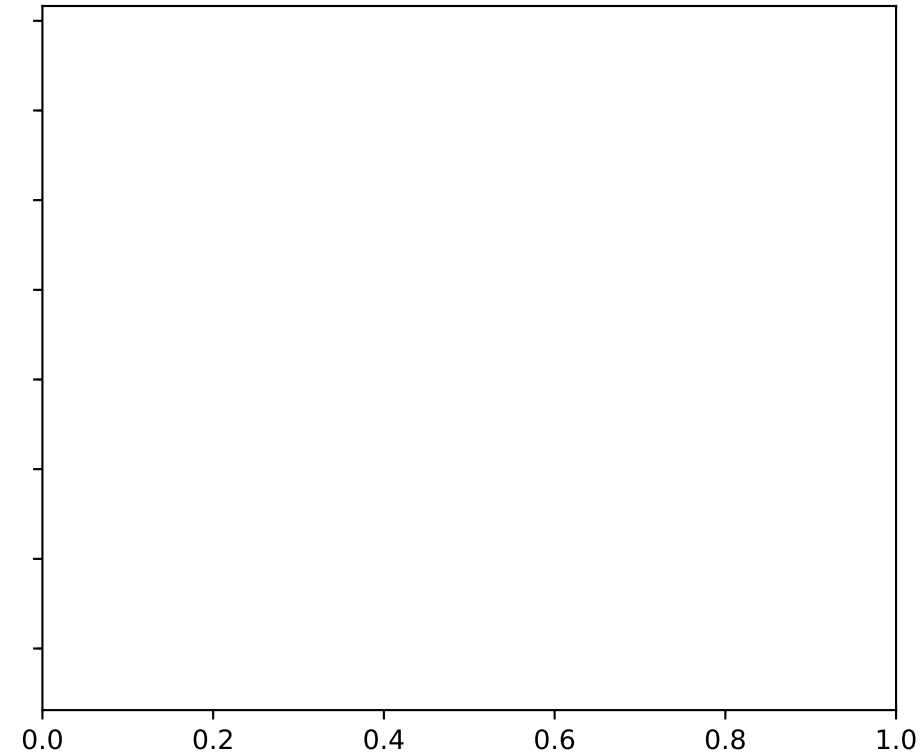
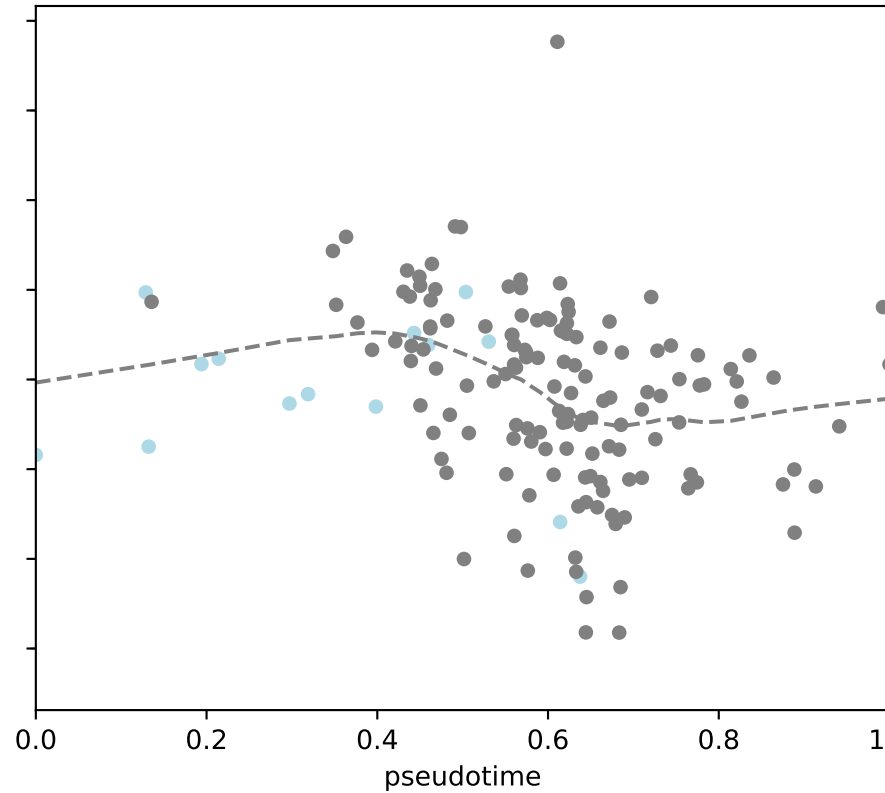


ENST00000510729 PPIAP11 (peptidylprolyl isomerase A pseudogene 11)

733 0spearman=-0.56



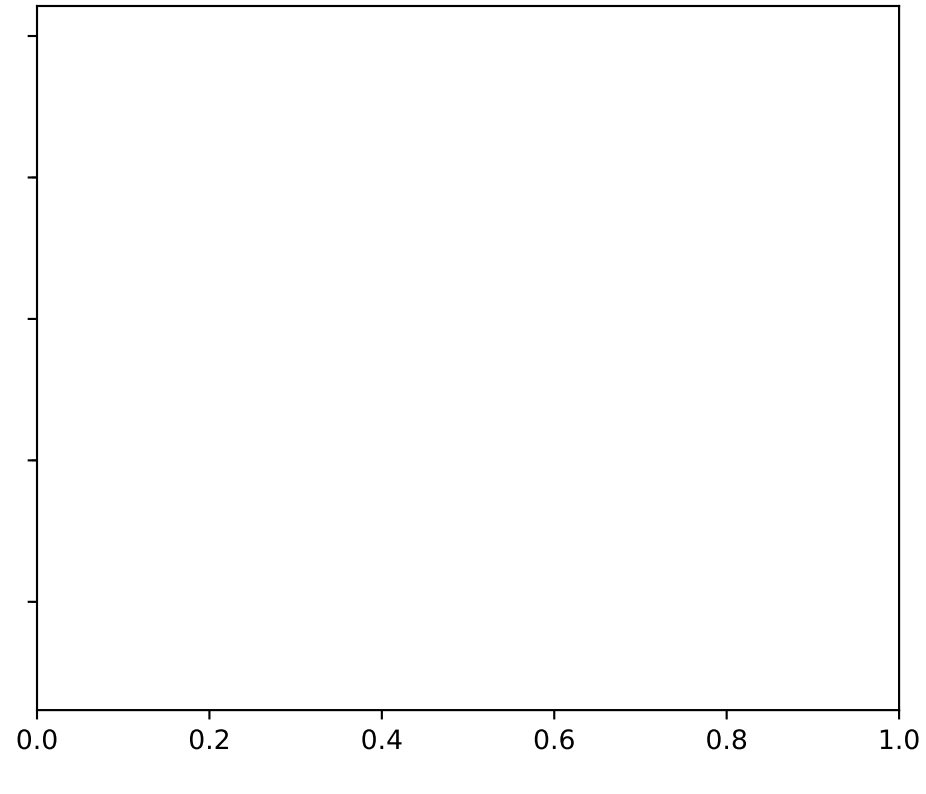
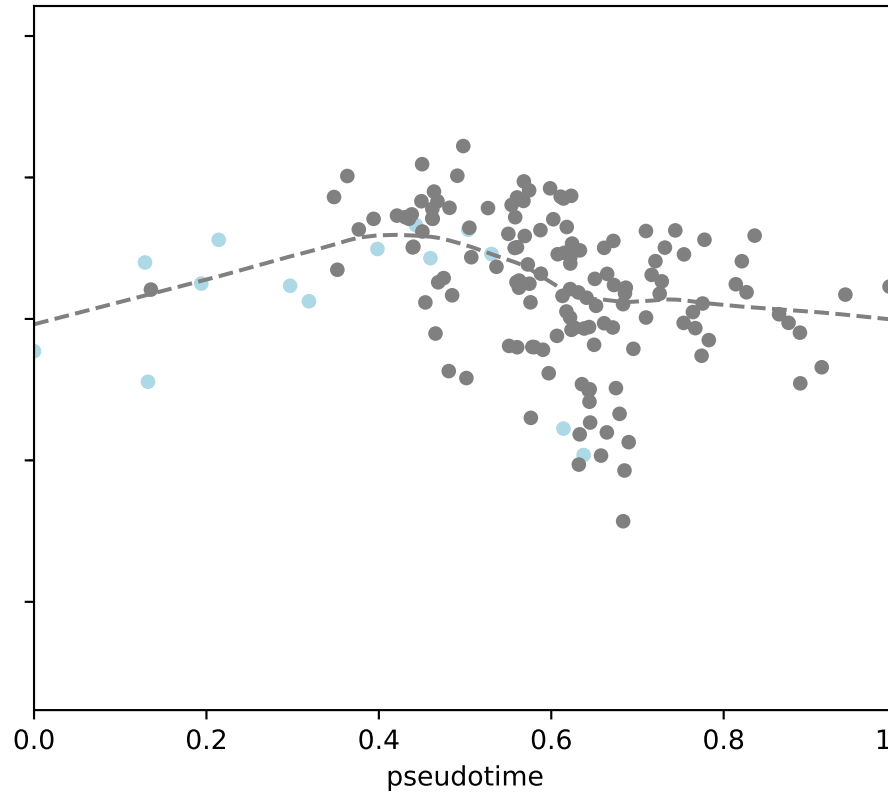
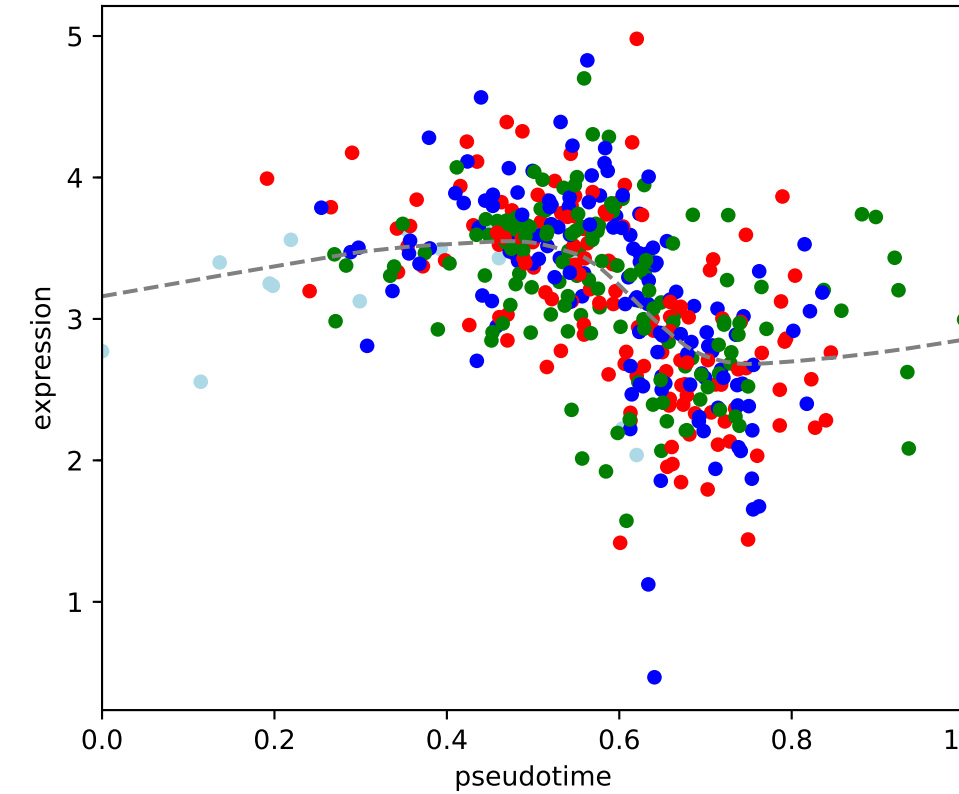
733 1spearman=-0.42



ENST00000414949 PPIAP29 (peptidylprolyl isomerase A pseudogene 29)

733 0spearman=-0.56

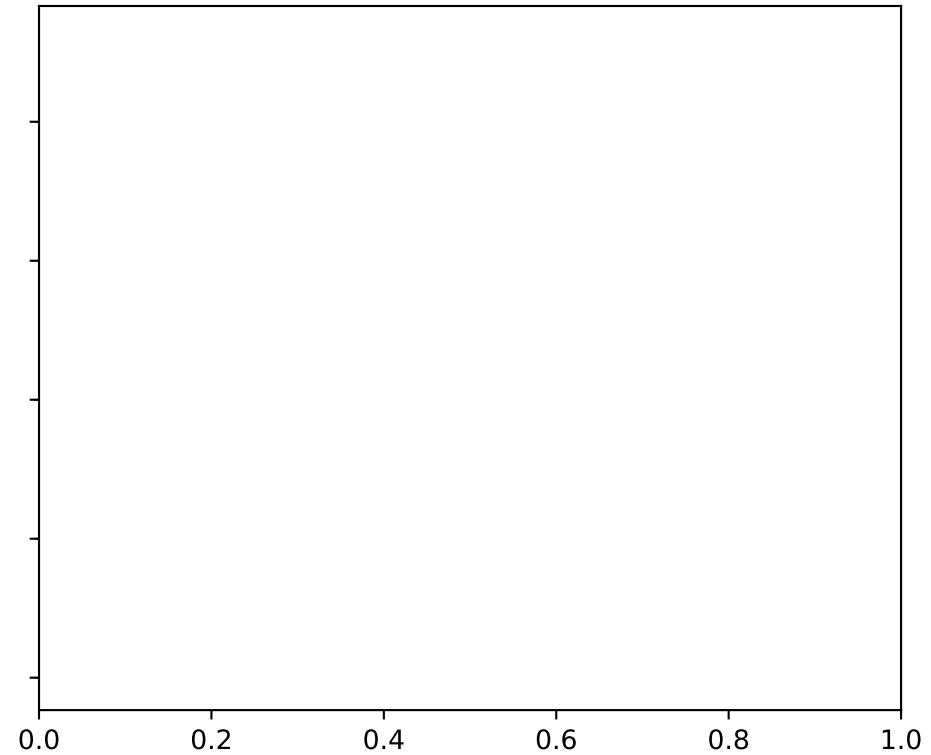
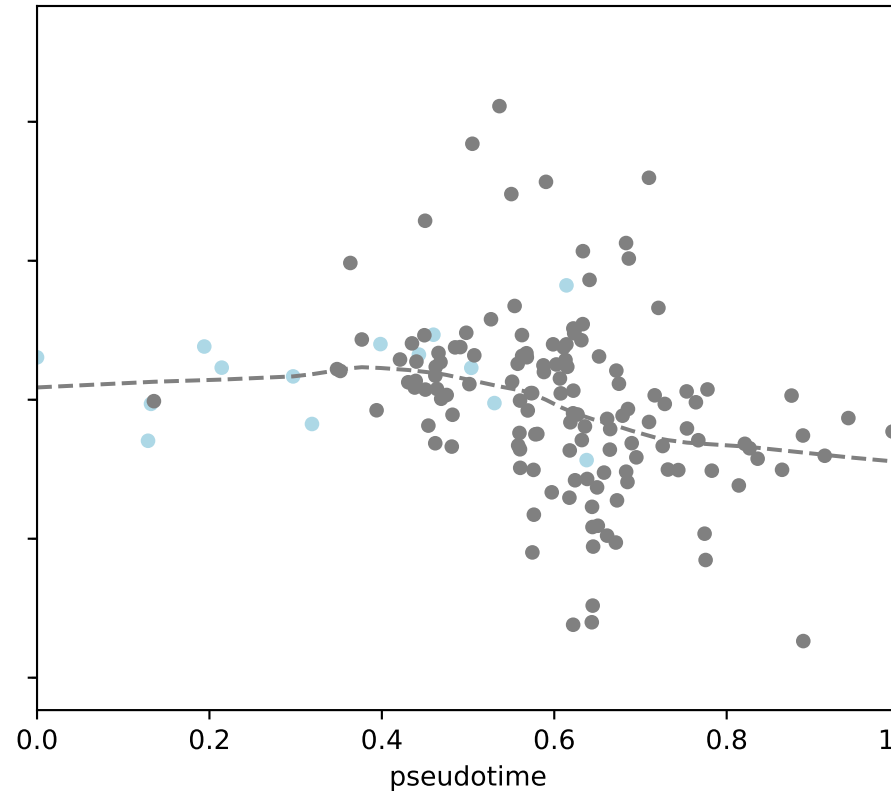
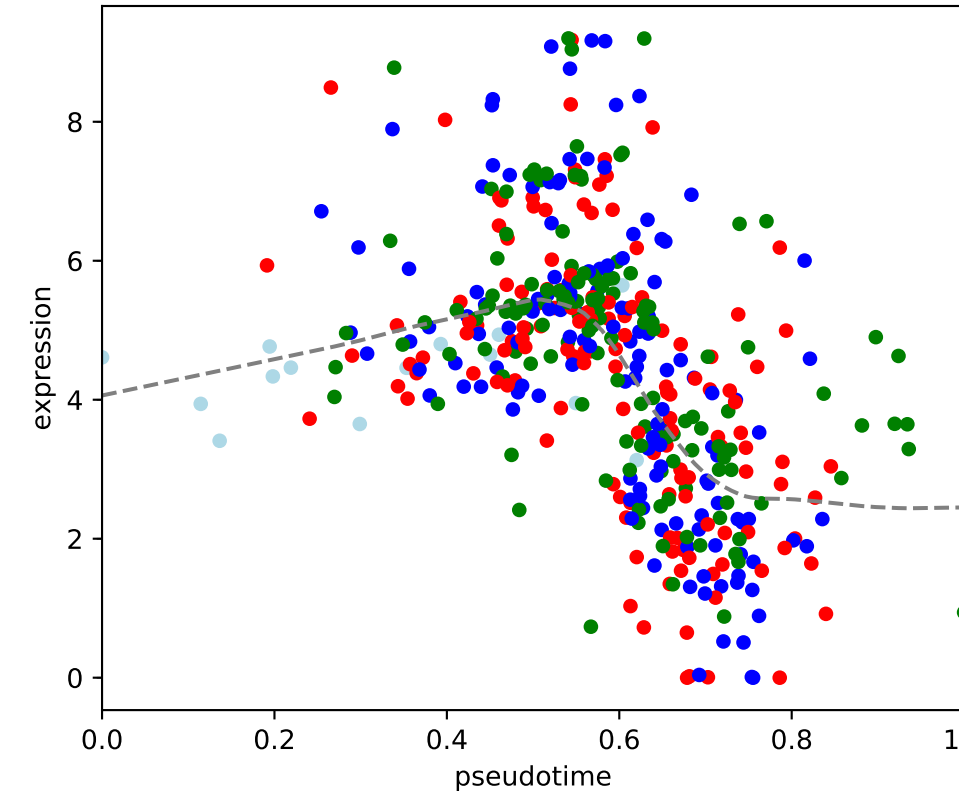
733 1spearman=-0.43



ENST00000336023 TUBA1B (tubulin alpha 1b)

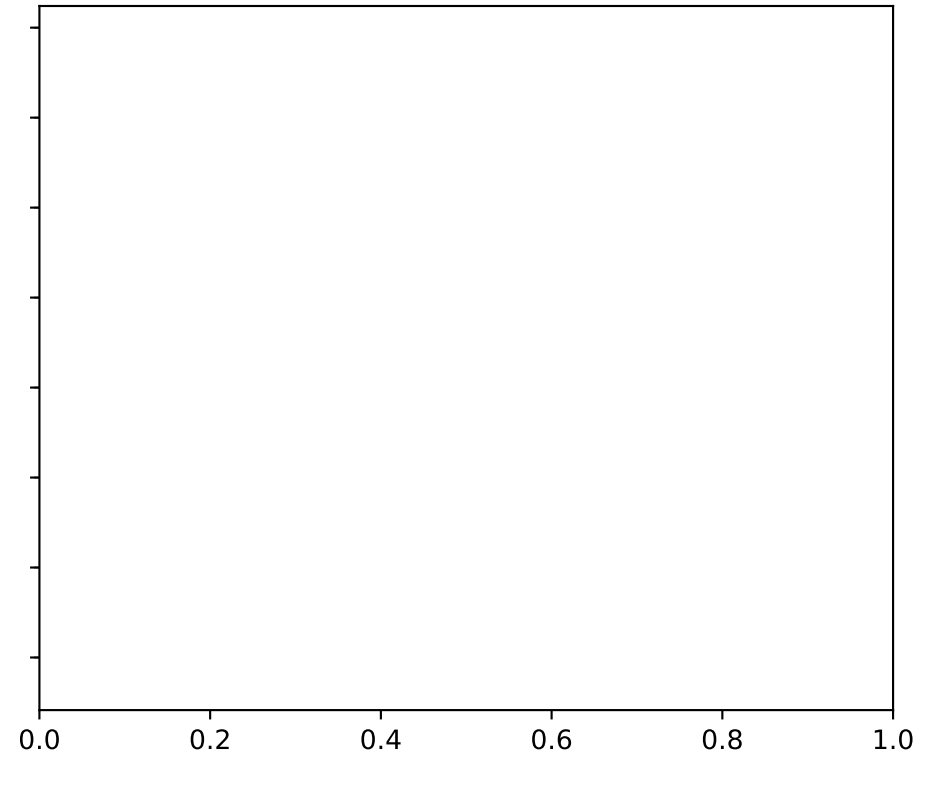
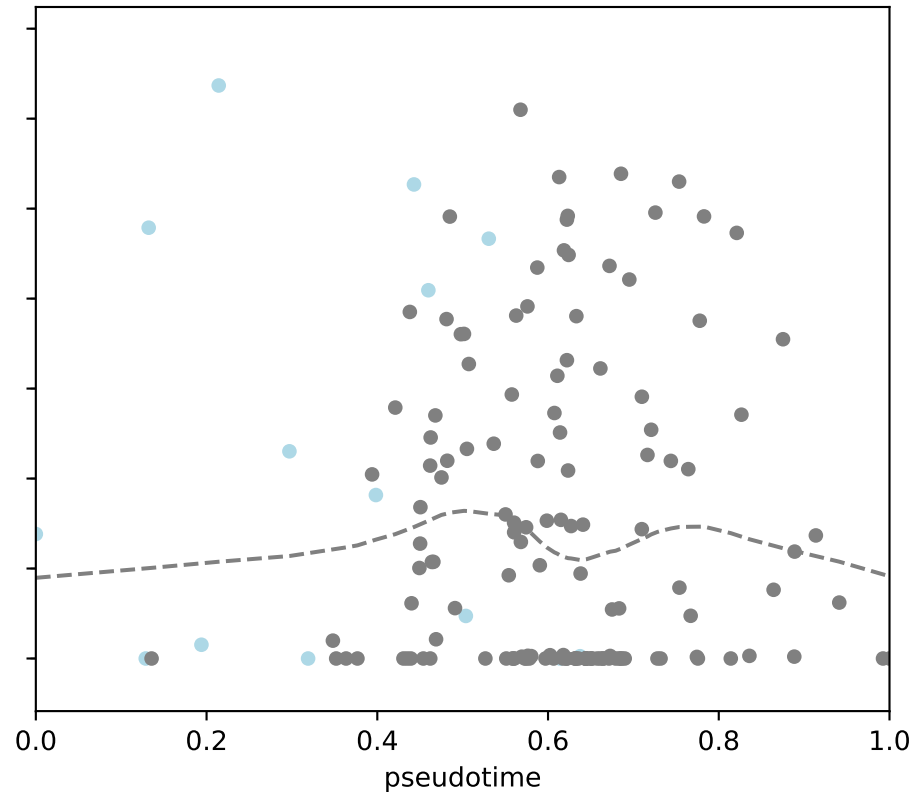
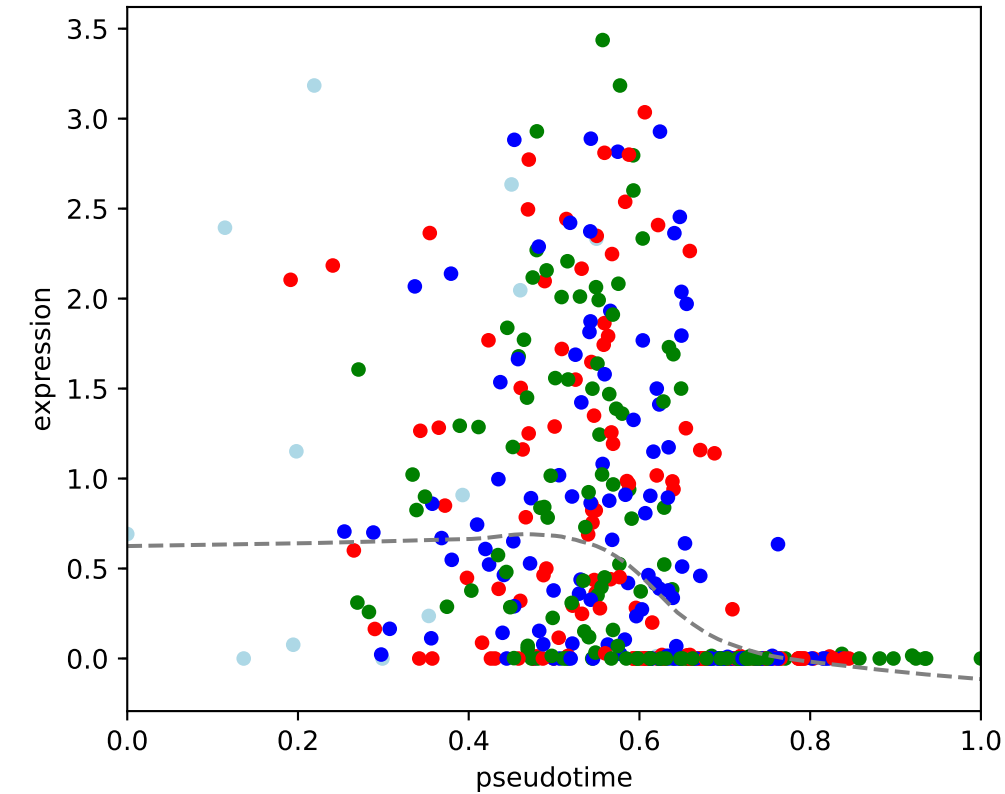
733 0spearman=-0.56

733 1spearman=-0.44



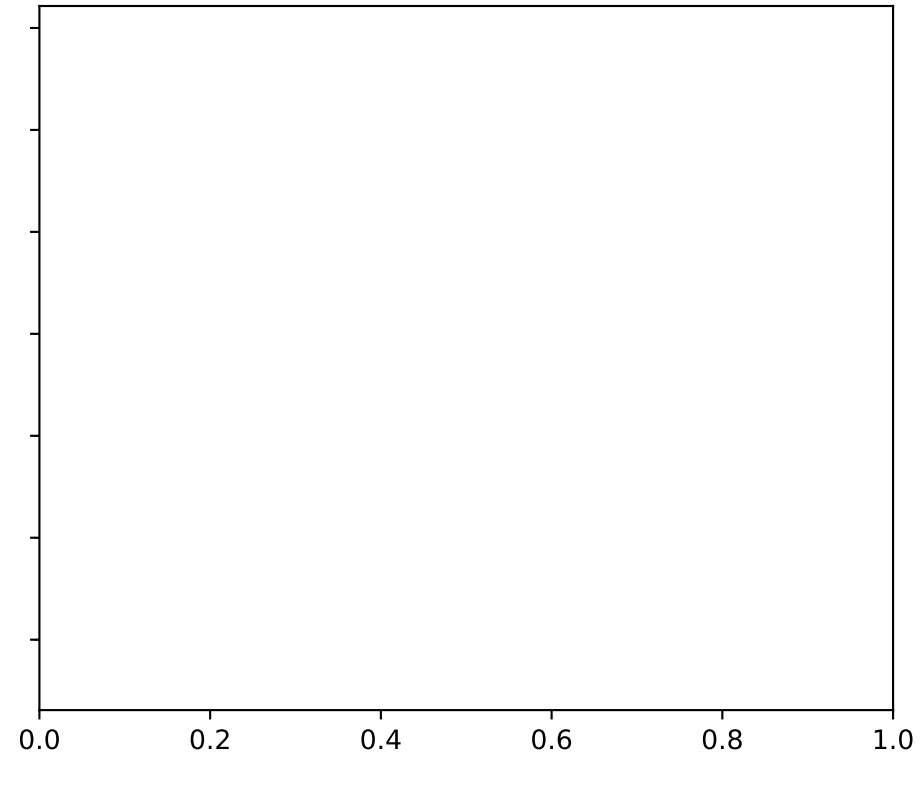
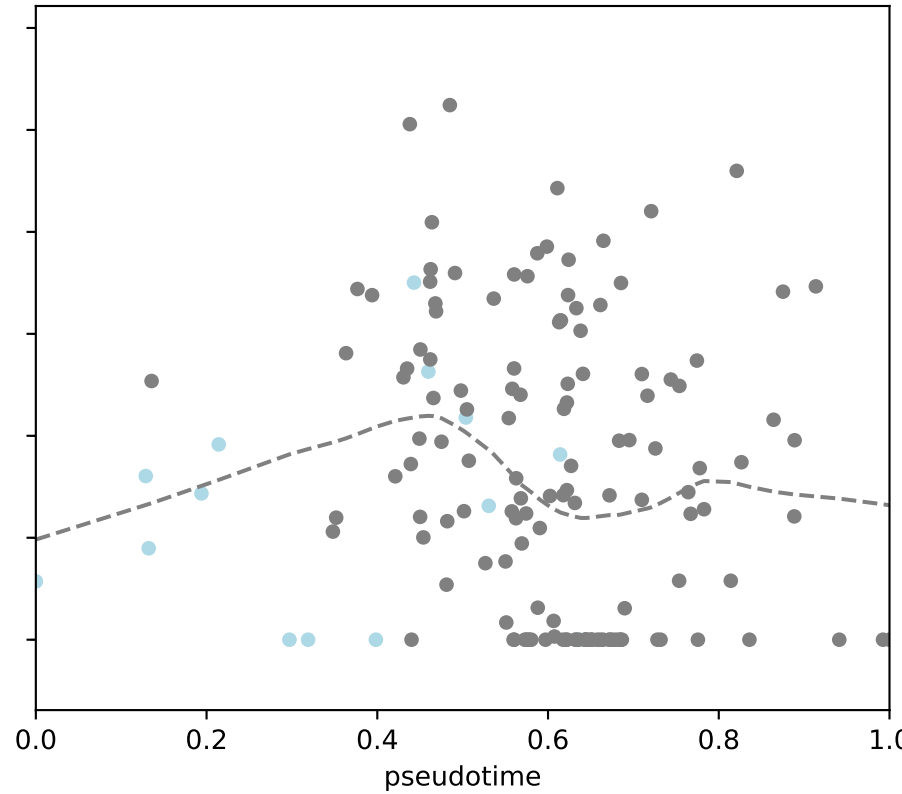
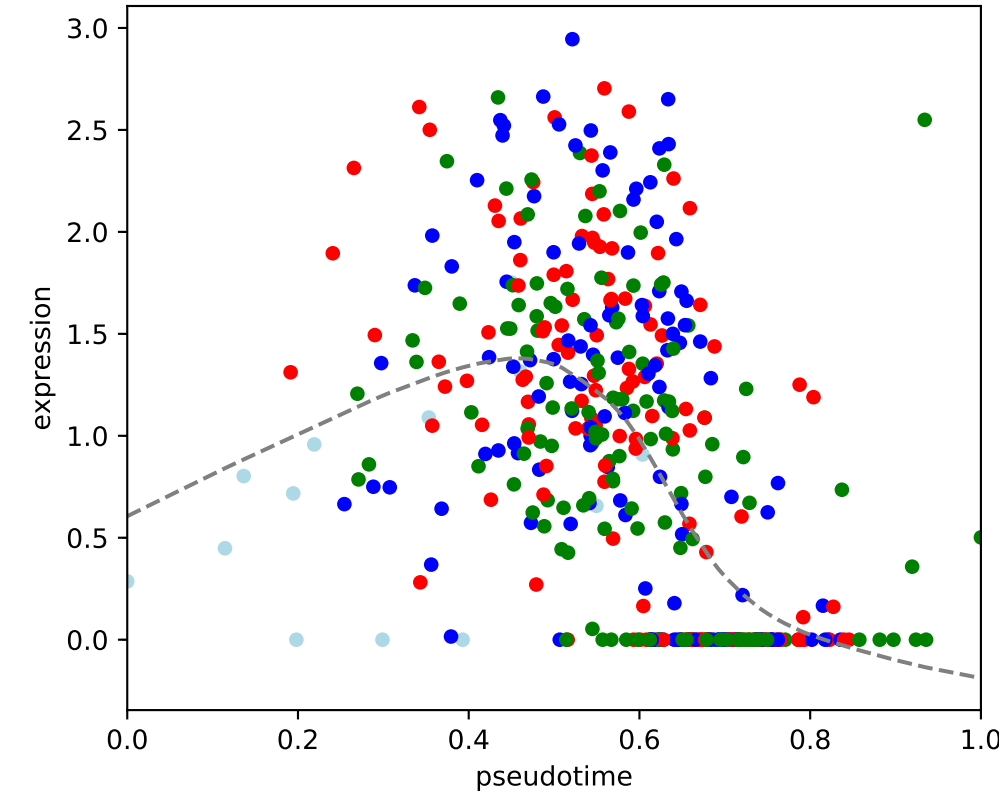
733 0spearman=-0.56

733 1spearman=-0.07



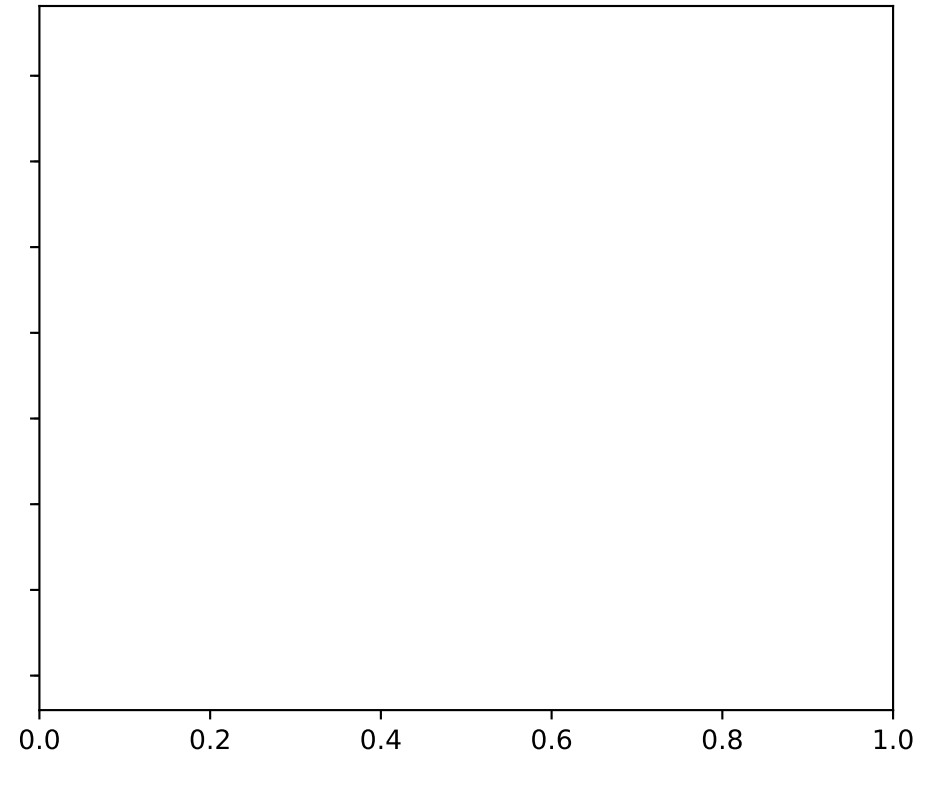
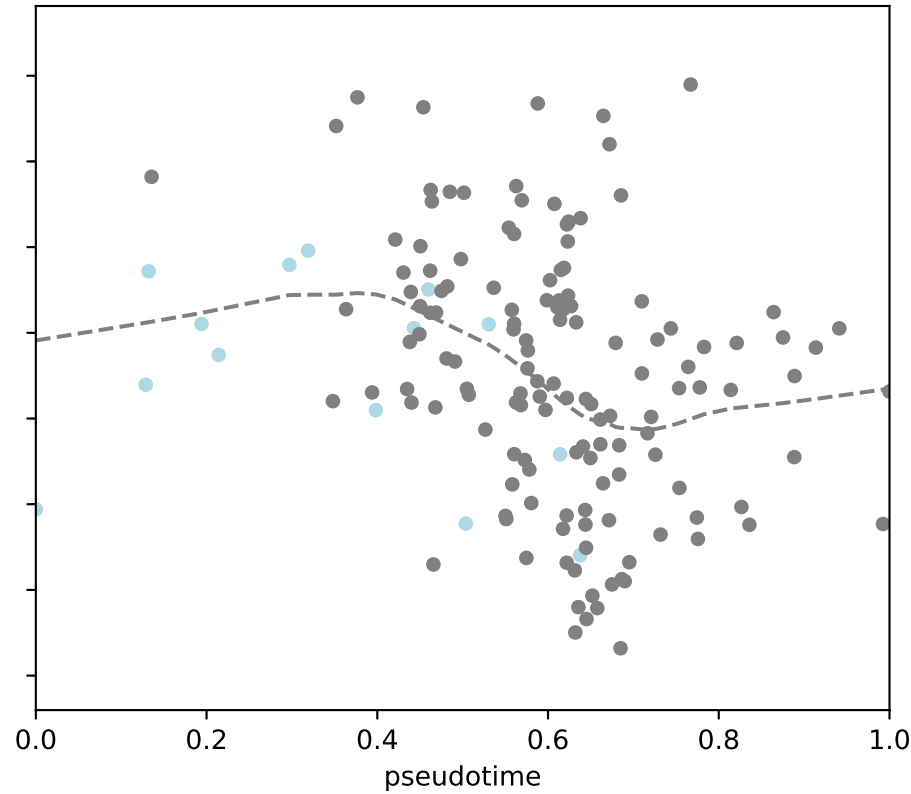
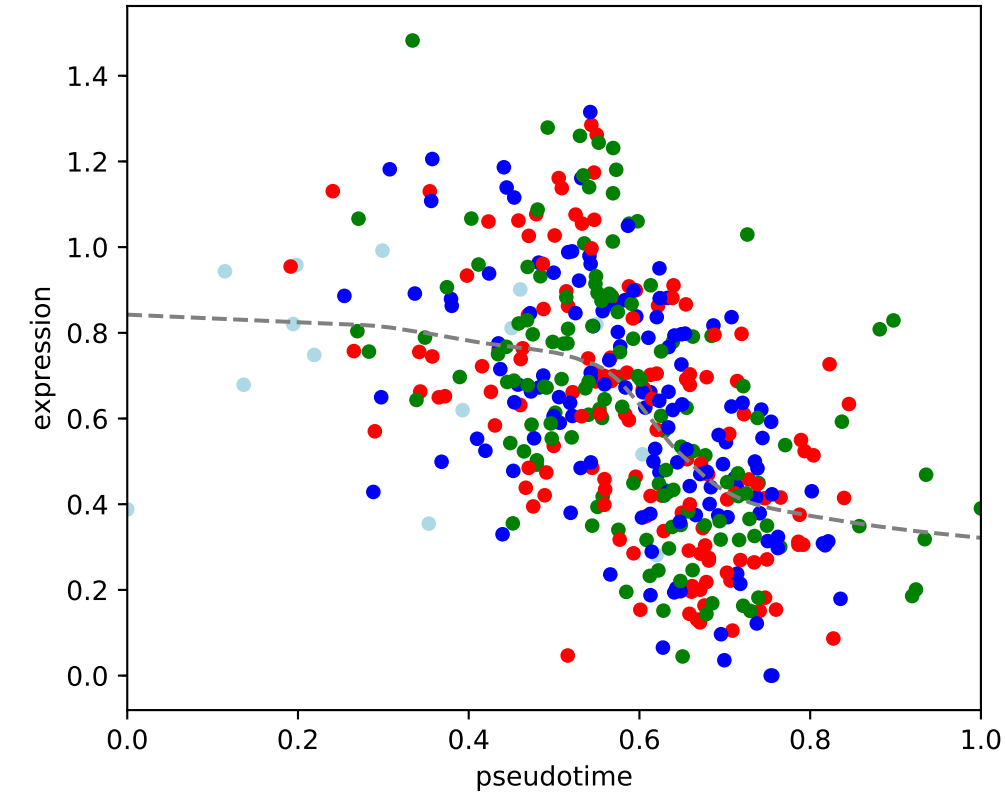
733 0spearman=-0.56

733 1spearman=-0.20



733 0spearman=-0.56

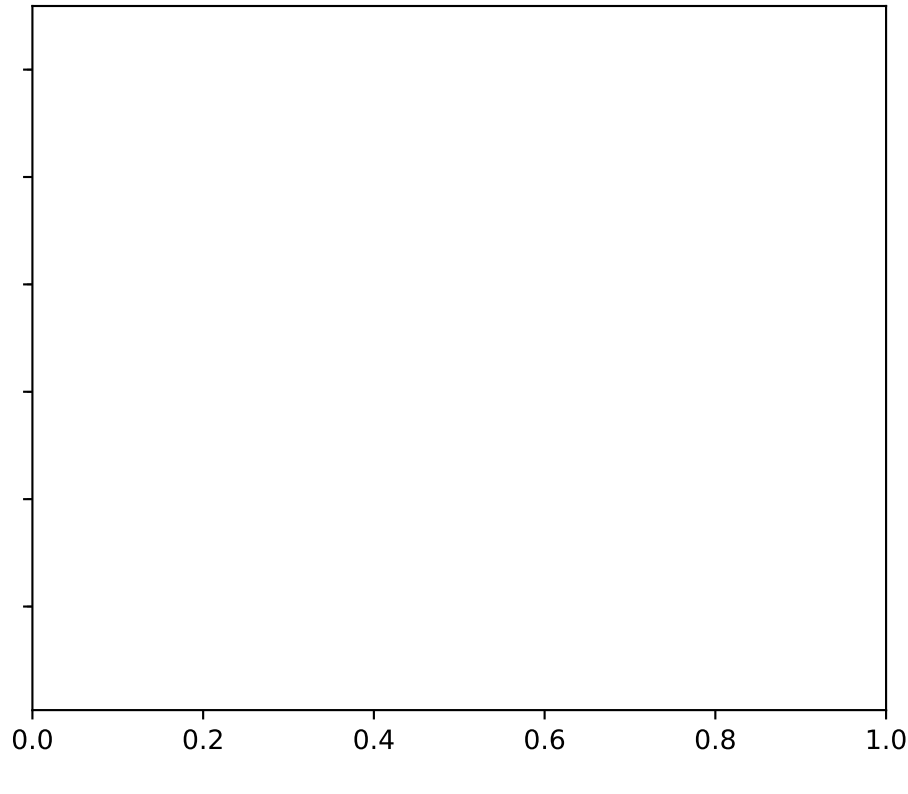
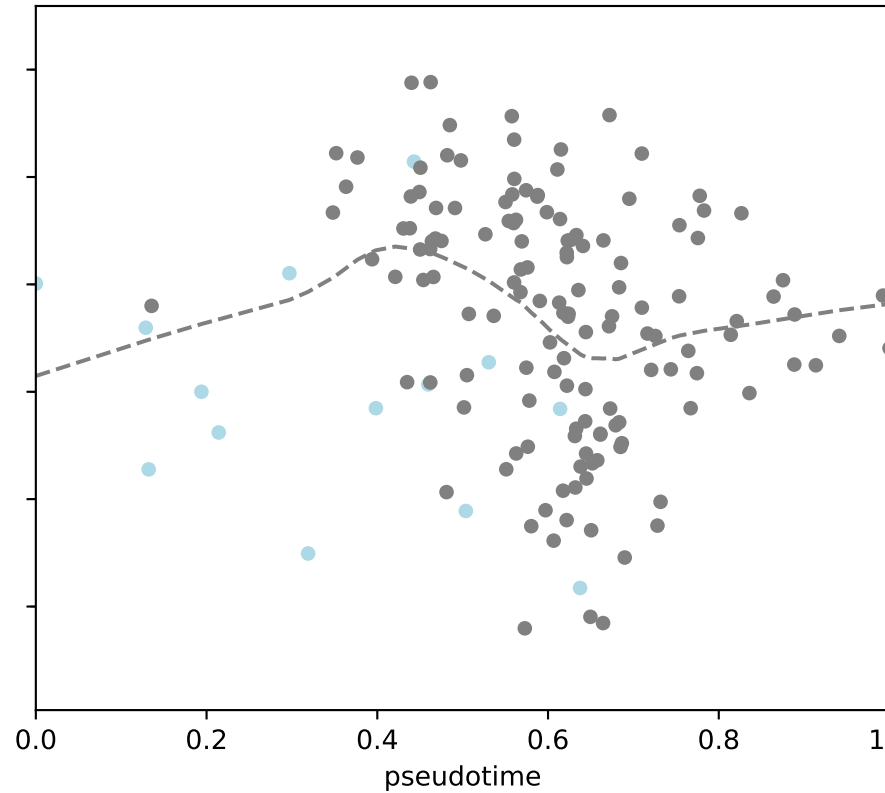
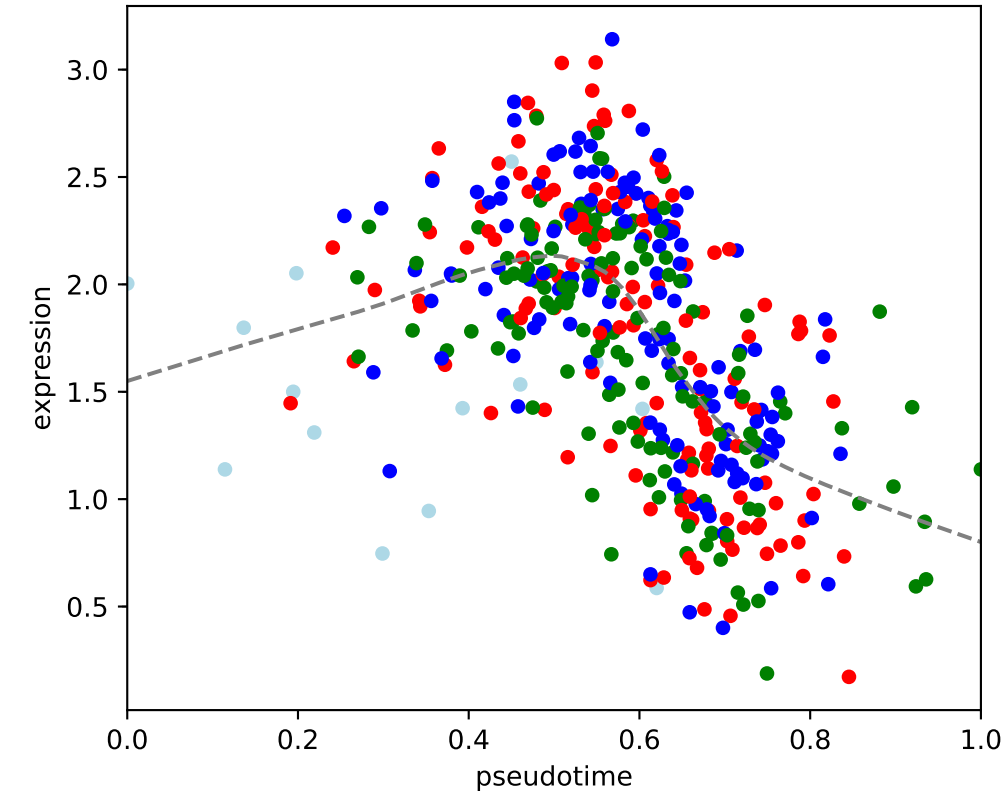
733 1spearman=-0.36



ENST00000461726 CHCHD2P9 (coiled-coil-helix-coiled-coil-helix domain containing 2 pseudogene 9)

733 0spearman=-0.56

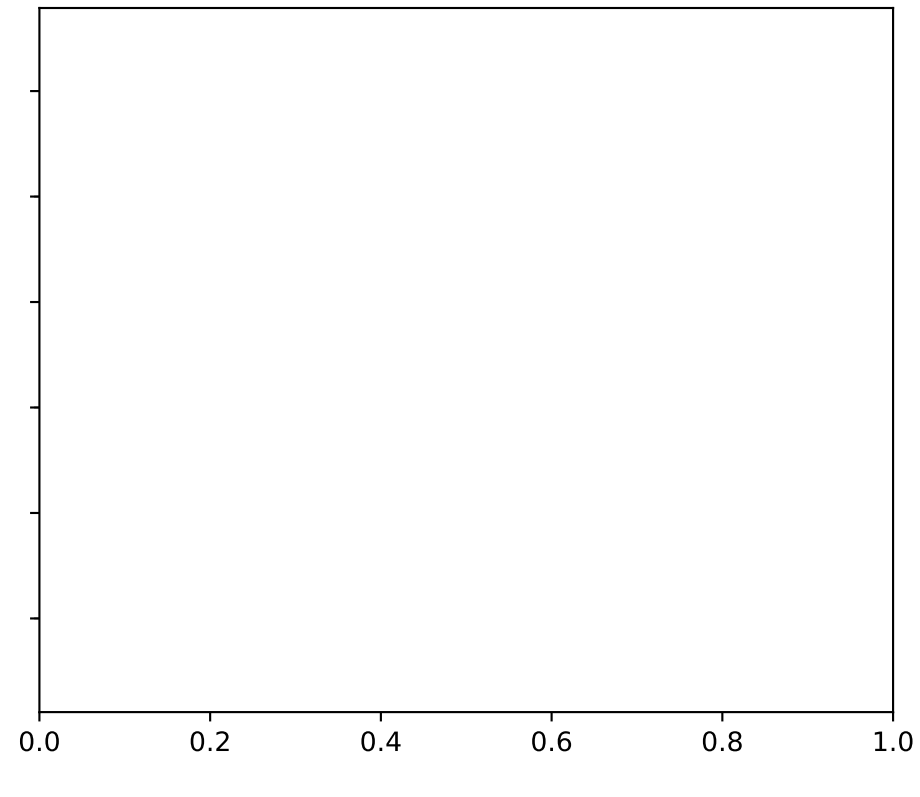
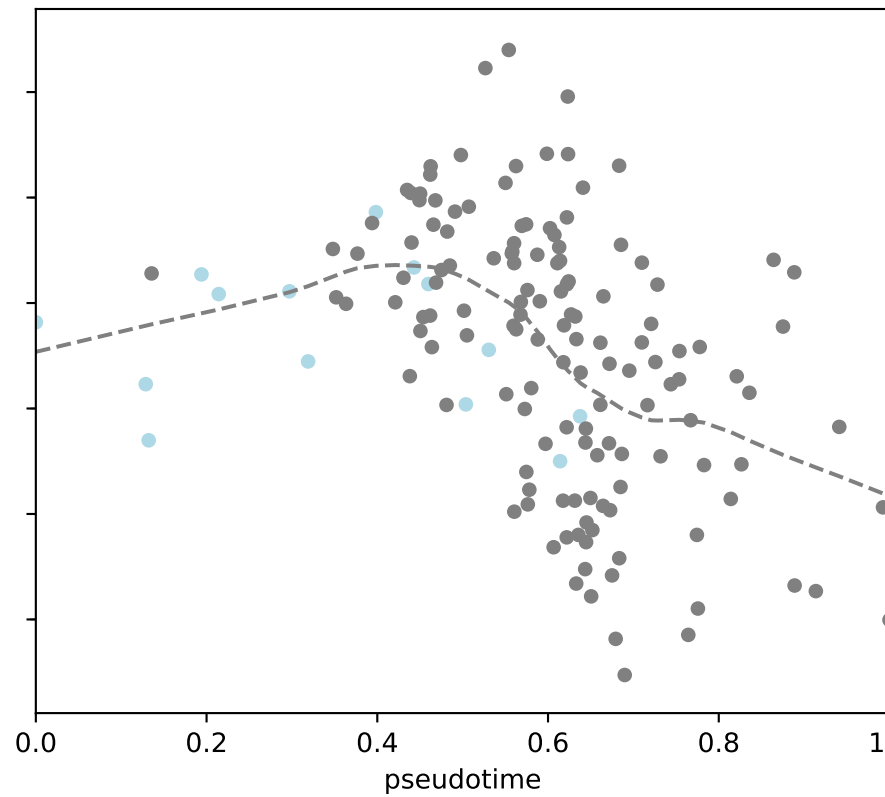
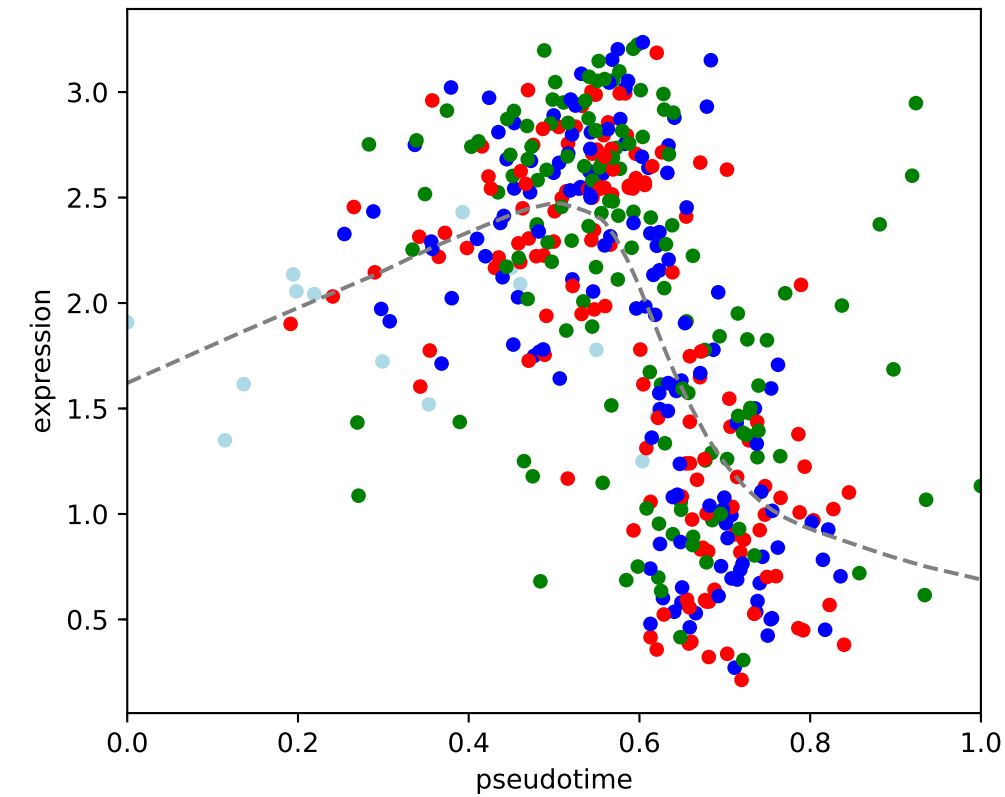
733 1spearman=-0.26



ENST00000295766 TUBA1A (tubulin alpha 1a)

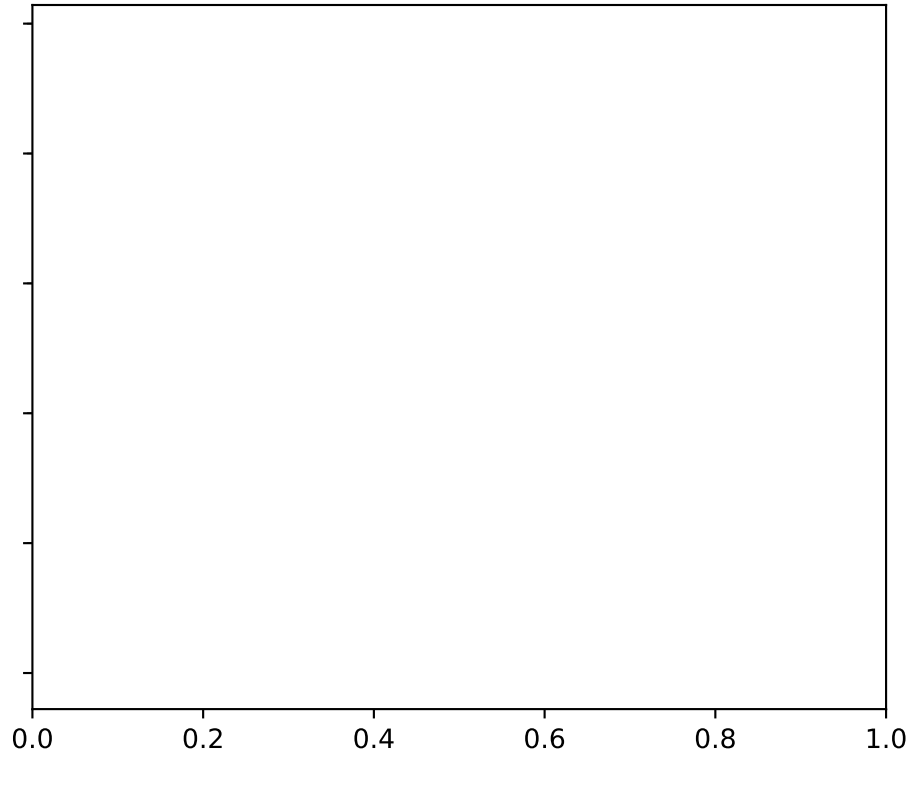
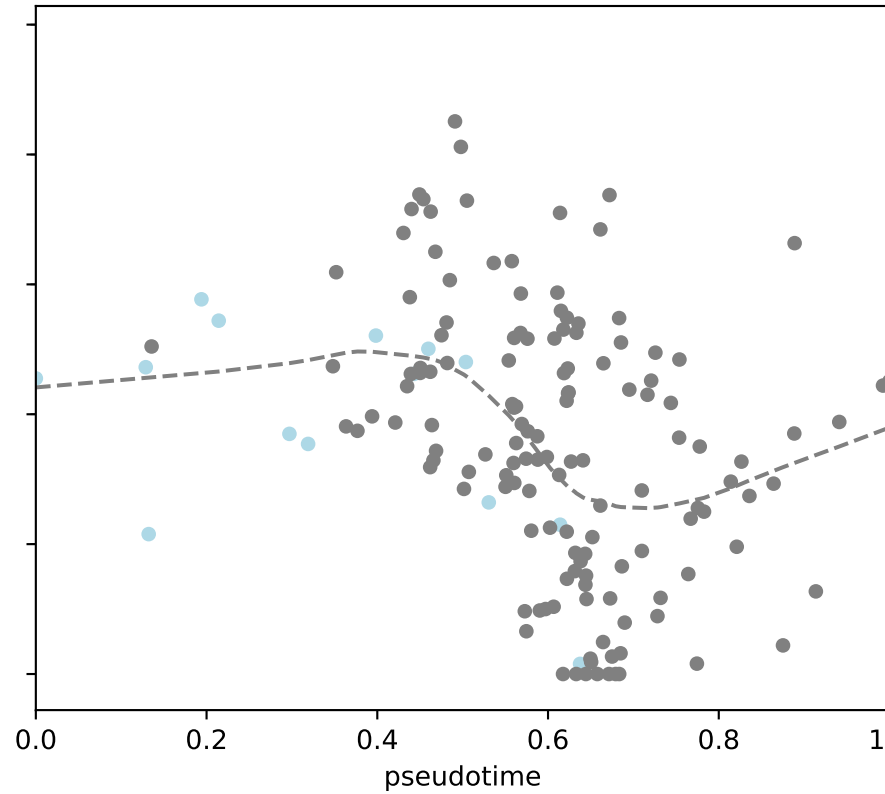
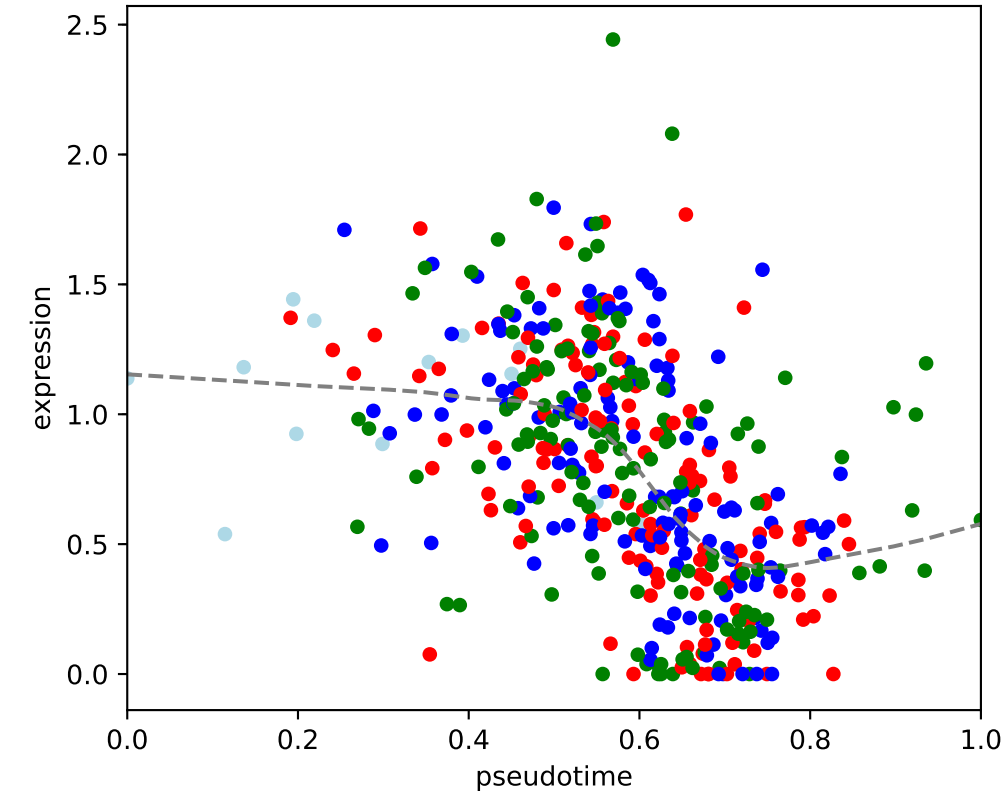
733 0spearman=-0.56

733 1spearman=-0.51



733 0spearman=-0.56

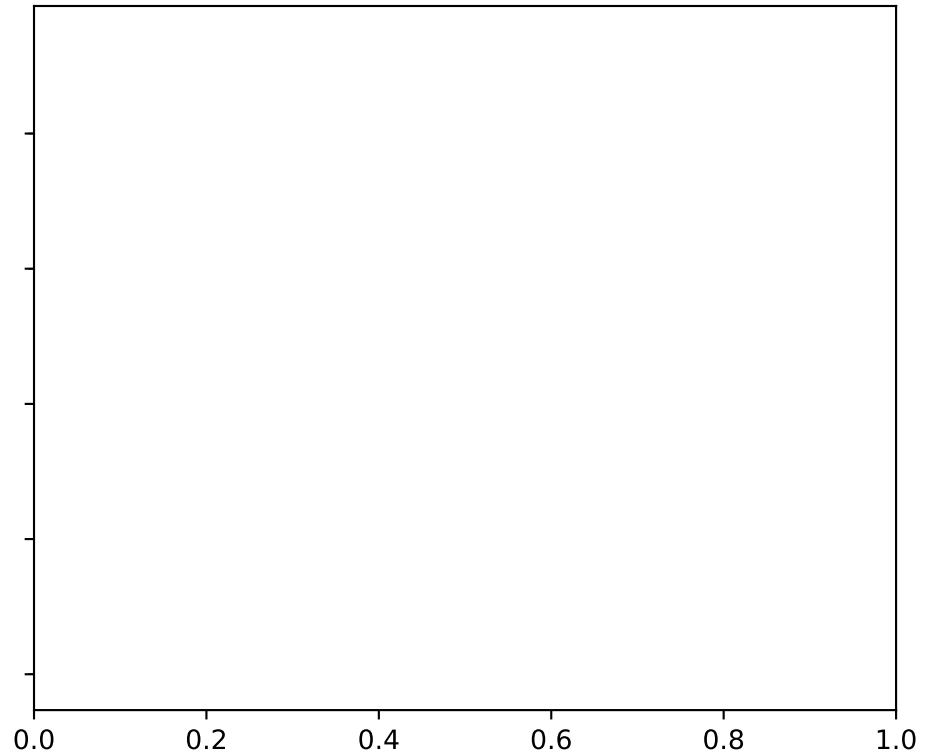
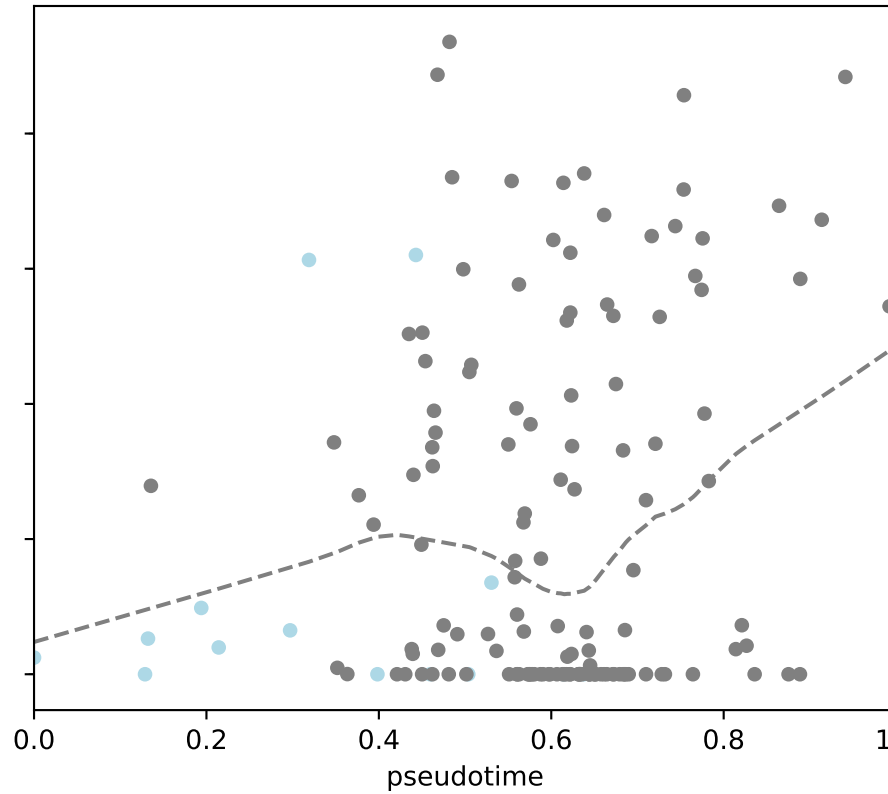
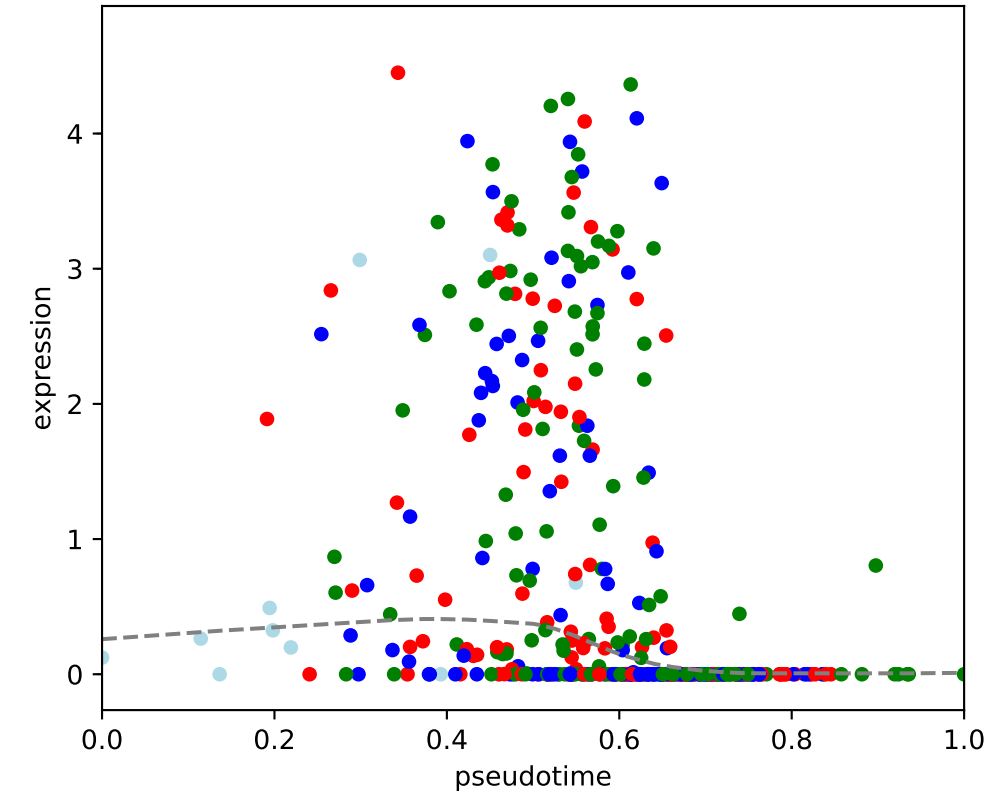
733 1spearman=-0.43



ENST00000588758 MIR7-3HG (MIR7-3 host gene)

733 0spearman=-0.56

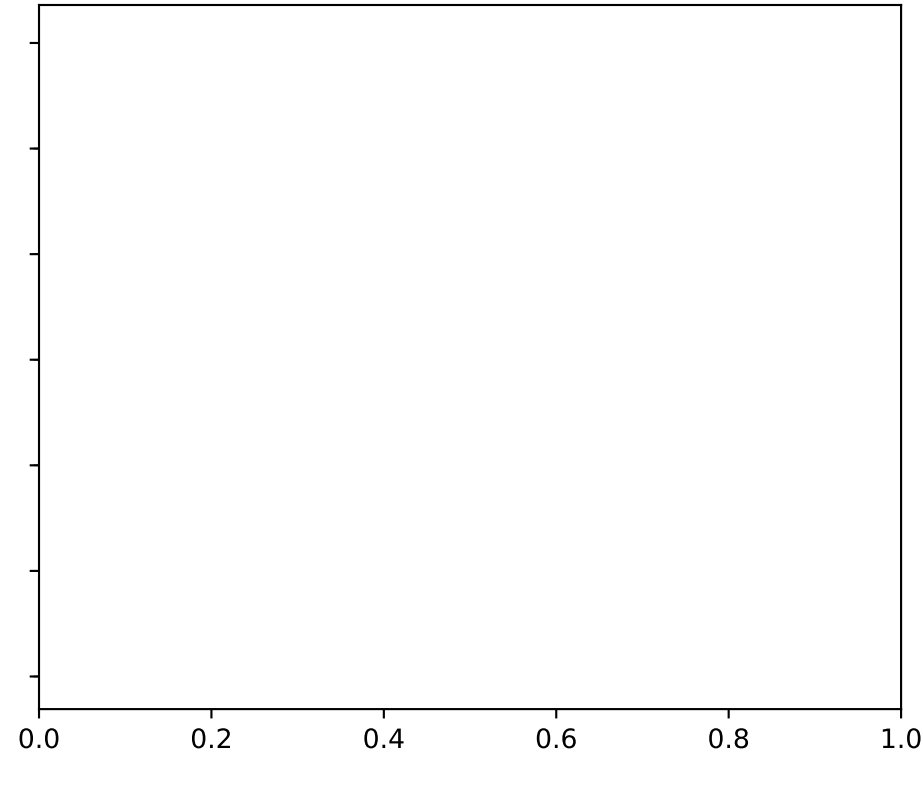
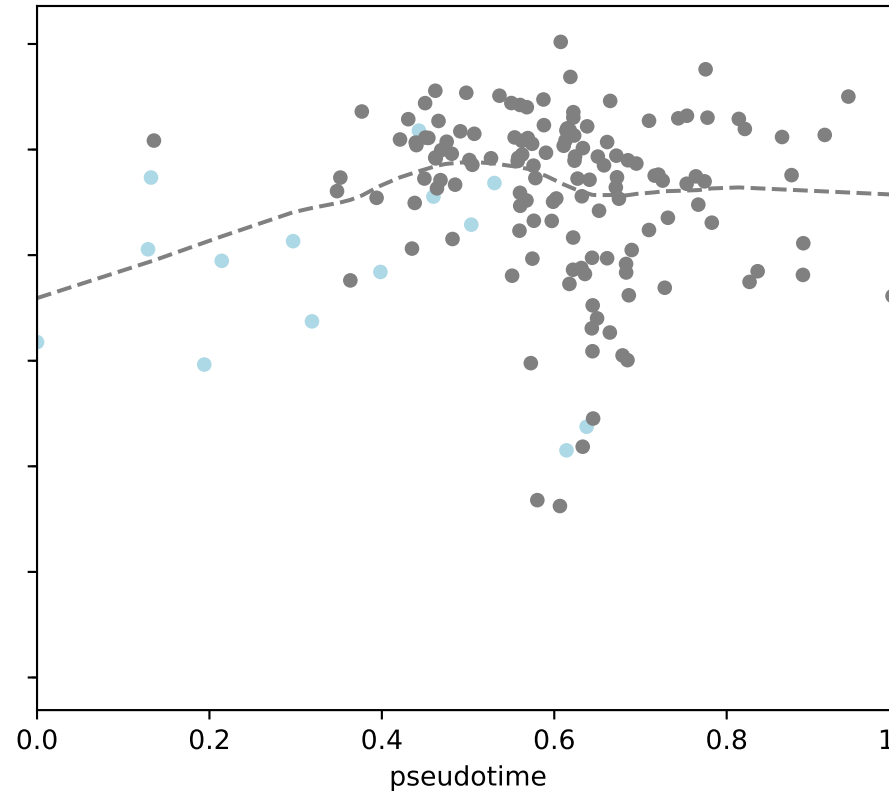
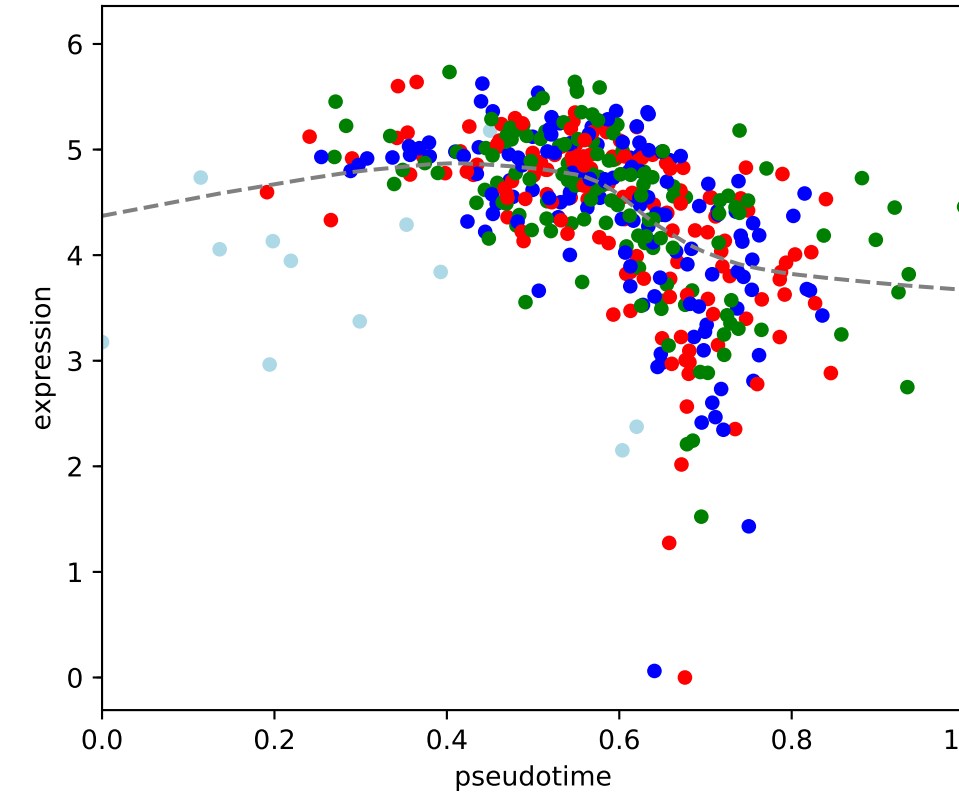
733 1spearman=-0.00



ENST00000281031 NDUFC2 (NADH:ubiquinone oxidoreductase subunit C2)

733 0spearman=-0.56

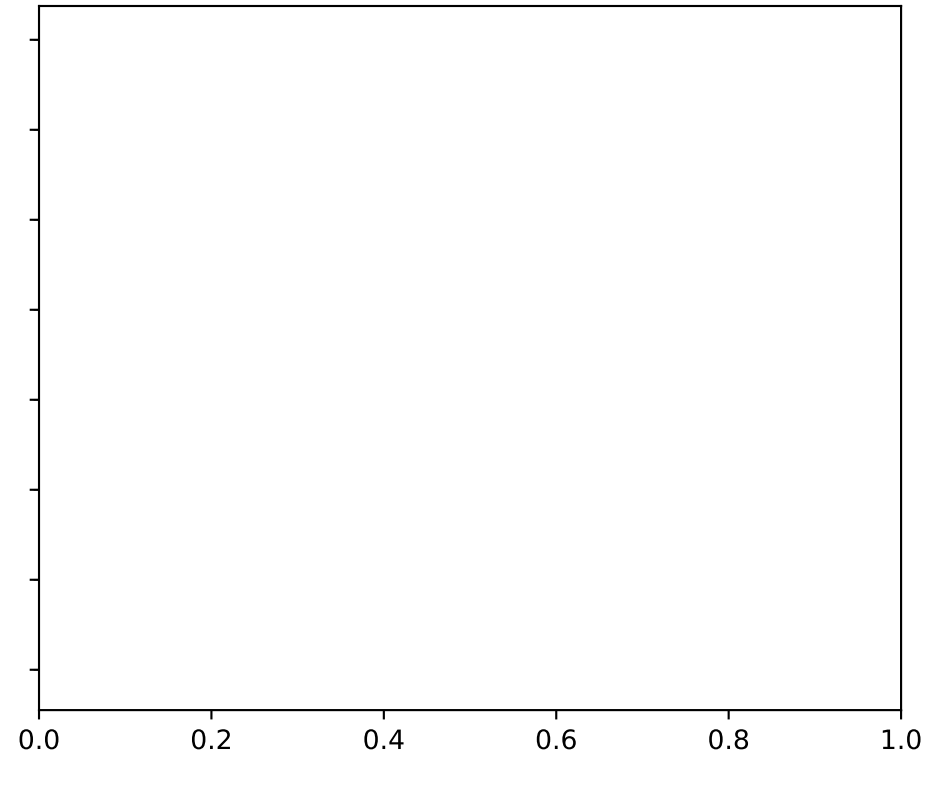
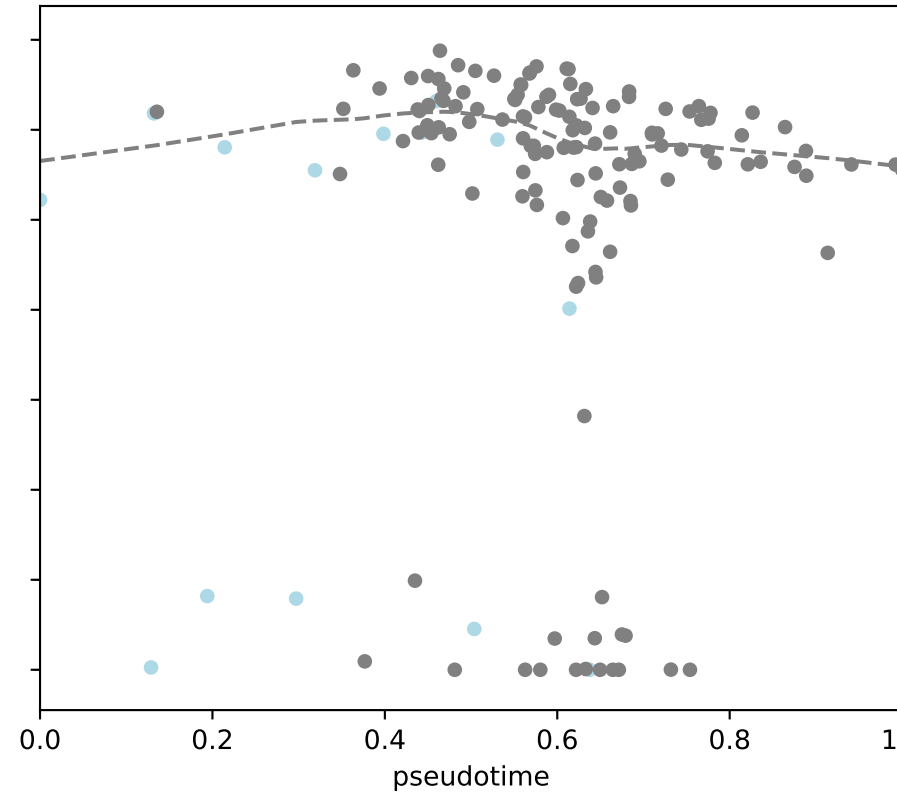
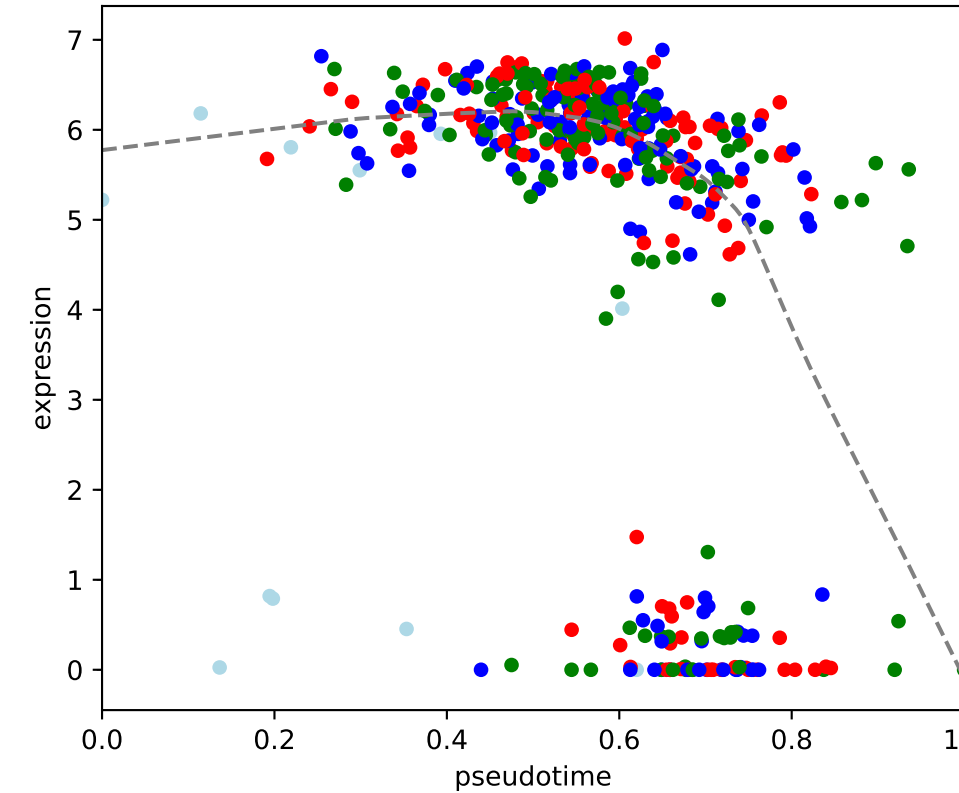
733 1spearman=-0.09



ENST00000401406 UQCR10 (ubiquinol-cytochrome c reductase, complex III subunit X)

733 0spearman=-0.56

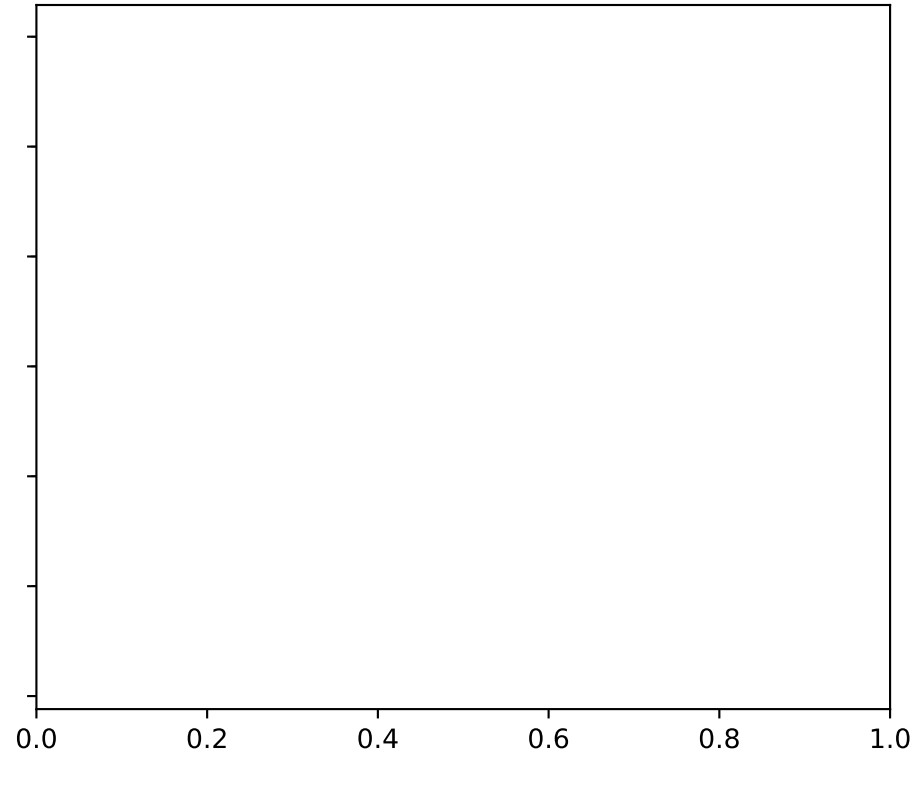
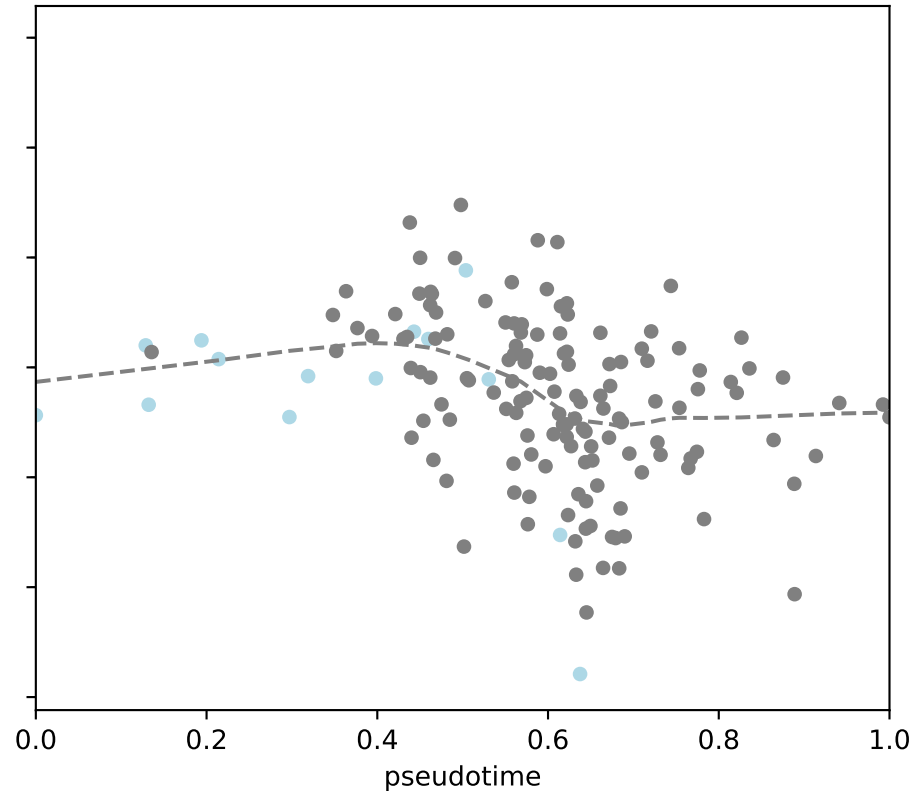
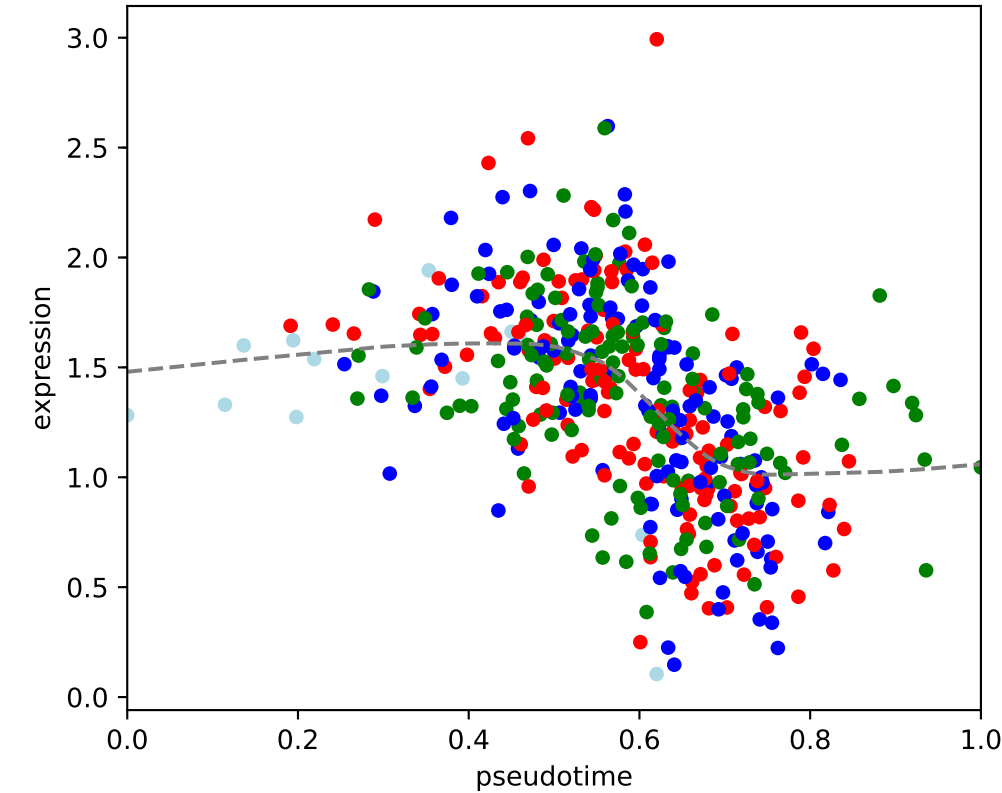
733 1spearman=-0.26



ENST00000557078 PPIAP6 (peptidylprolyl isomerase A pseudogene 6)

733 0spearman=-0.56

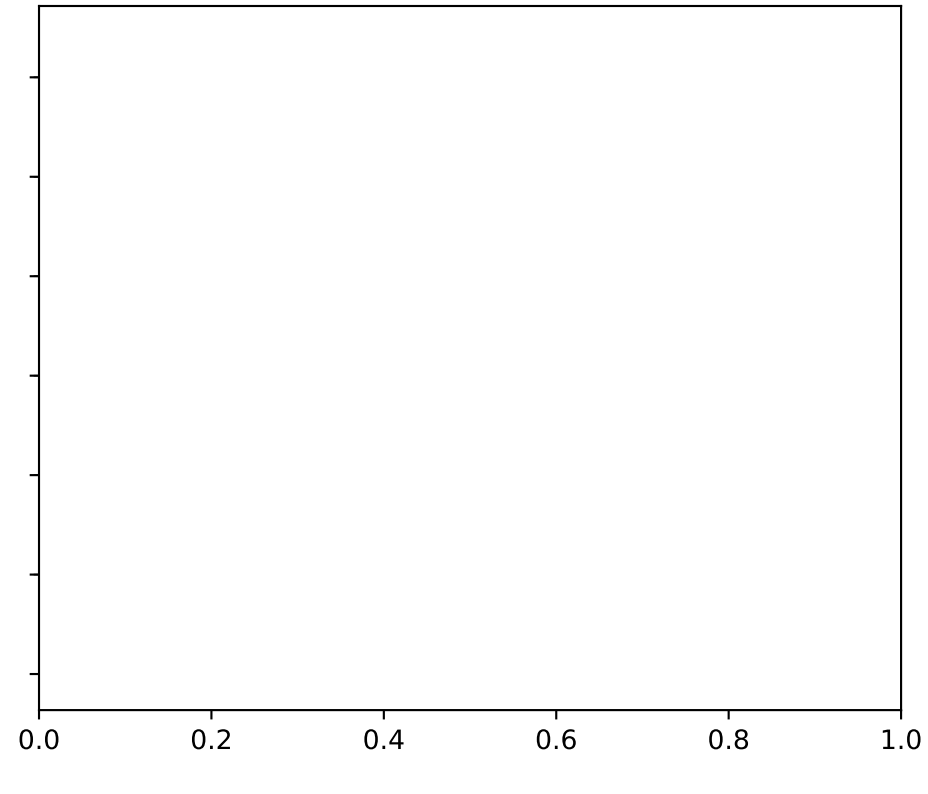
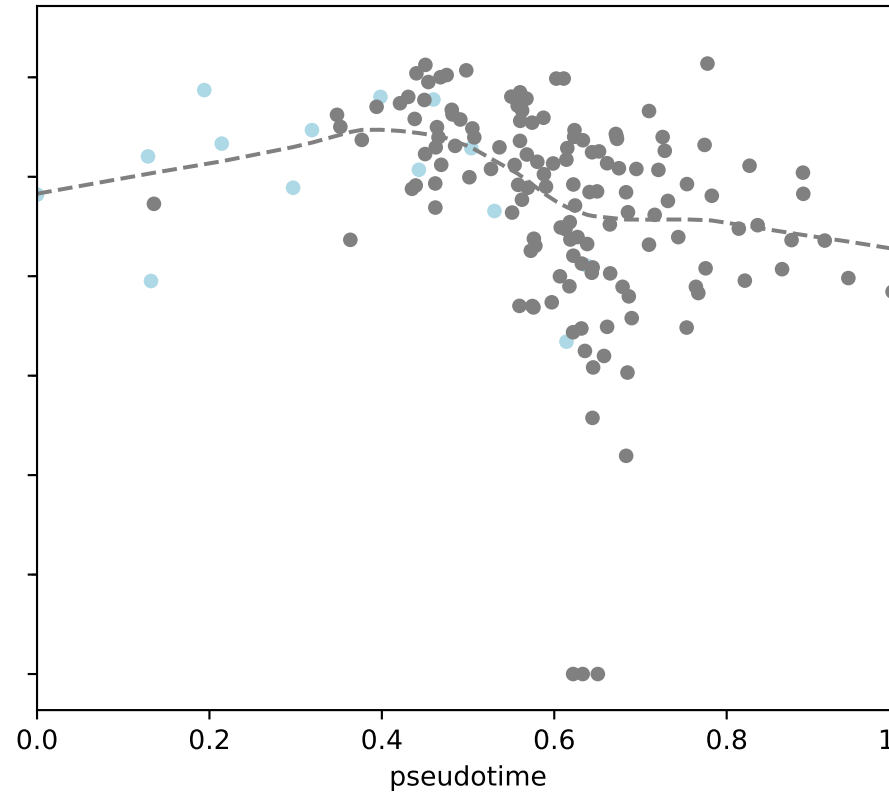
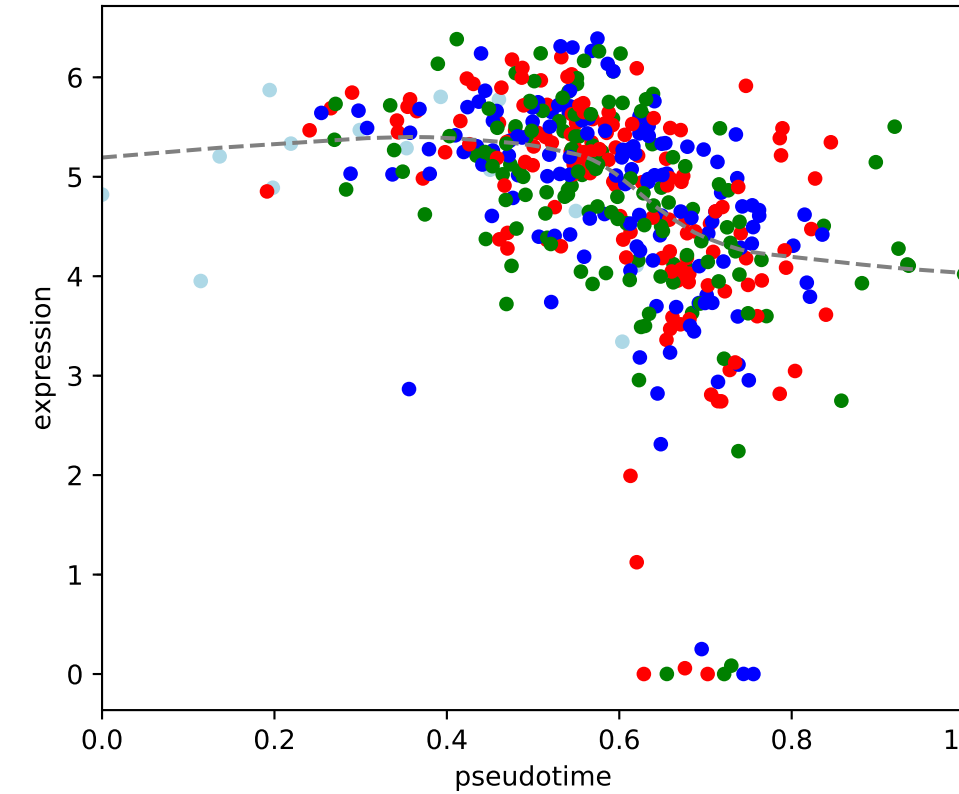
733 1spearman=-0.43



ENST00000233893 HSPE1 (heat shock protein family E (Hsp10) member 1)

733 0spearman=-0.56

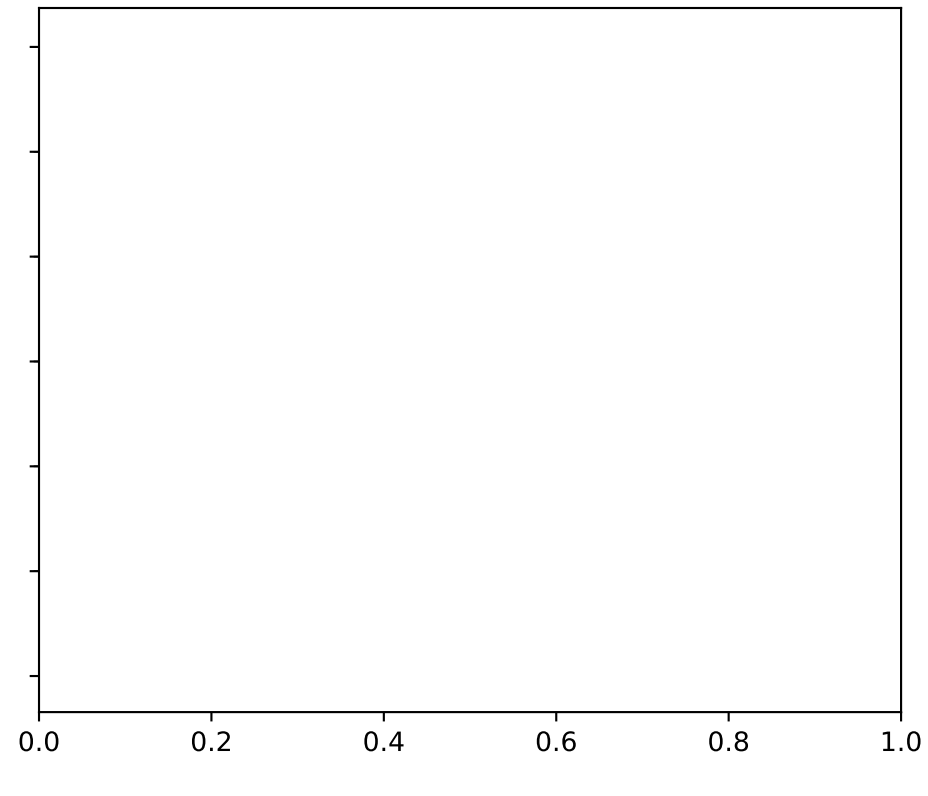
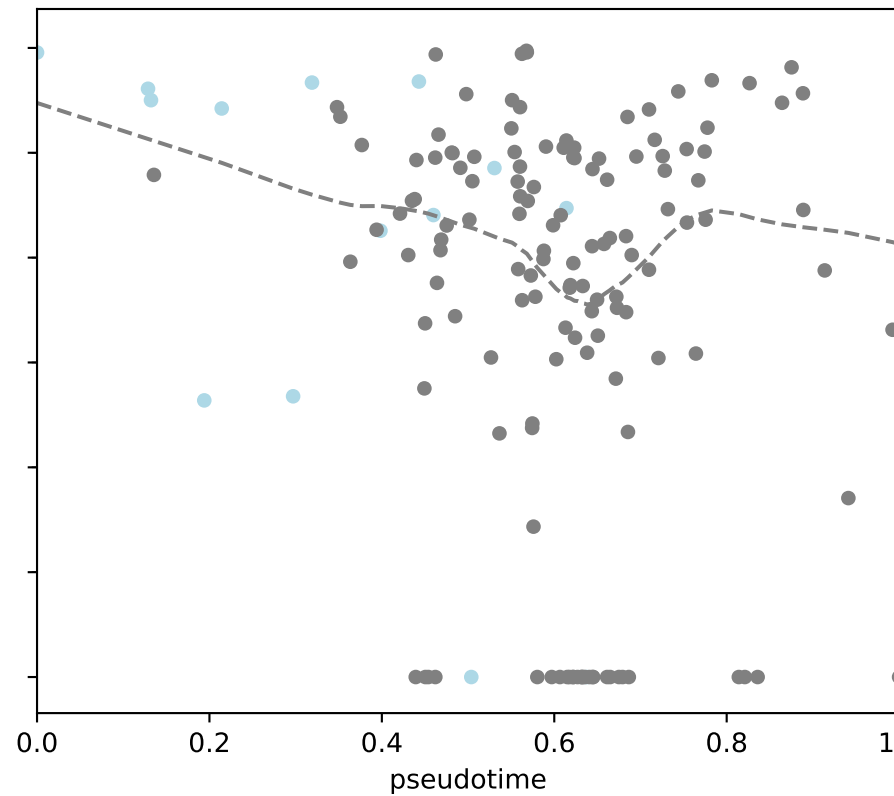
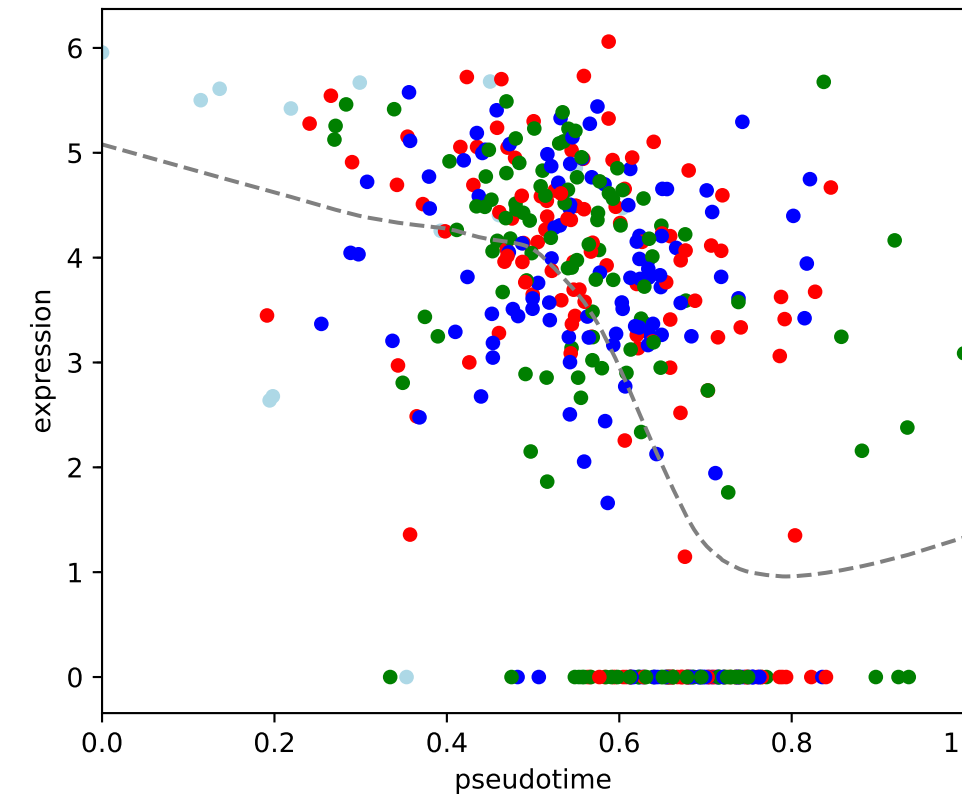
733 1spearman=-0.48



ENST00000338702 MAOA (monoamine oxidase A)

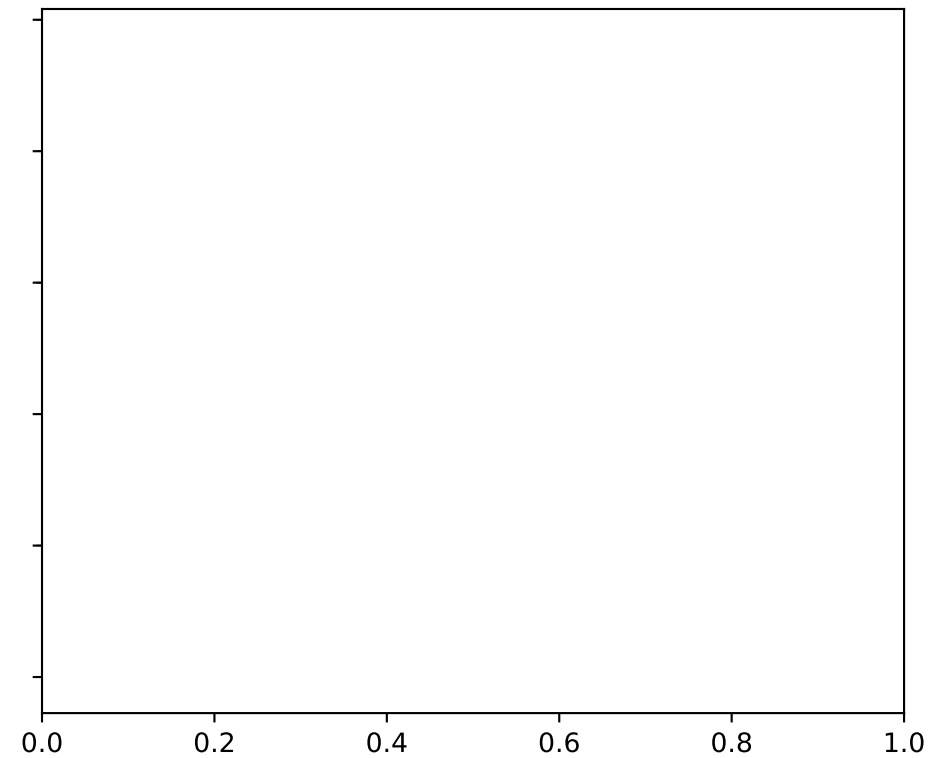
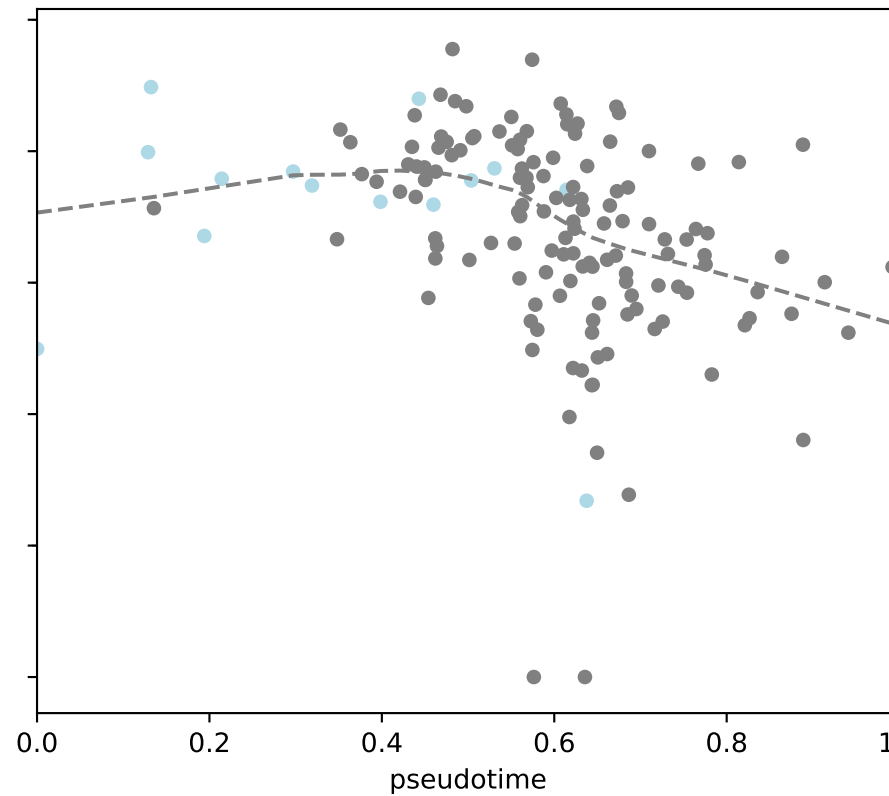
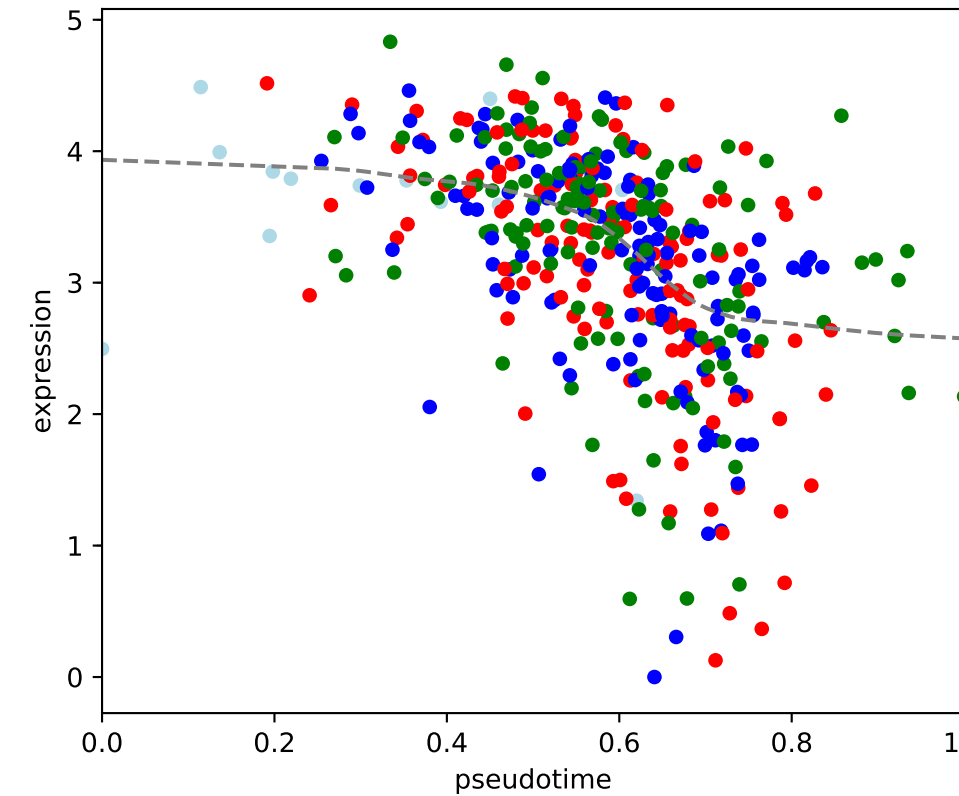
733 0spearman=-0.56

733 1spearman=-0.17



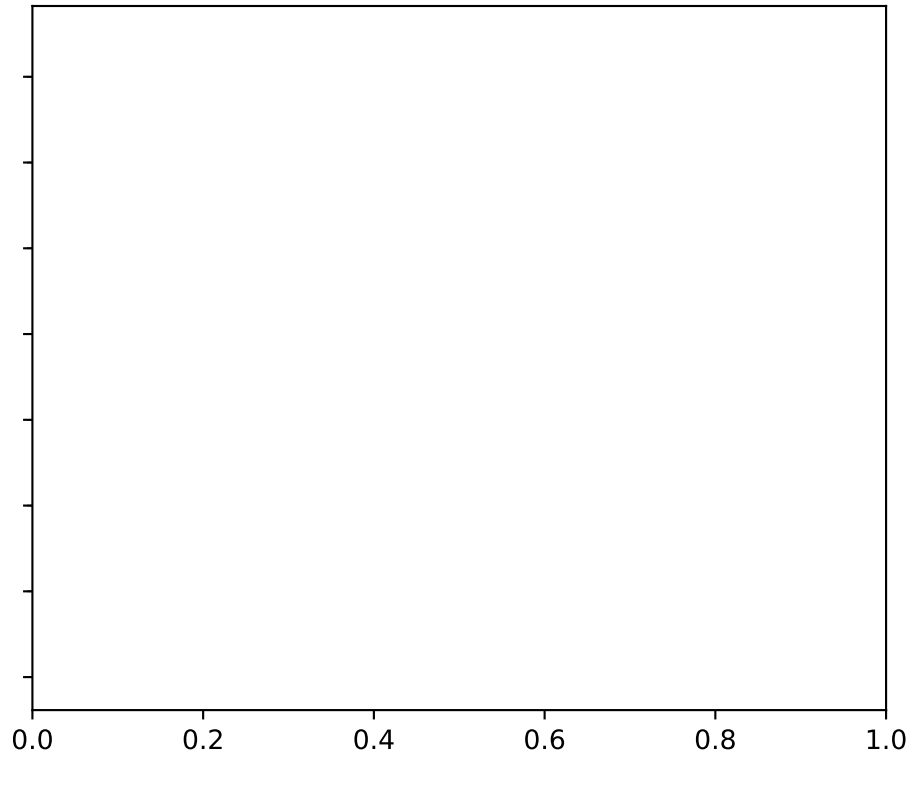
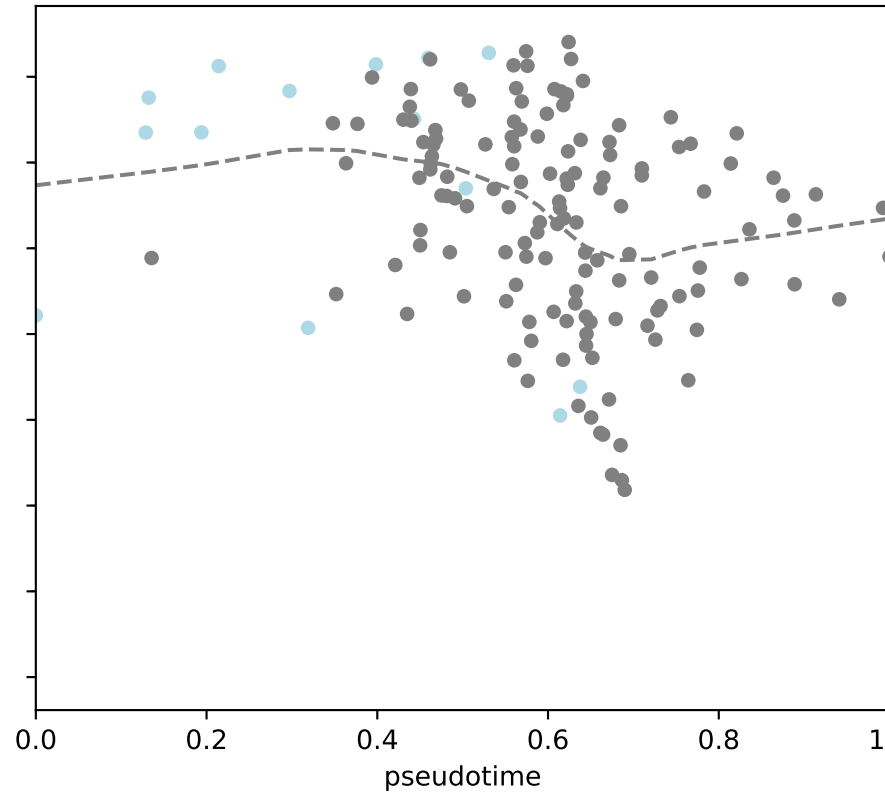
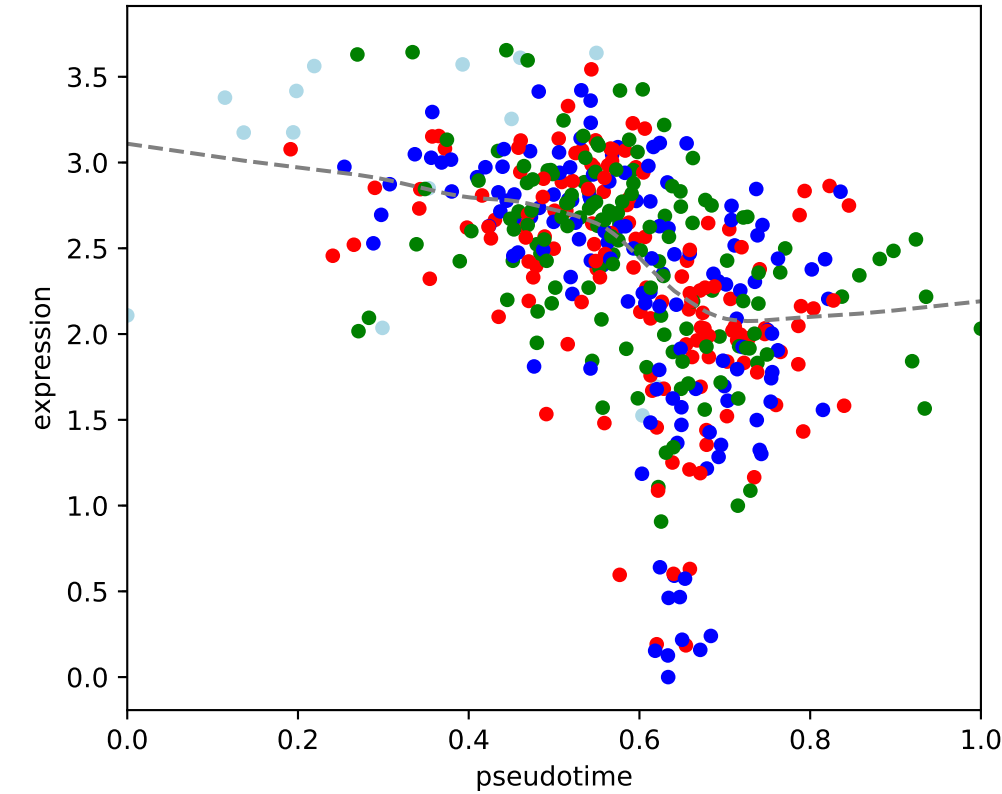
733 0spearman=-0.56

733 1spearman=-0.47



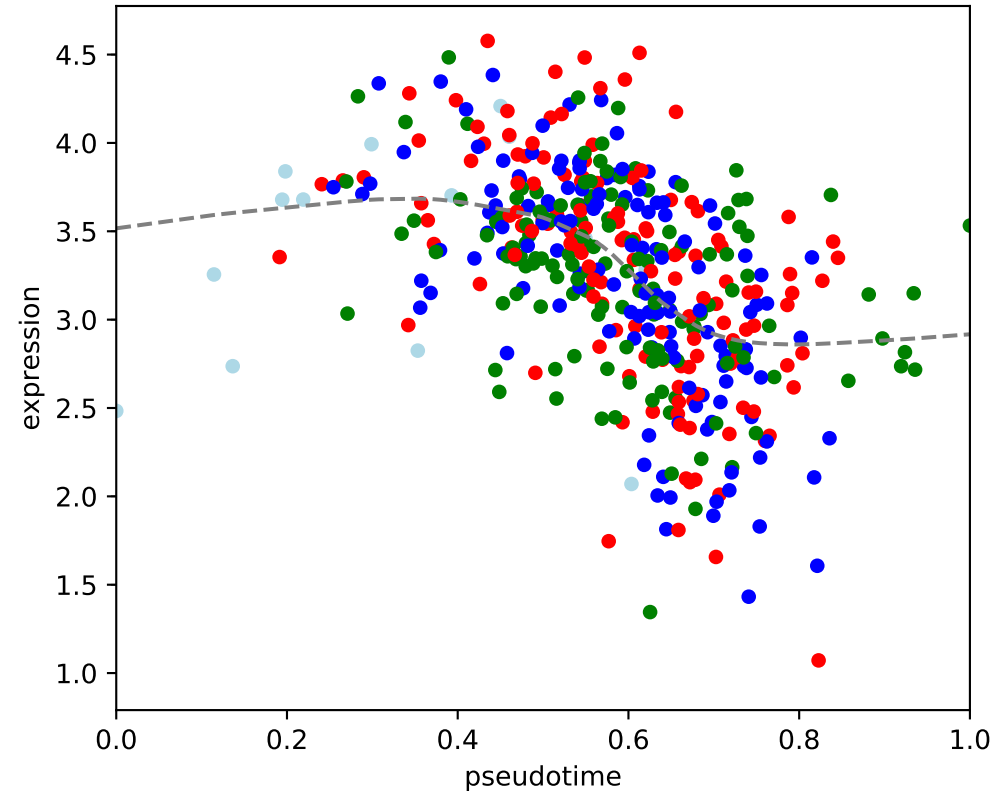
733 0spearman=-0.56

733 1spearman=-0.38

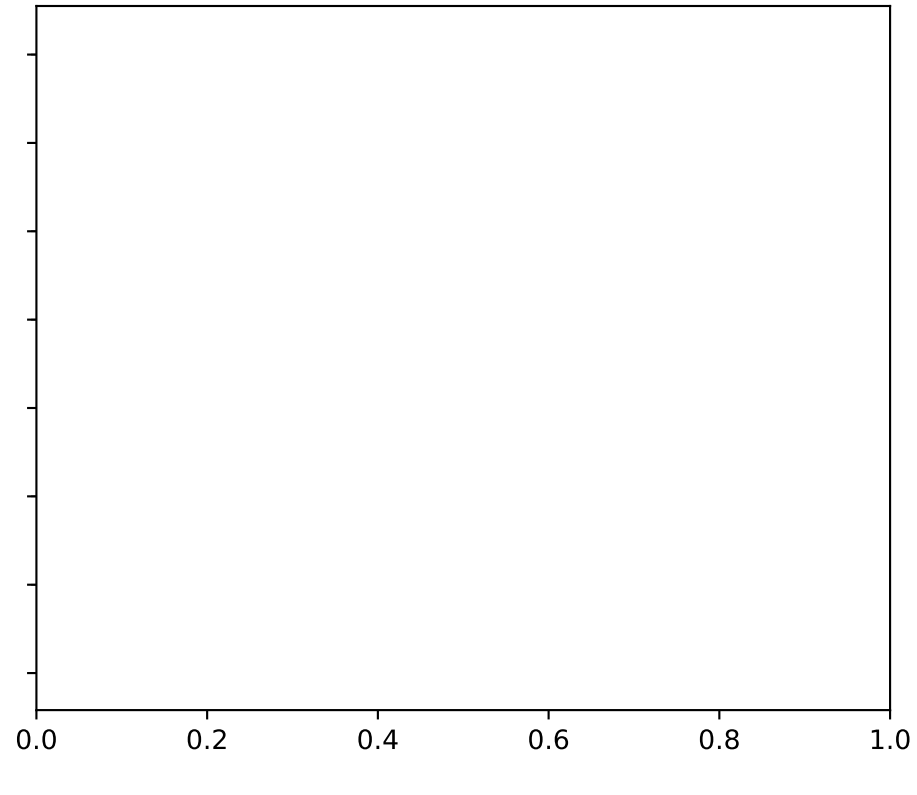
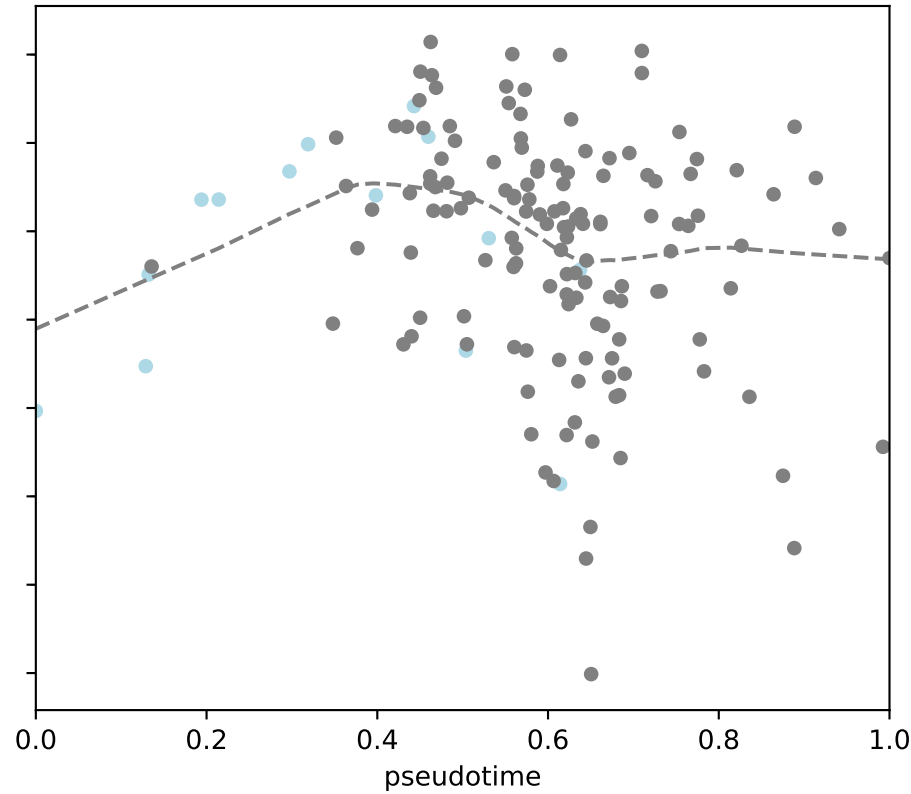


ENST00000433675 AC007969.1

733 0spearman=-0.56



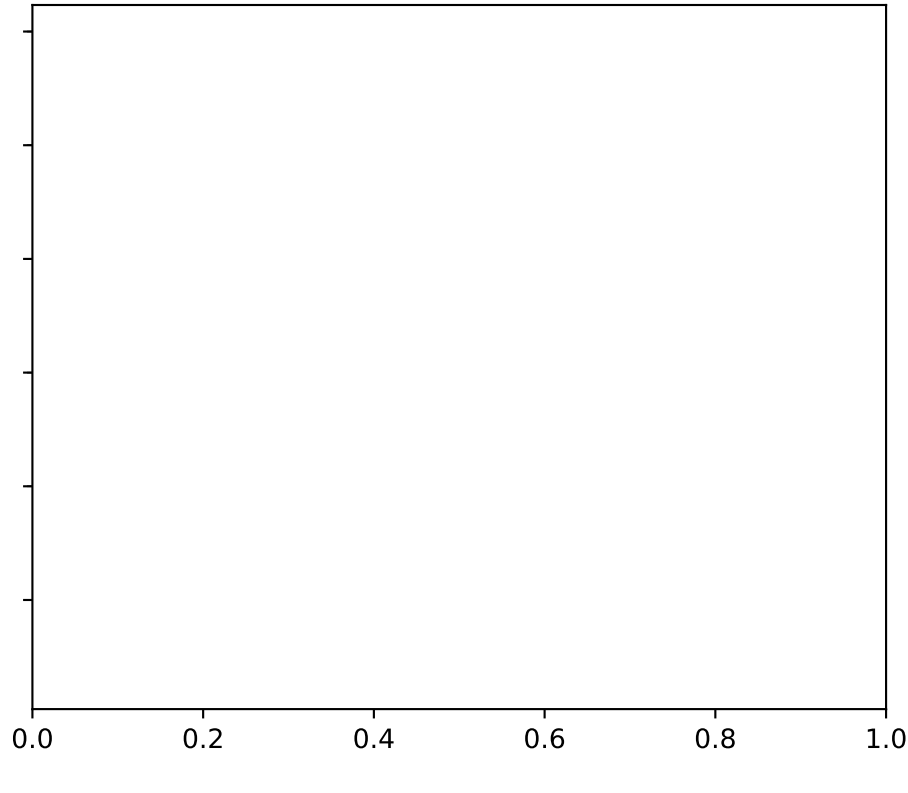
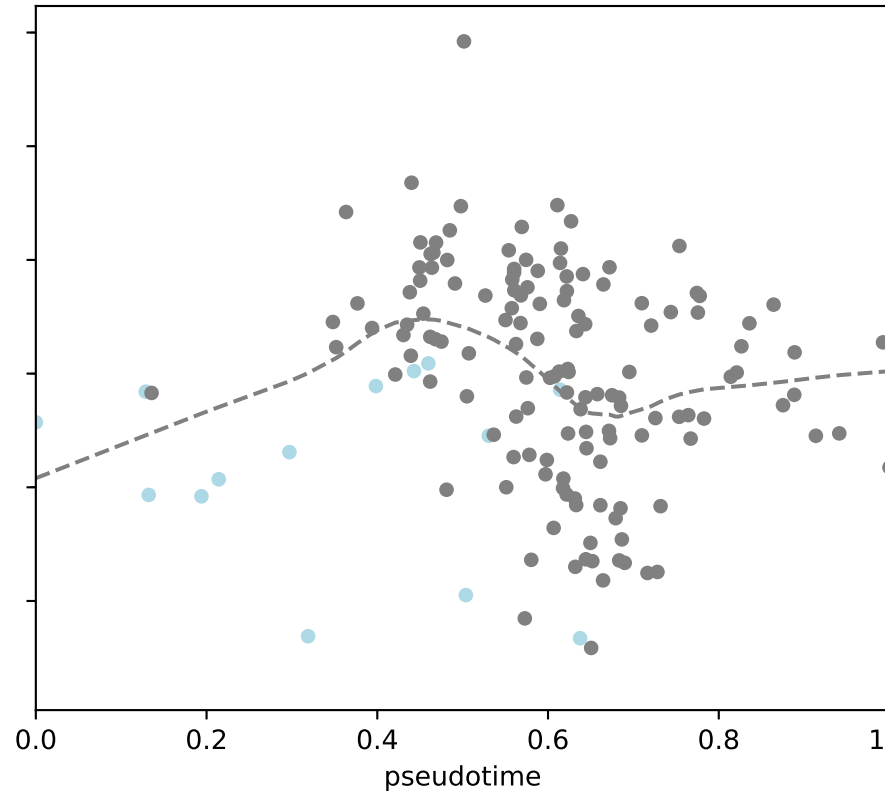
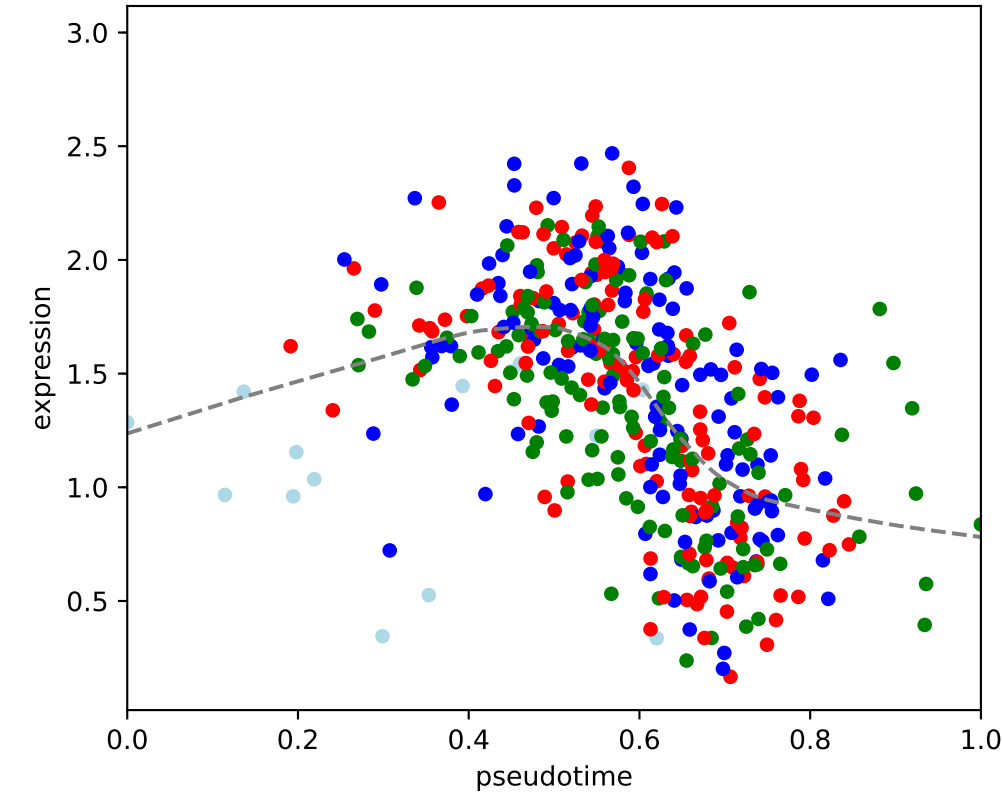
733 1spearman=-0.27



ENST00000503744 CHCHD2P2 (coiled-coil-helix-coiled-coil-helix domain containing 2 pseudogene 2)

733 0spearman=-0.55

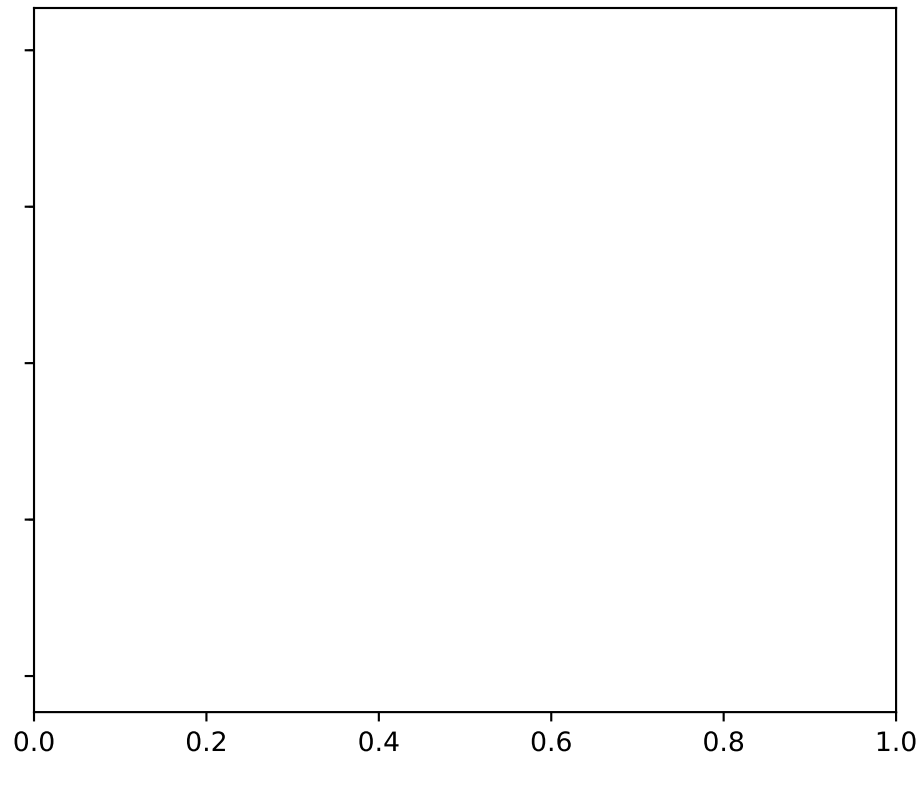
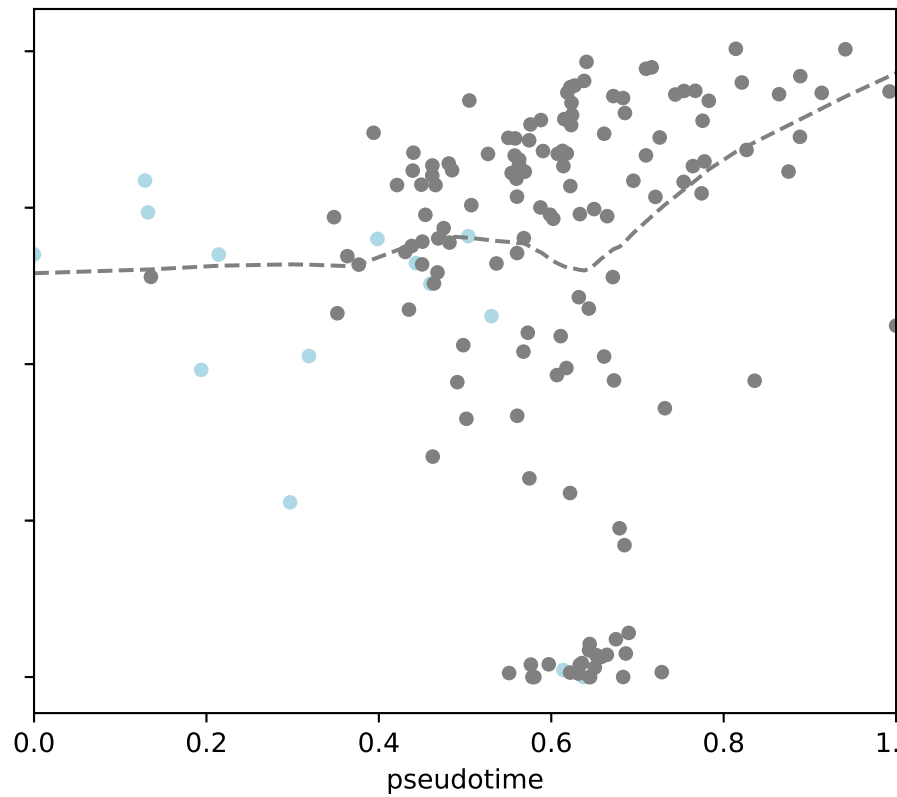
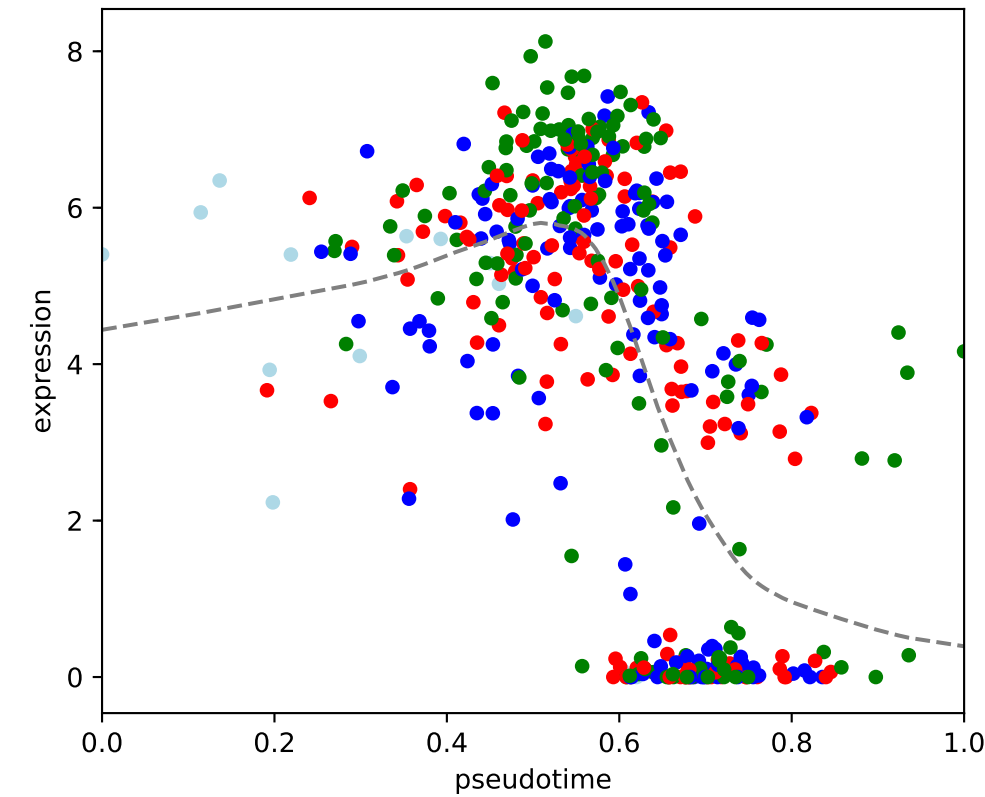
733 1spearman=-0.26



ENST00000252854 OLFM1 (olfactomedin 1)

733 0spearman=-0.55

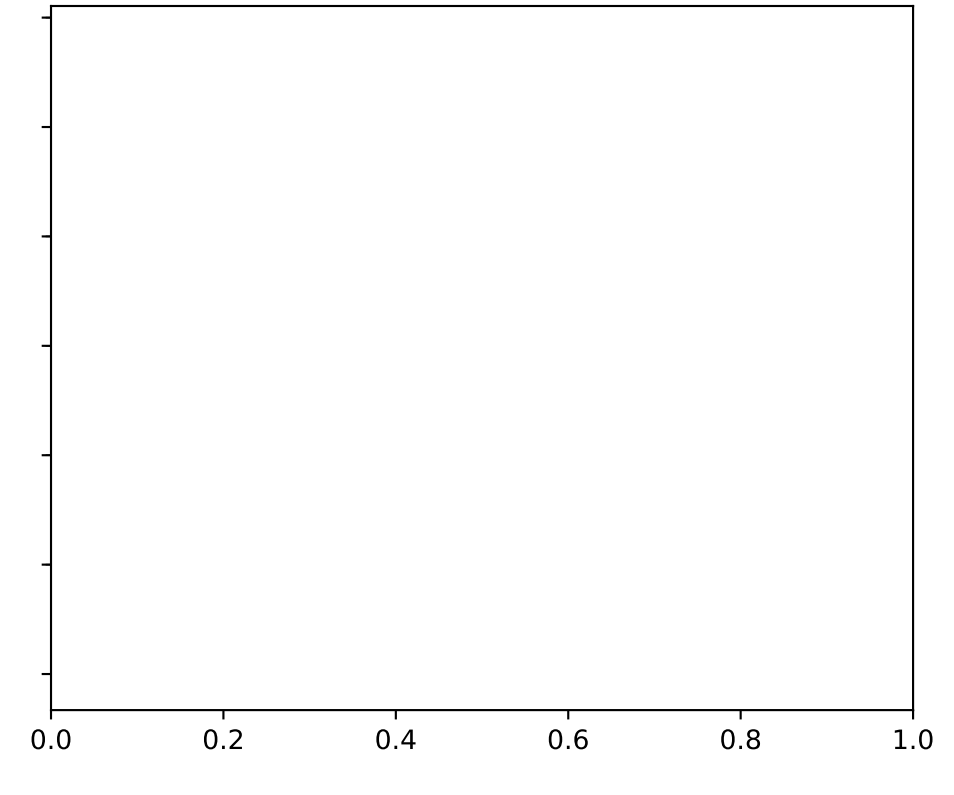
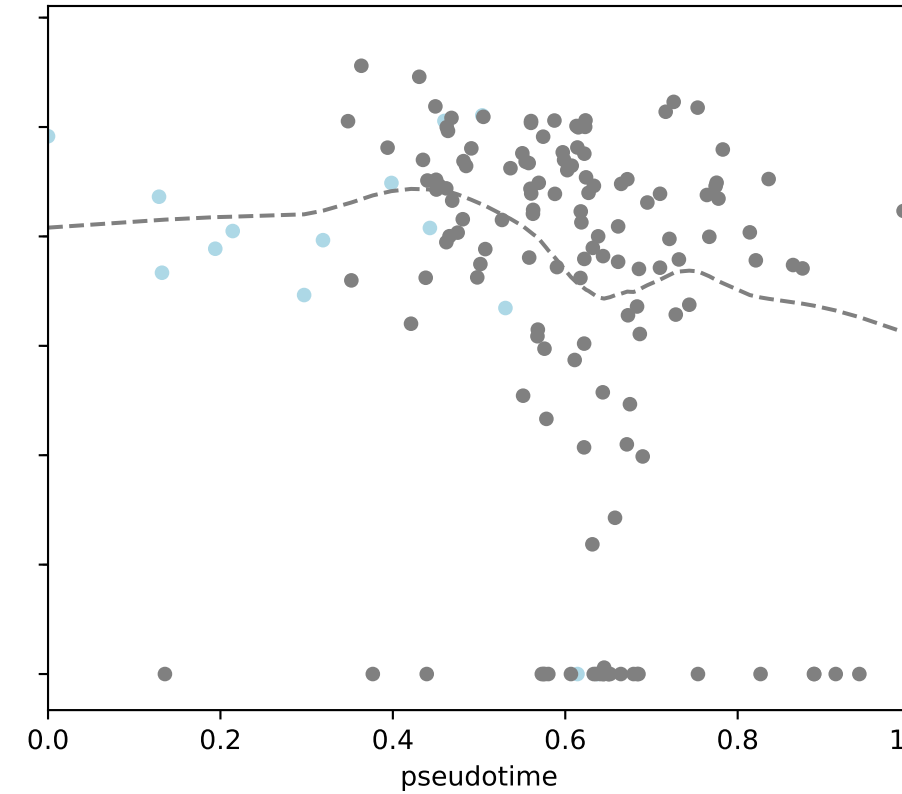
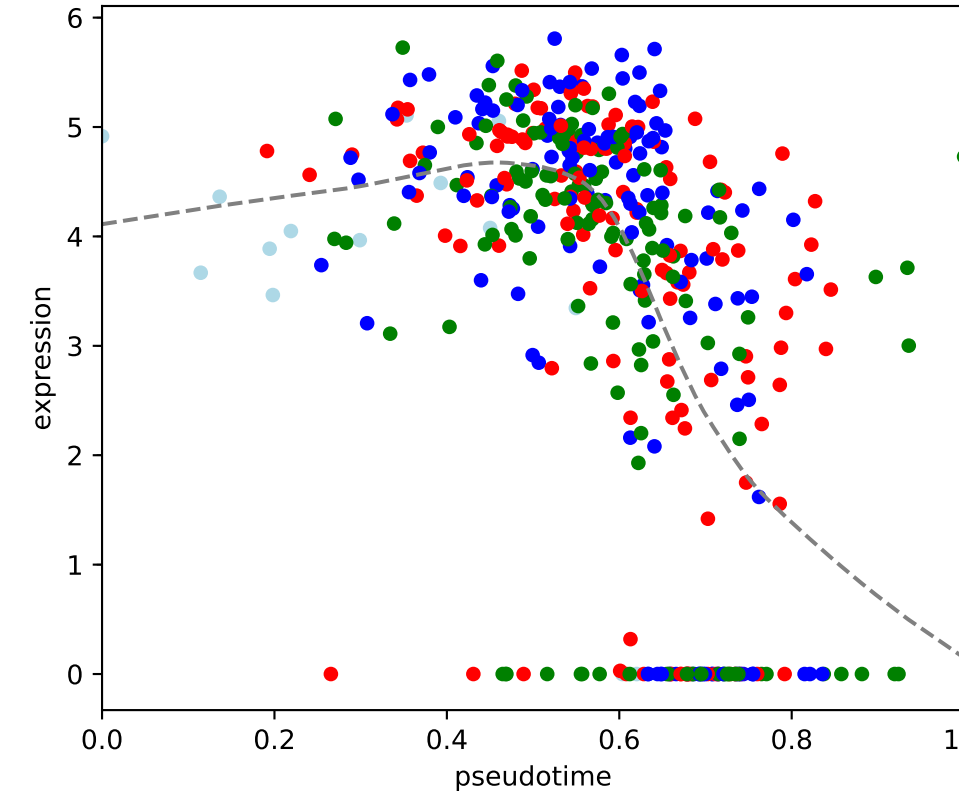
733 1spearman=0.20



ENST00000432020 MDH2 (malate dehydrogenase 2)

733 0spearman=-0.55

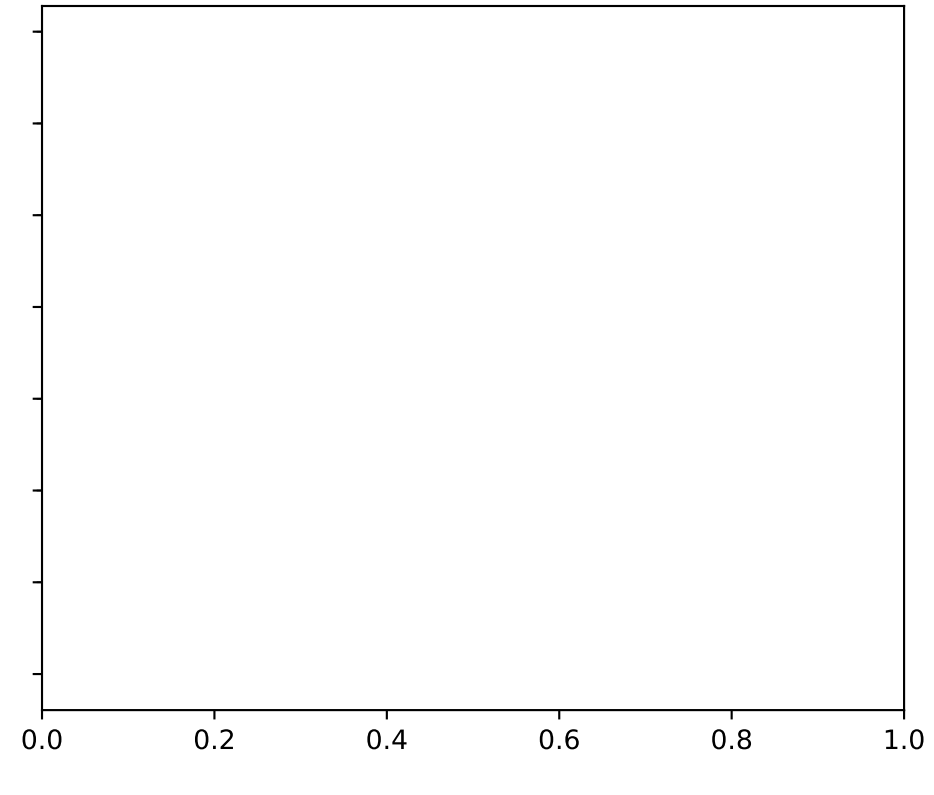
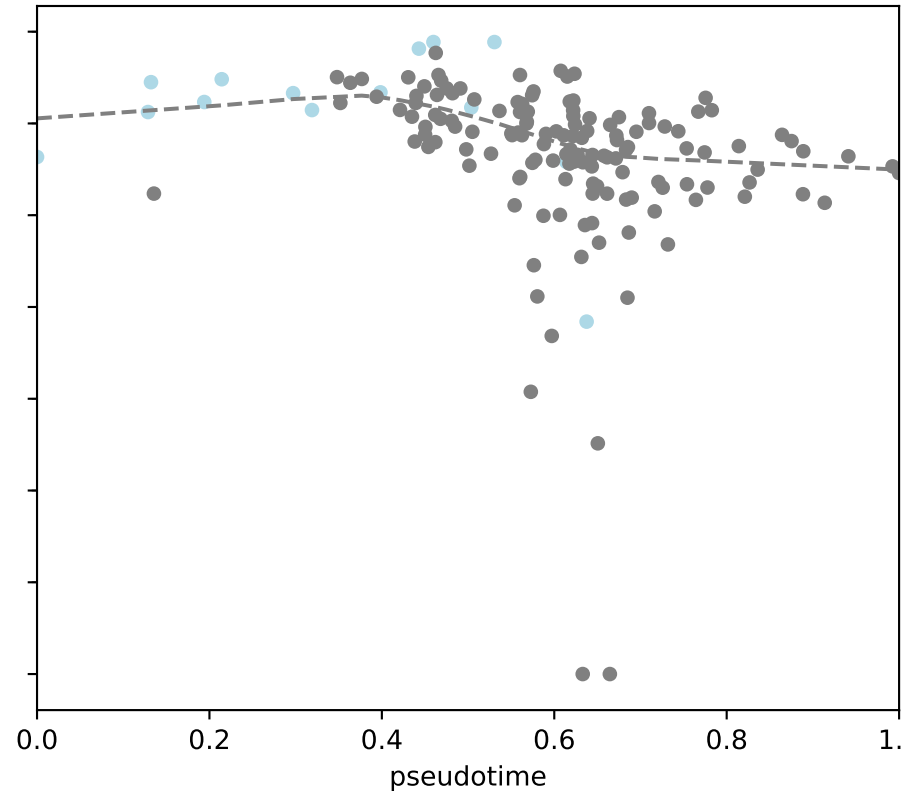
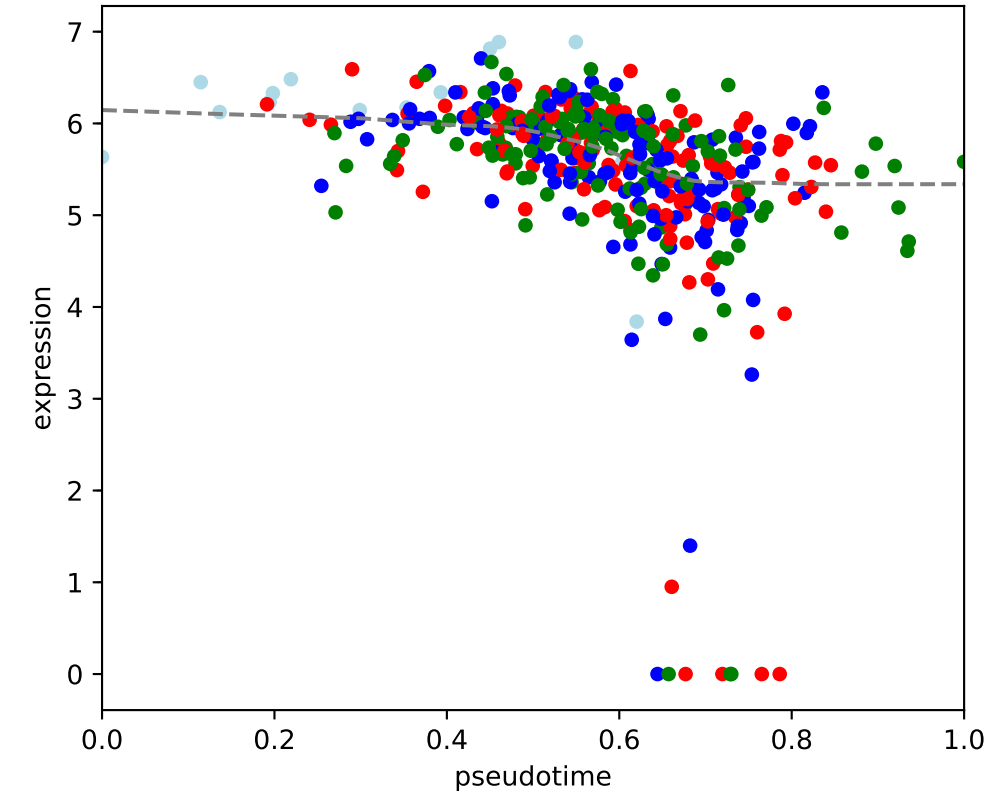
733 1spearman=-0.30



ENST00000527673 RPS25 (ribosomal protein S25)

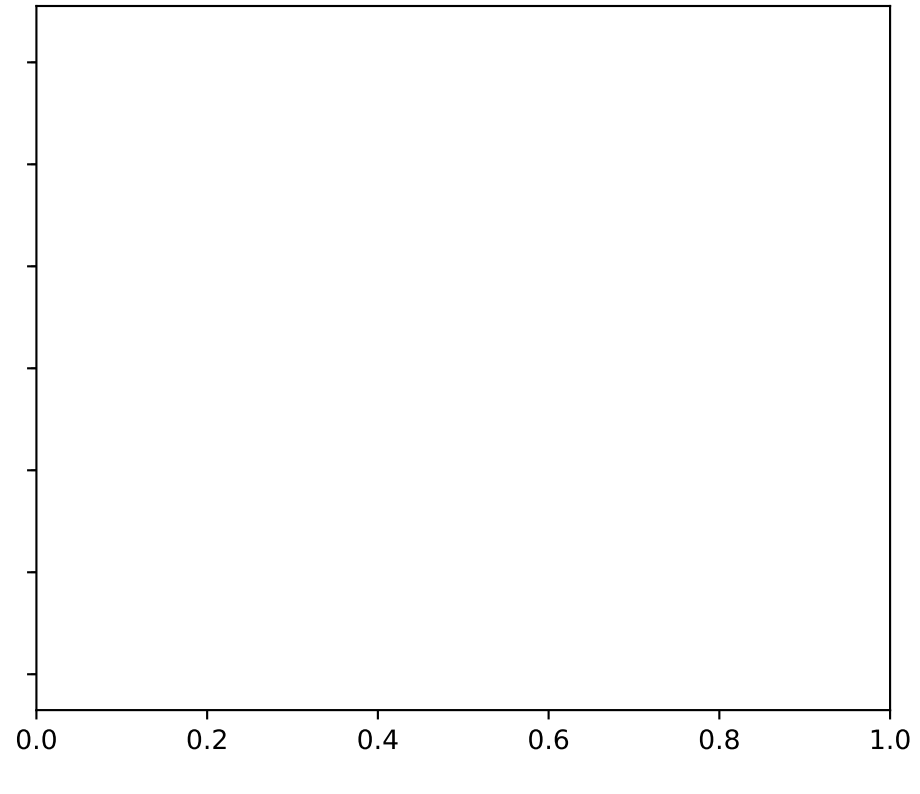
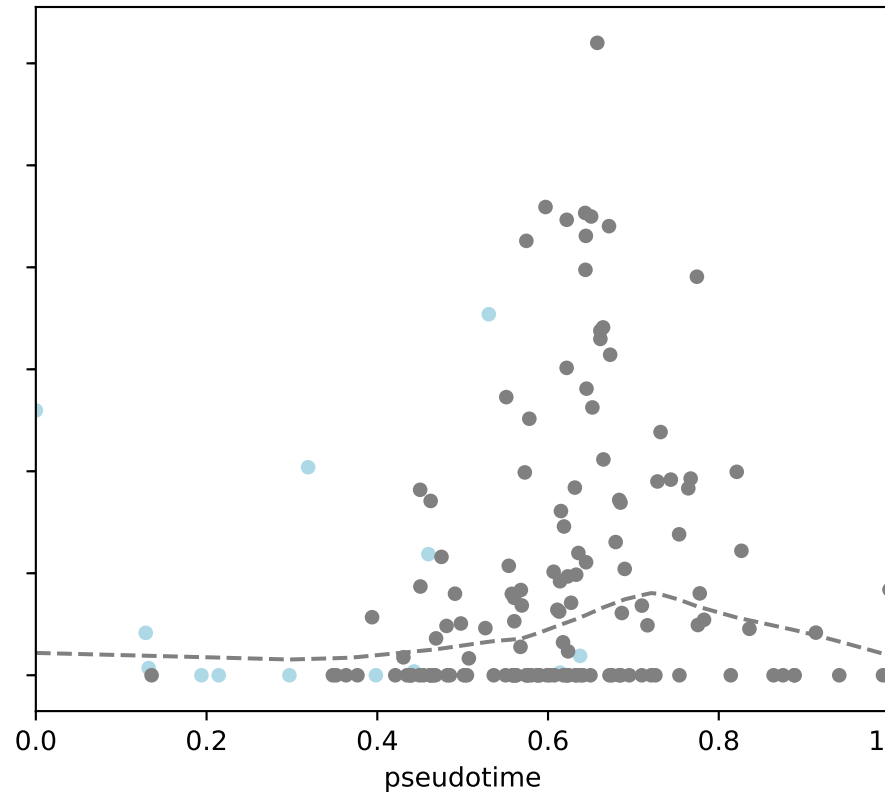
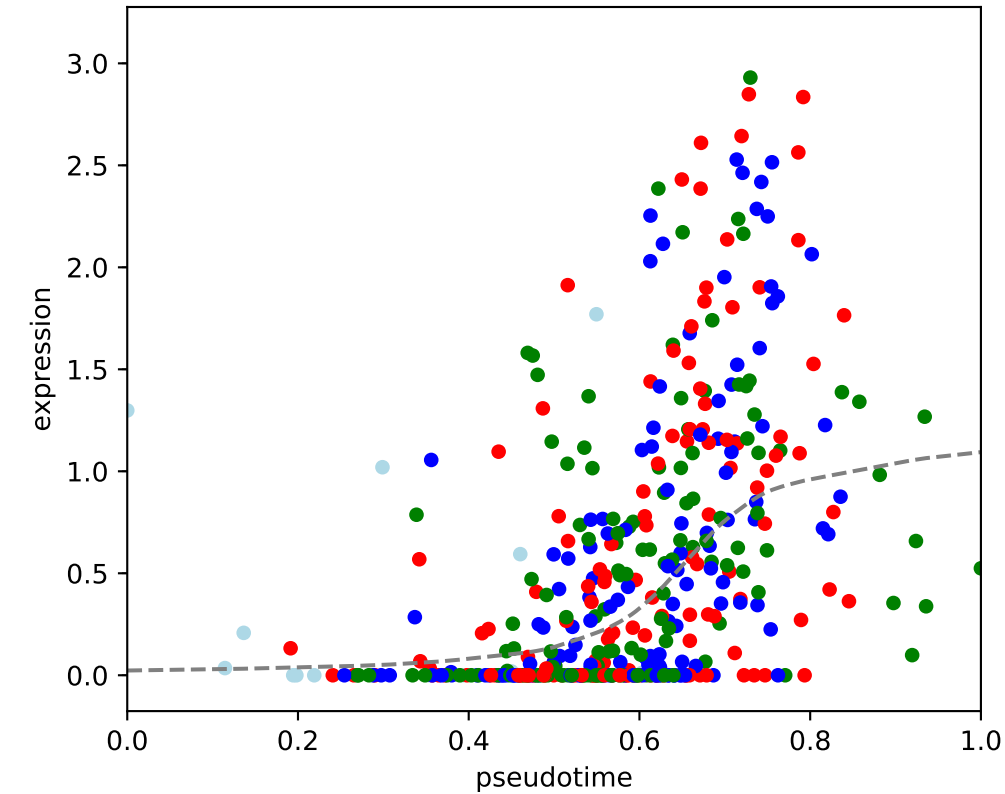
733 0spearman=-0.55

733 1spearman=-0.53



733 0spearman=0.55

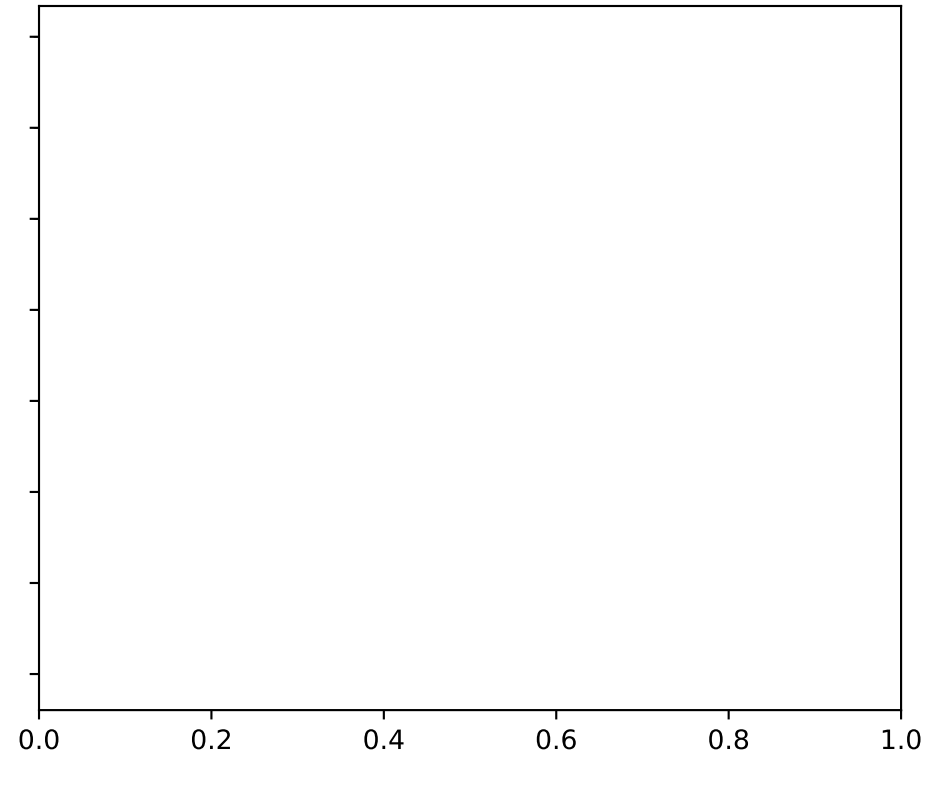
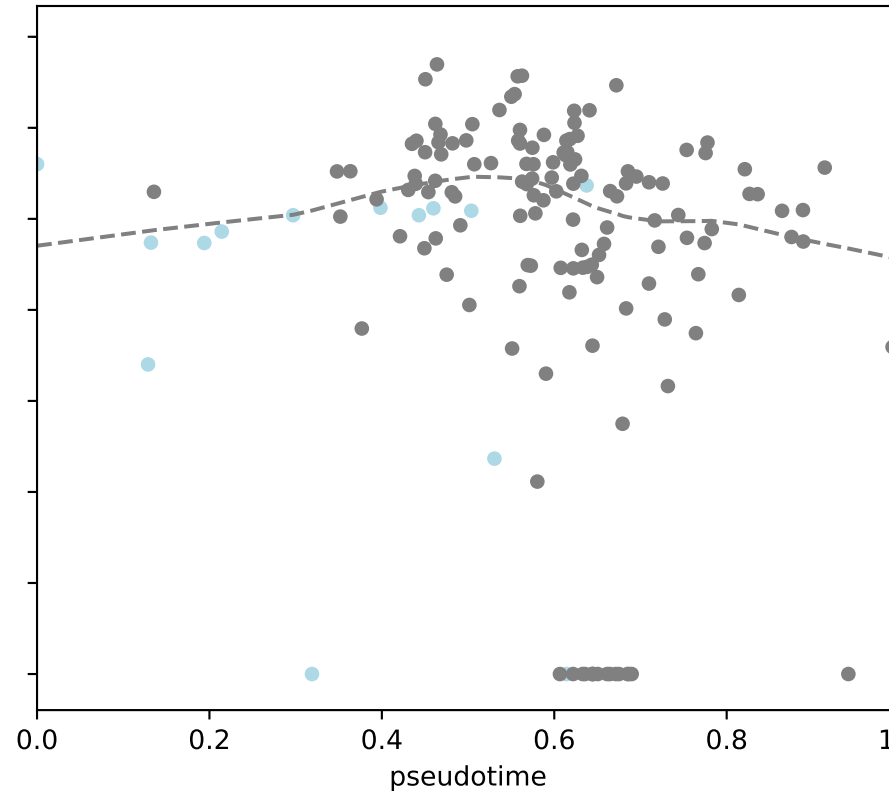
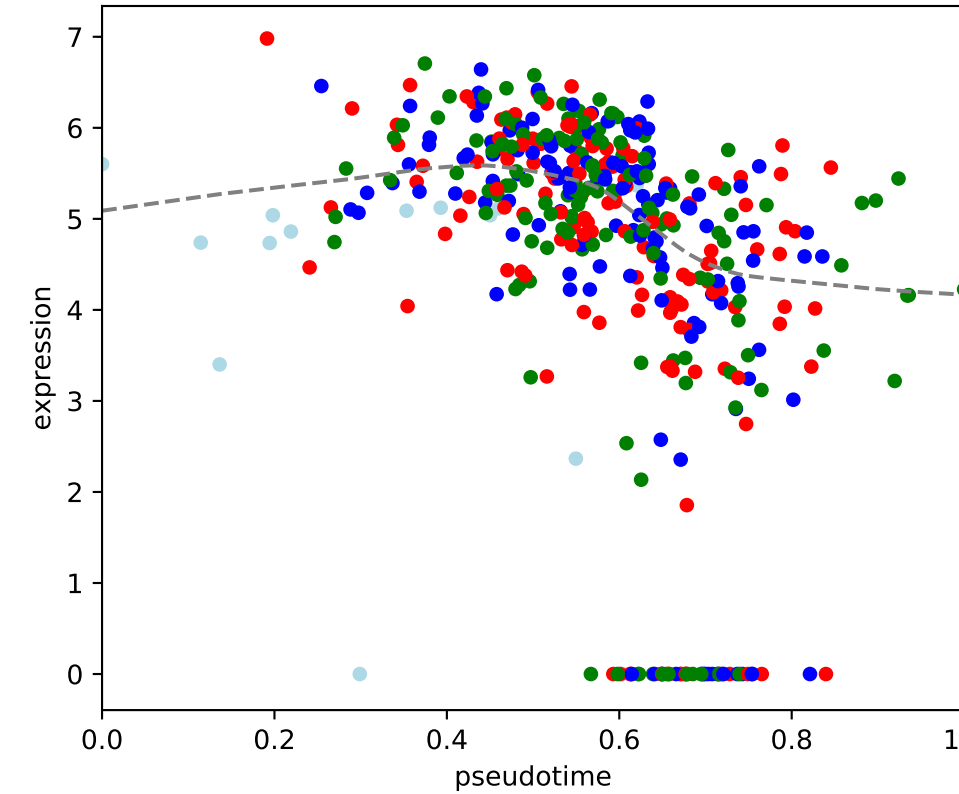
733 1spearman=0.25



ENST00000313115 ATOX1 (antioxidant 1 copper chaperone)

733 0spearman=-0.55

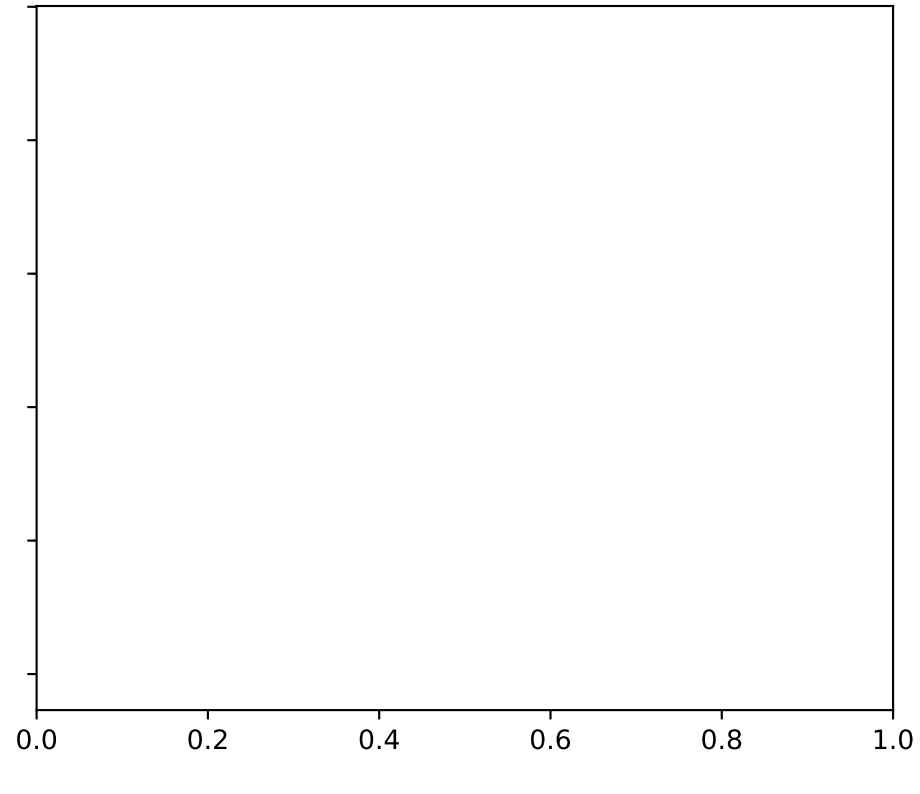
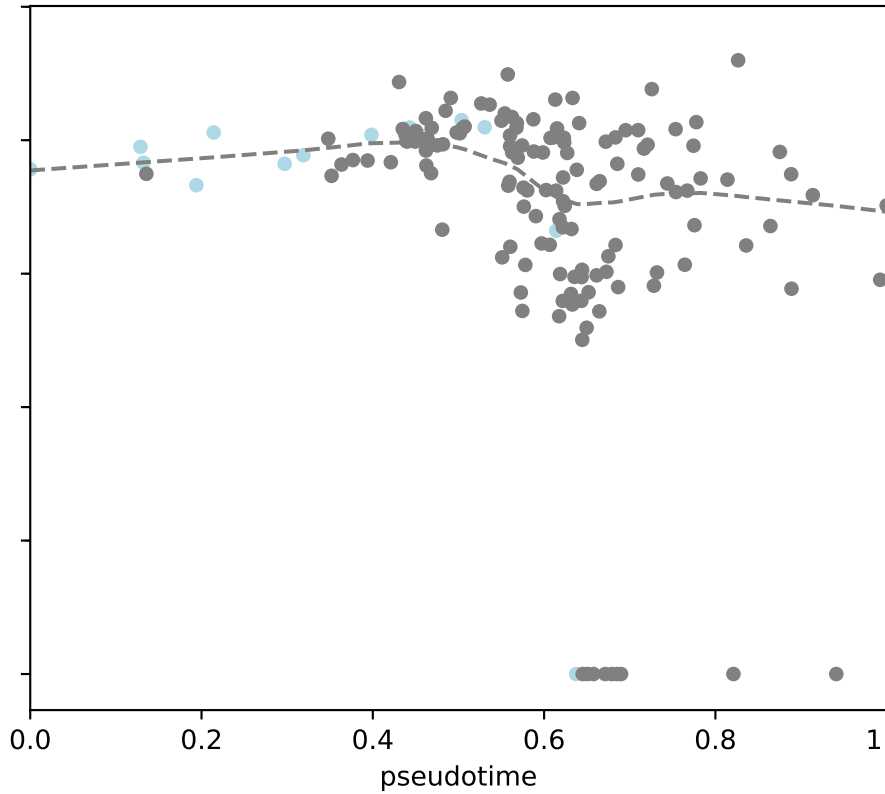
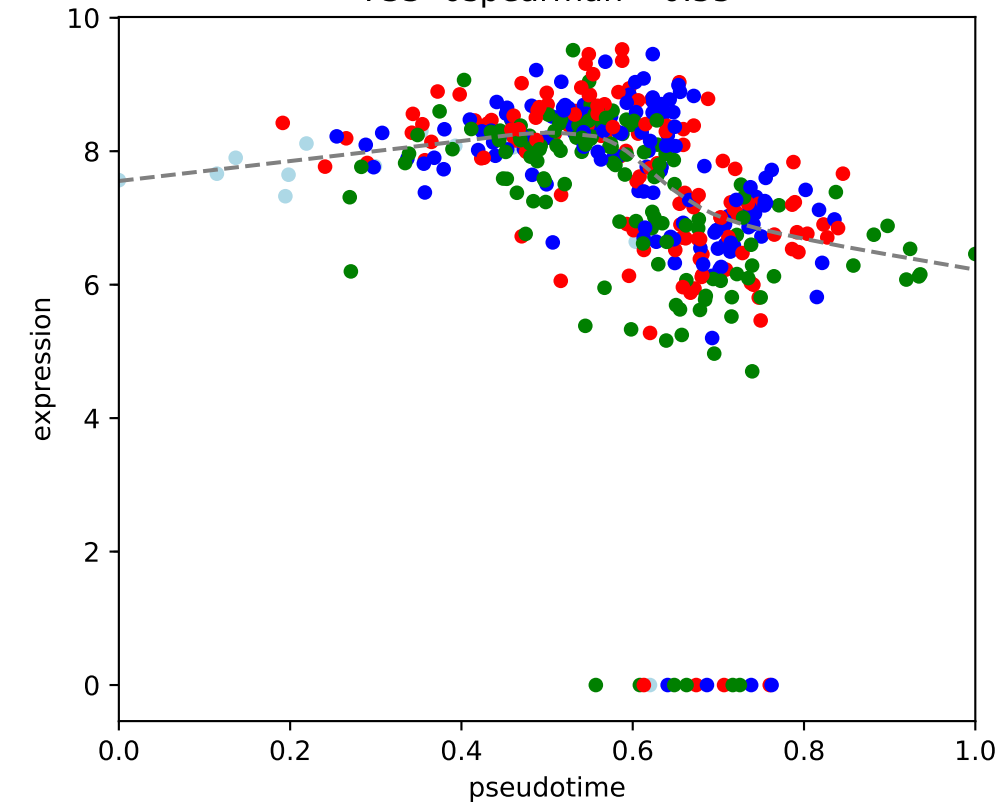
733 1spearman=-0.28



ENST00000553652 CKB (creatine kinase B)

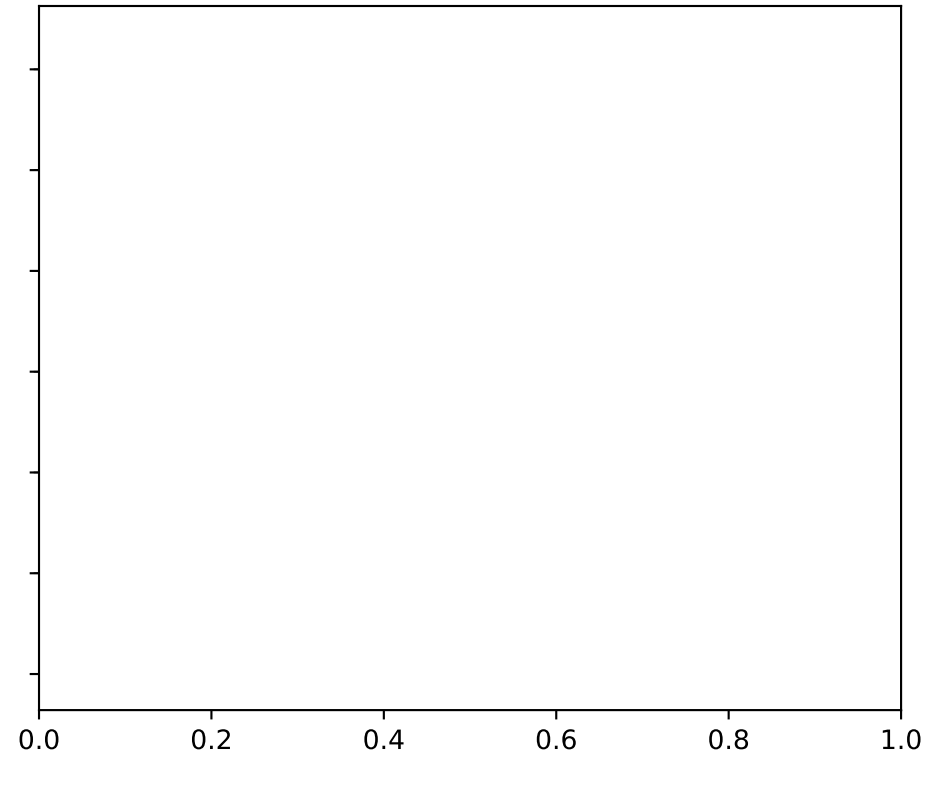
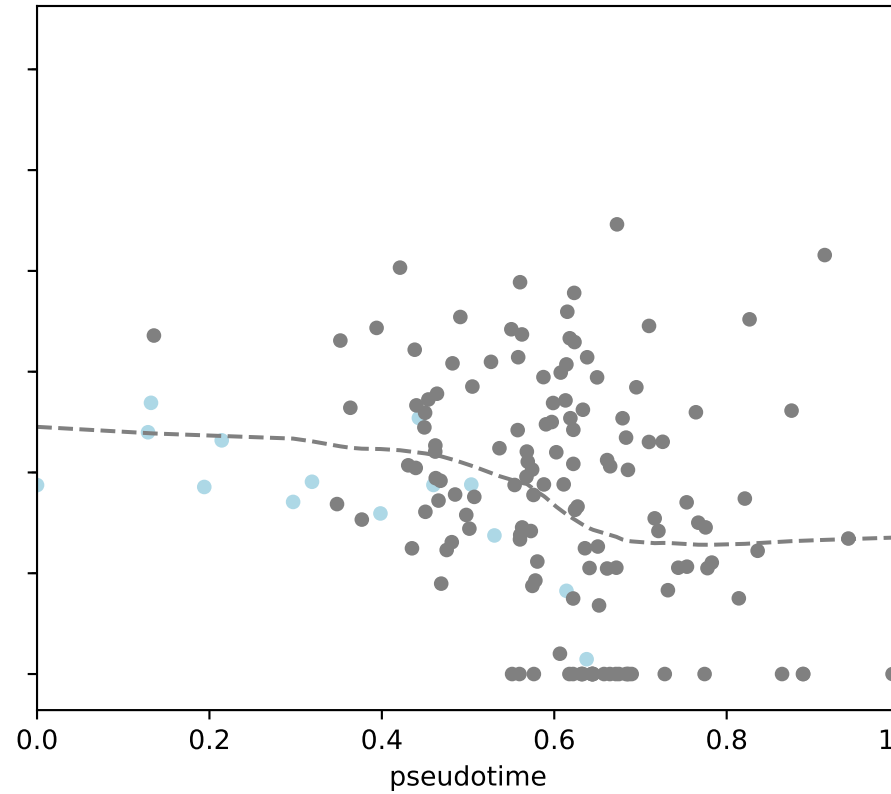
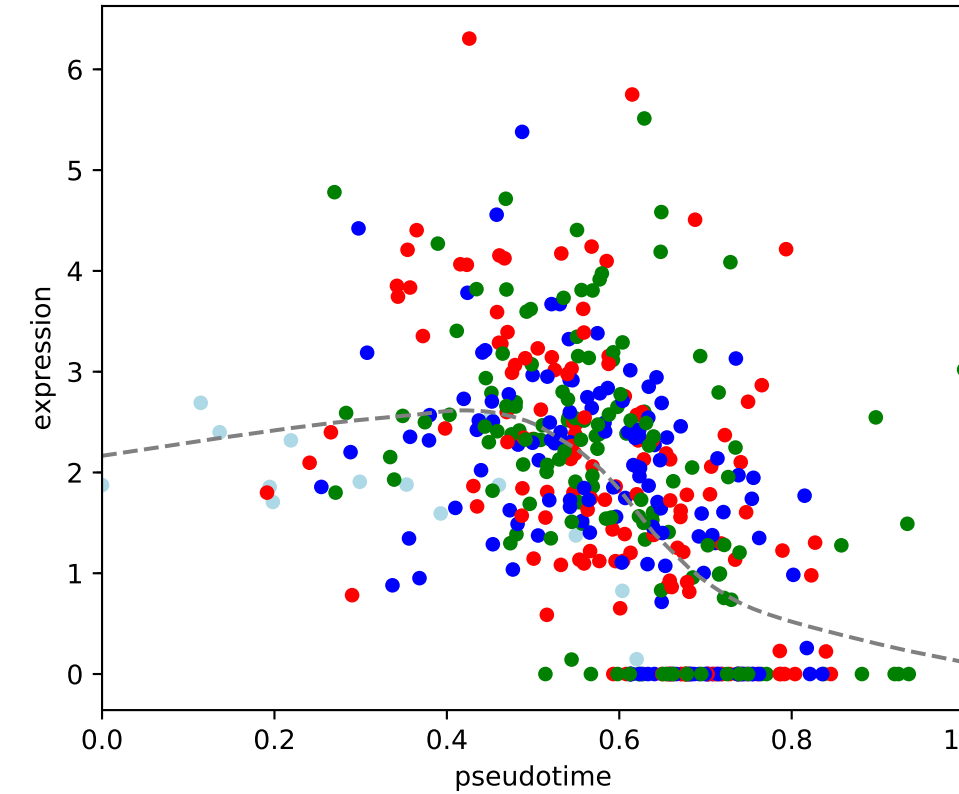
733 0spearman=-0.55

733 1spearman=-0.43



733 0spearman=-0.55

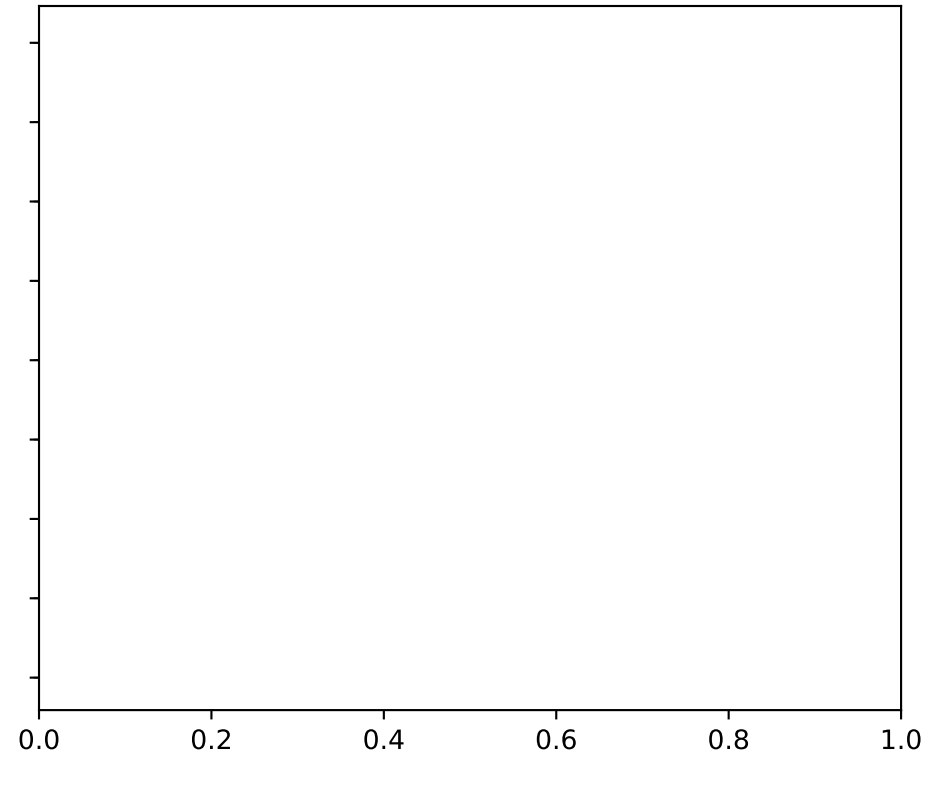
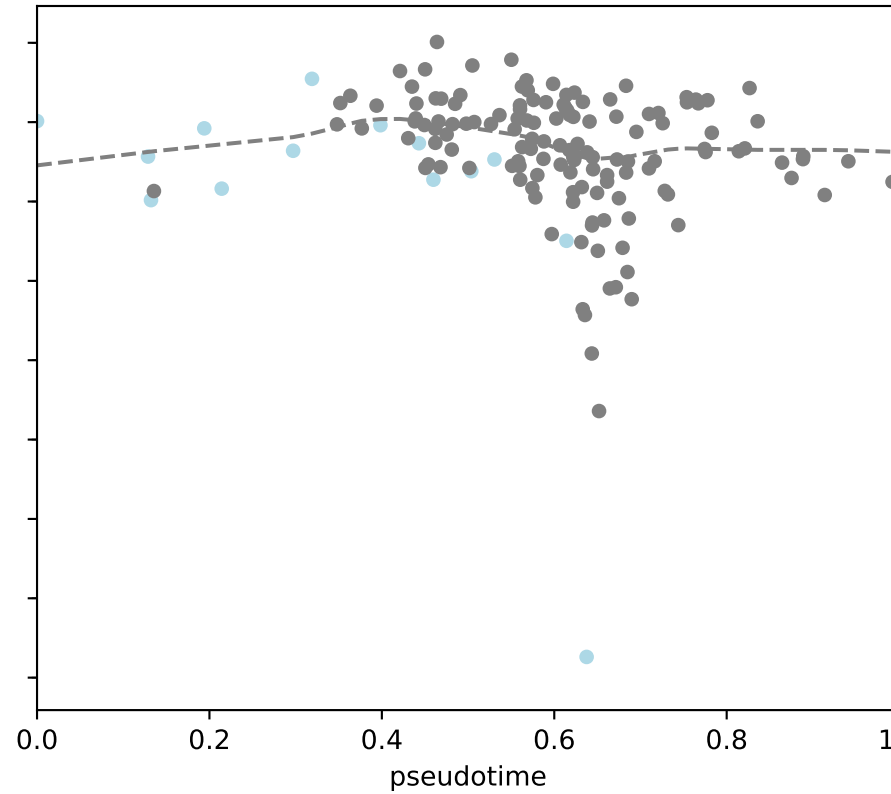
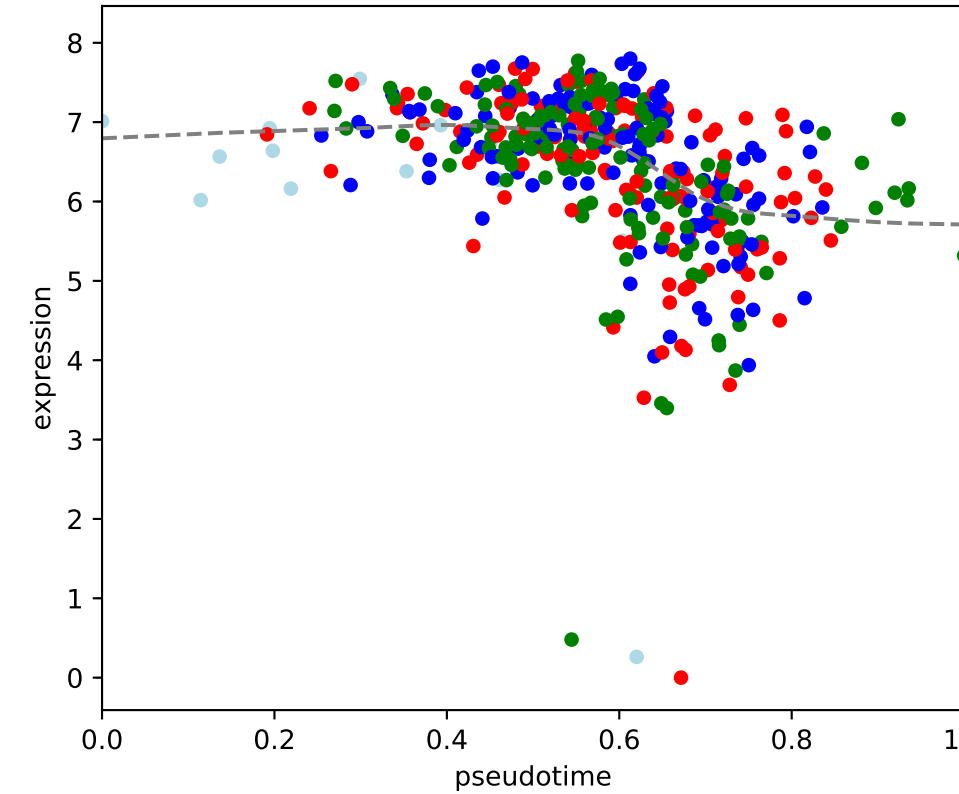
733 1spearman=-0.33



ENST00000258424 COX5B (cytochrome c oxidase subunit 5B)

733 0spearman=-0.55

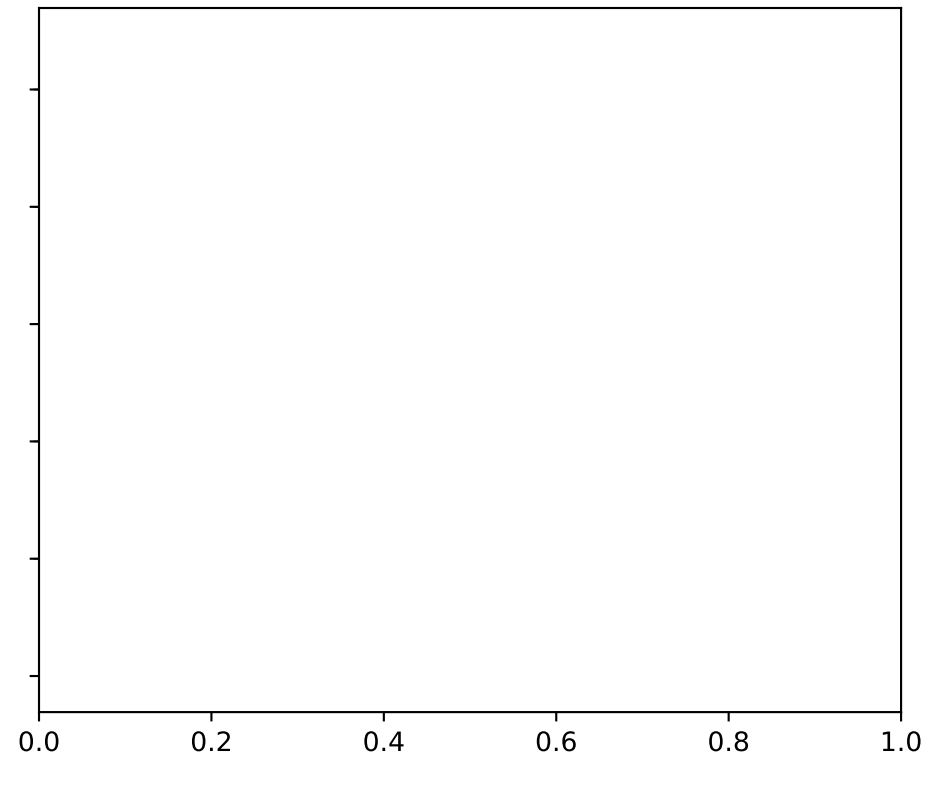
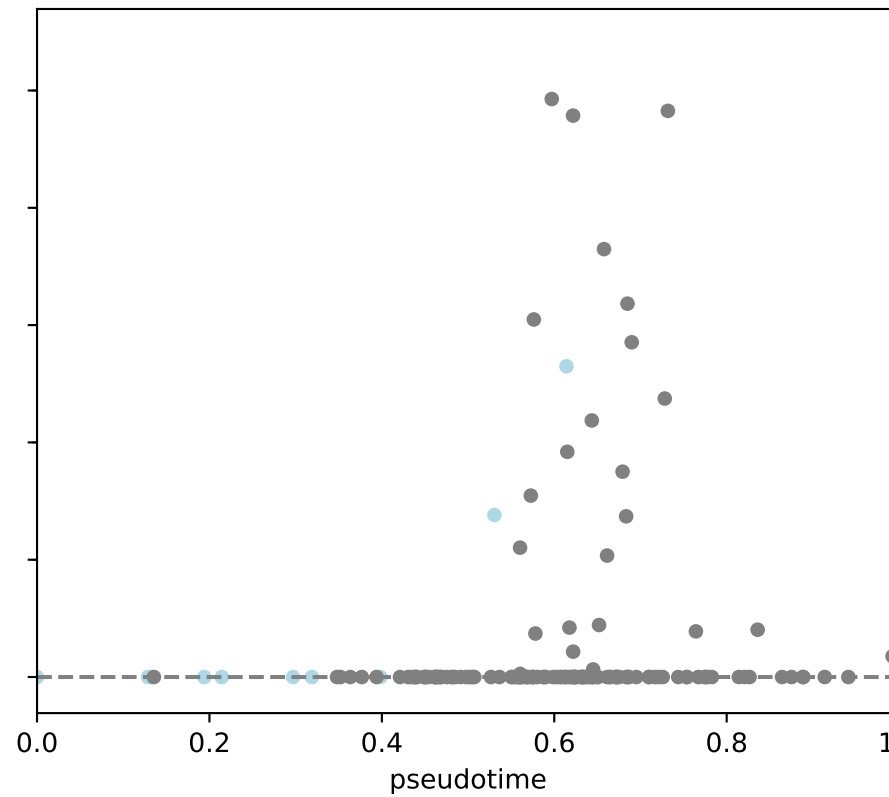
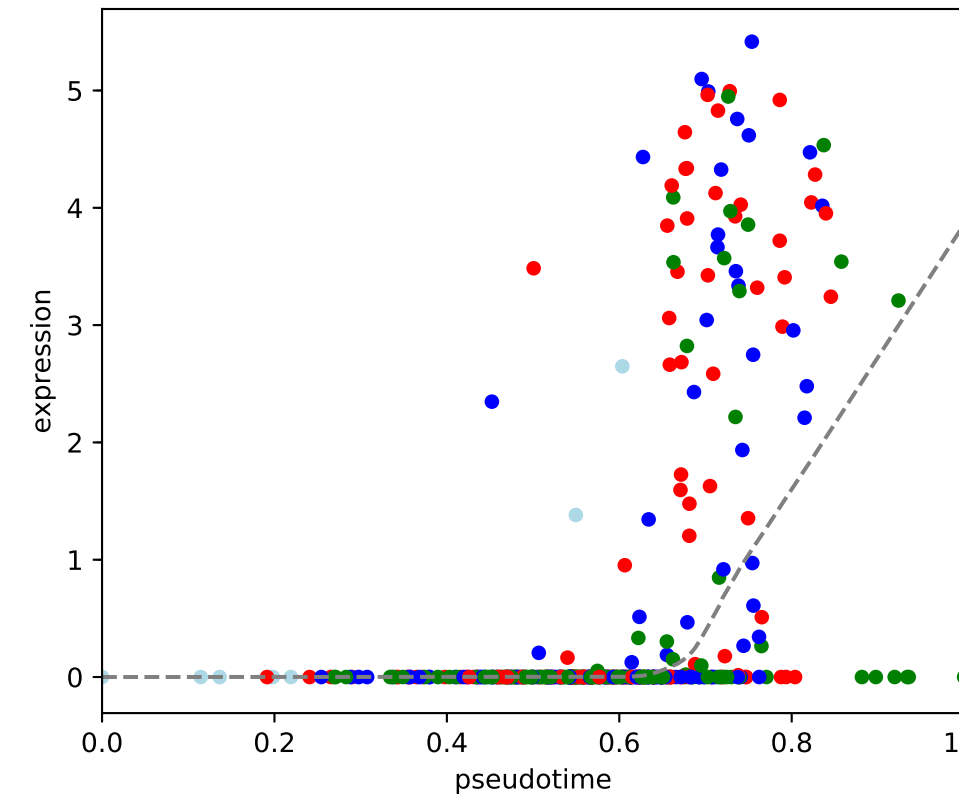
733 1spearman=-0.28



ENST00000563041 AC104938.1

733 0spearman=0.55

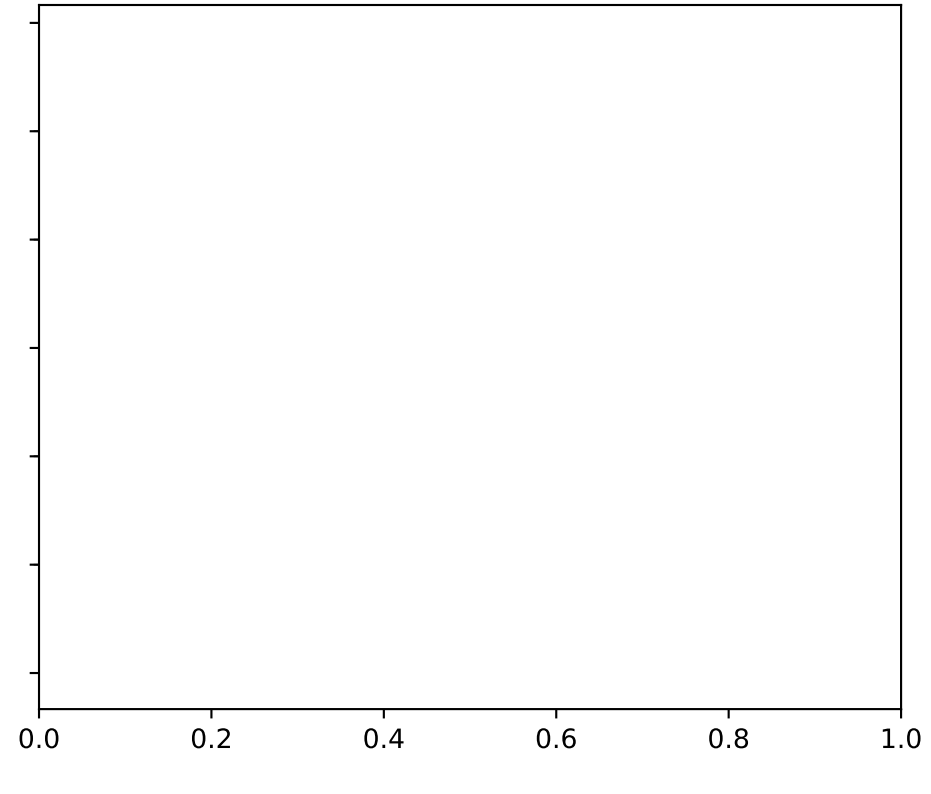
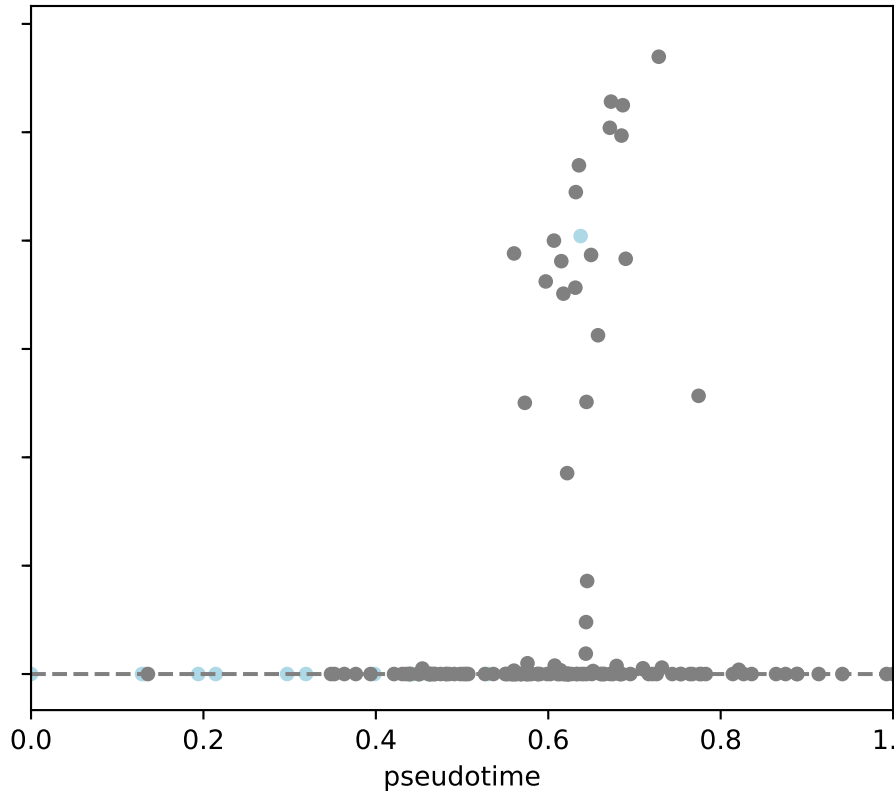
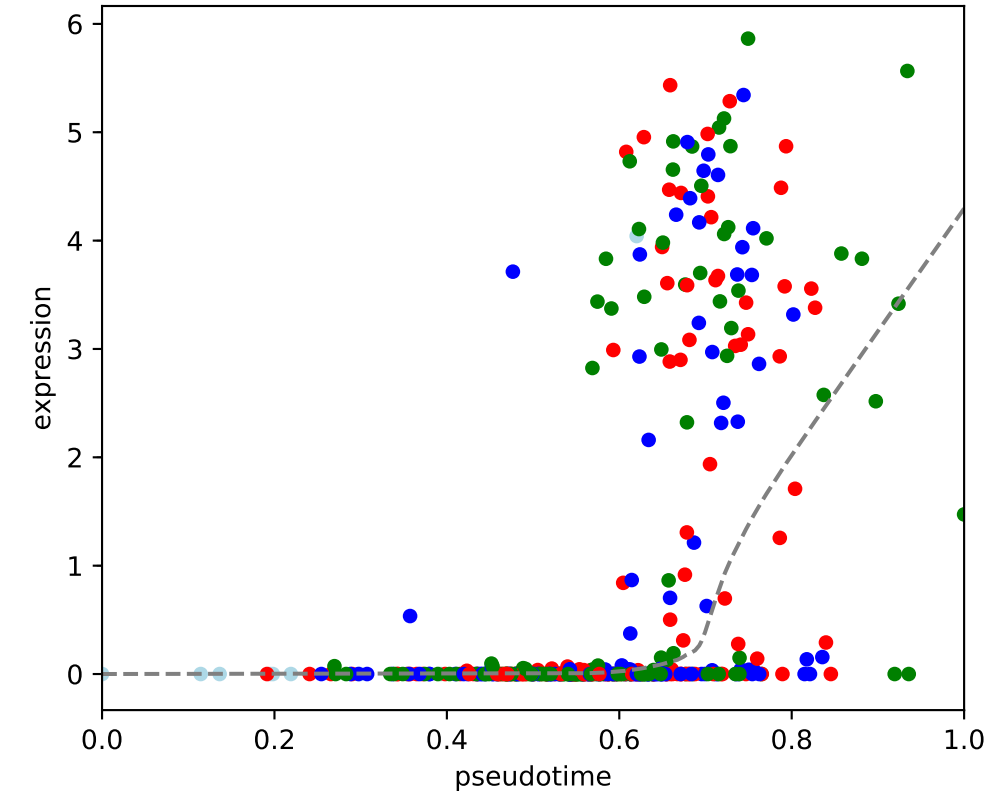
733 1spearman=0.21



ENST00000290271 STC1 (stanniocalcin 1)

733 0spearman=0.55

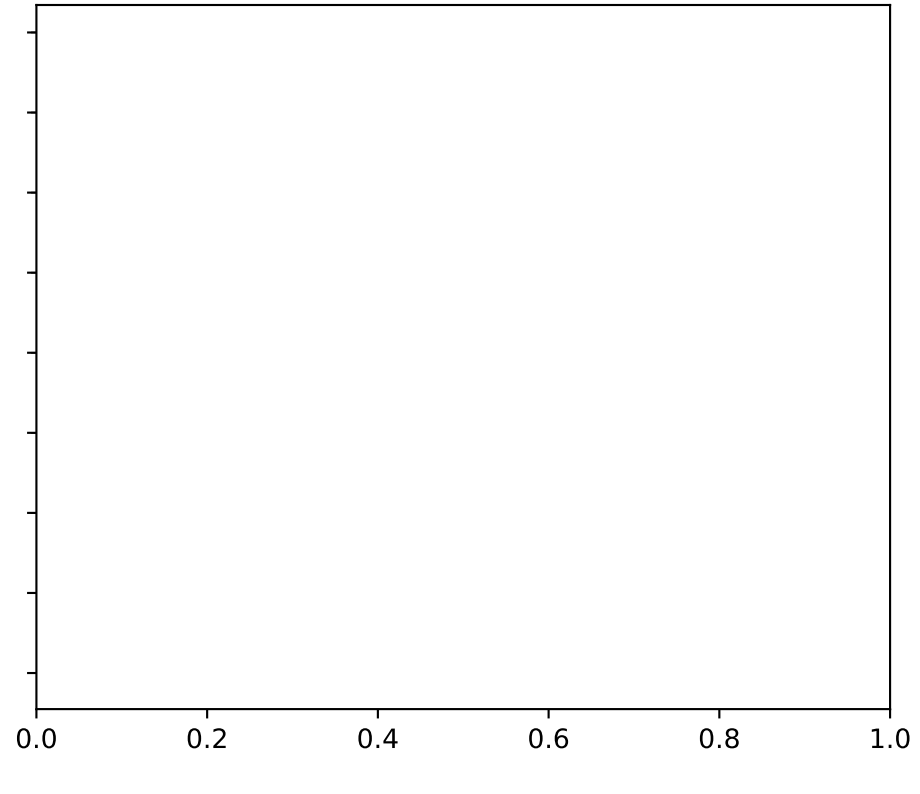
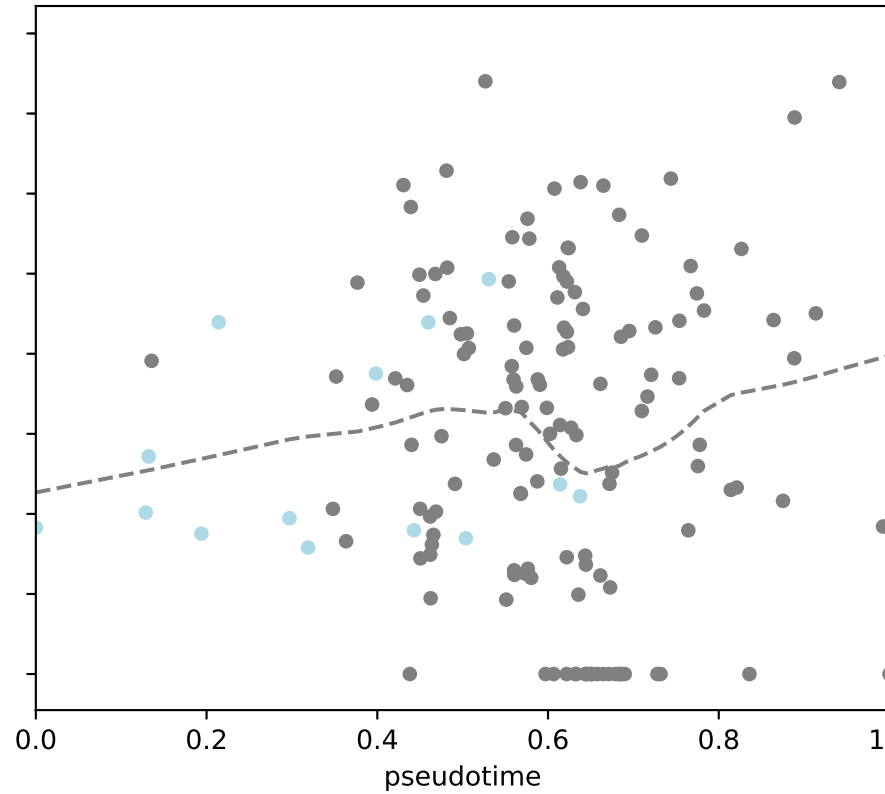
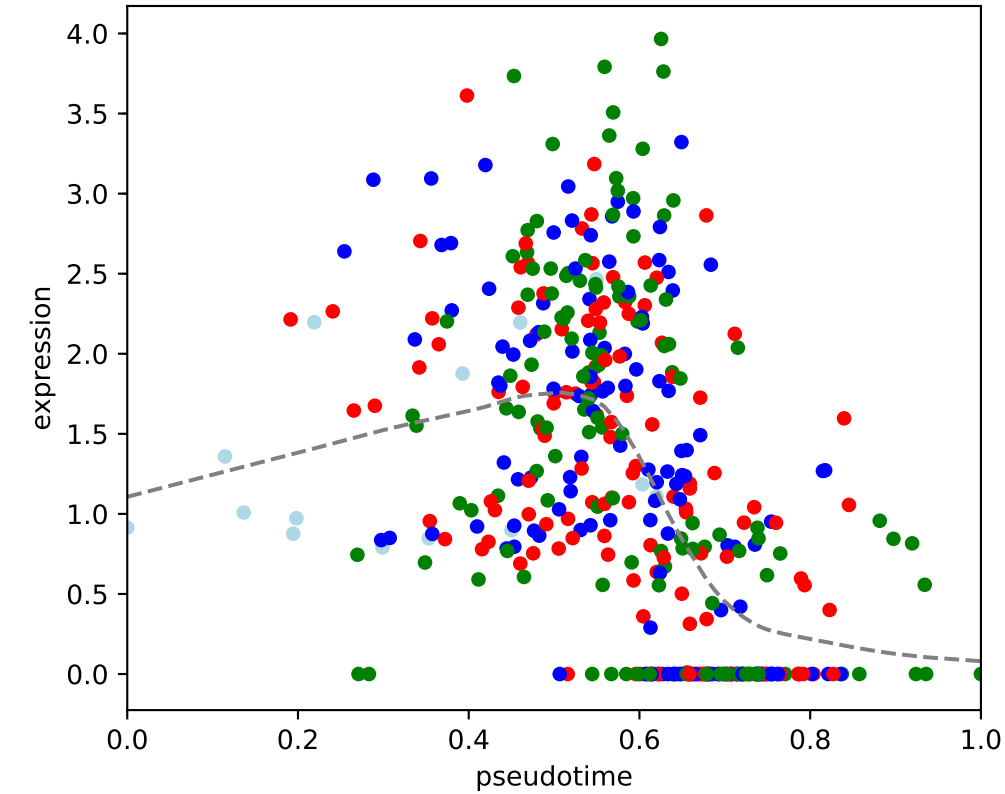
733 1spearman=0.26



ENST00000409574 HES6 (hes family bHLH transcription factor 6)

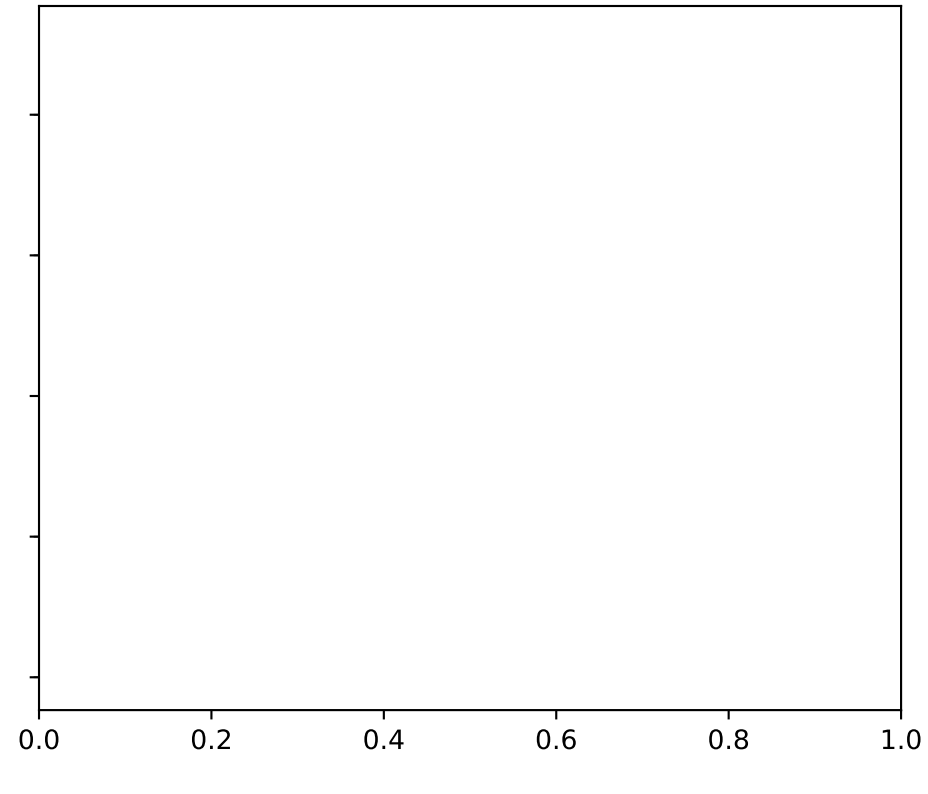
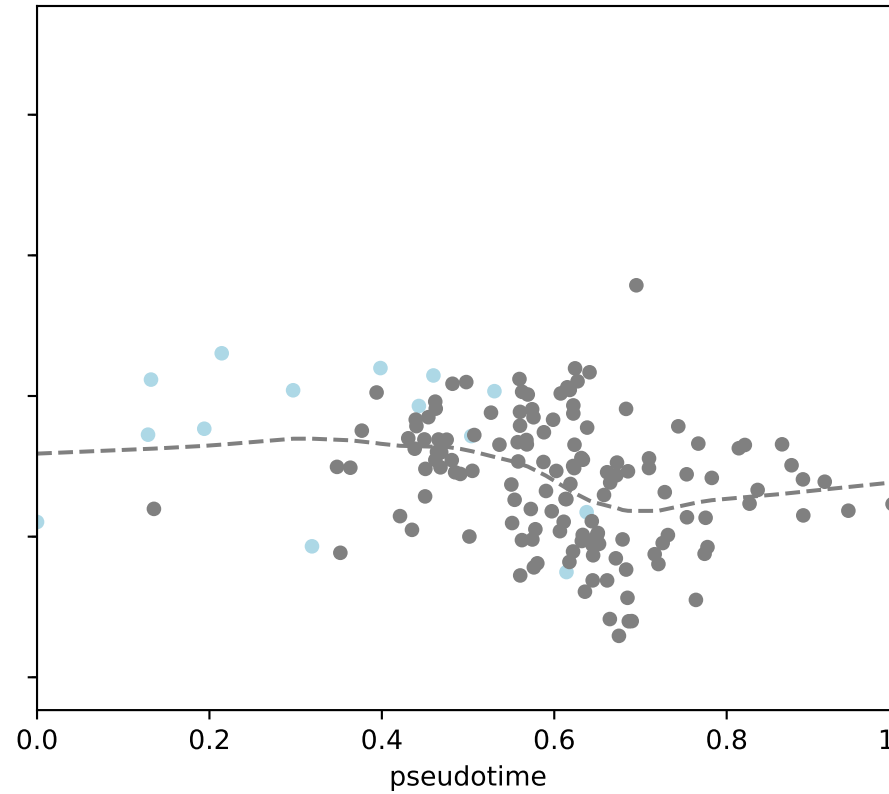
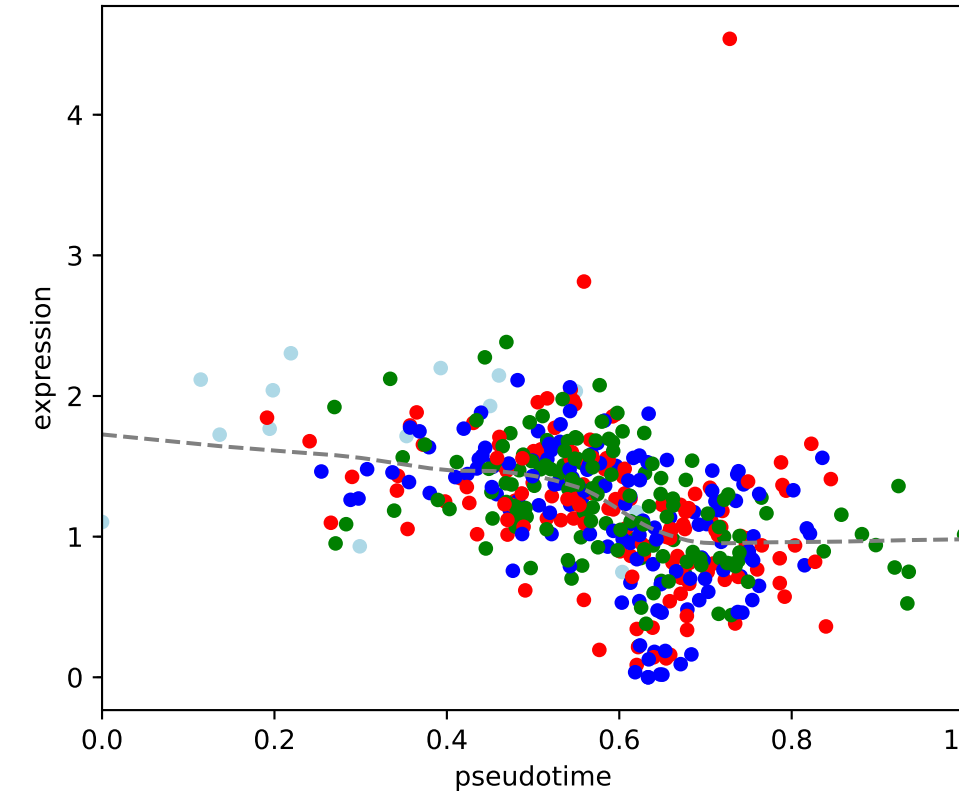
733 0spearman=-0.55

733 1spearman=-0.06



733 0spearman=-0.55

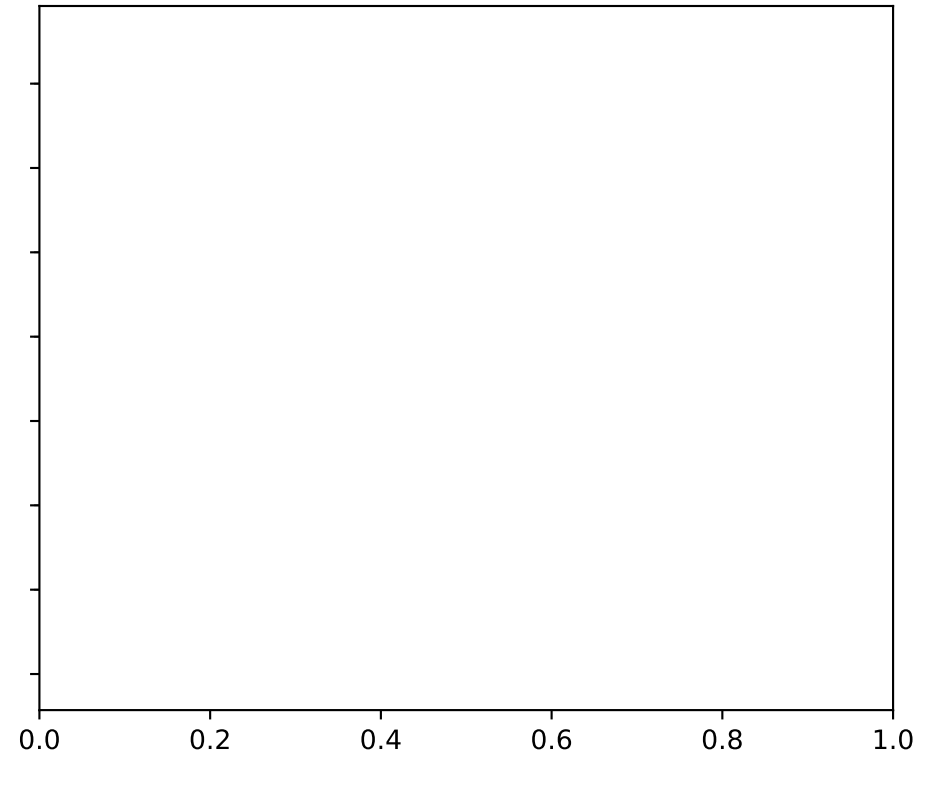
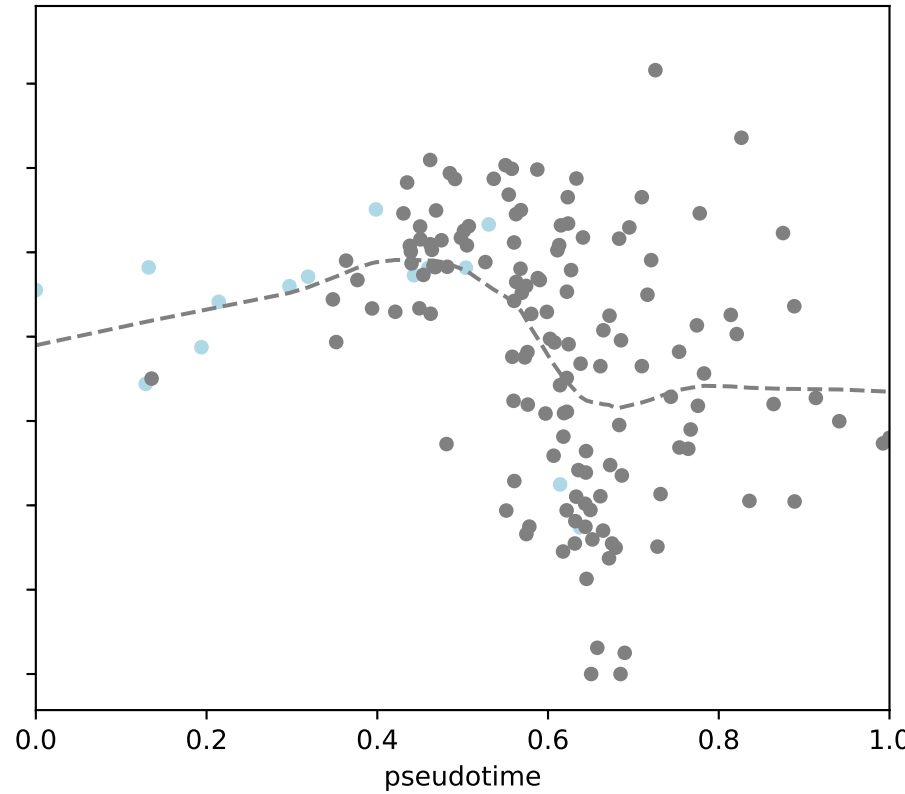
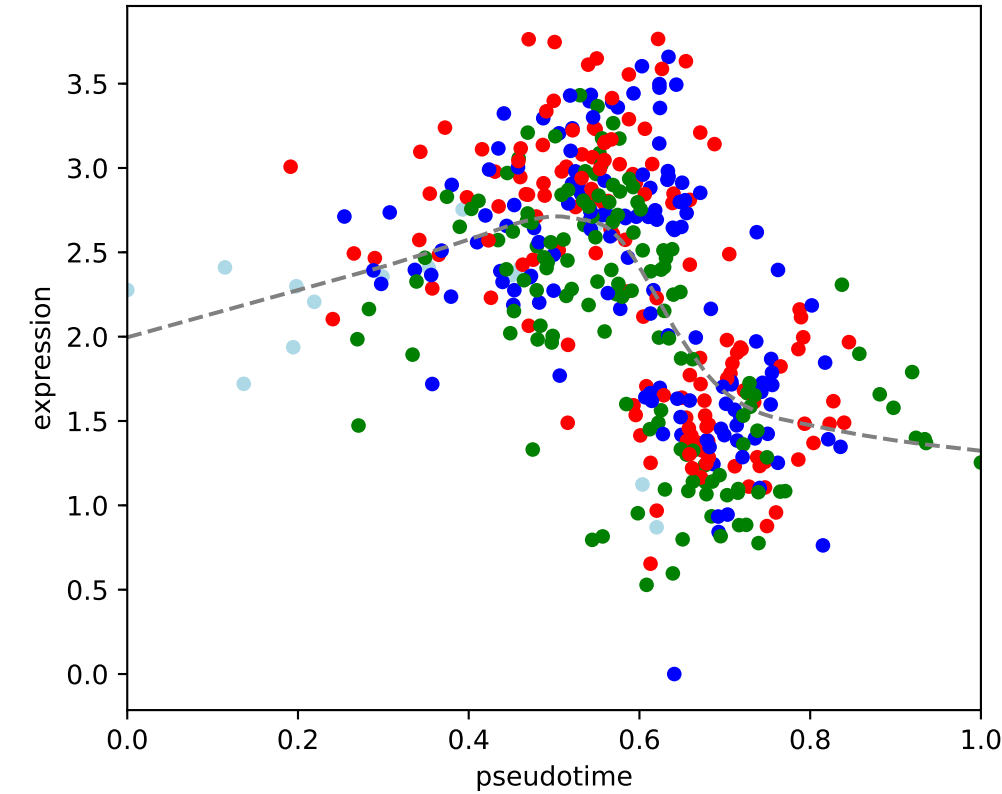
733 1spearman=-0.39



ENST00000555039 CKB (creatine kinase B)

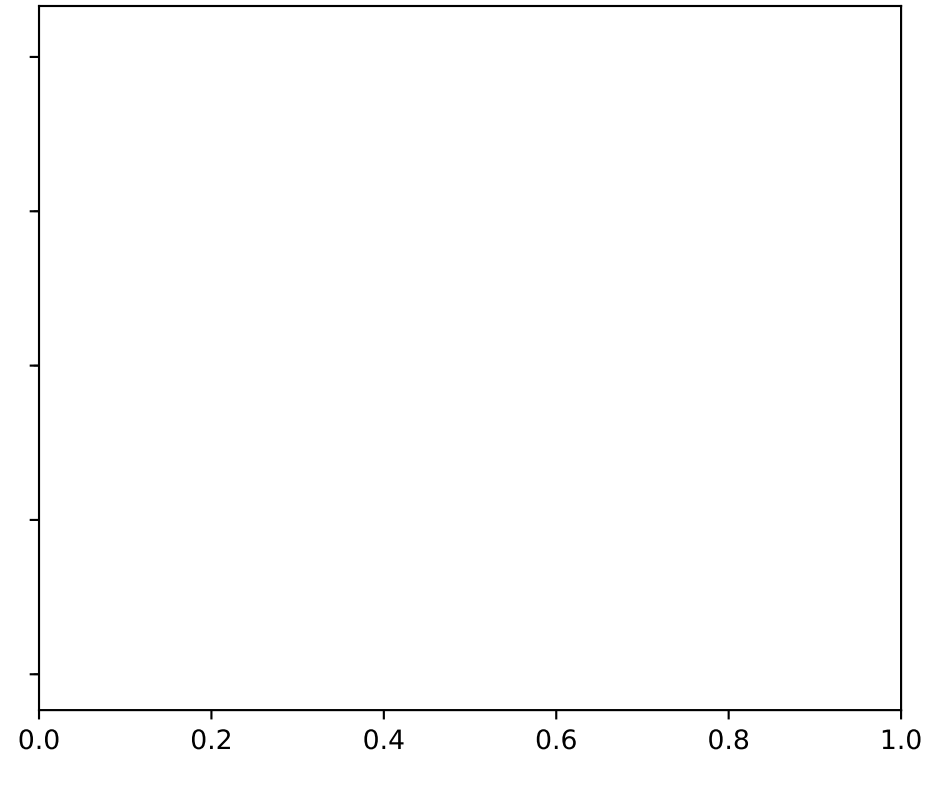
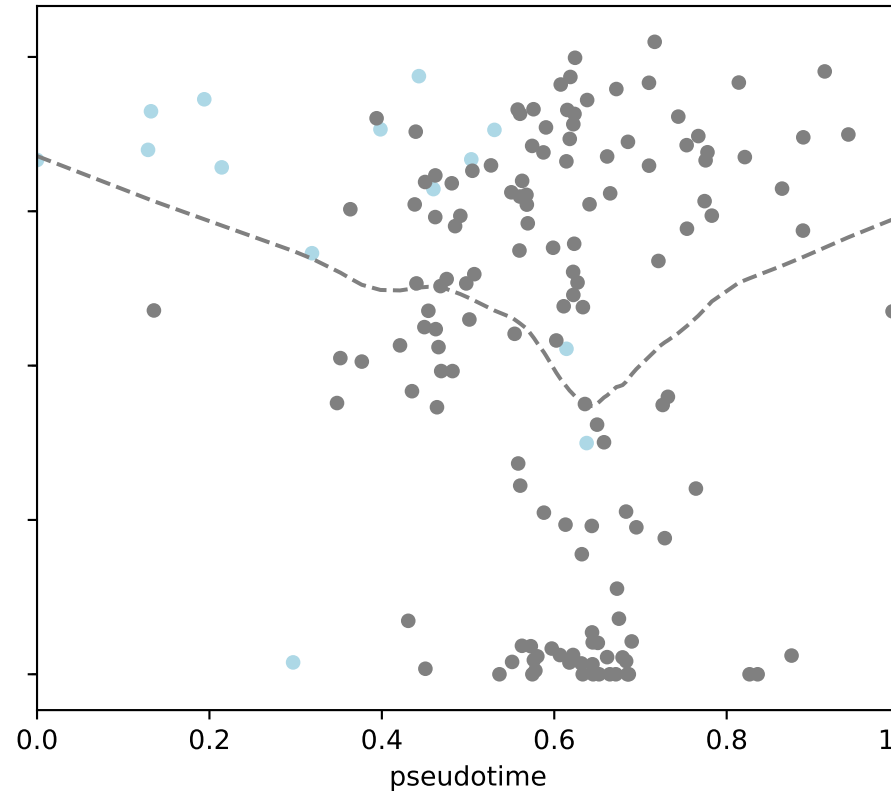
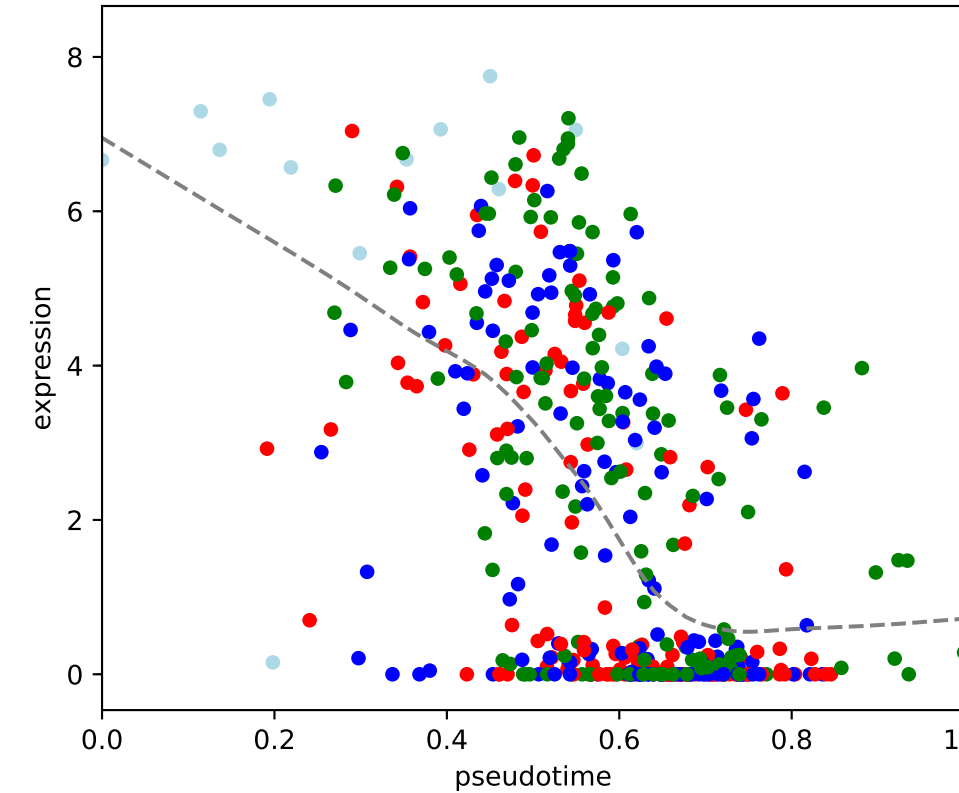
733 0spearman=-0.55

733 1spearman=-0.42



733 0spearman=-0.55

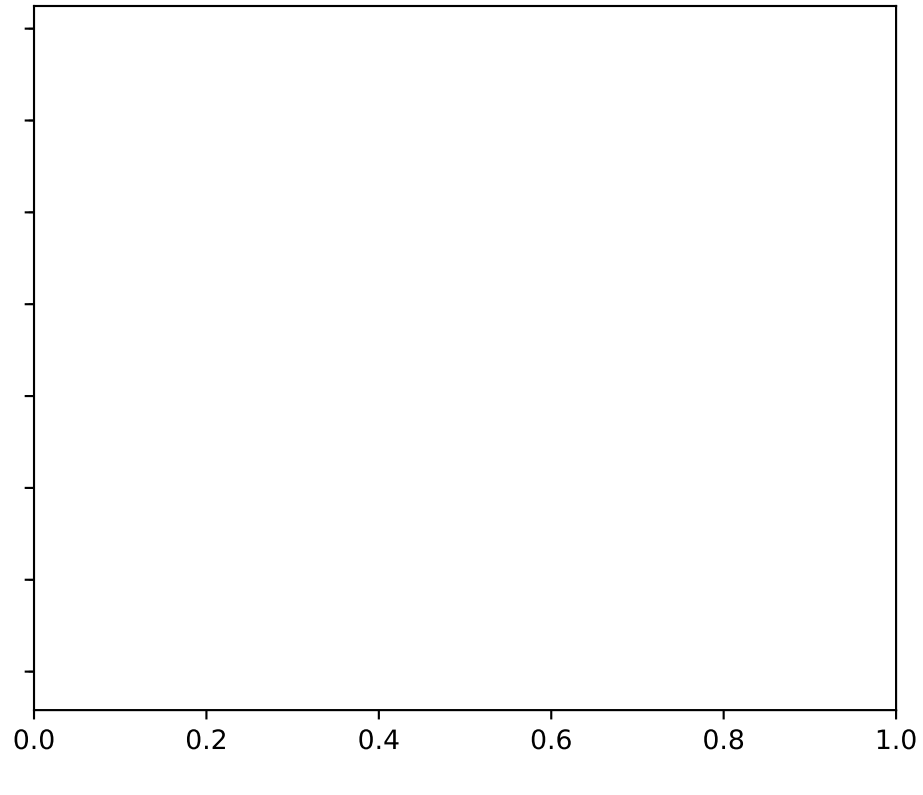
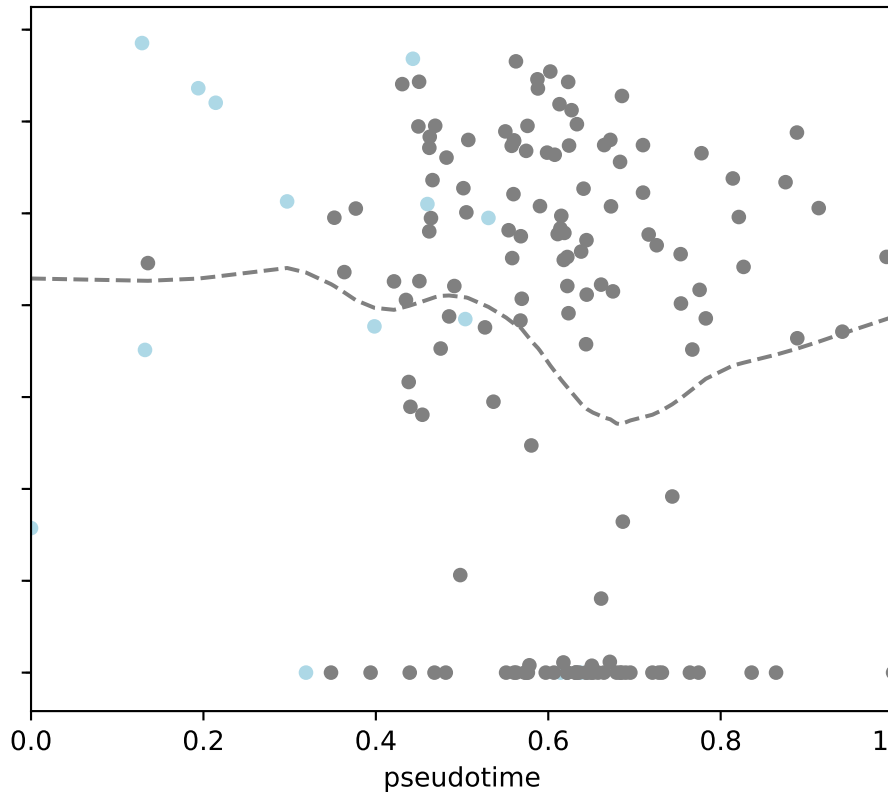
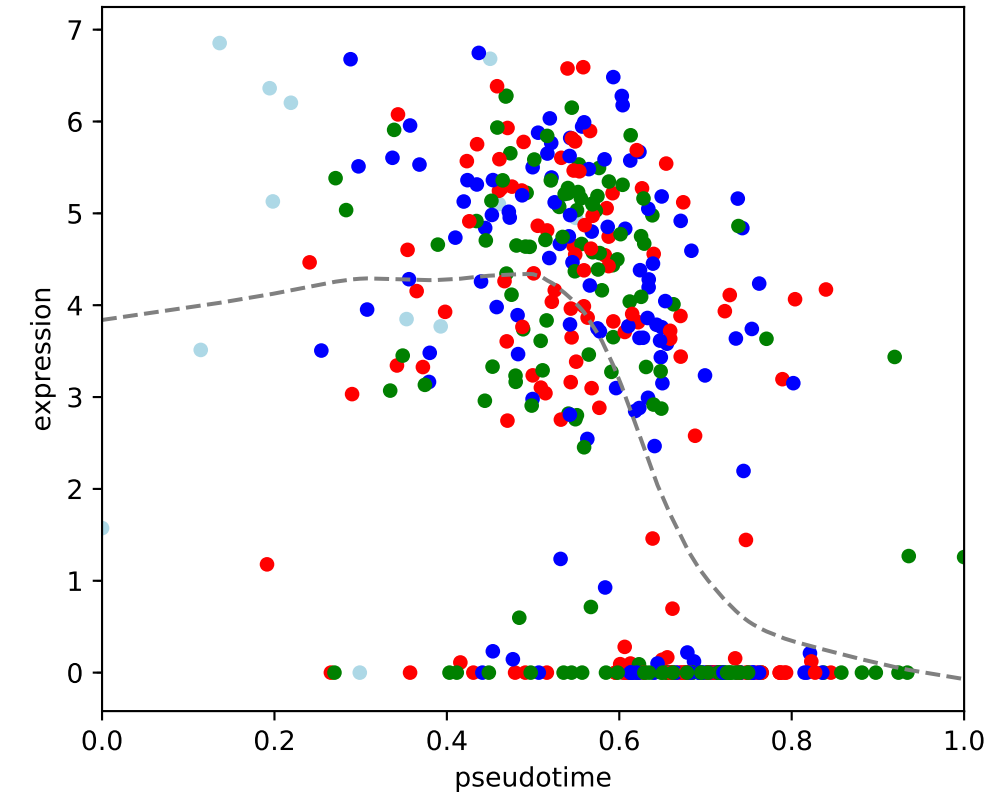
733 1spearman=-0.10



ENST00000264735 HRASLS (HRAS like suppressor)

733 0spearman=-0.55

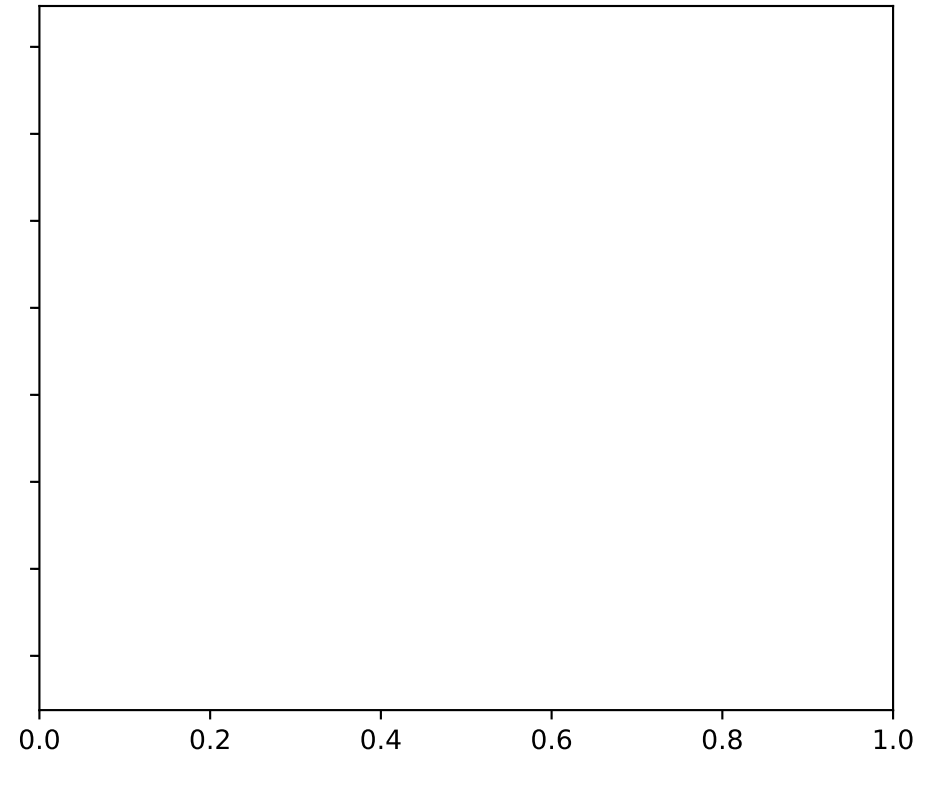
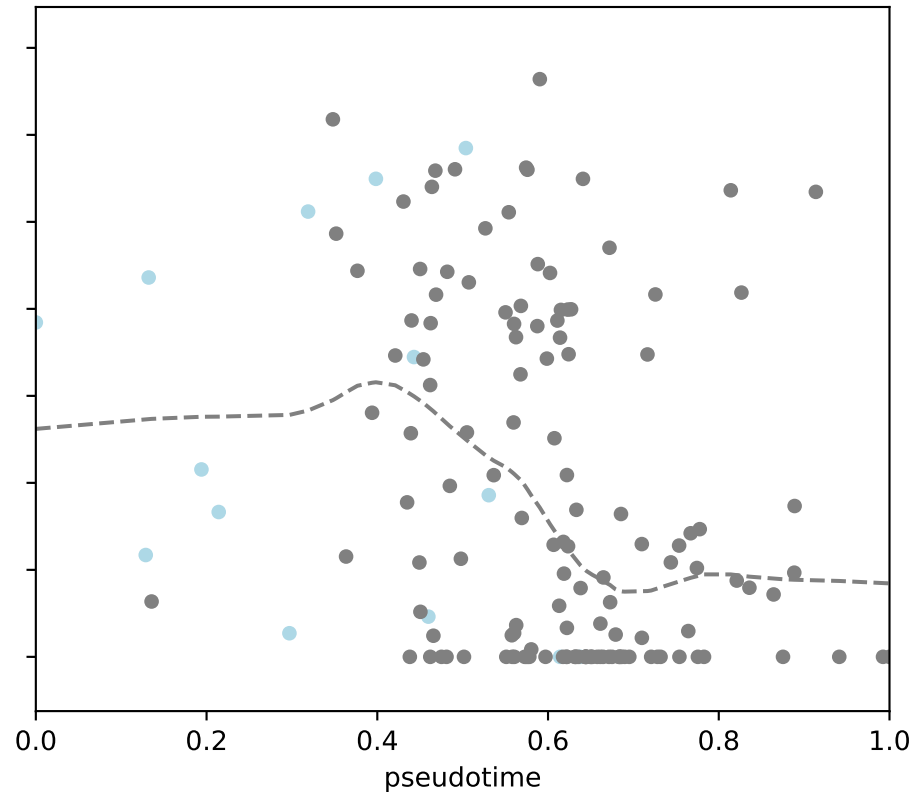
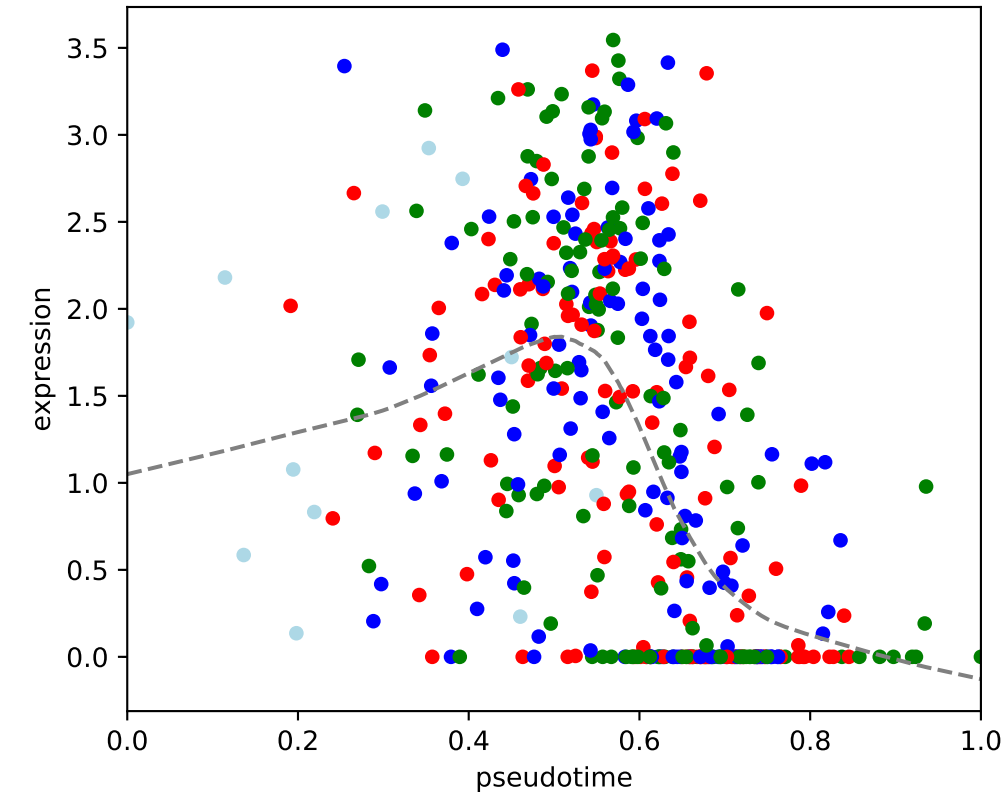
733 1spearman=-0.20



ENST00000304062 CPLX1 (complexin 1)

733 0spearman=-0.54

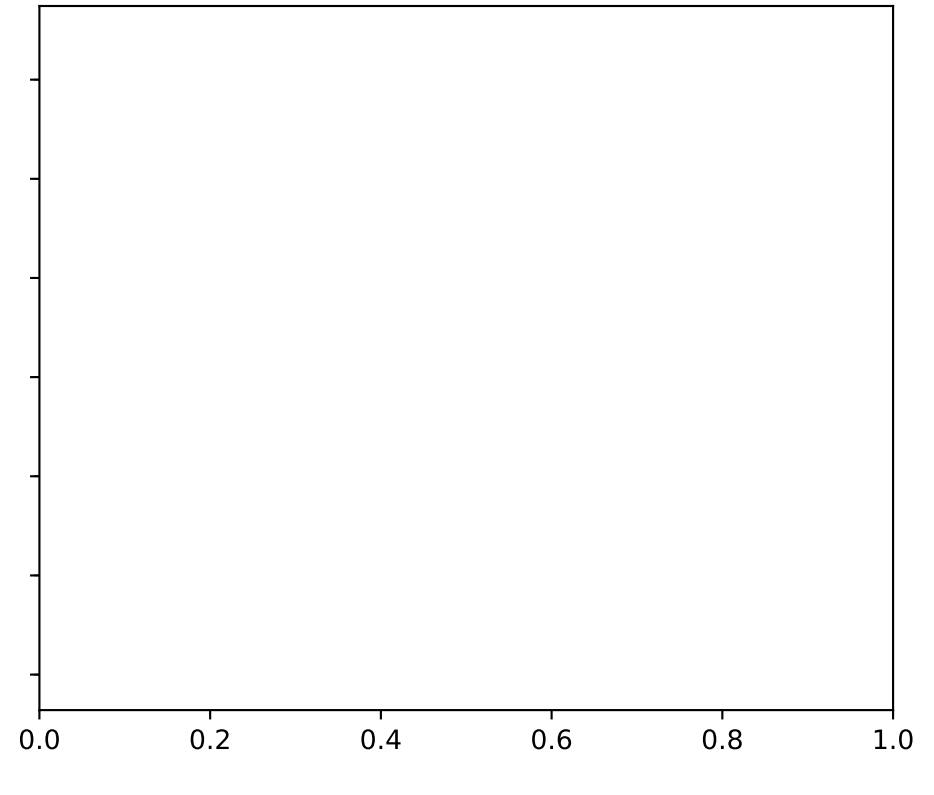
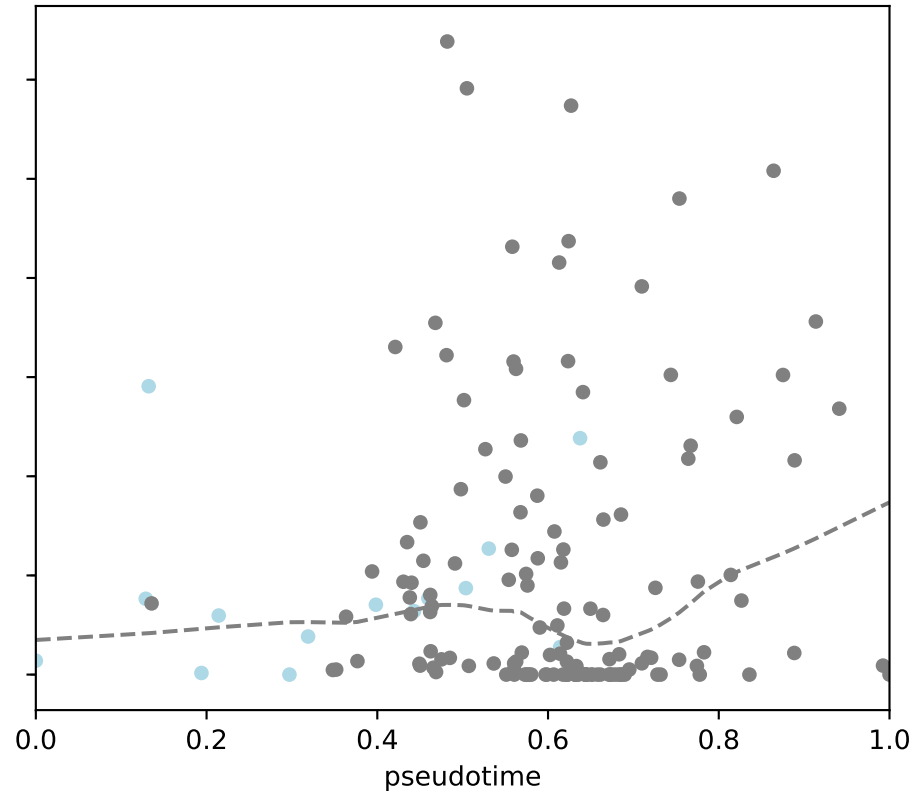
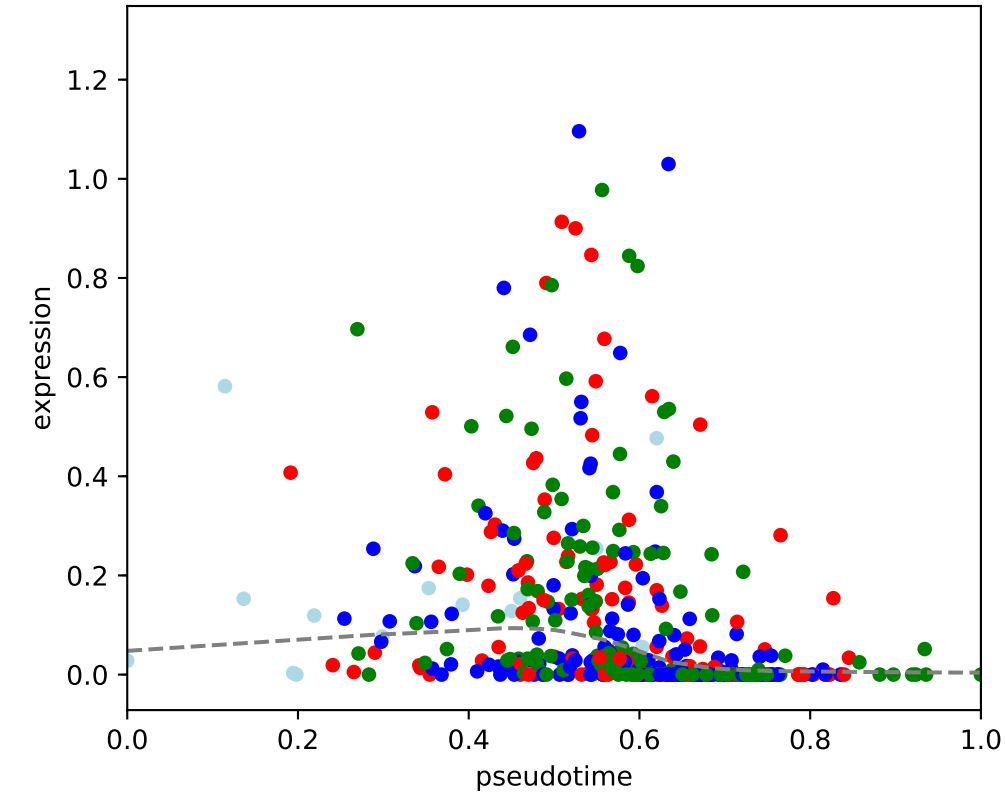
733 1spearman=-0.39



ENST00000512888 SPON2 (spodin 2)

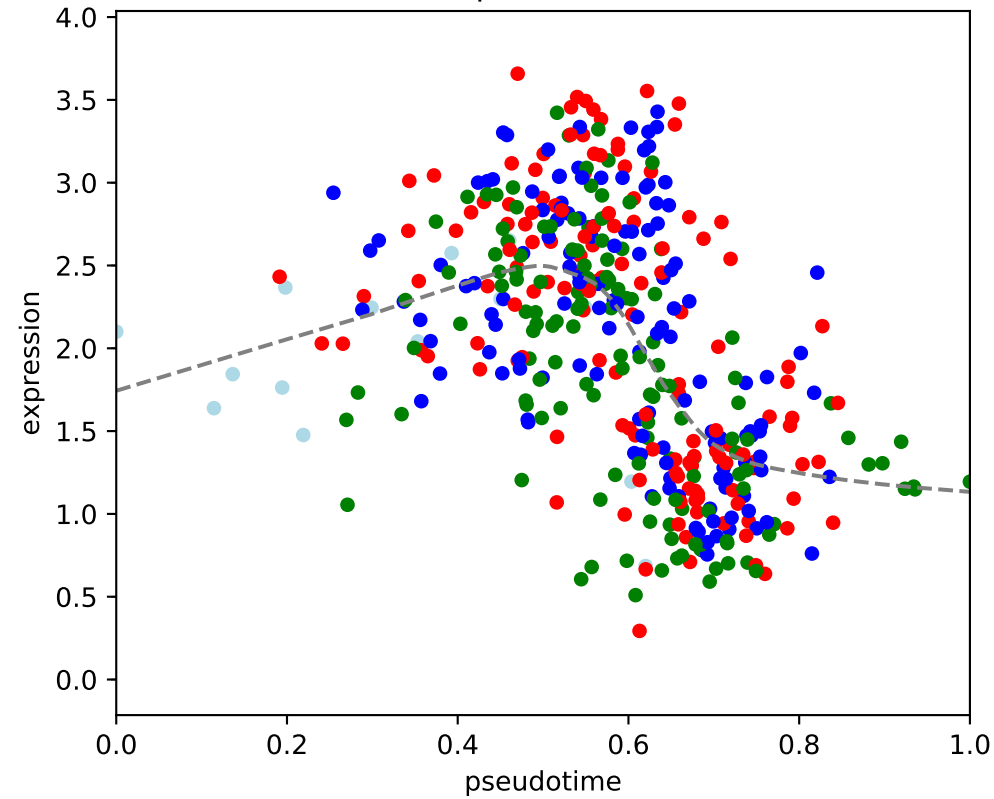
733 0spearman=-0.54

733 1spearman=-0.13

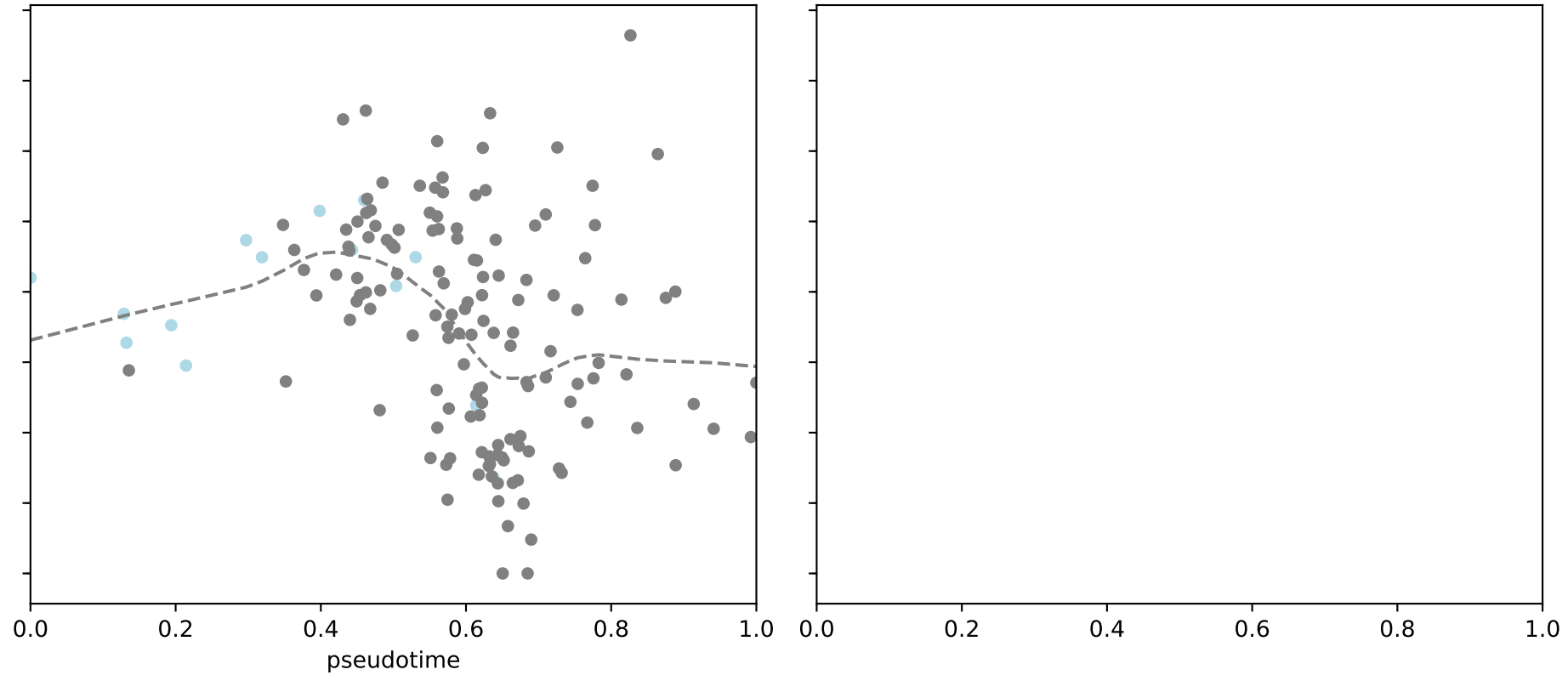


ENST00000553610 CKB (creatine kinase B)

733 0spearman=-0.54



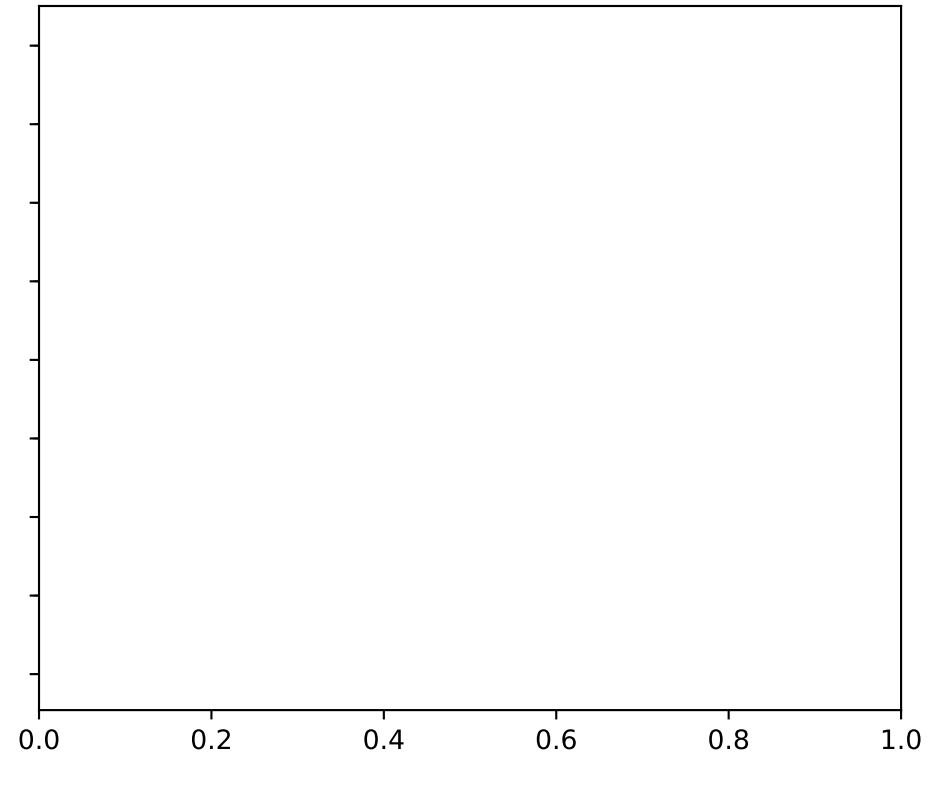
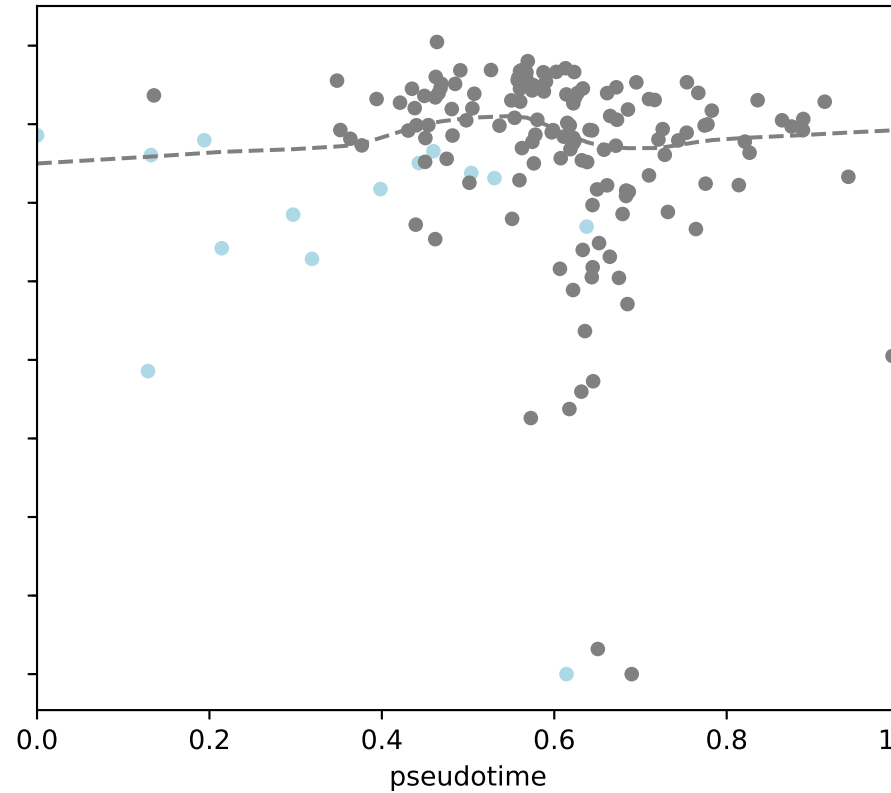
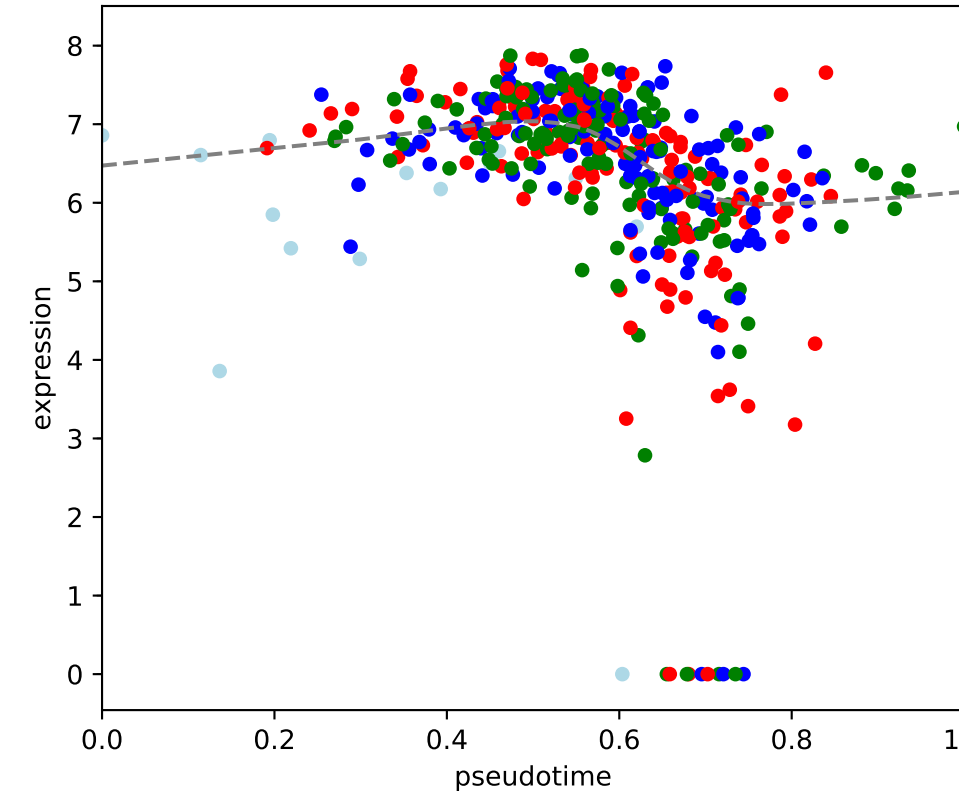
733 1spearman=-0.38



ENST00000309579 ATP5MD (ATP synthase membrane subunit DAPIT)

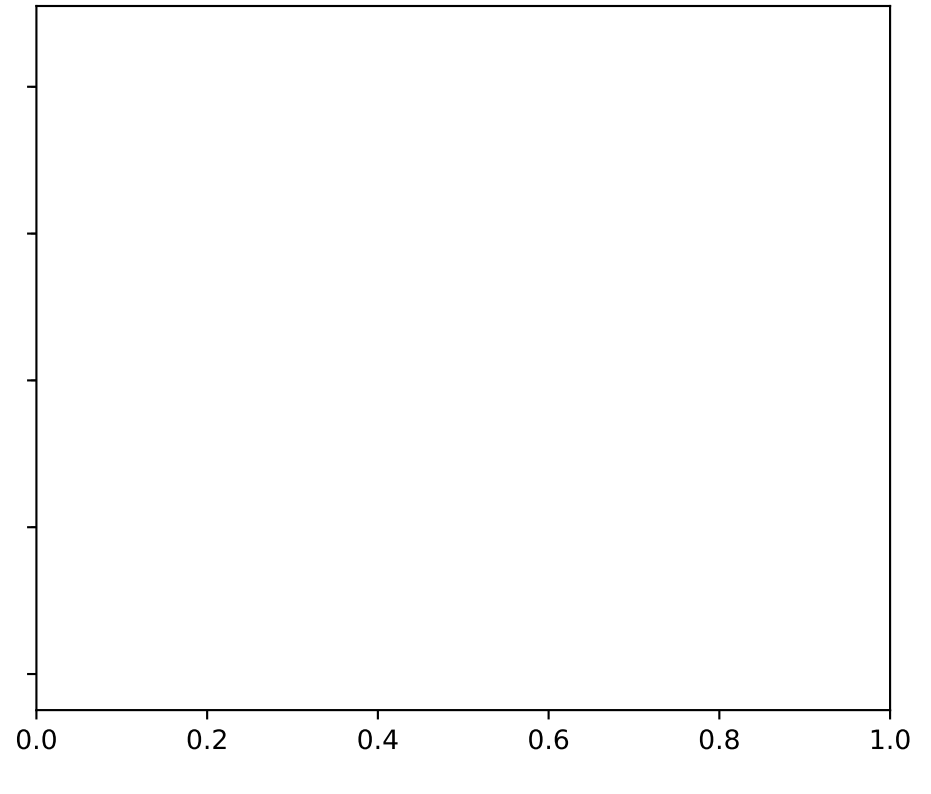
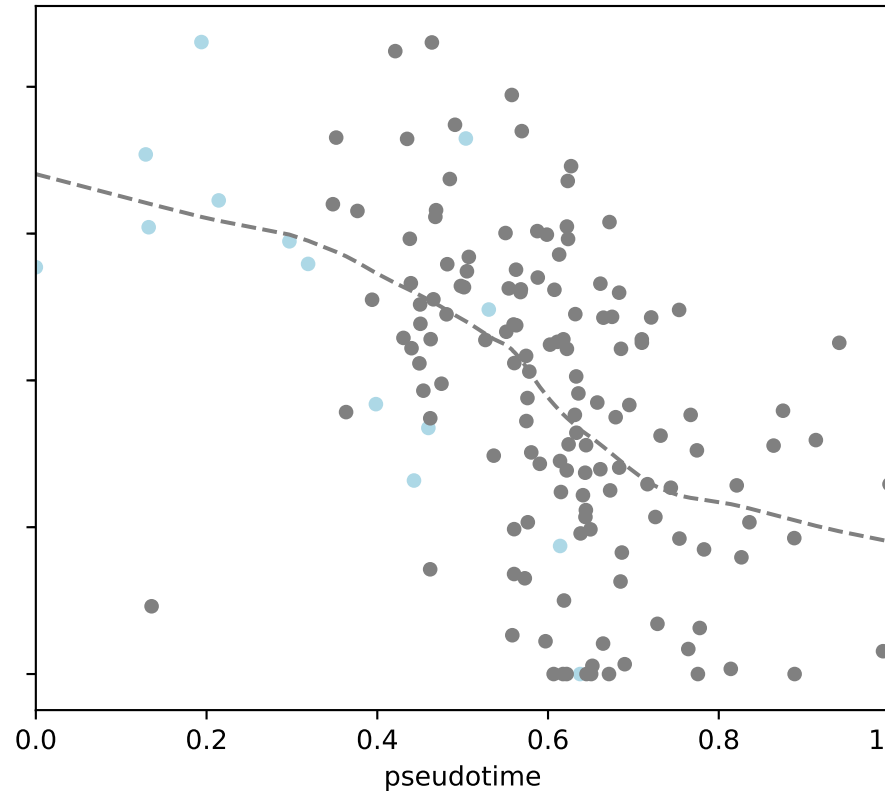
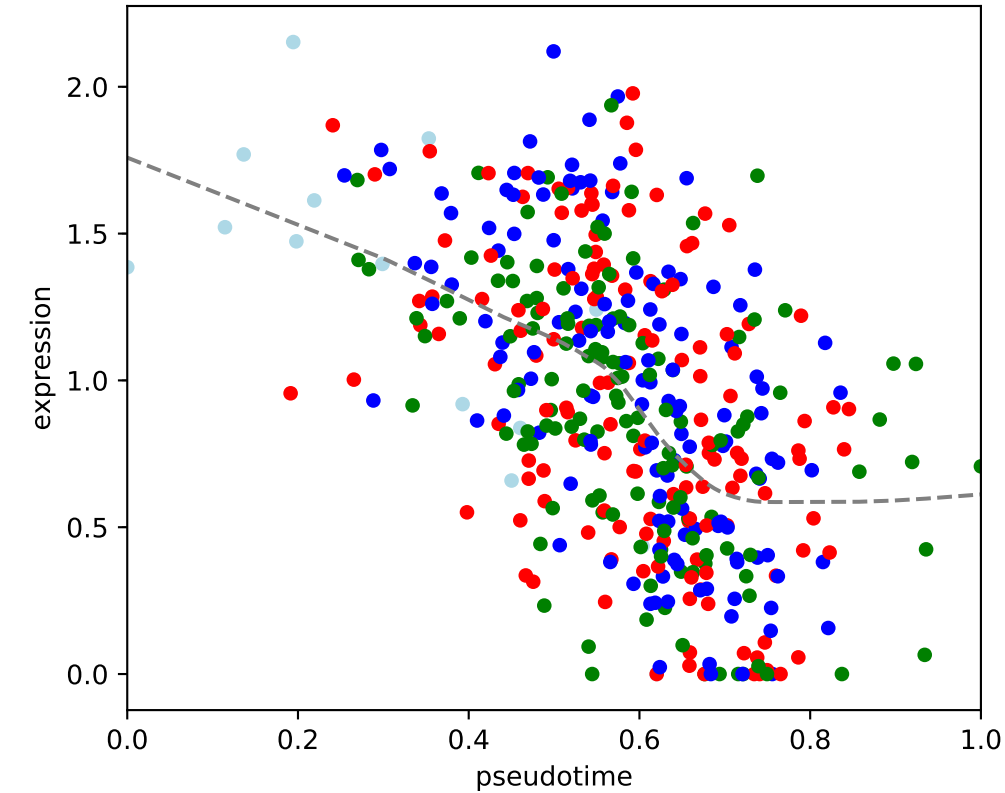
733 0spearman=-0.54

733 1spearman=-0.18



733 0spearman=-0.54

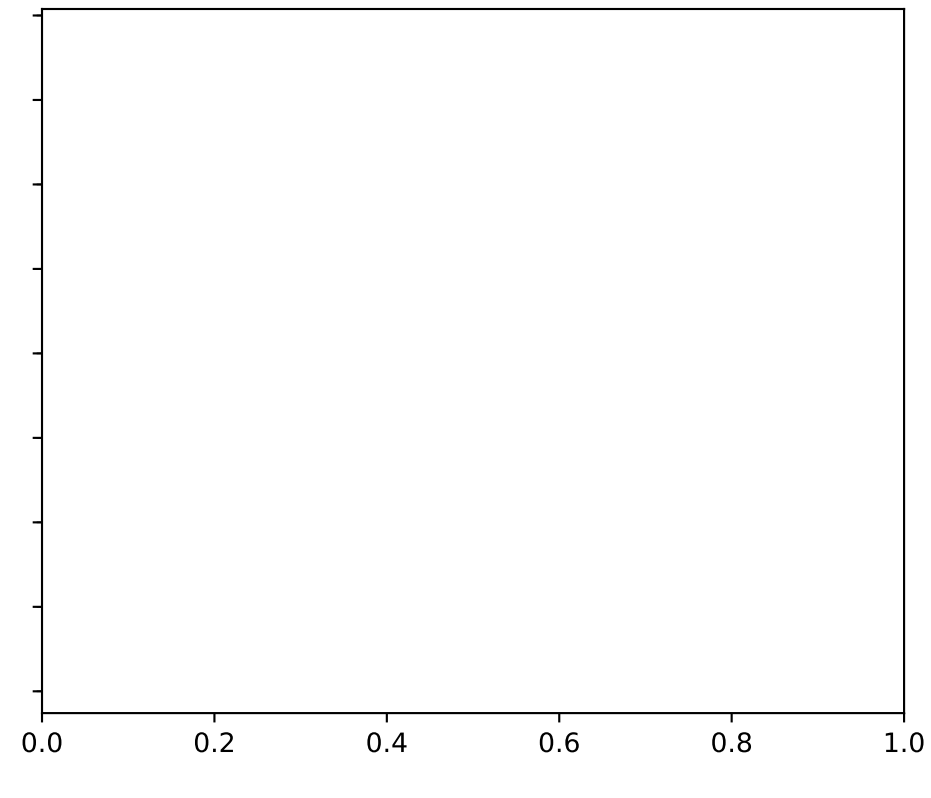
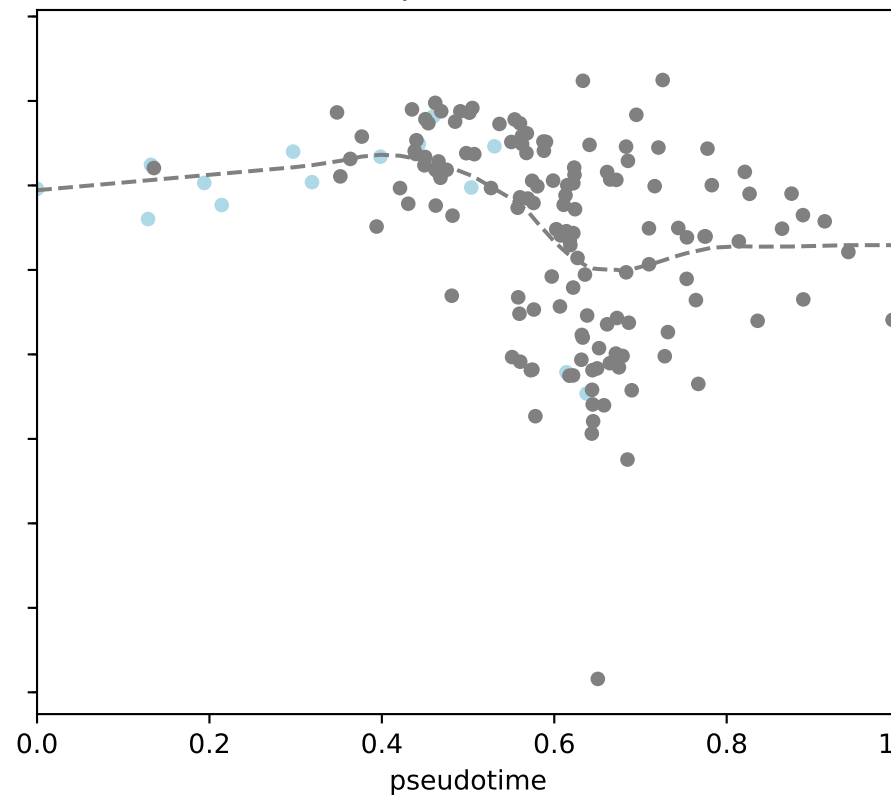
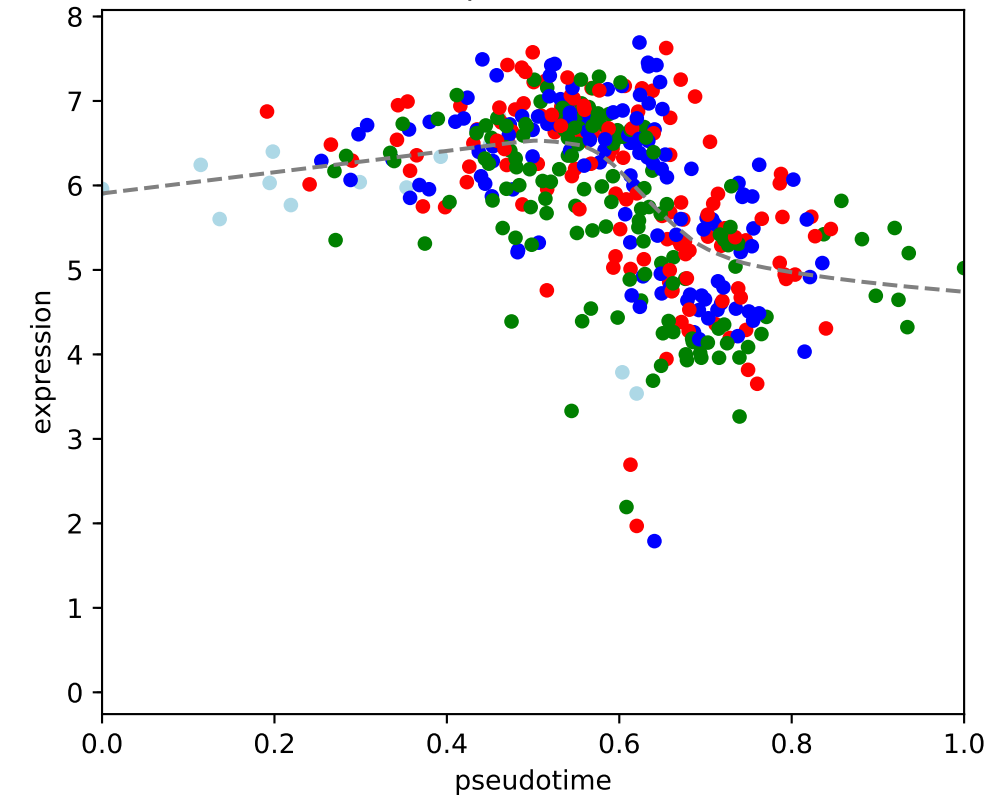
733 1spearman=-0.55



ENST00000348956 CKB (creatine kinase B)

733 0spearman=-0.54

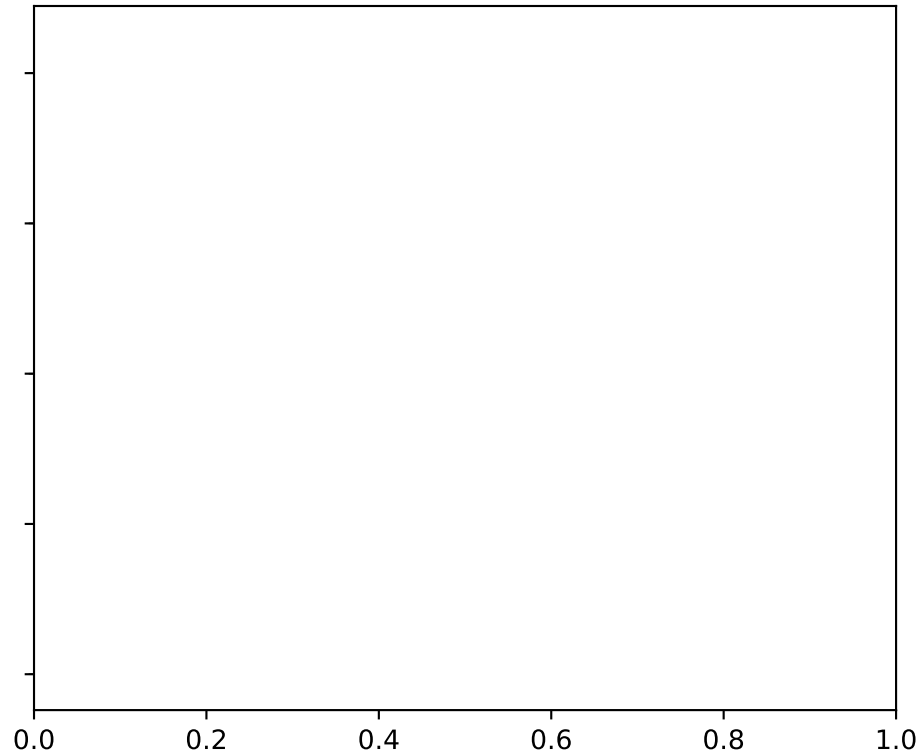
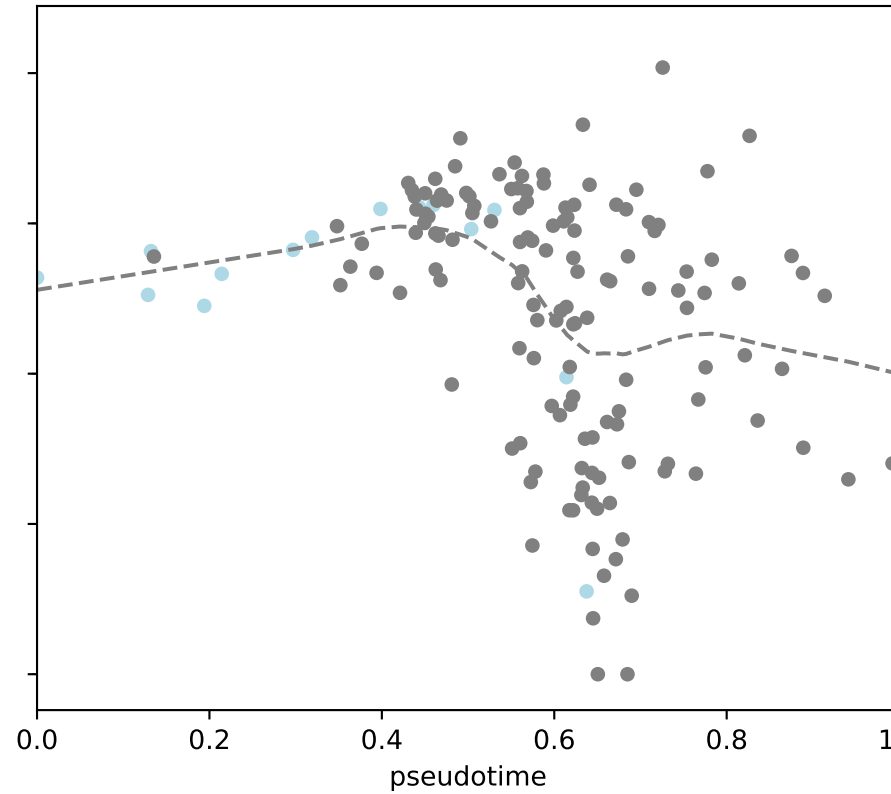
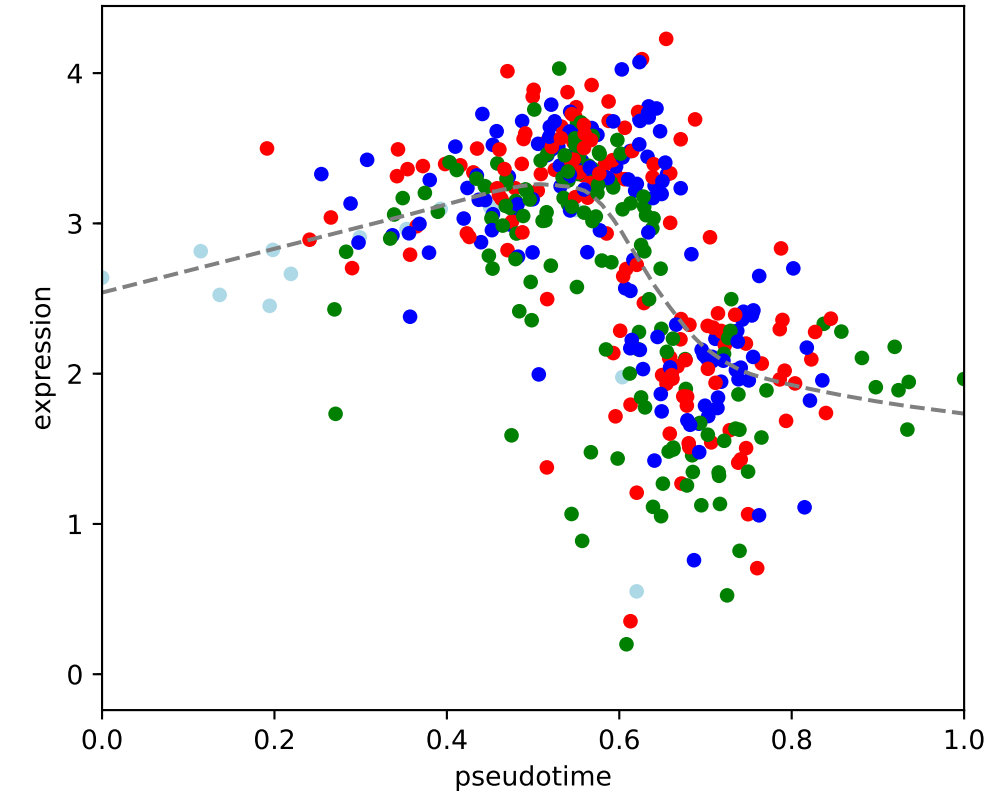
733 1spearman=-0.47



ENST00000554282 CKB (creatine kinase B)

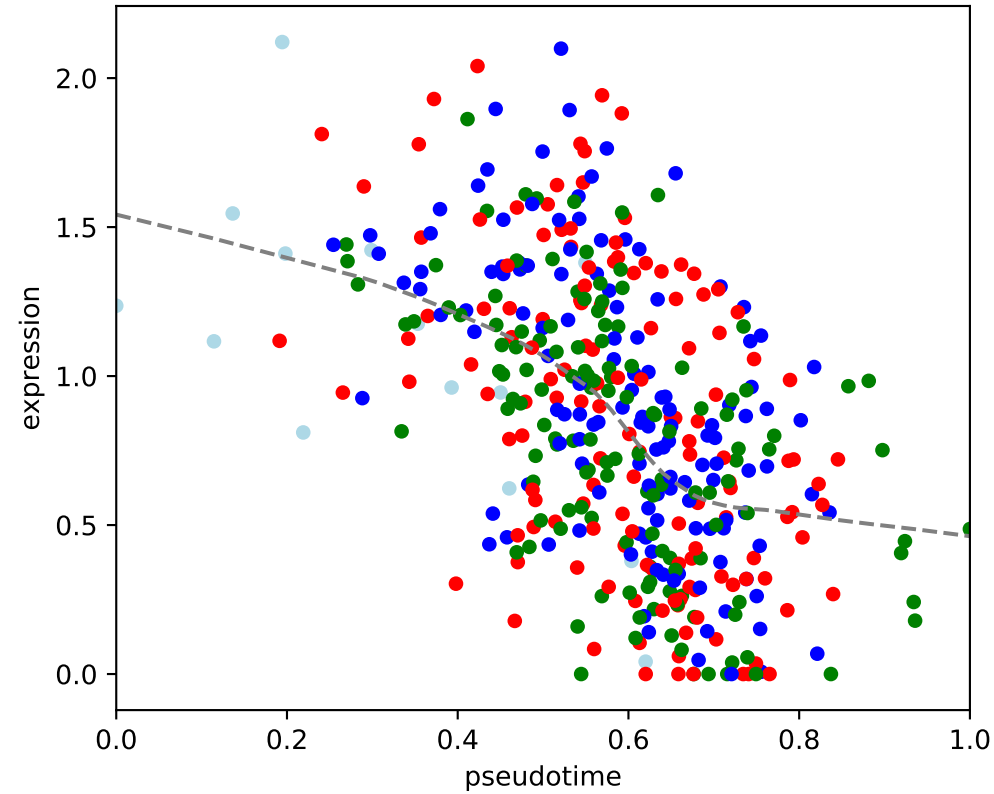
733 0spearman=-0.54

733 1spearman=-0.42

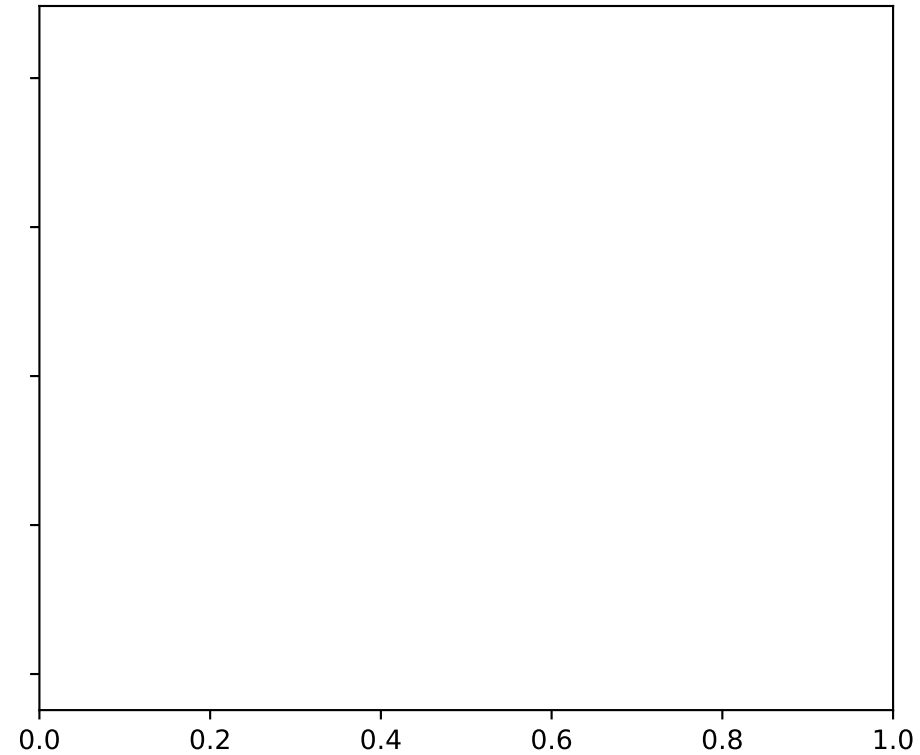
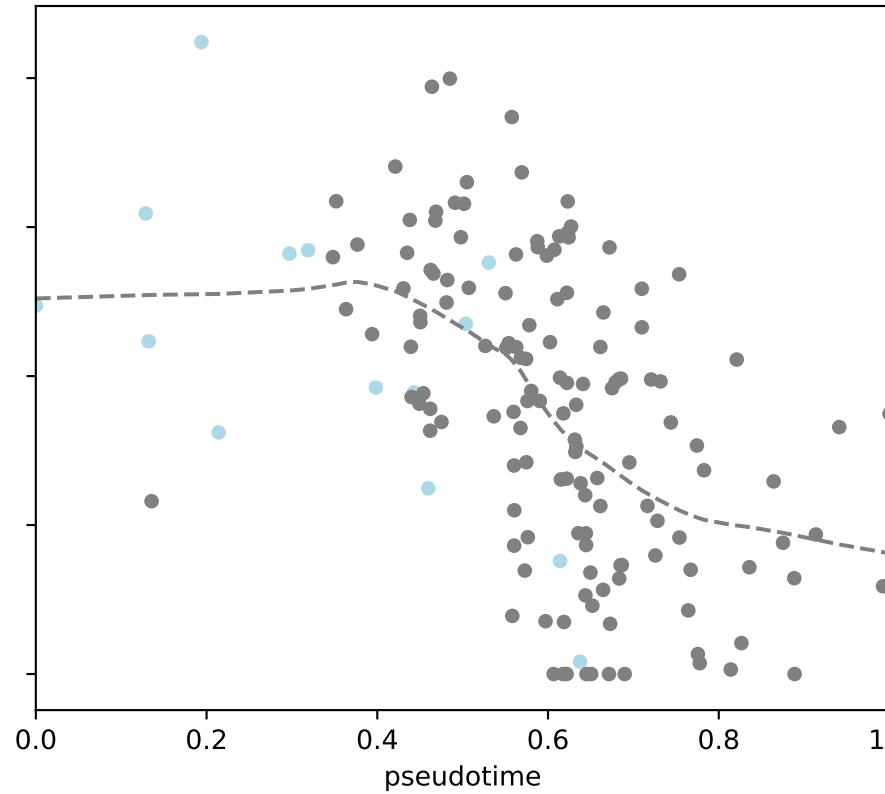


ENST00000431241 RPSAP19 (ribosomal protein SA pseudogene 19)

733 0spearman=-0.54



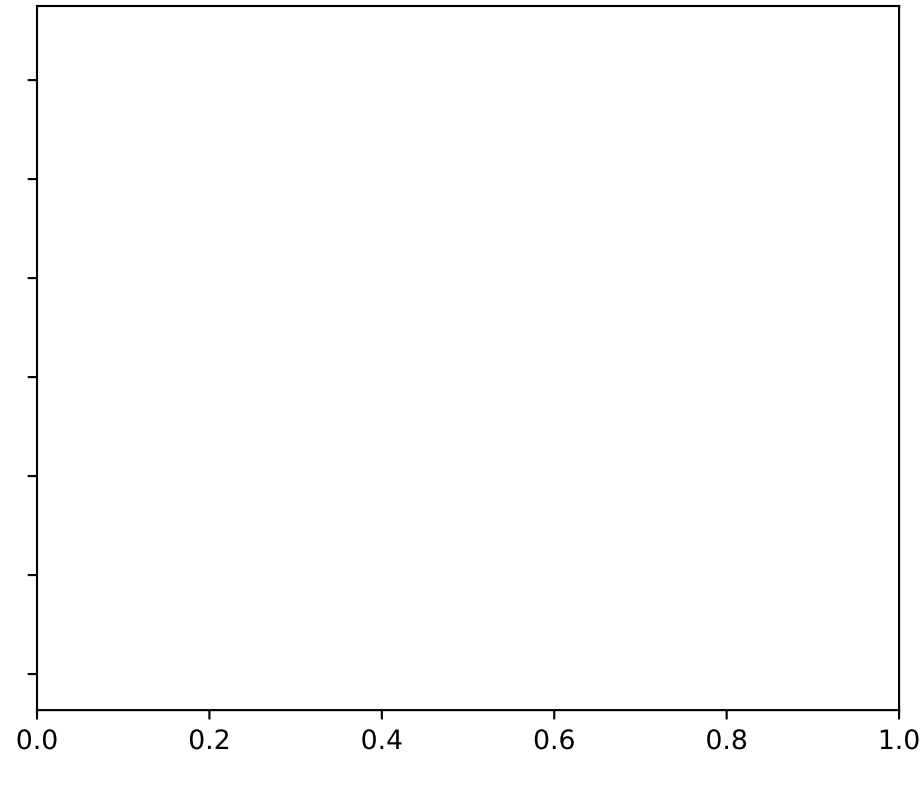
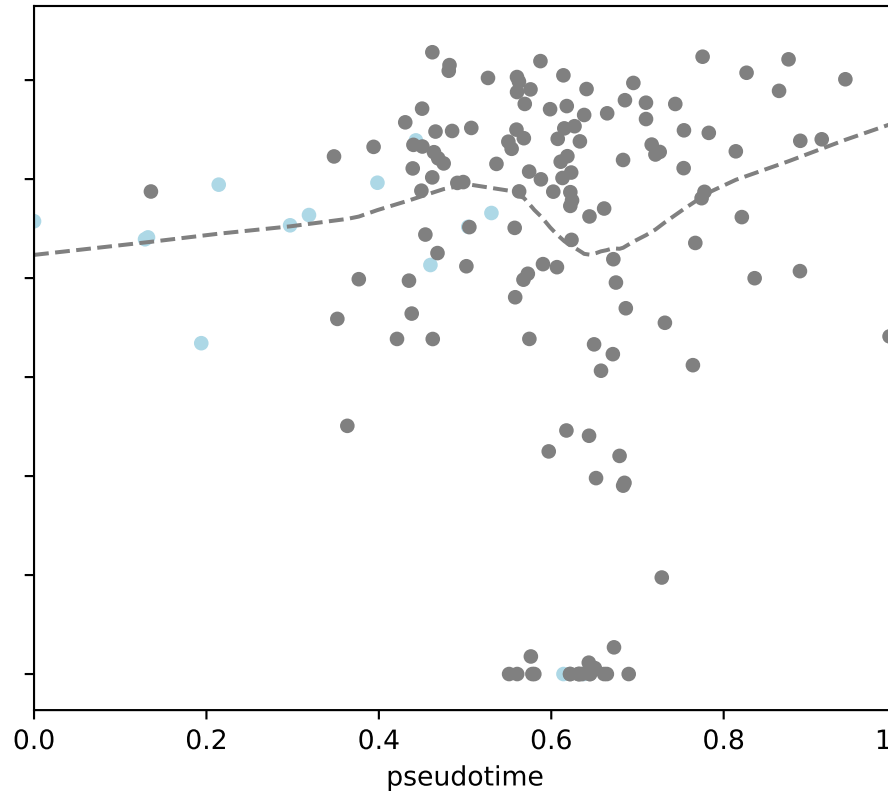
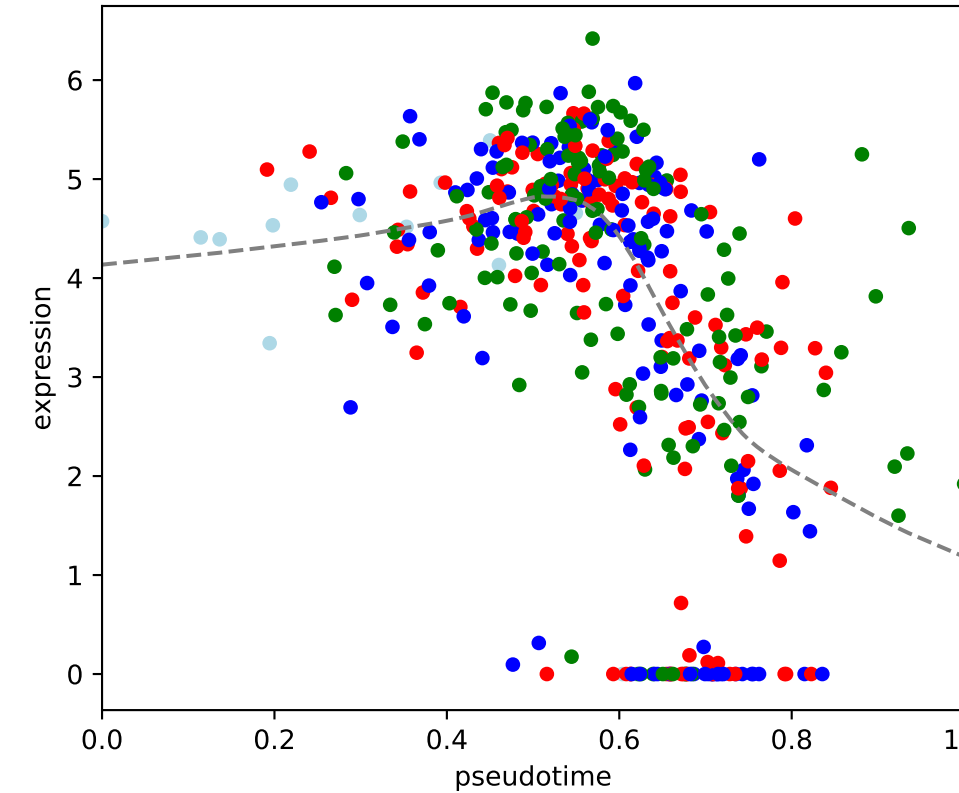
733 1spearman=-0.55



ENST00000218230 PCSK1N (proprotein convertase subtilisin/kexin type 1 inhibitor)

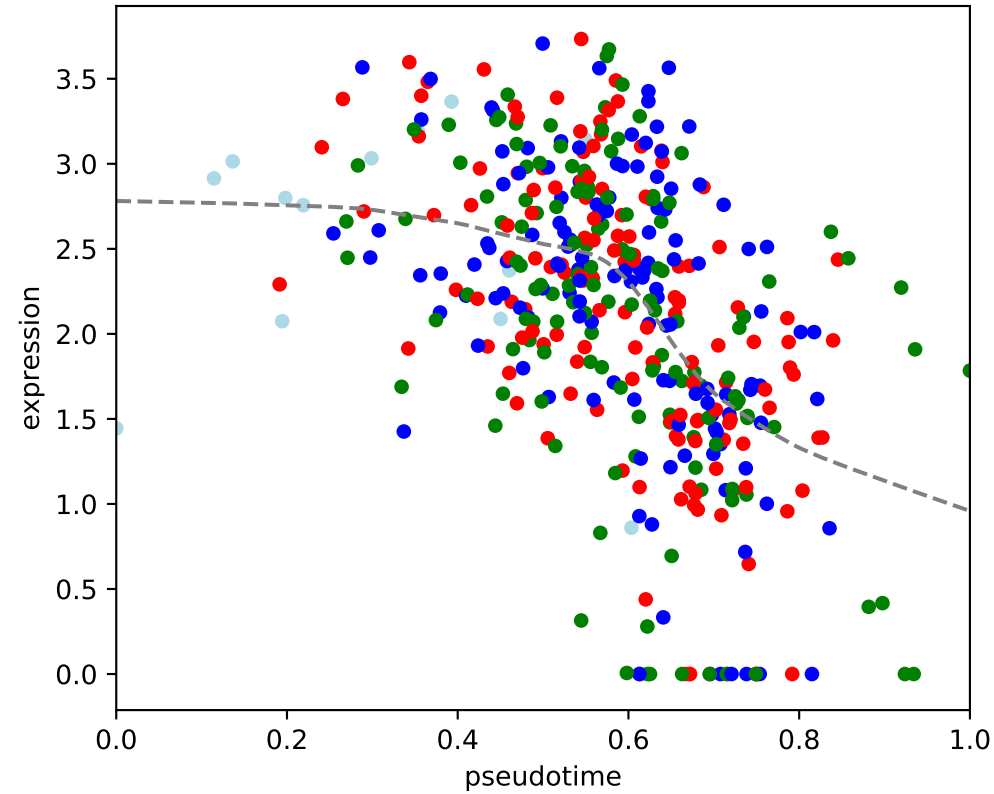
733 0spearman=-0.54

733 1spearman=-0.02

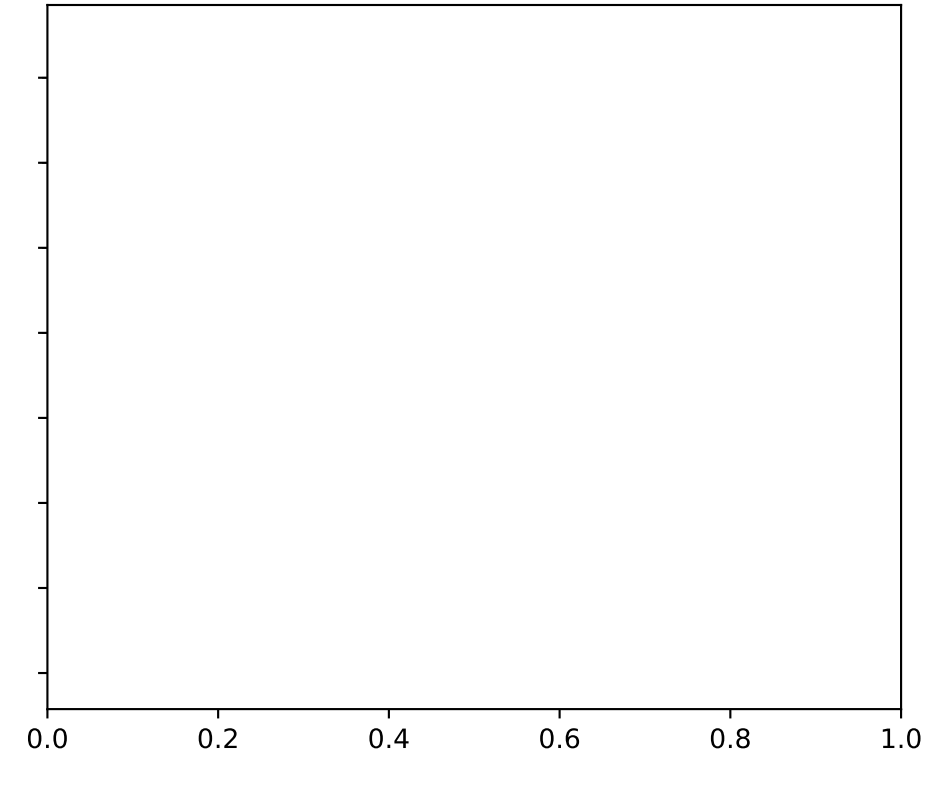
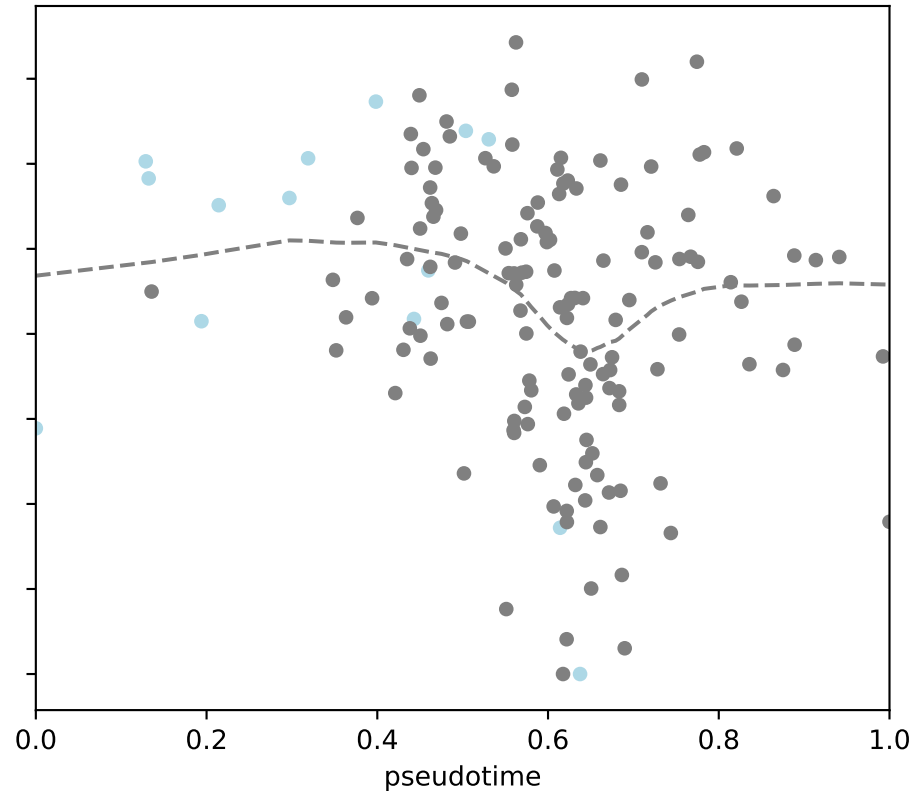


ENST00000492871 GUK1 (guanylate kinase 1)

733 0spearman=-0.54



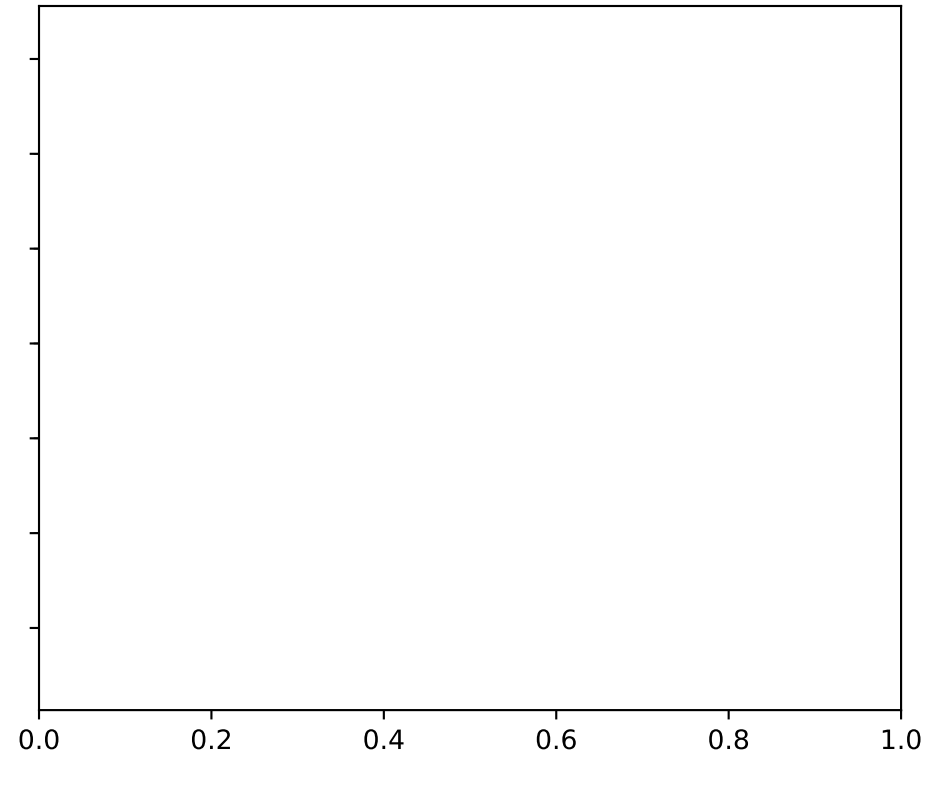
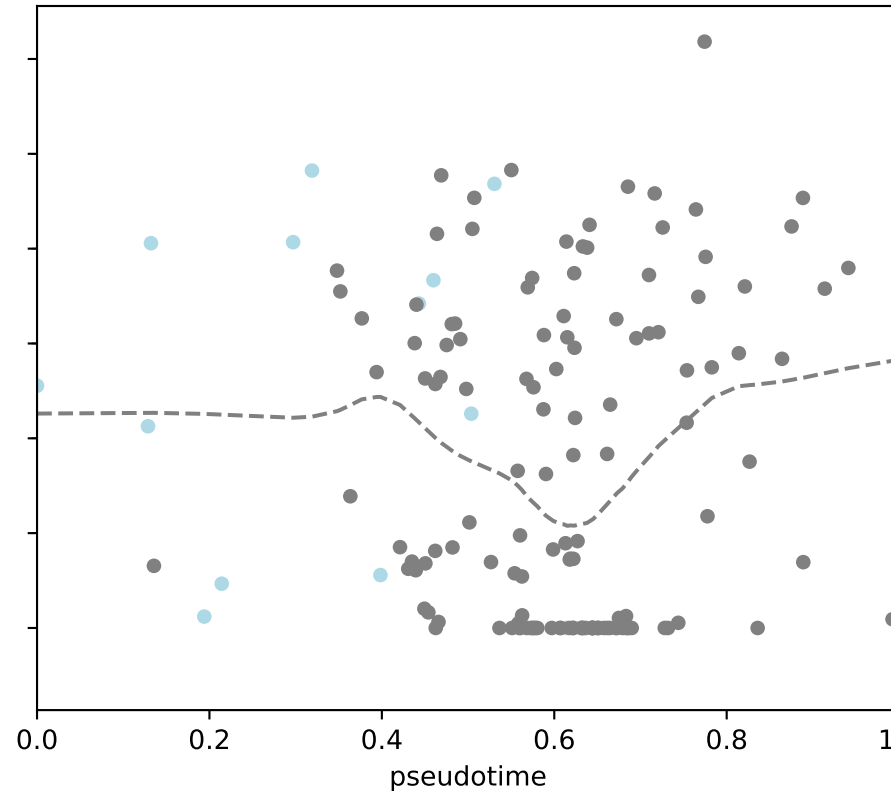
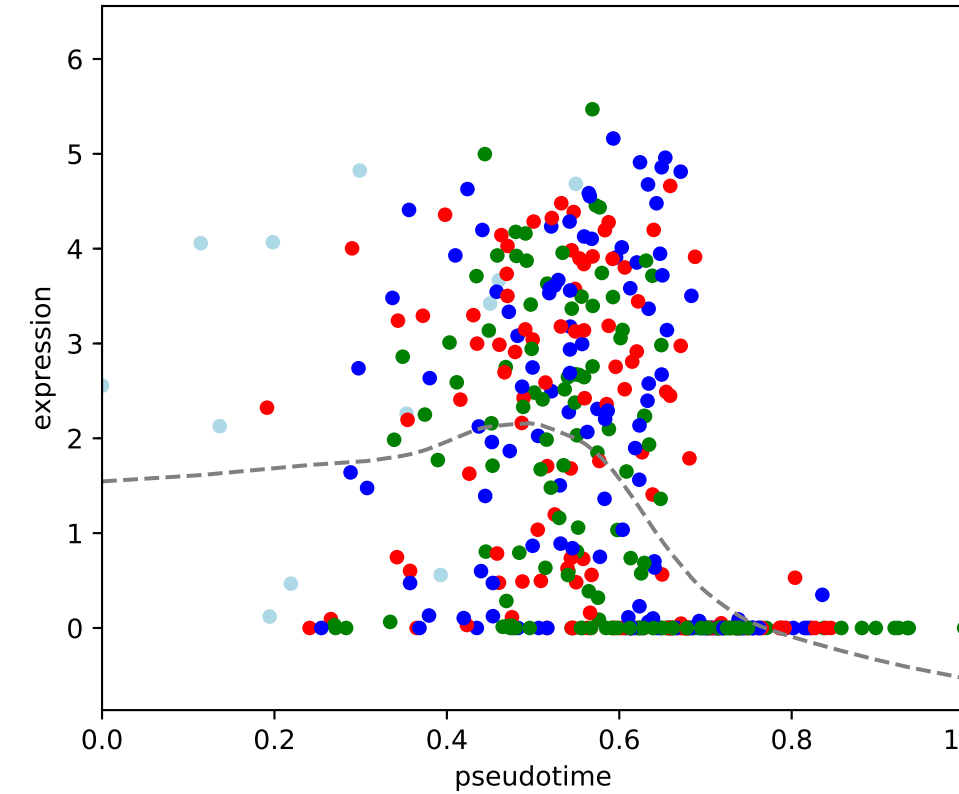
733 1spearman=-0.23



ENST00000280696 THRB (thyroid hormone receptor beta)

733 0spearman=-0.54

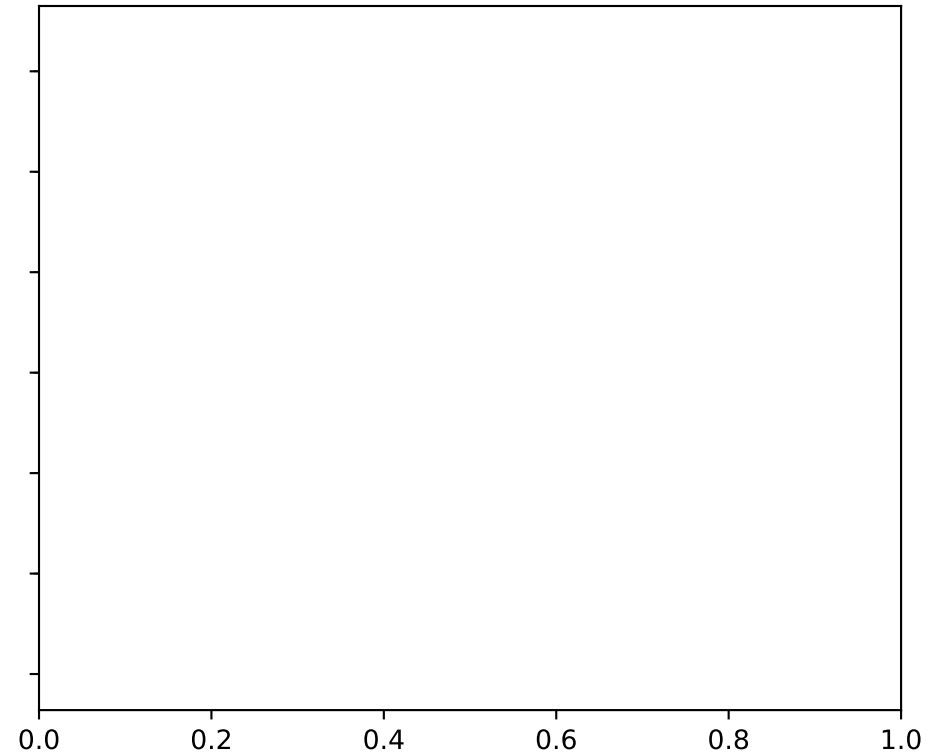
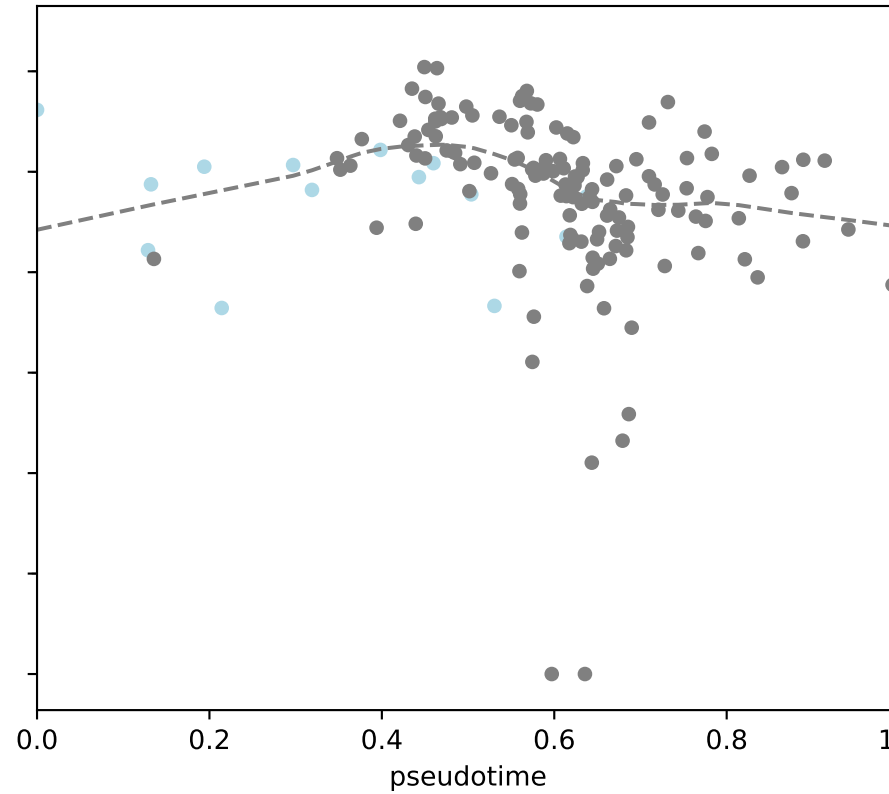
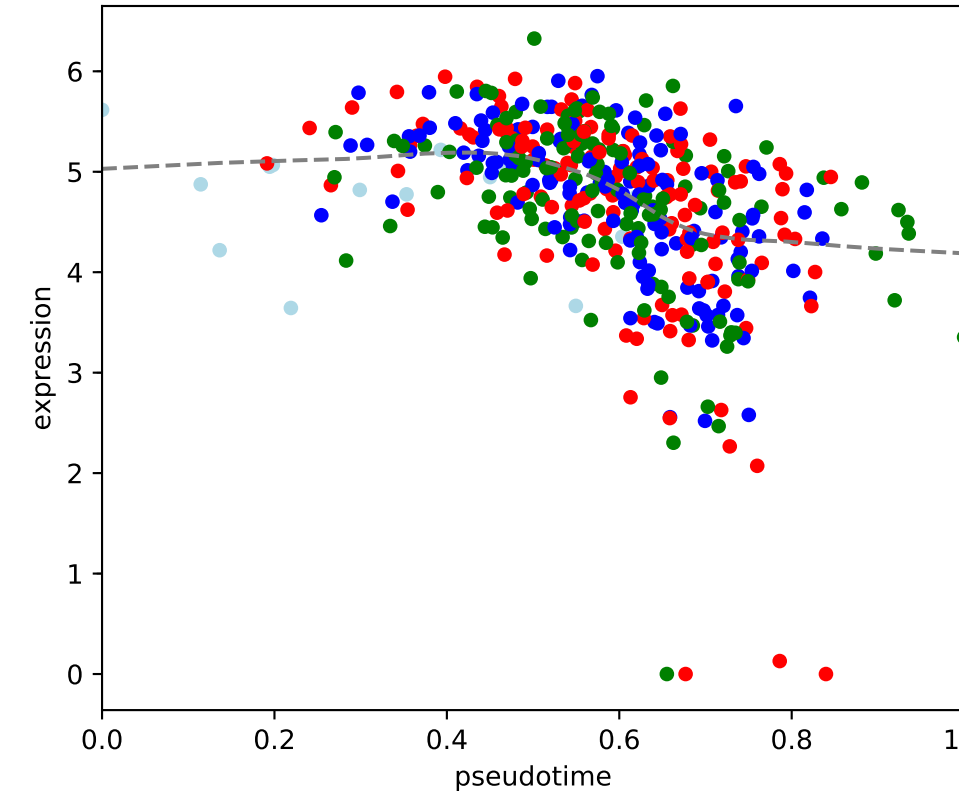
733 1spearman=-0.09



ENST00000274137 NDUF56 (NADH:ubiquinone oxidoreductase subunit S6)

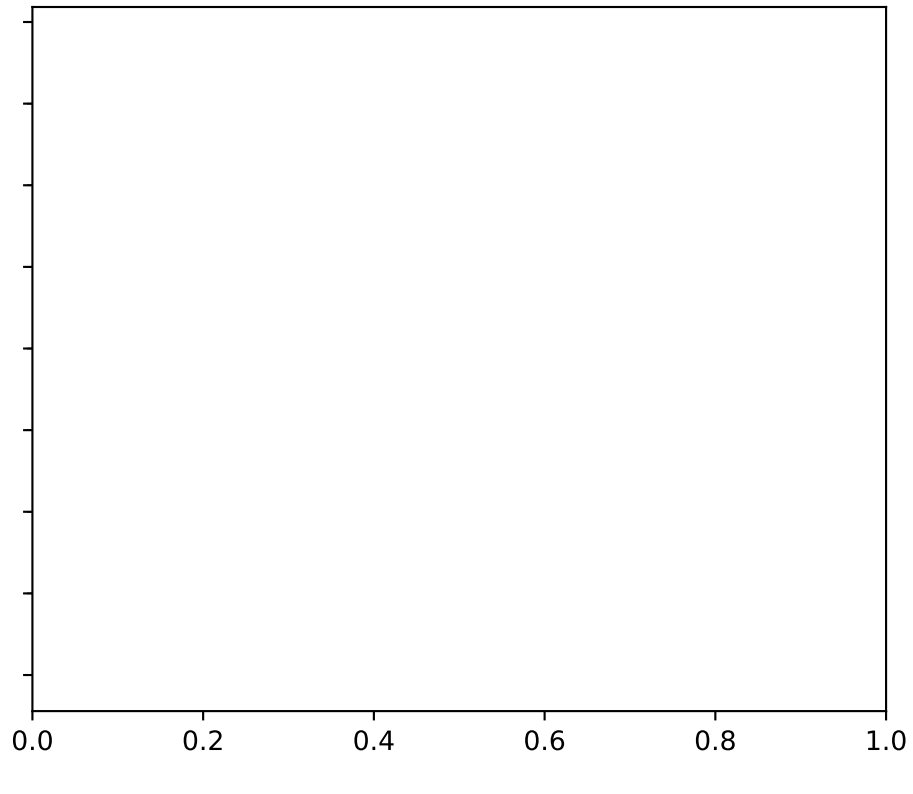
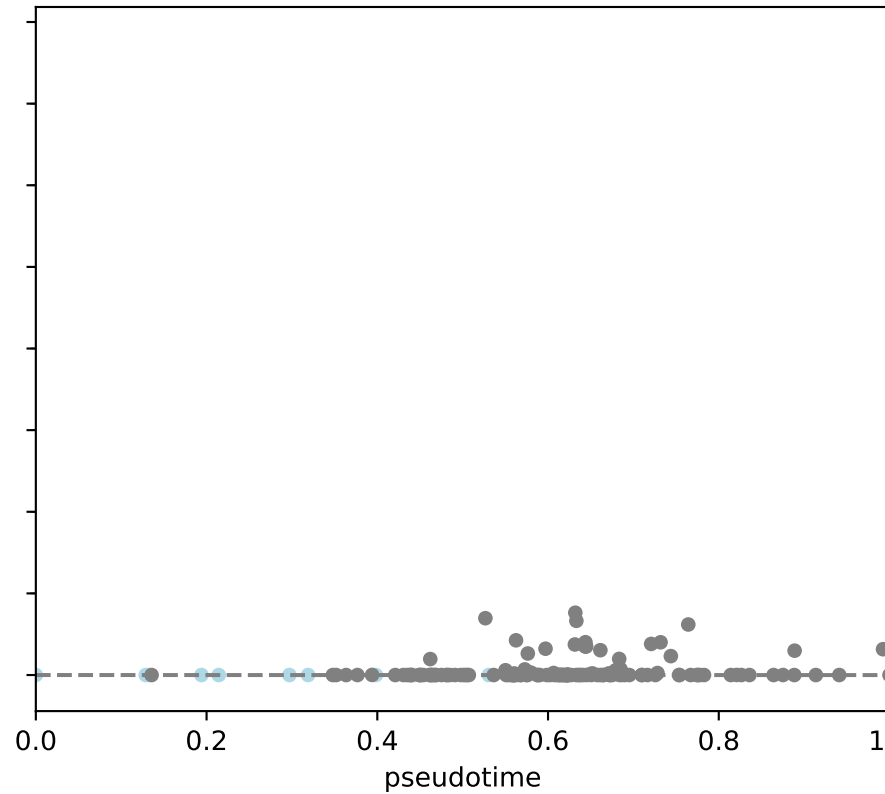
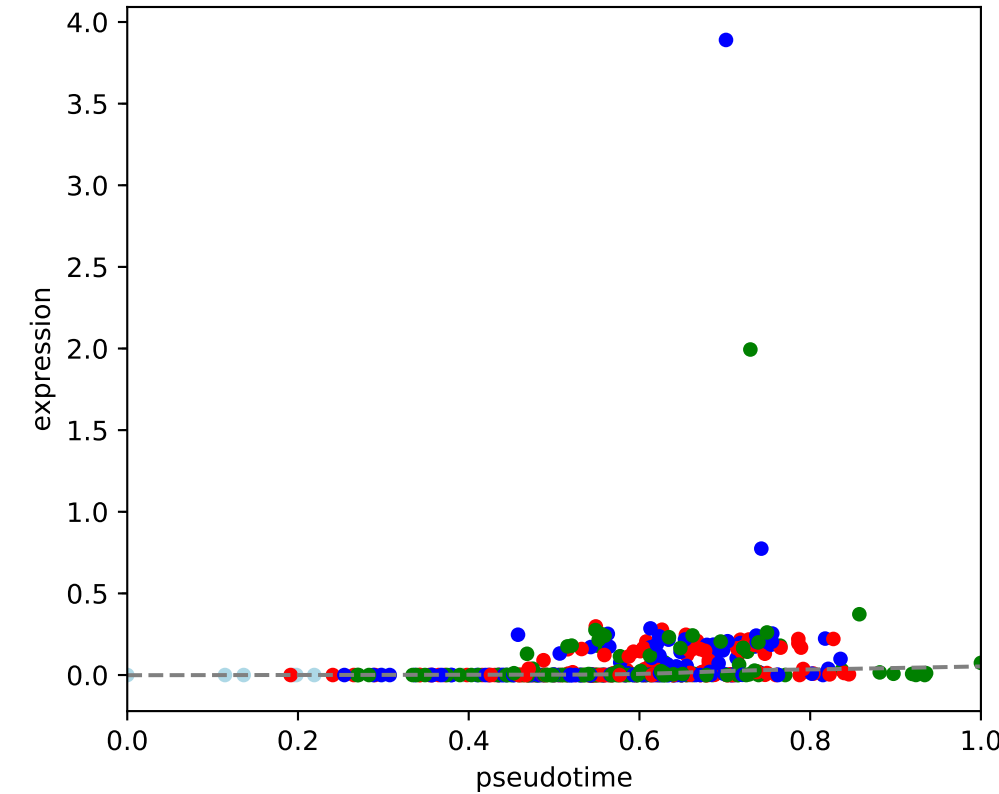
733 0spearman=-0.54

733 1spearman=-0.44



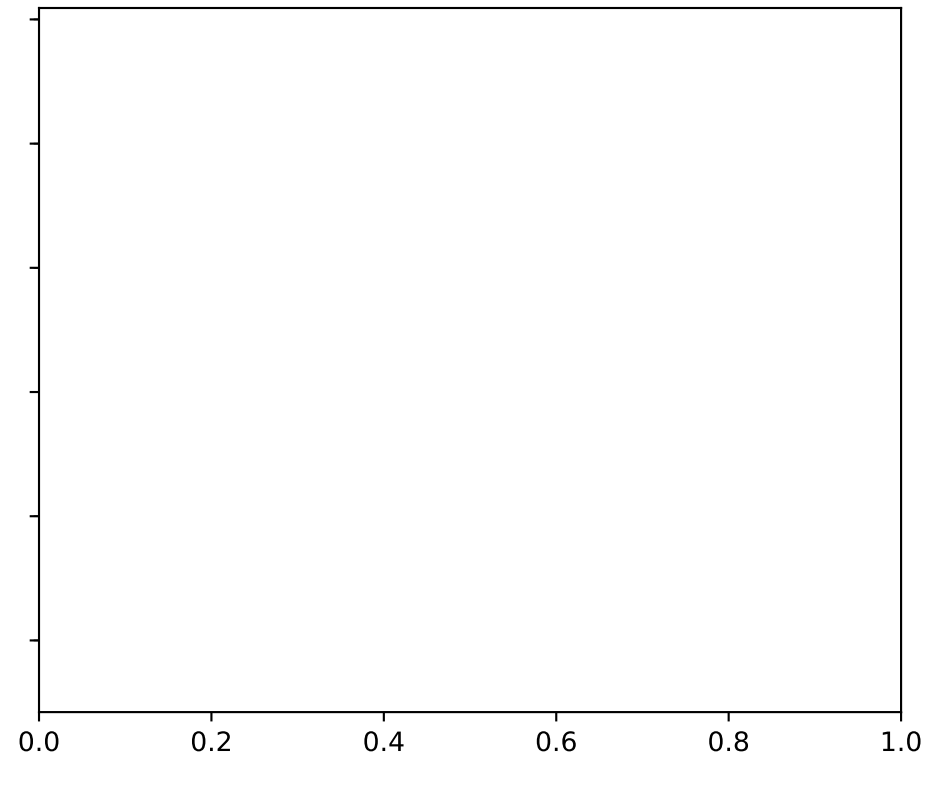
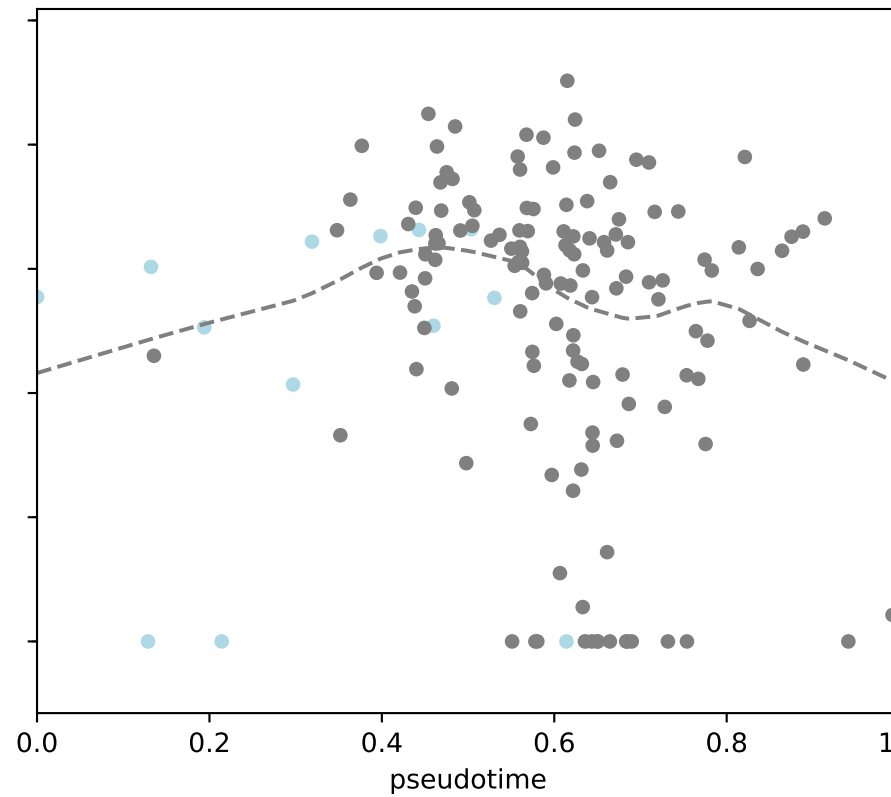
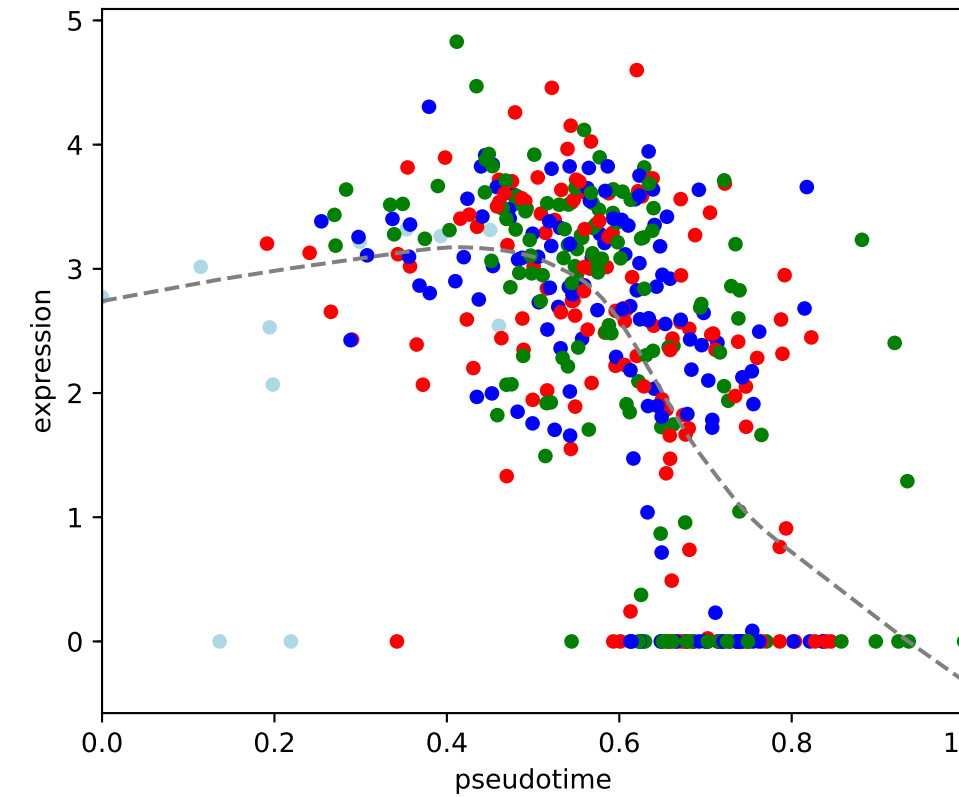
733 0spearman=0.54

733 1spearman=0.25



733 0spearman=-0.54

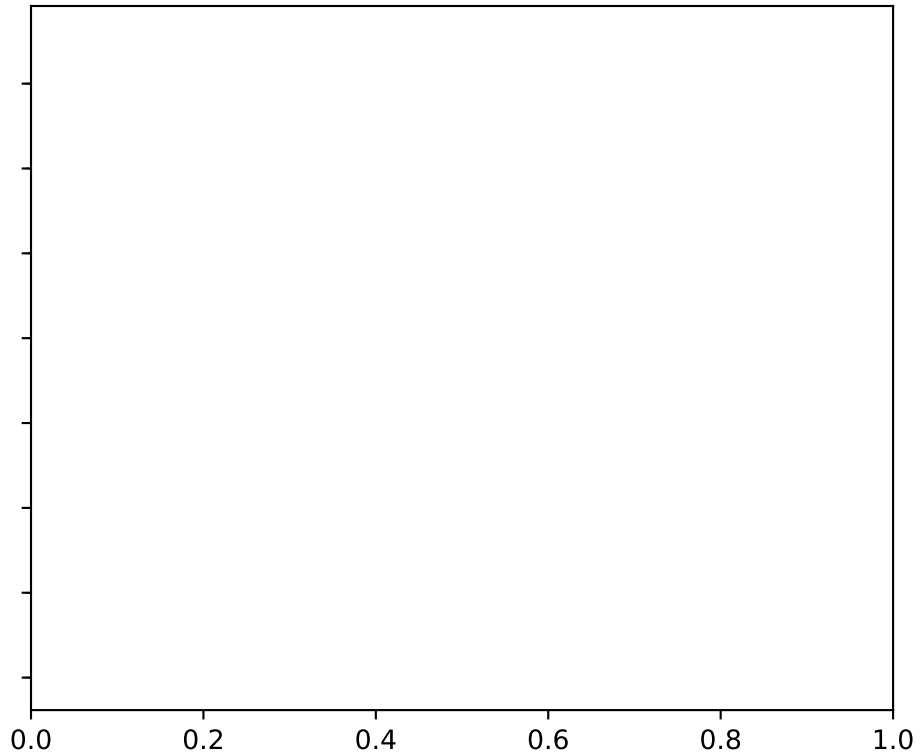
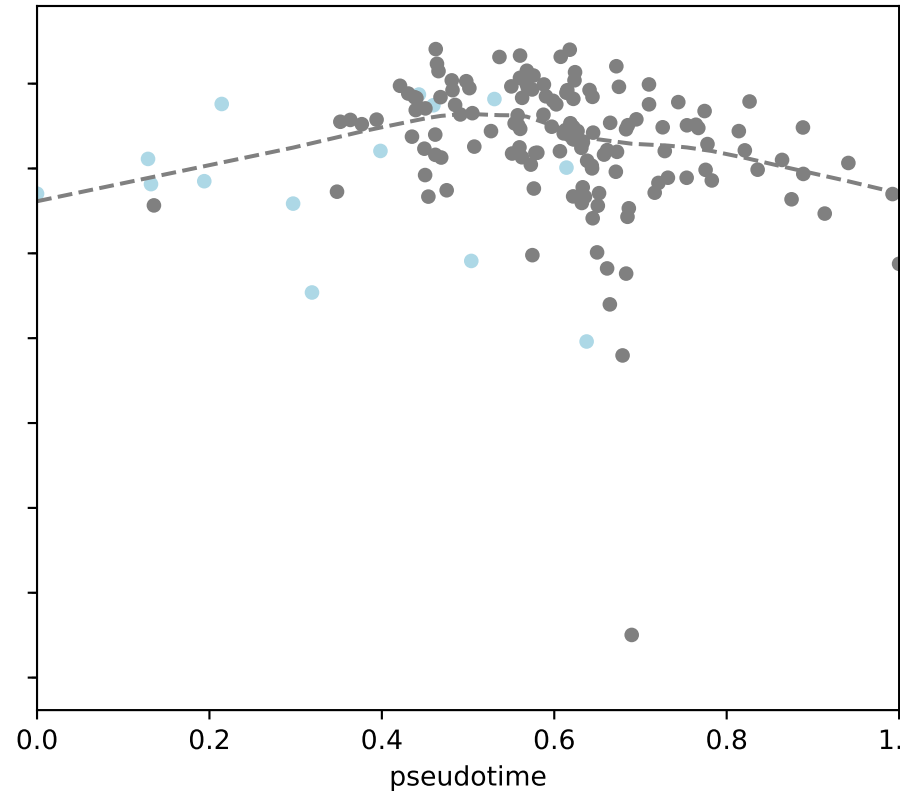
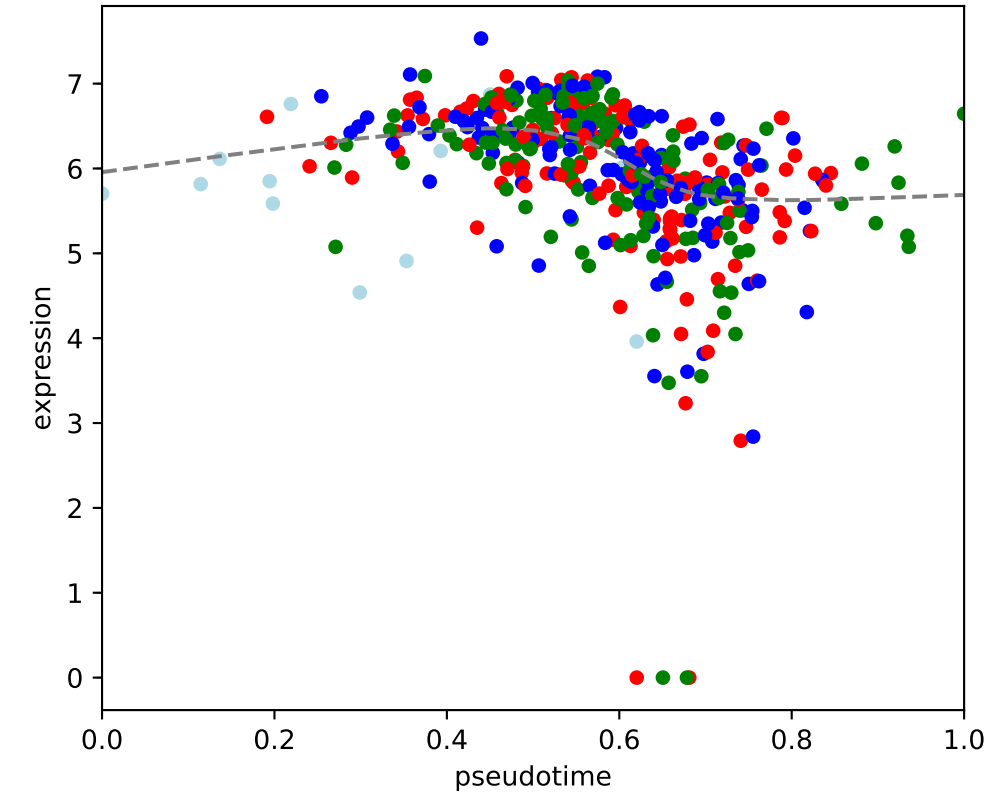
733 1spearman=-0.22



ENST00000245458 RPS29 (ribosomal protein S29)

733 0spearman=-0.54

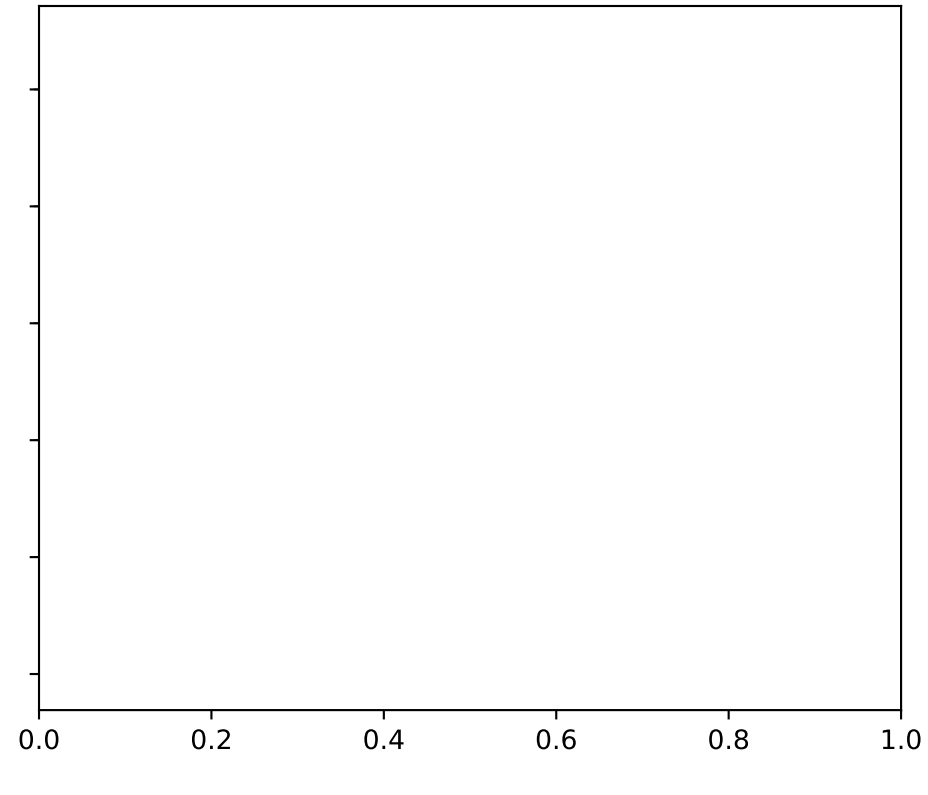
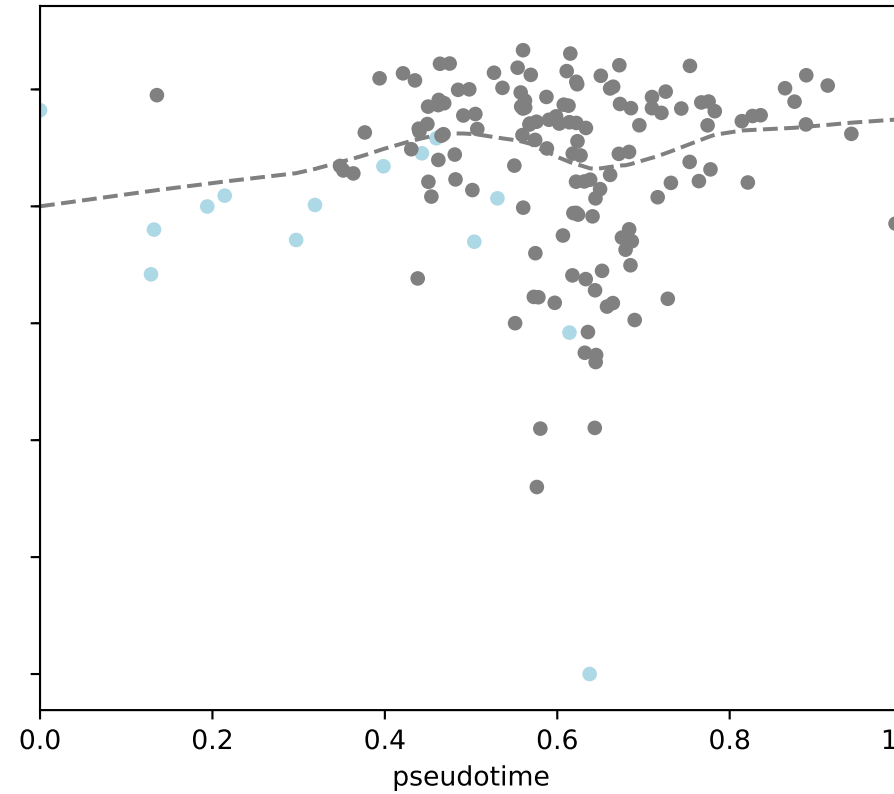
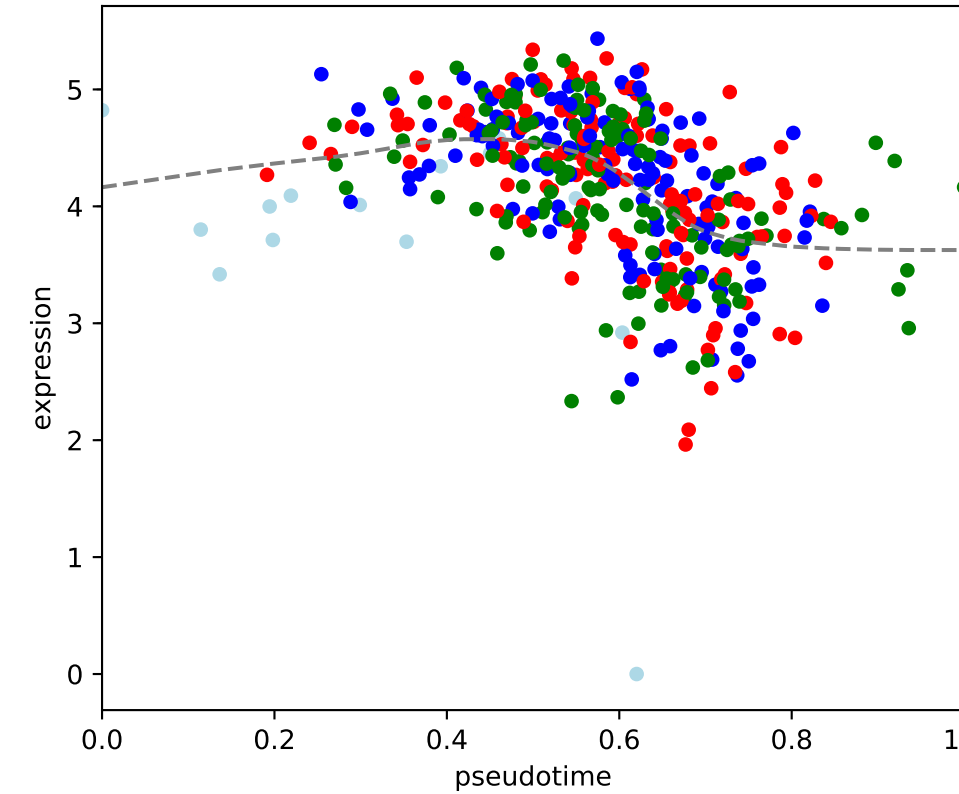
733 1spearman=-0.25



ENST00000243997 ATP5F1E (ATP synthase F1 subunit epsilon)

733 0spearman=-0.54

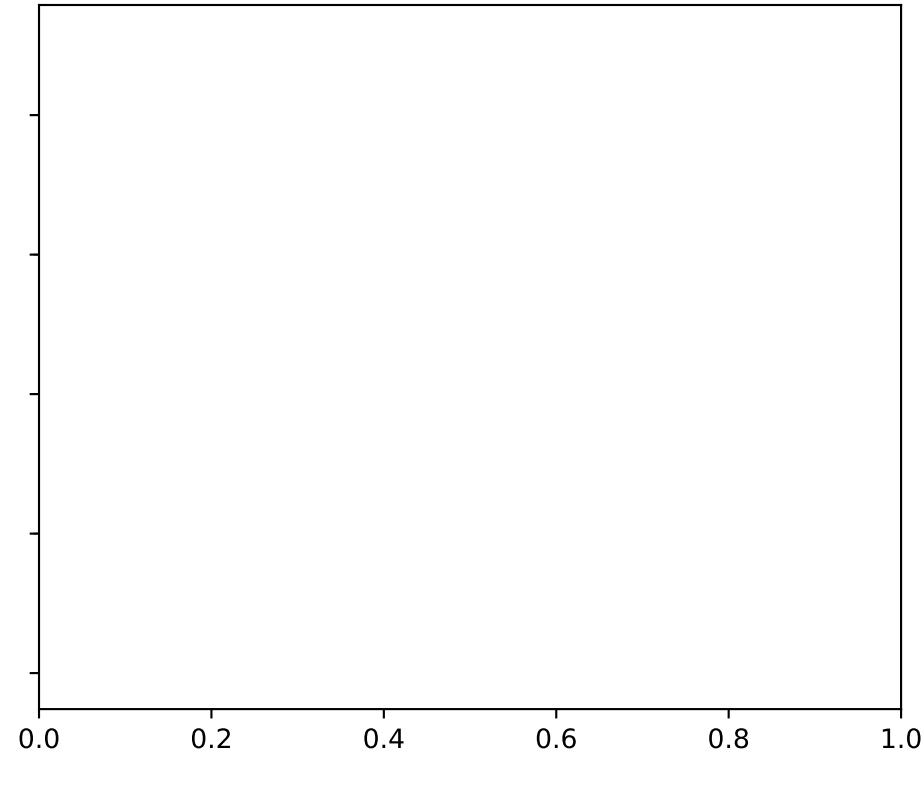
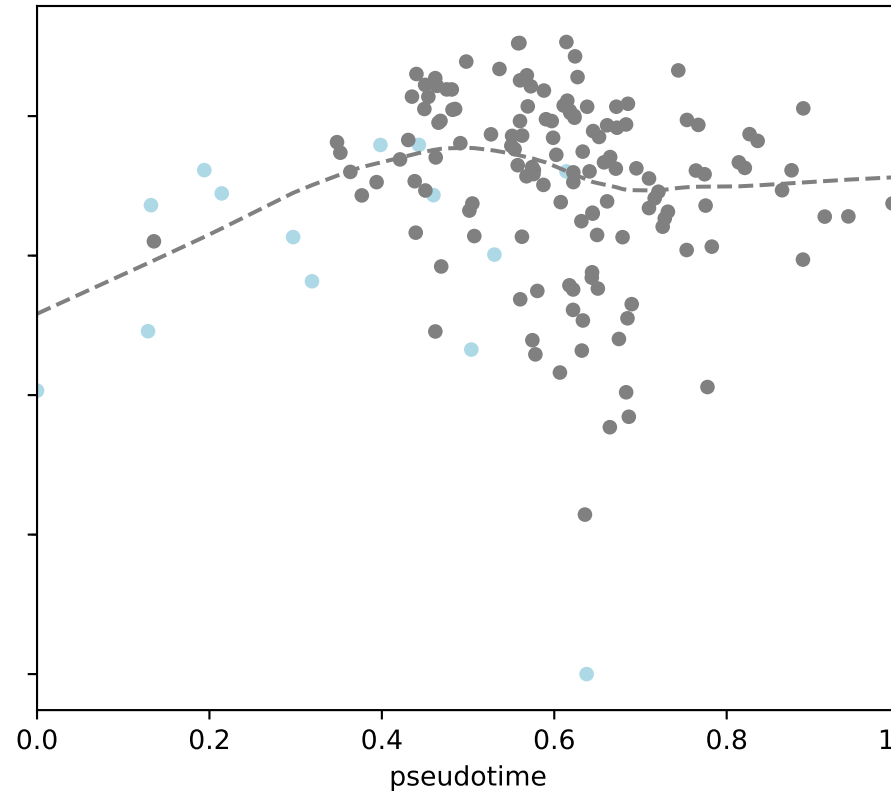
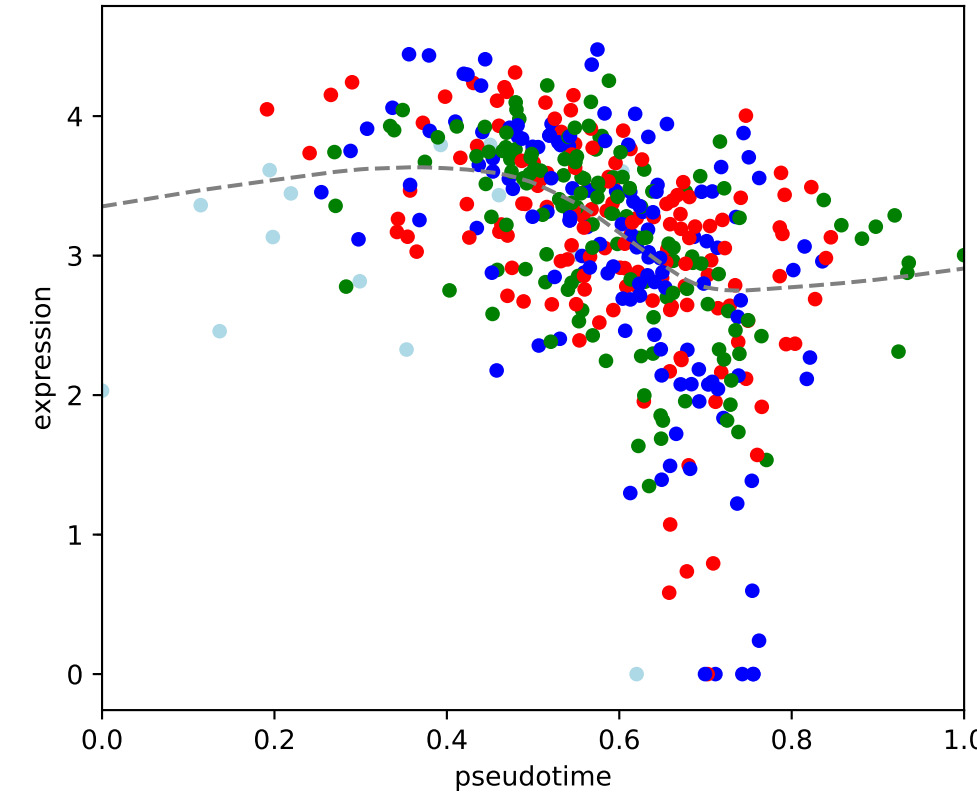
733 1spearman=-0.04



ENST00000491306 RPL37A (ribosomal protein L37a)

733 0spearman=-0.54

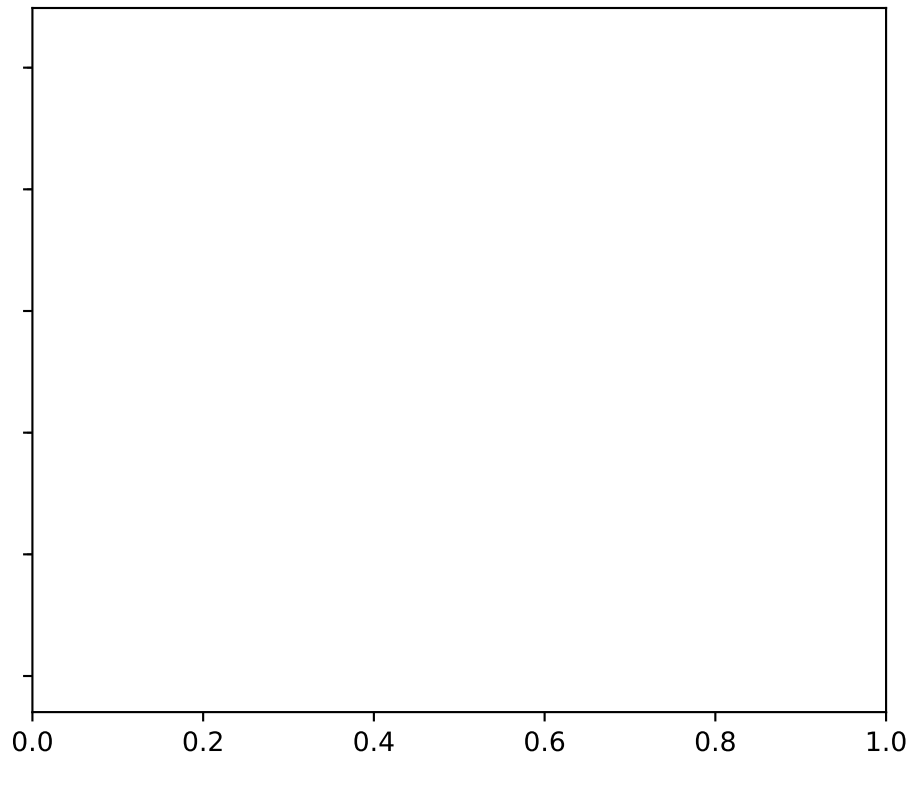
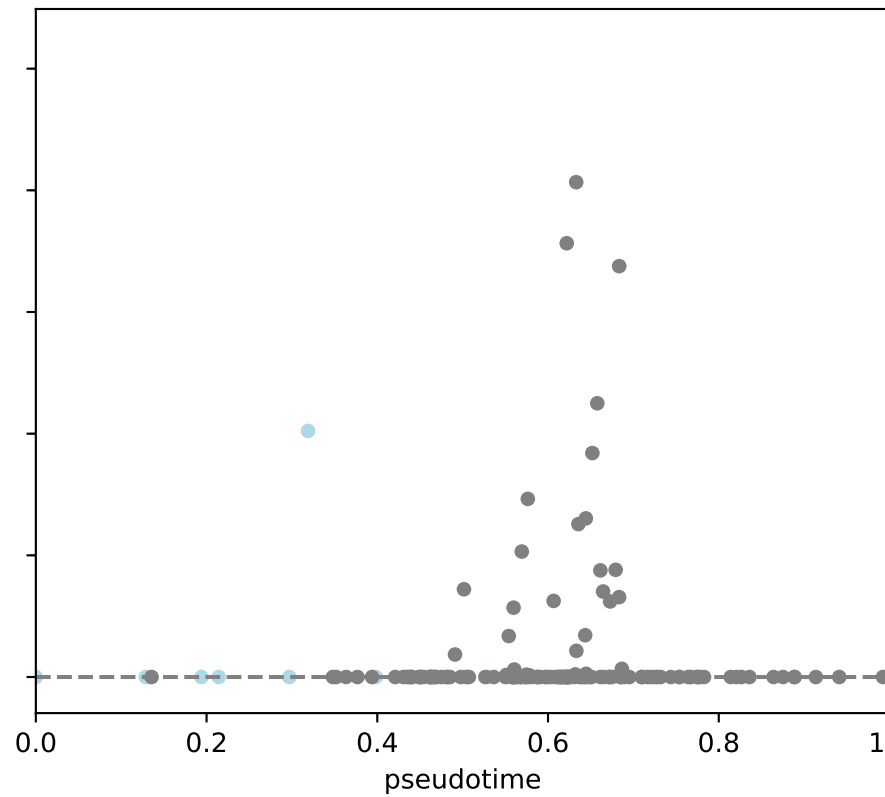
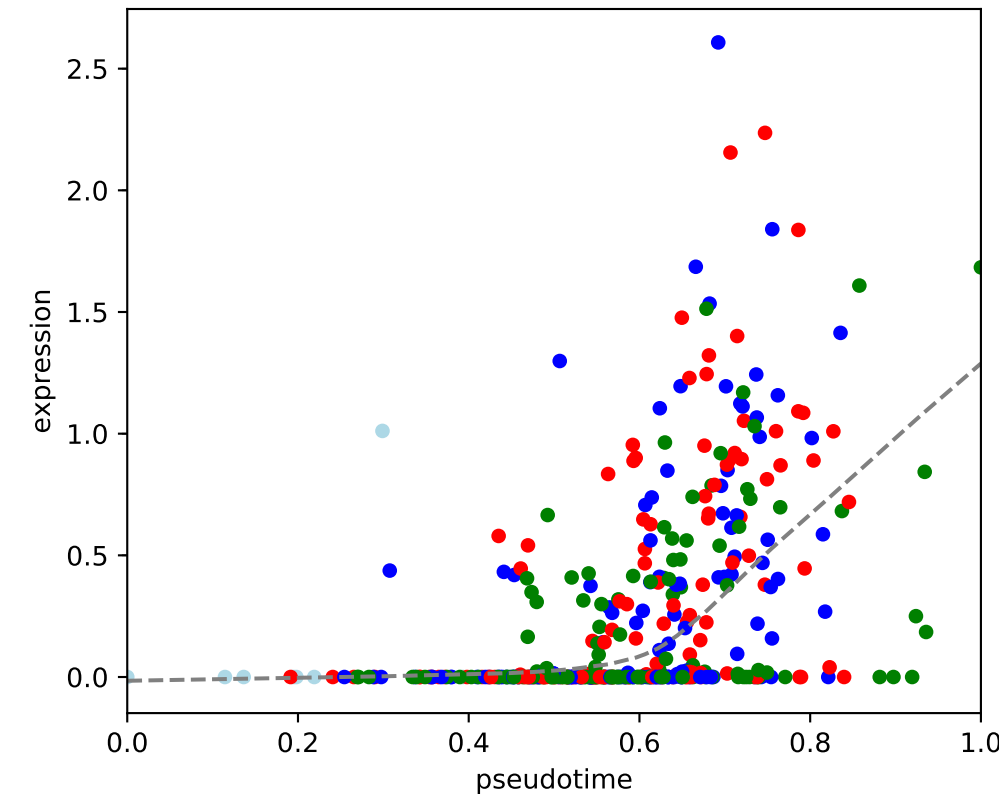
733 1spearman=-0.15



ENST00000564448 CNGB1 (cyclic nucleotide gated channel beta 1)

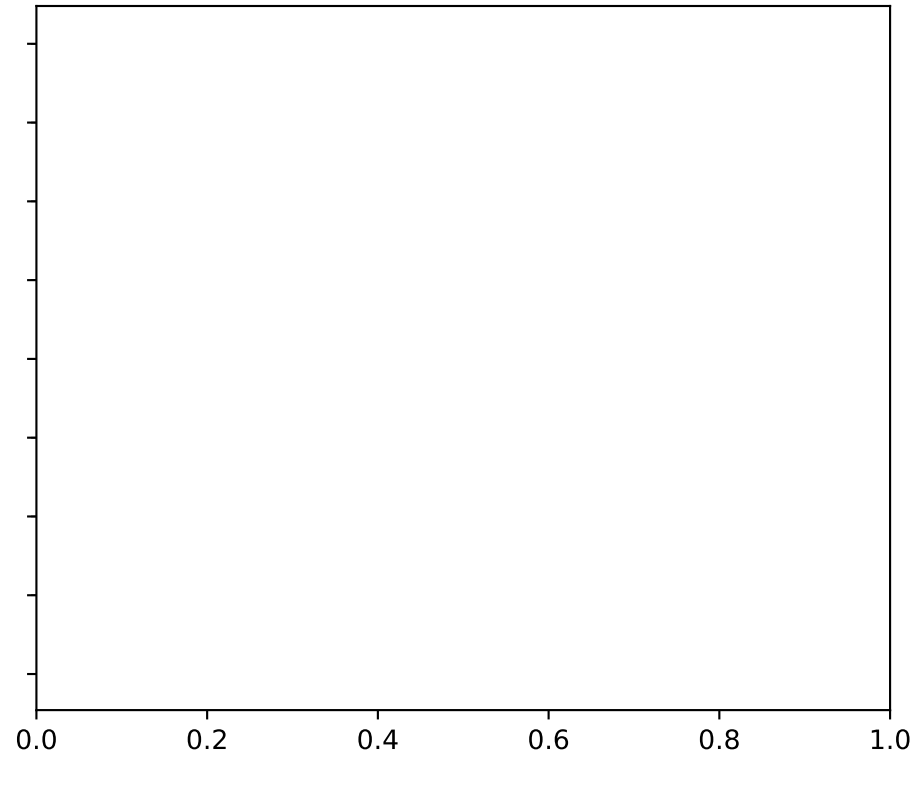
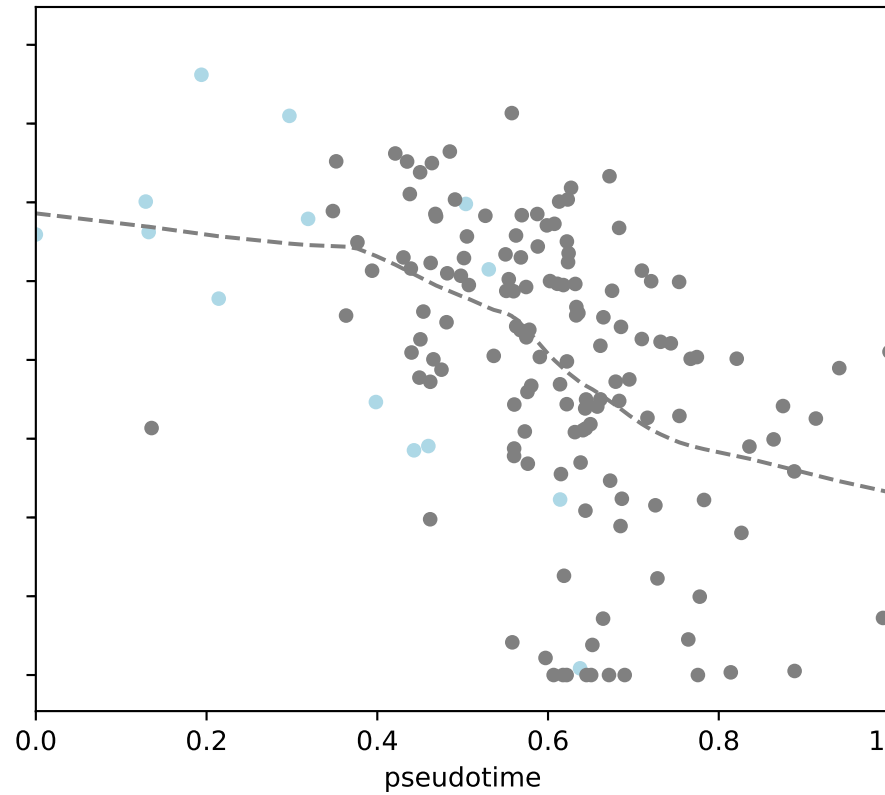
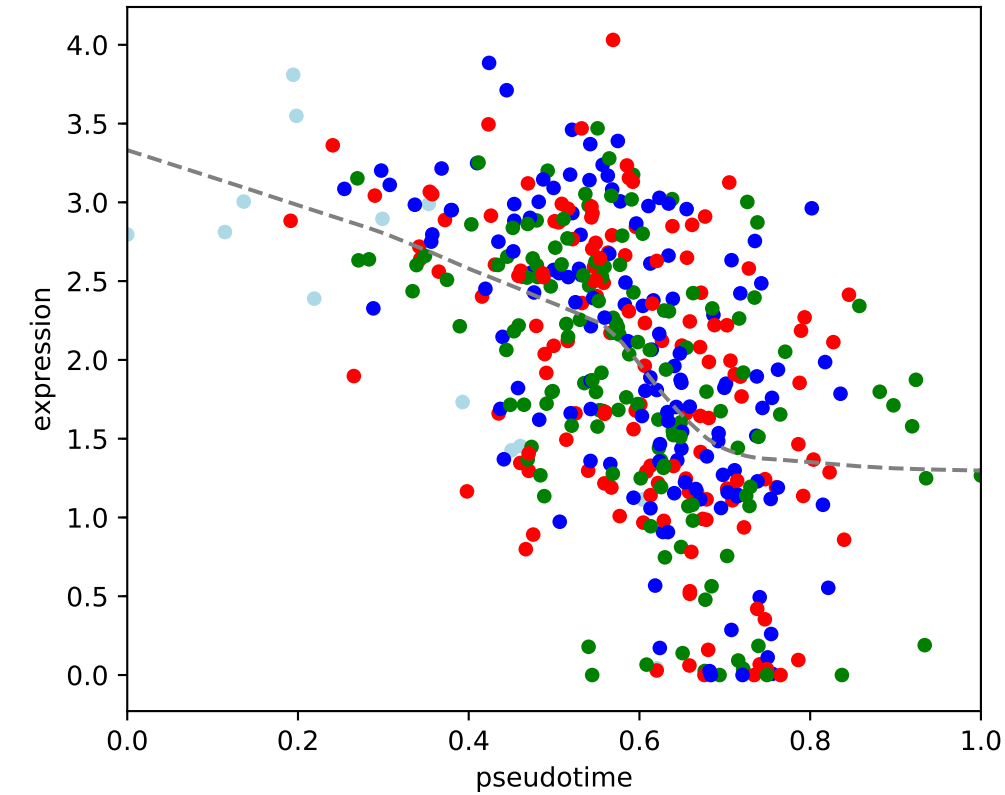
733 0spearman=0.54

733 1spearman=0.08



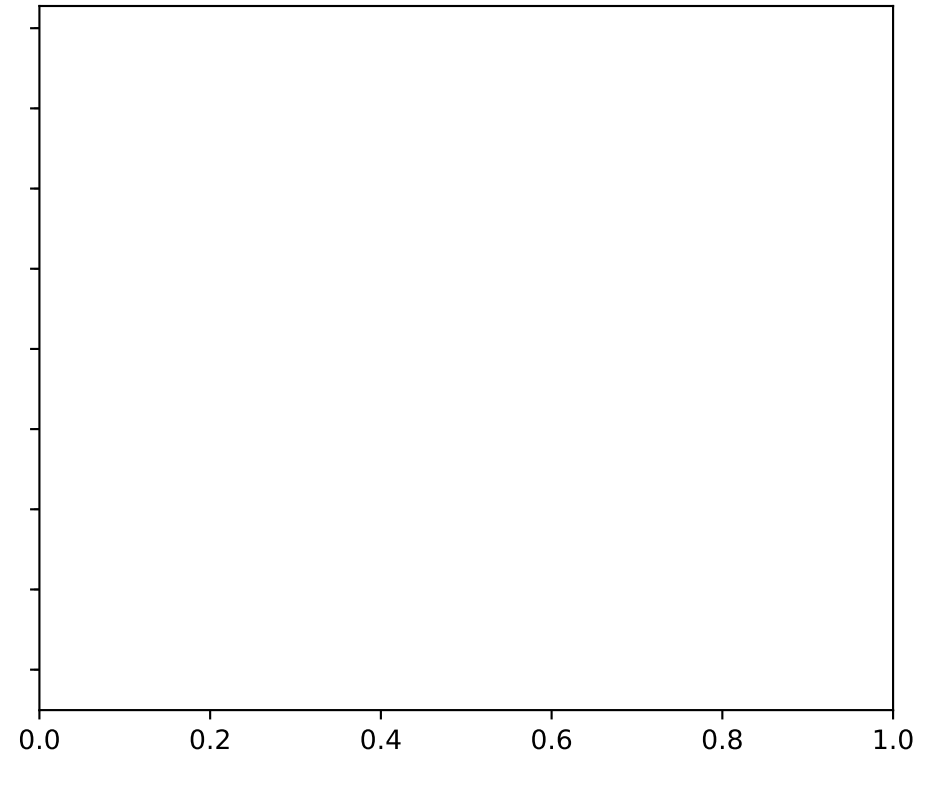
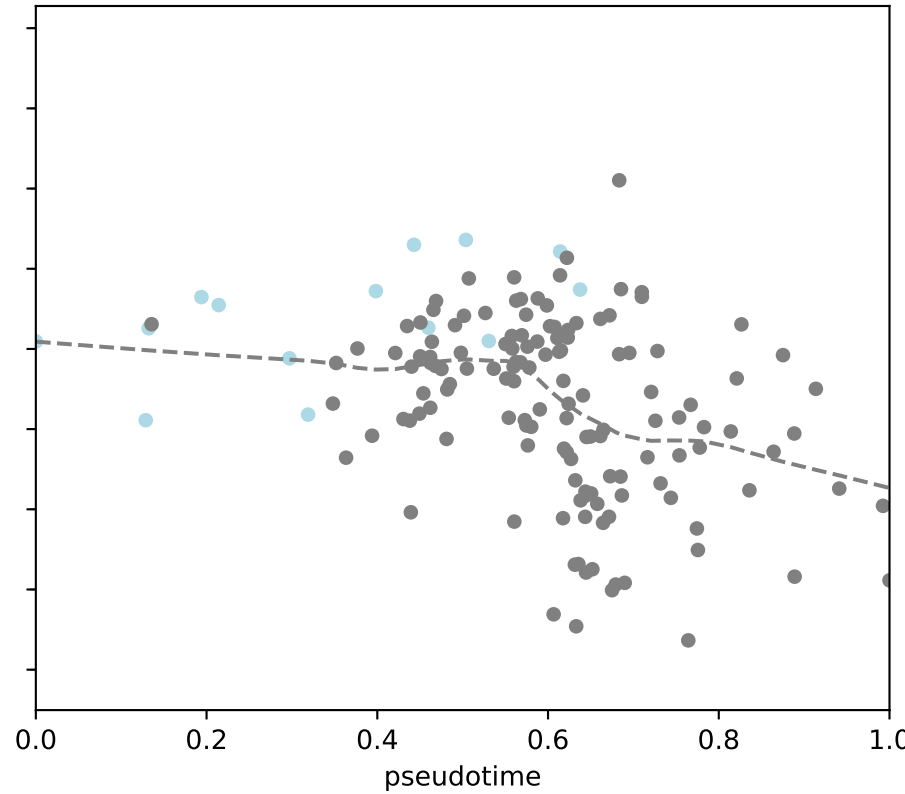
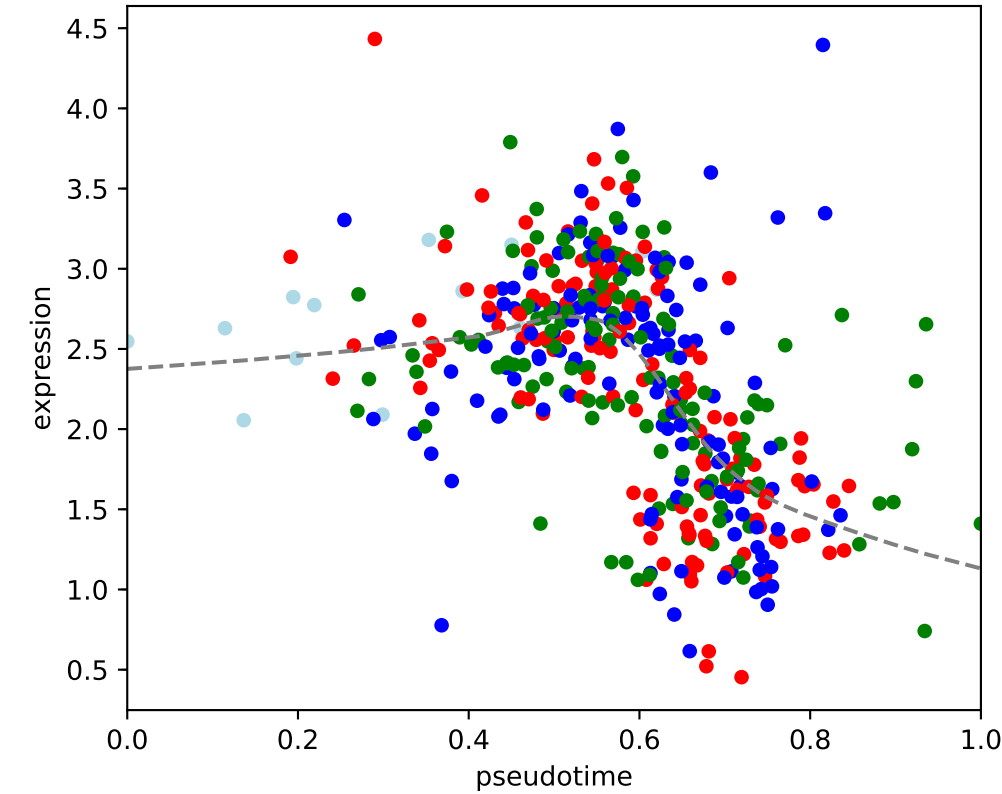
733 0spearman=-0.54

733 1spearman=-0.52



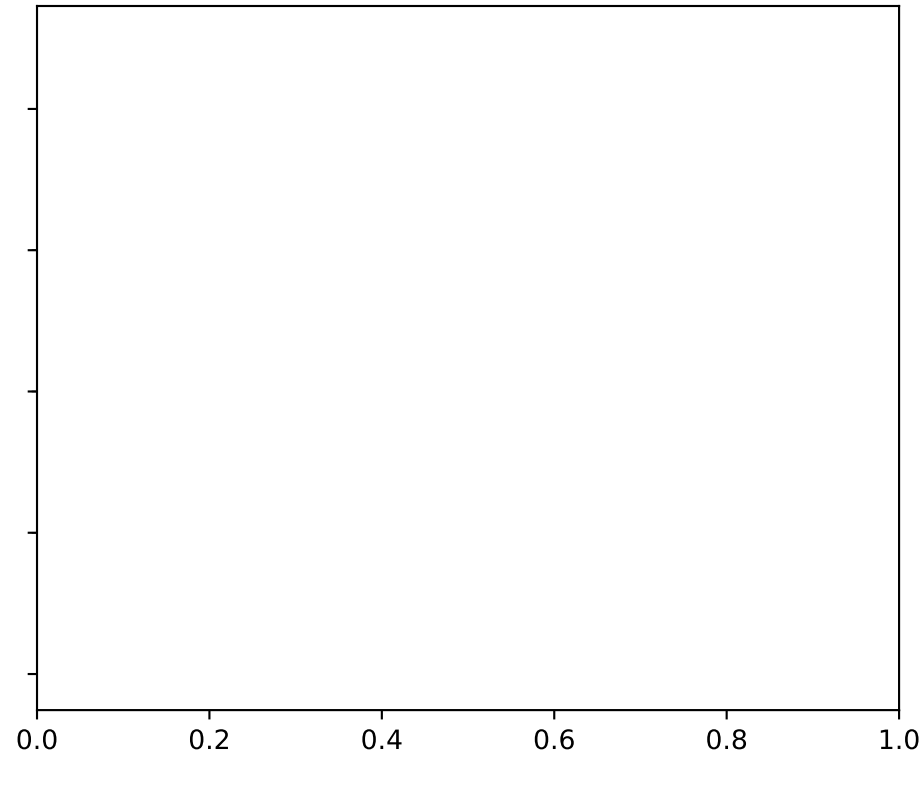
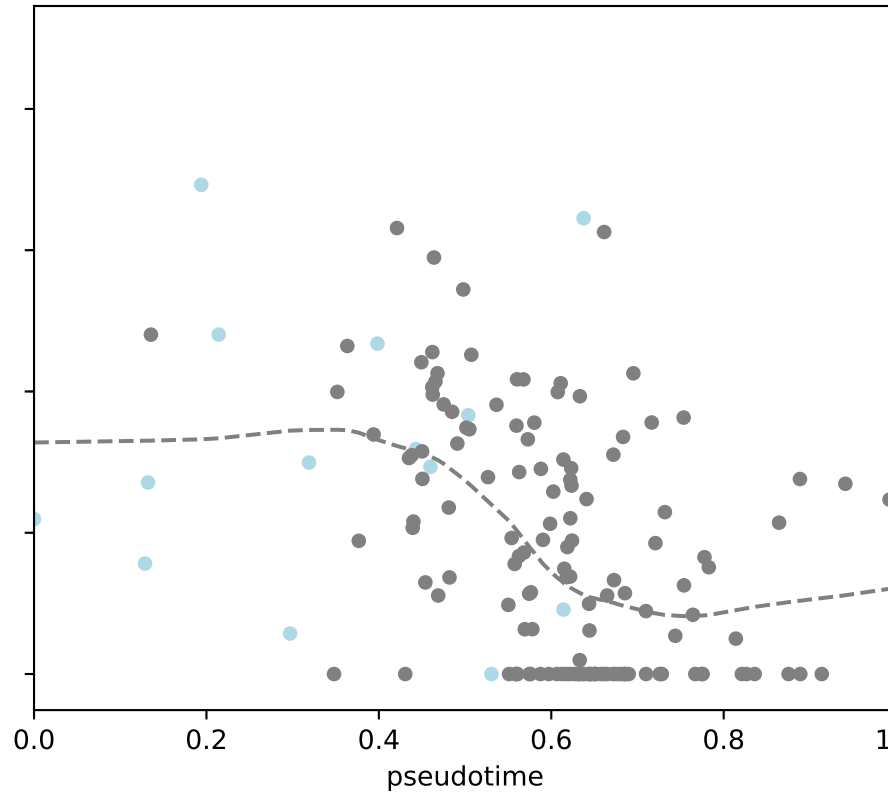
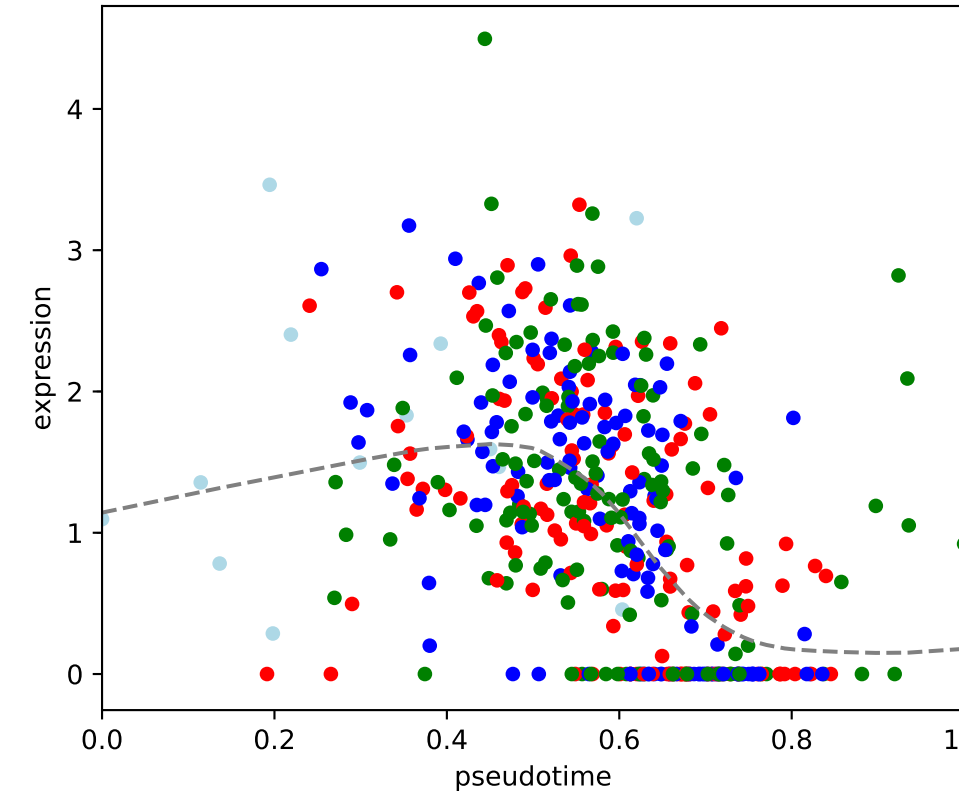
733 0spearman=-0.54

733 1spearman=-0.41



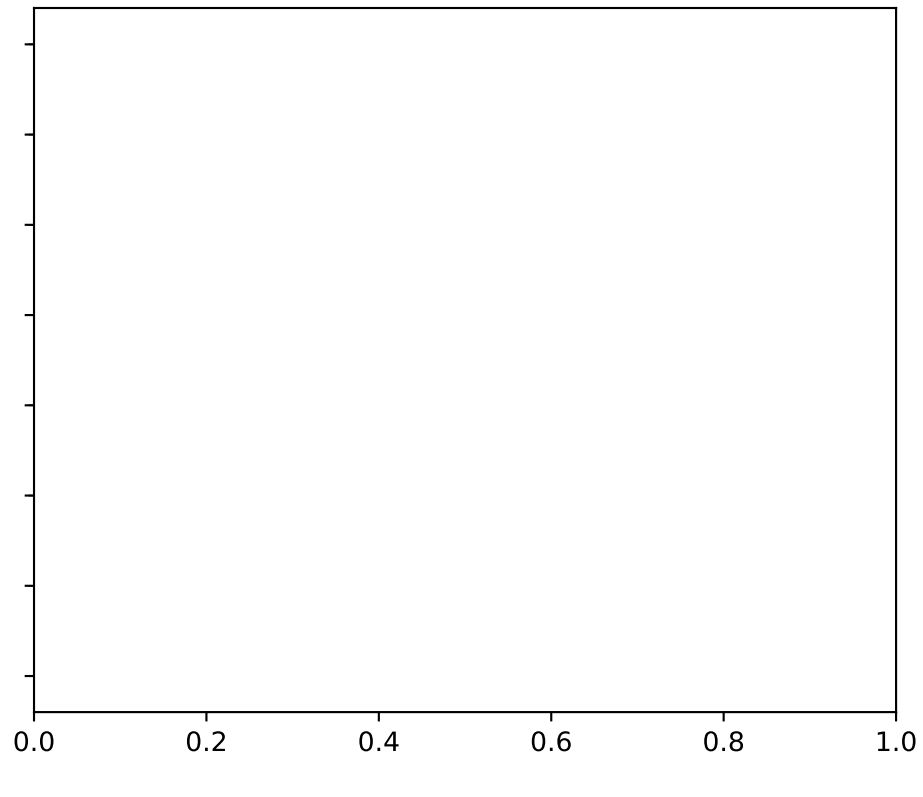
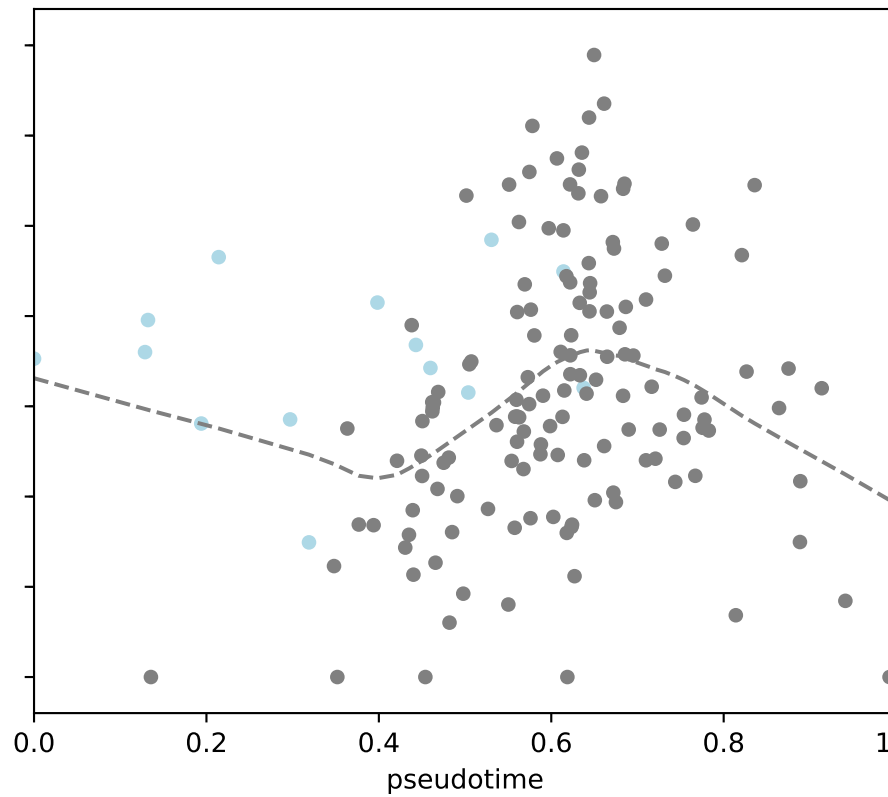
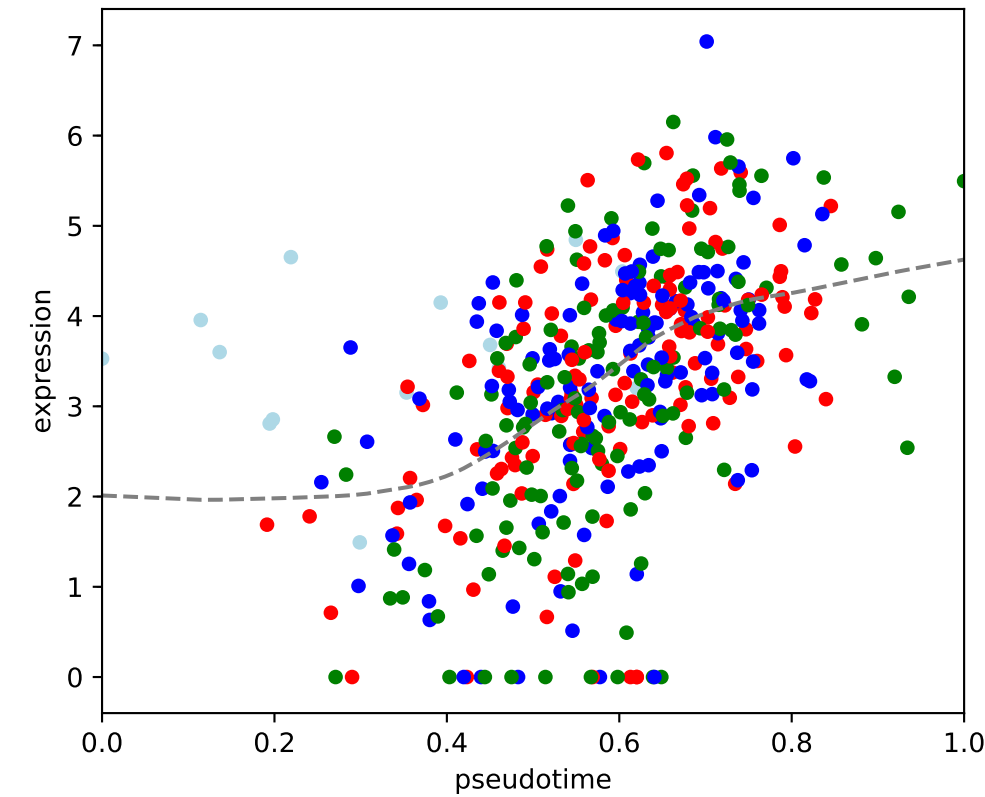
733 0spearman=-0.54

733 1spearman=-0.47



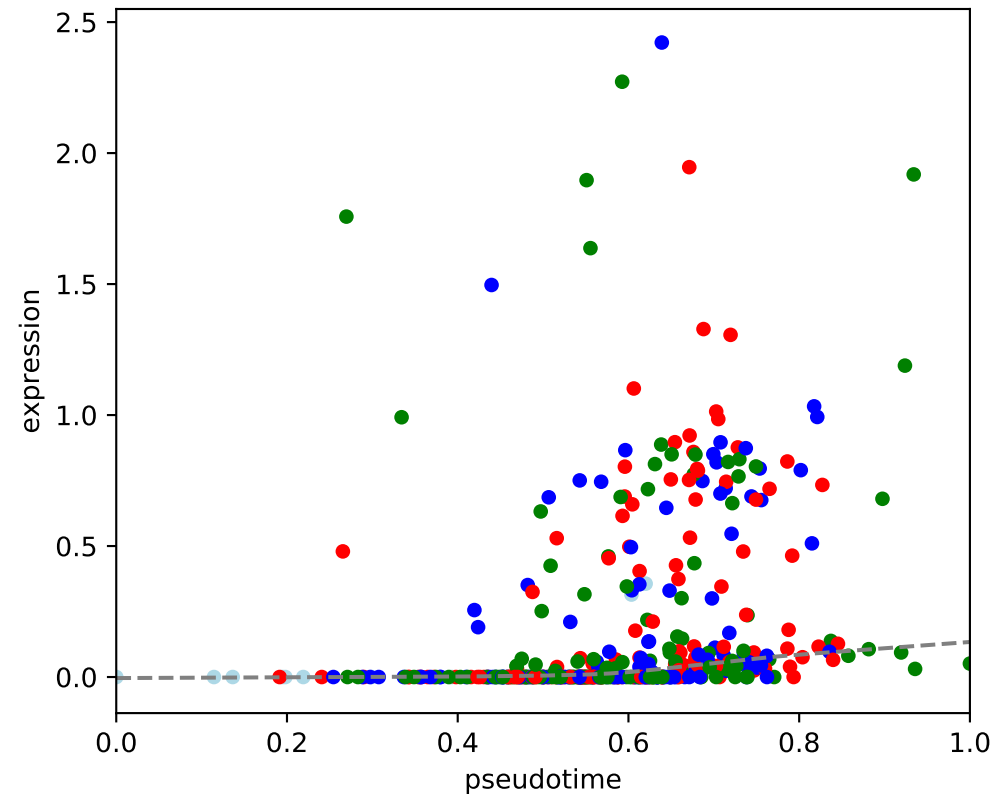
733 0spearman=0.54

733 1spearman=0.22

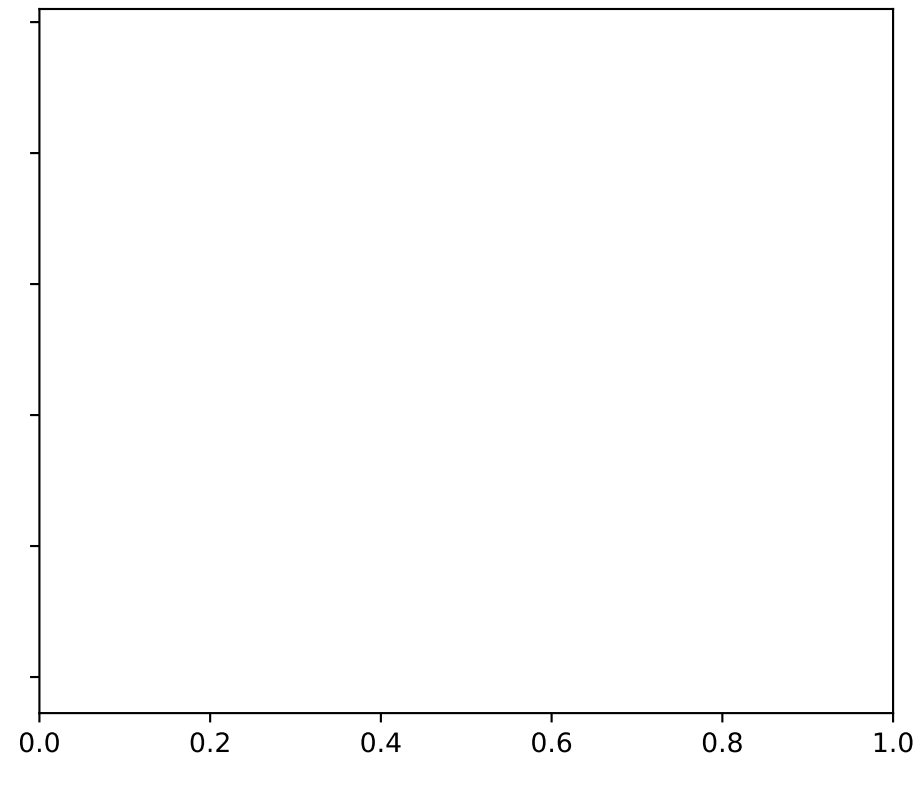
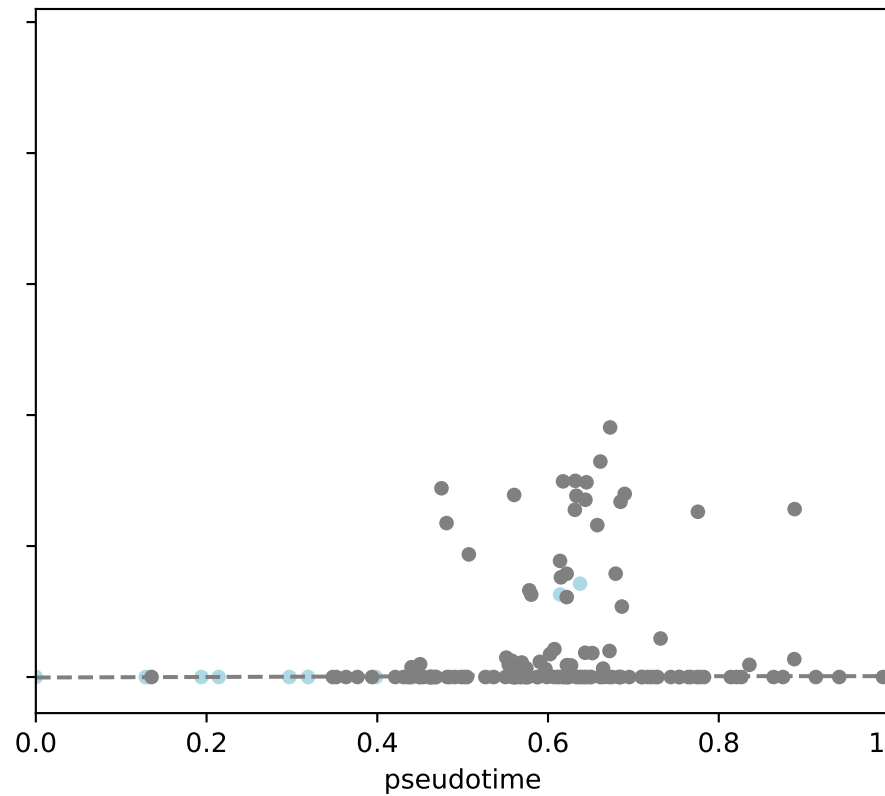


ENST00000618779 SAMD11 (sterile alpha motif domain containing 11)

733 0spearman=0.54

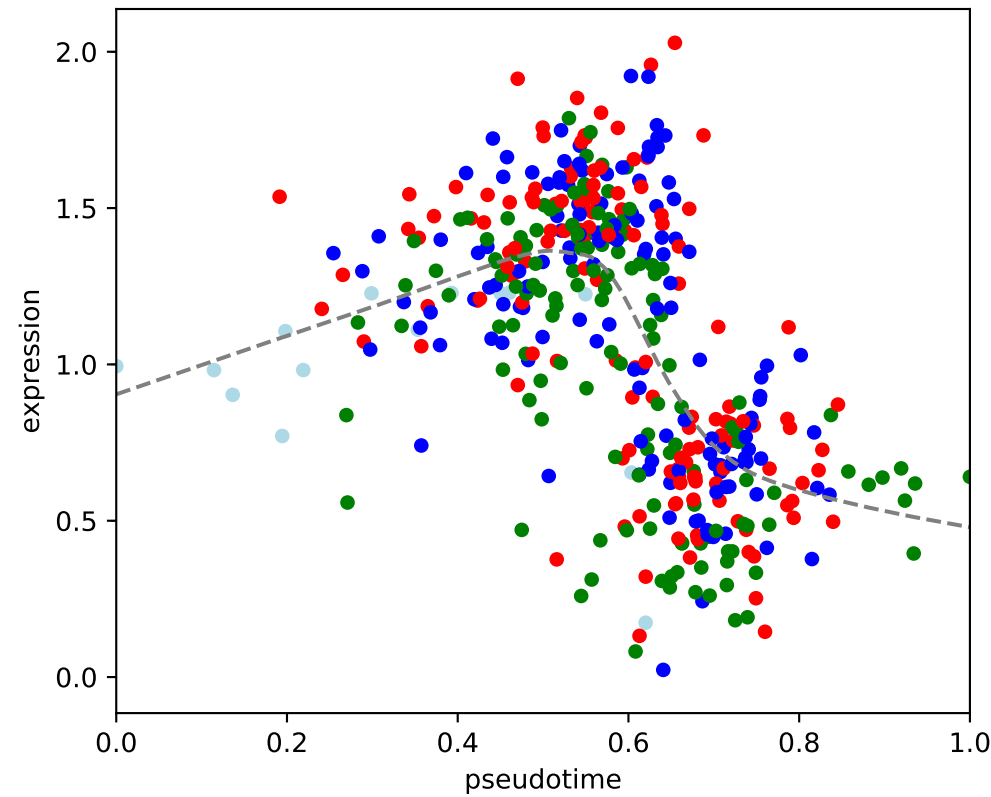


733 1spearman=0.18

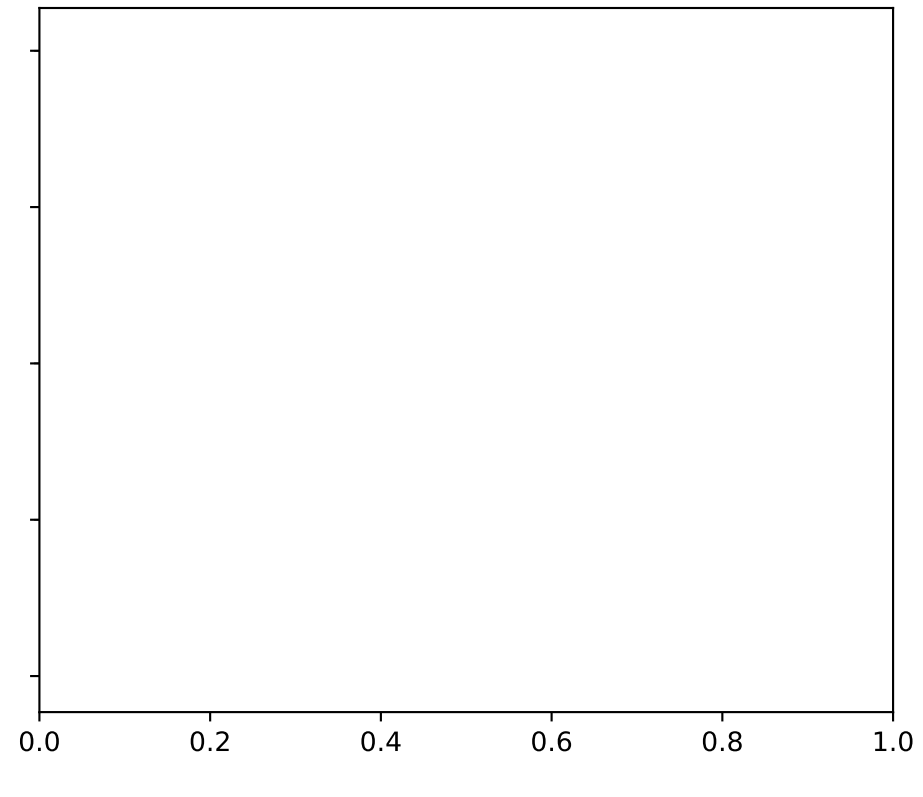
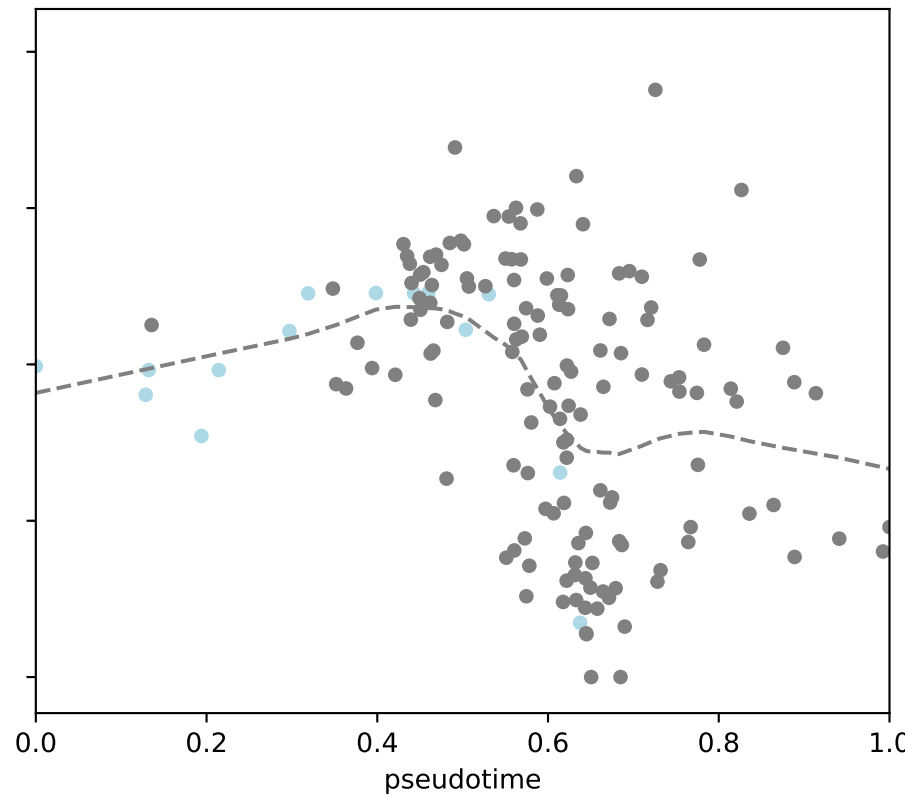


ENST00000554989 CKB (creatine kinase B)

733 0spearman=-0.54

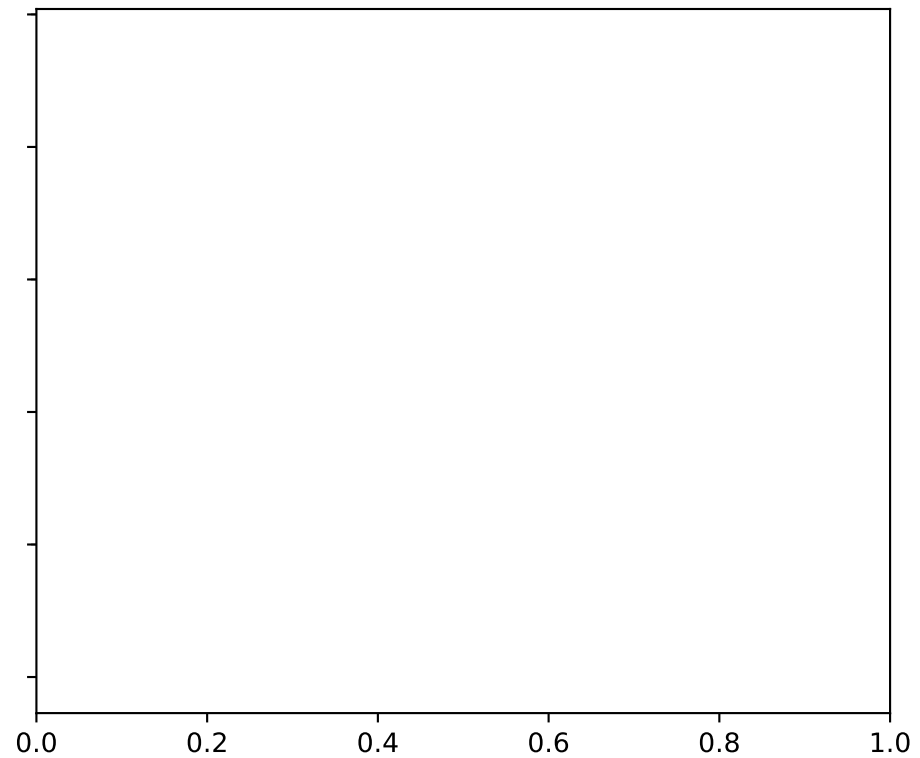
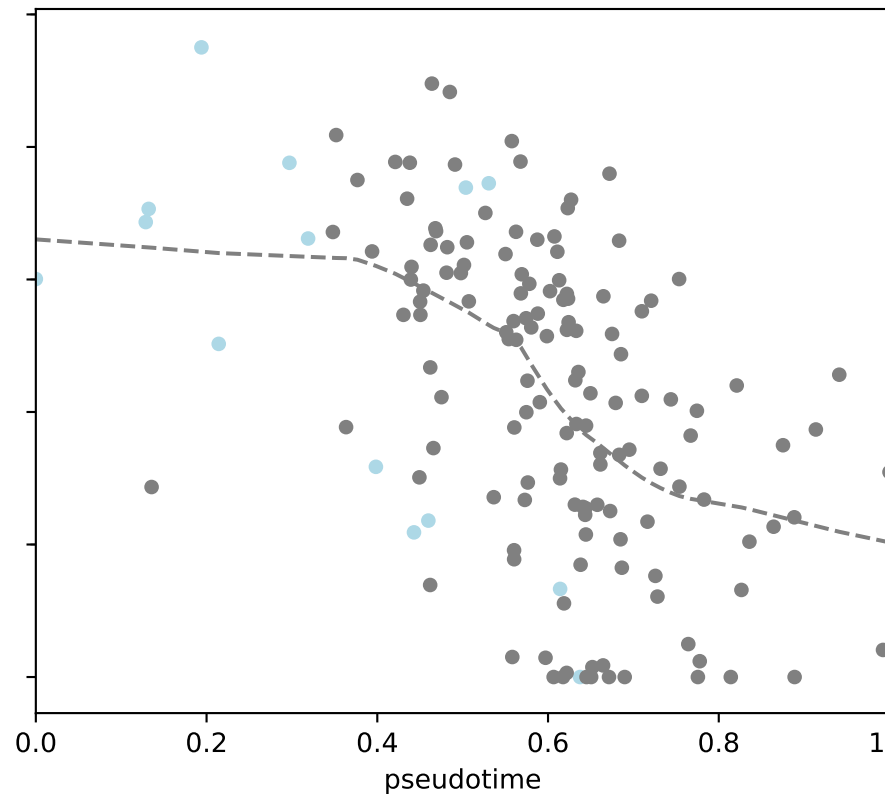
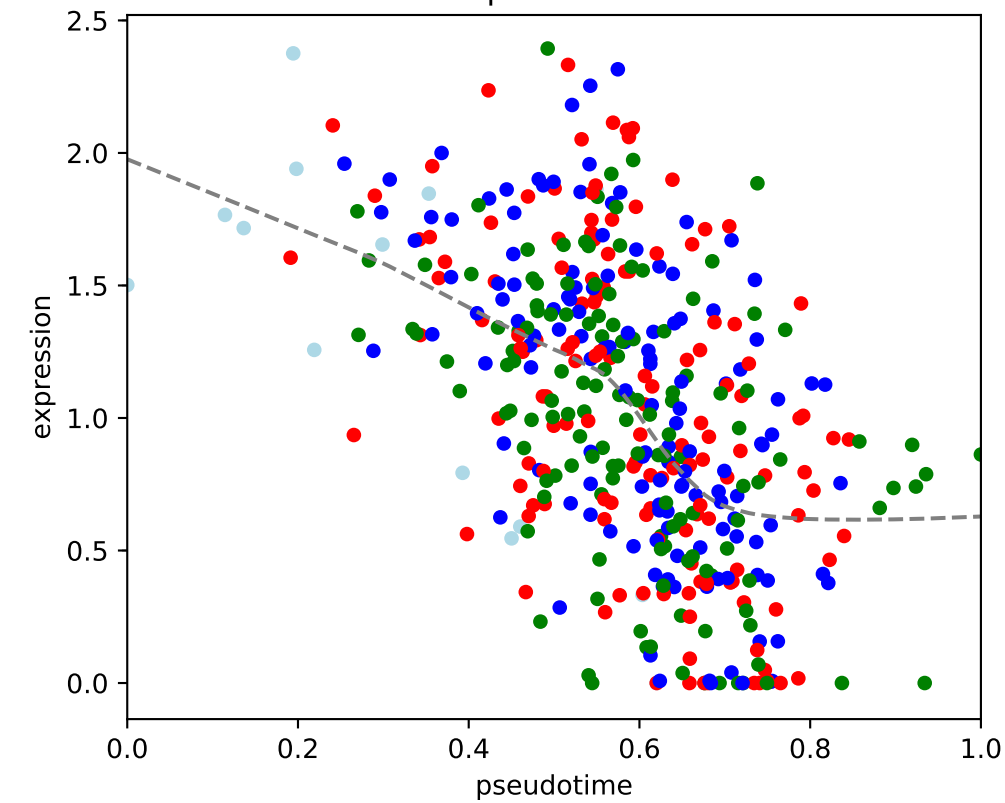


733 1spearman=-0.42



733 0spearman=-0.54

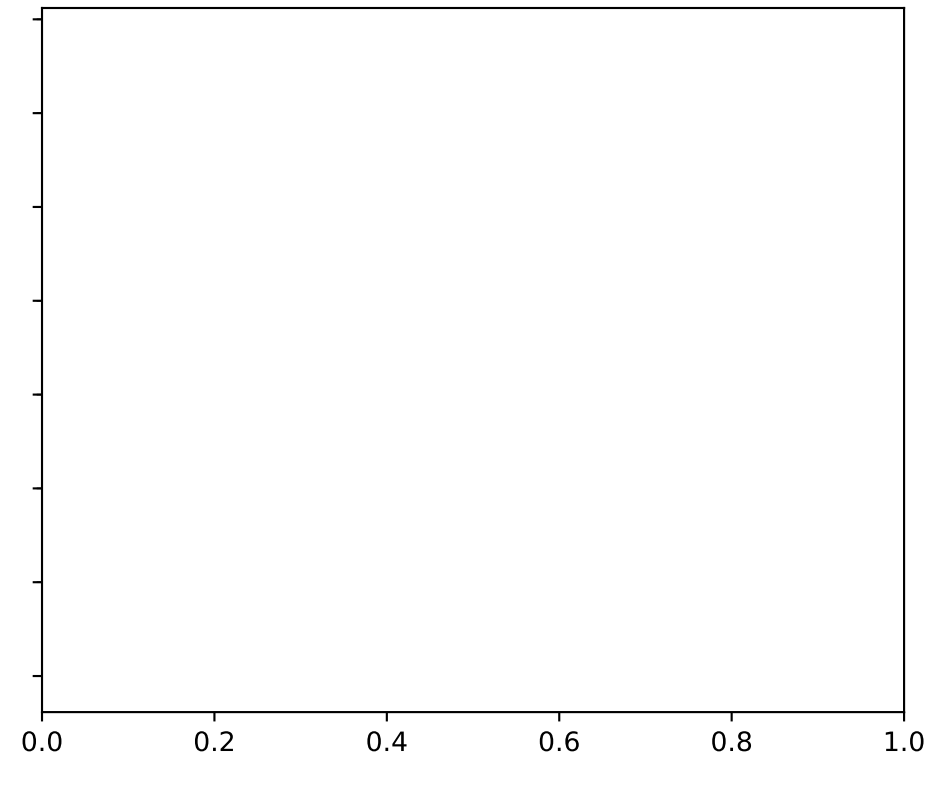
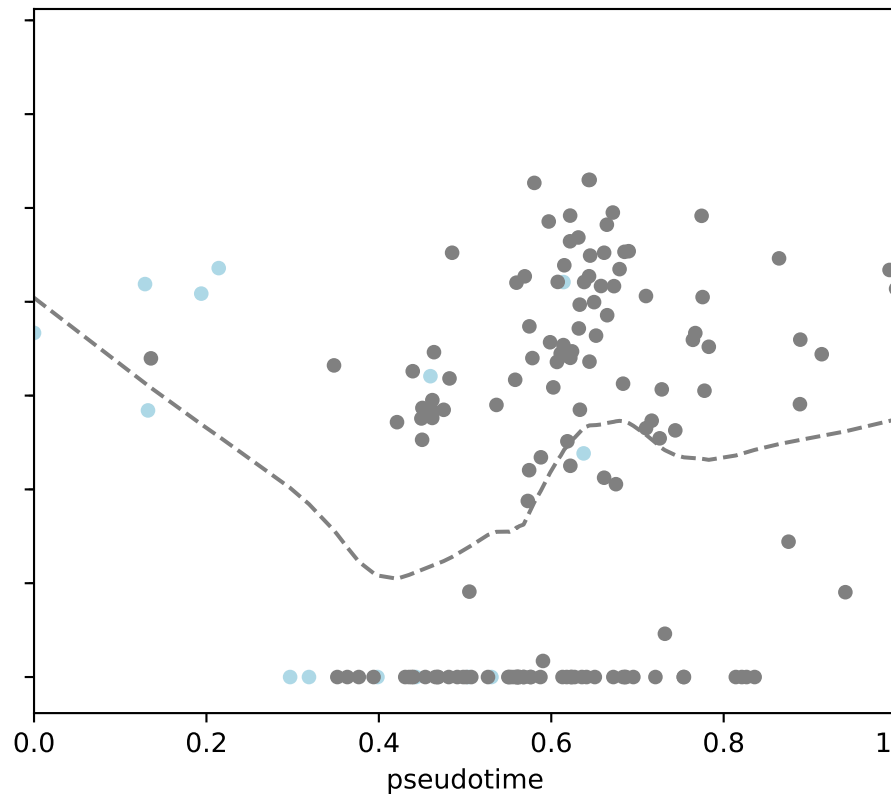
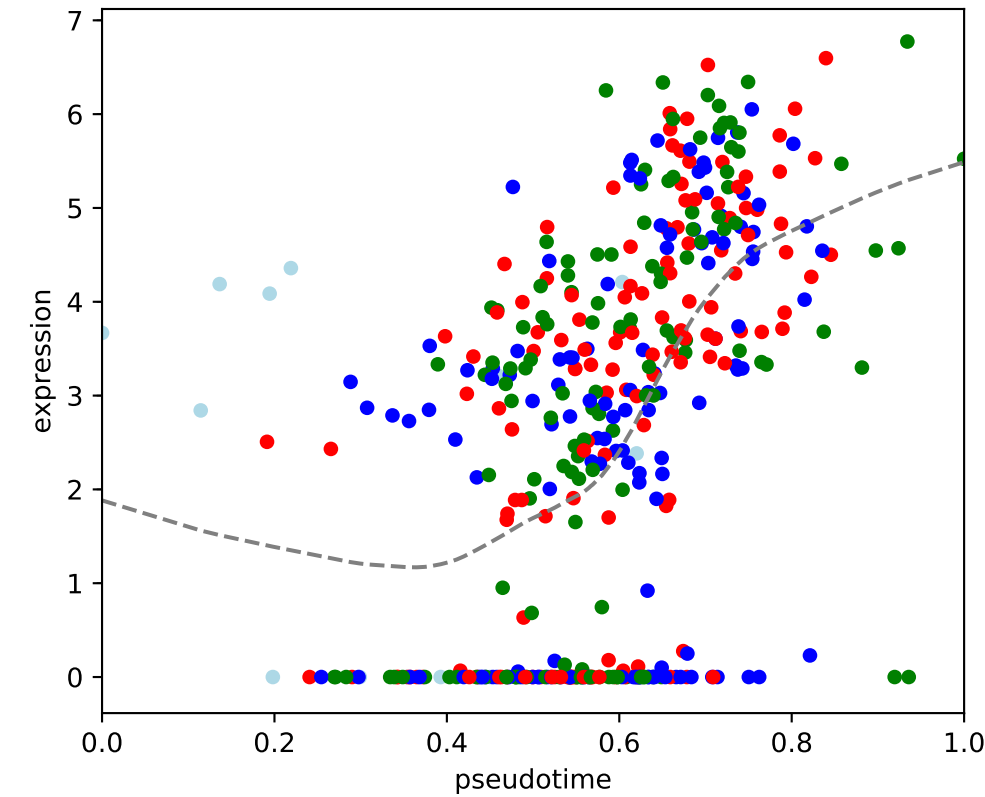
733 1spearman=-0.54



ENST00000374644 EPHA8 (EPH receptor A8)

733 0spearman=0.54

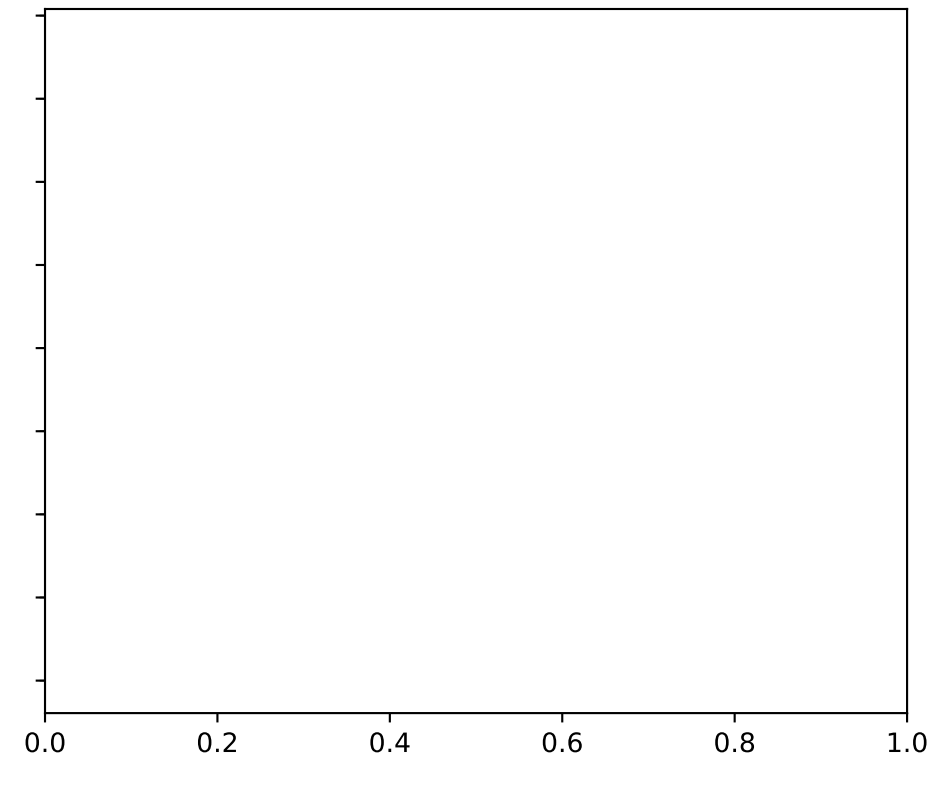
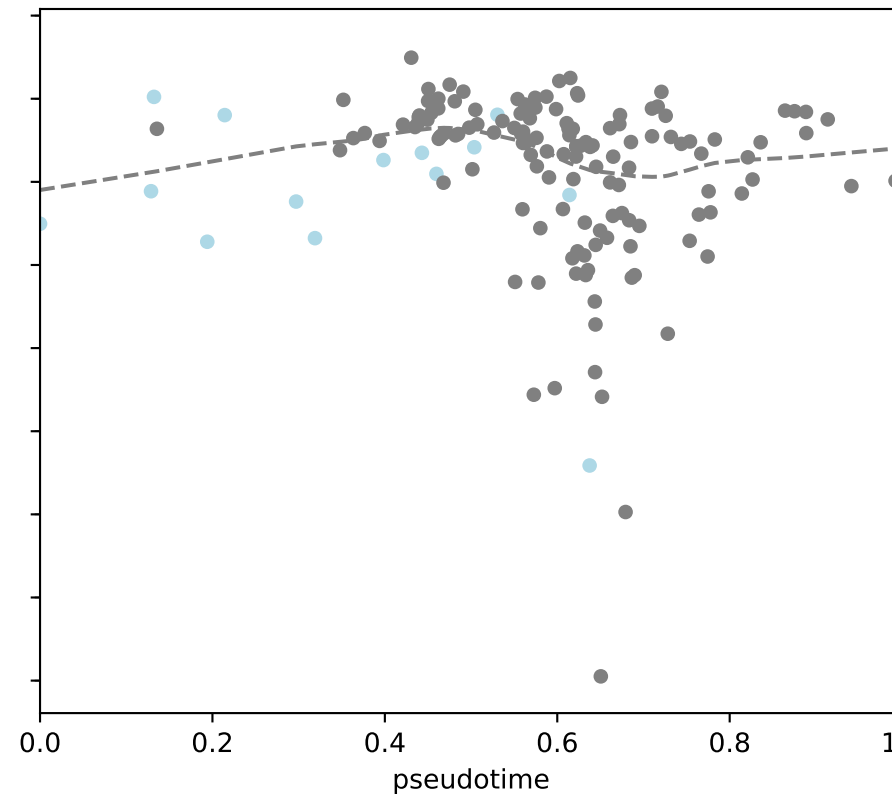
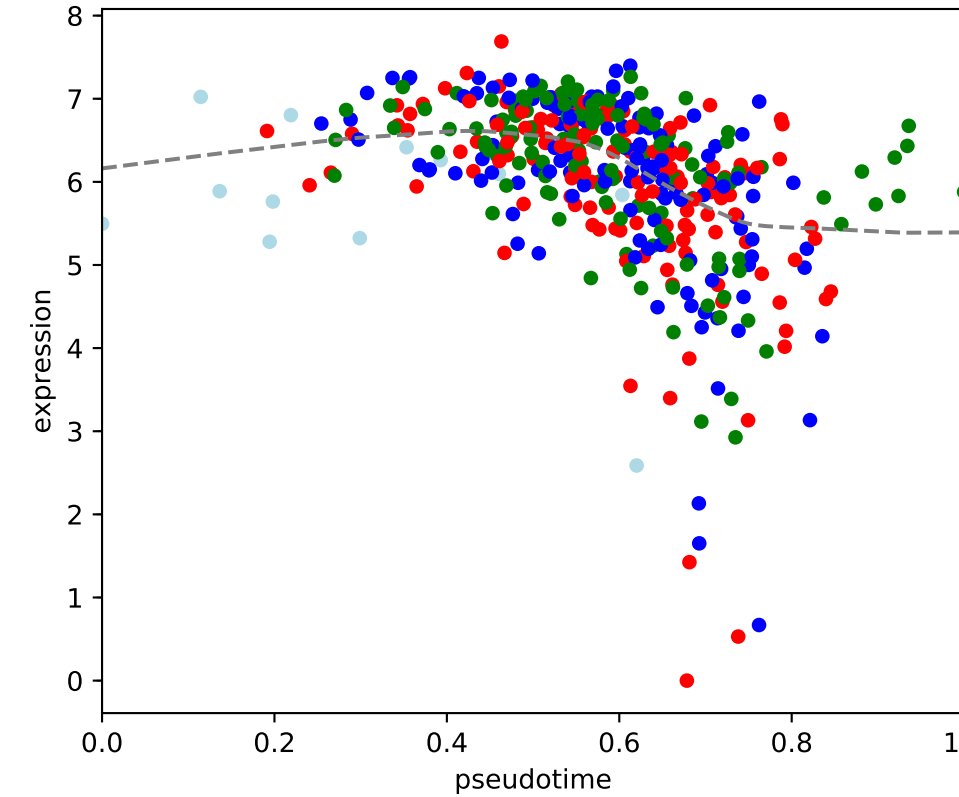
733 1spearman=0.25



ENST00000299166 NDUFB8 (NADH:ubiquinone oxidoreductase subunit B8)

733 0spearman=-0.54

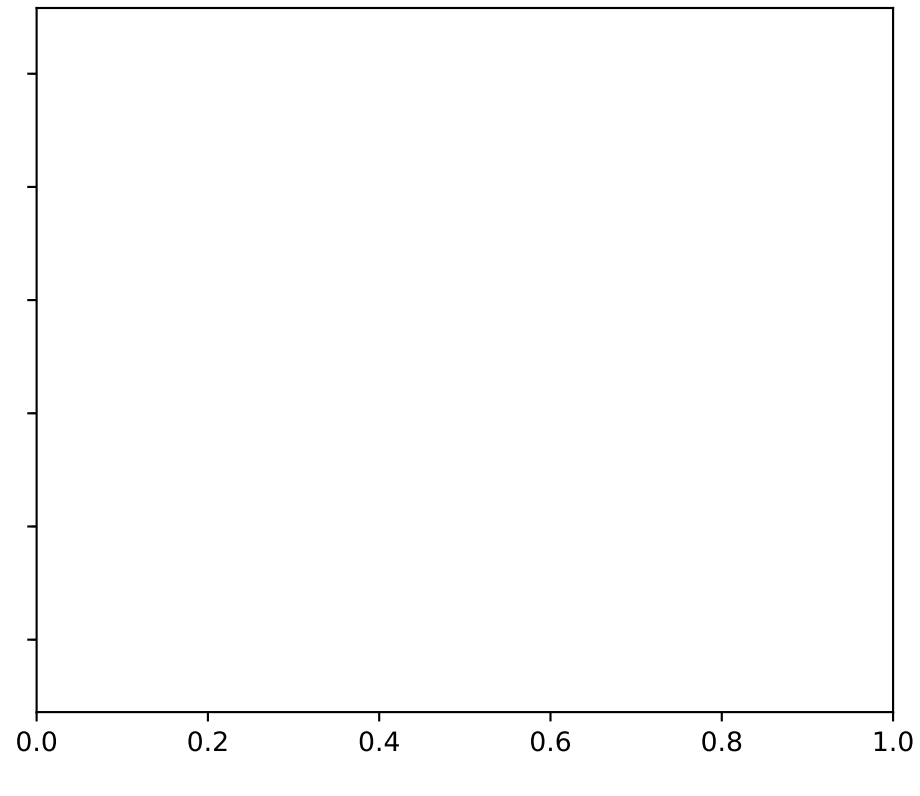
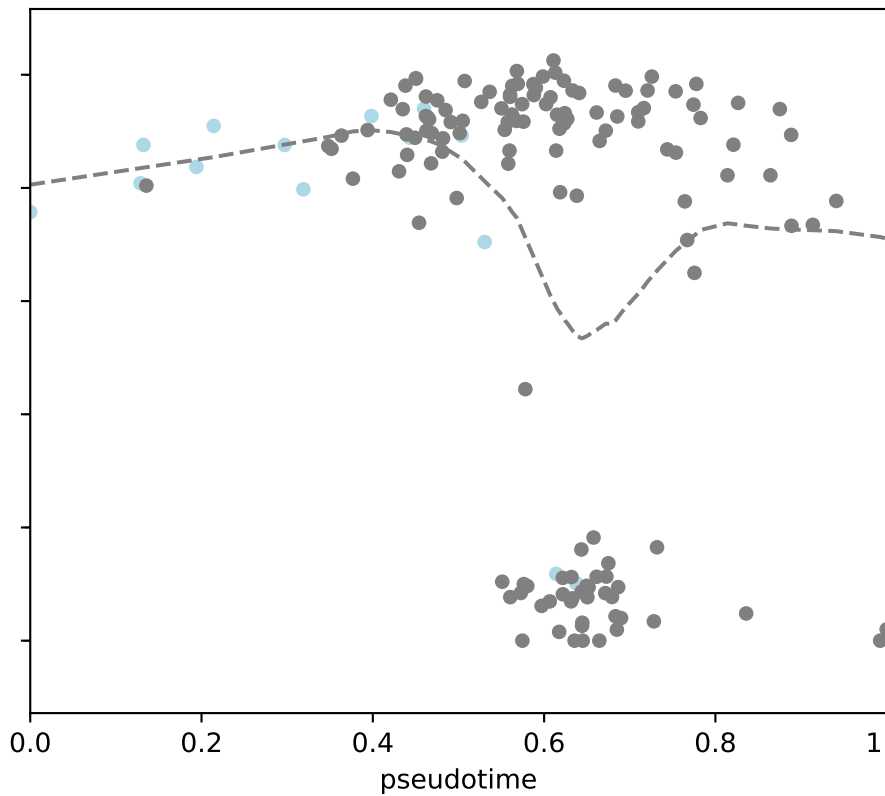
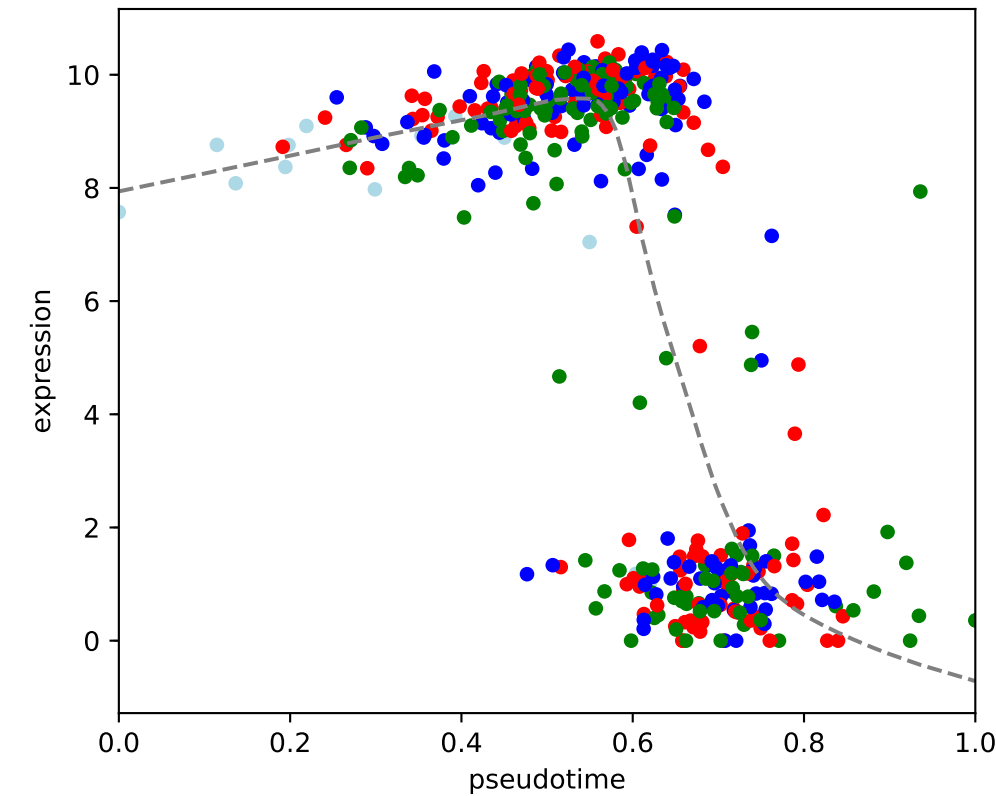
733 1spearman=-0.31



ENST00000266395 PDE6H (phosphodiesterase 6H)

733 0spearman=-0.54

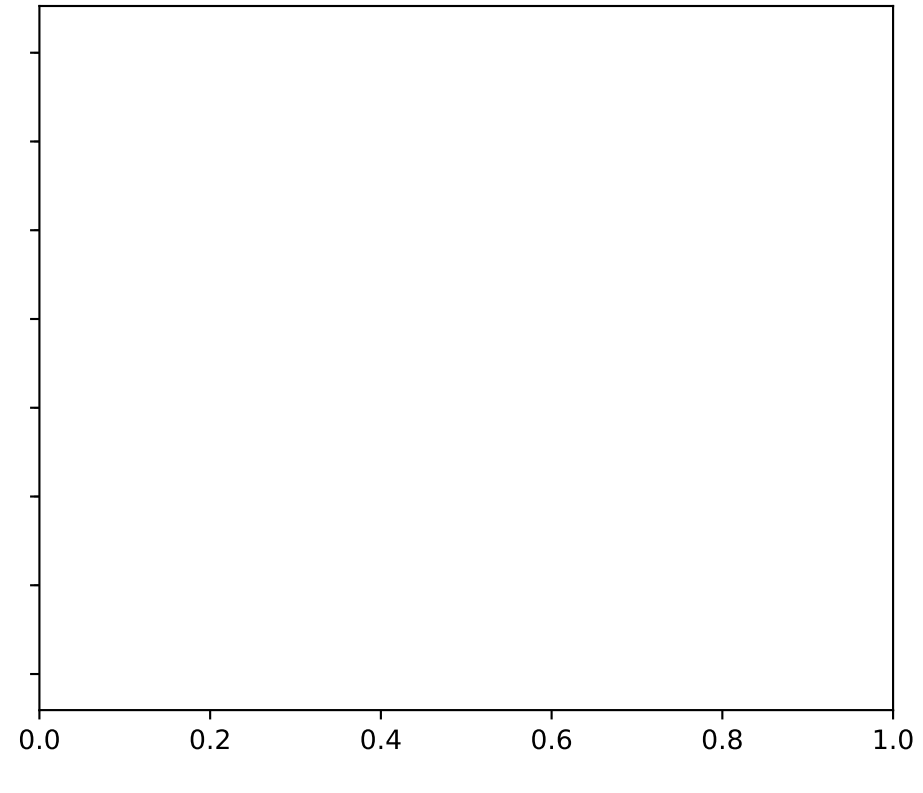
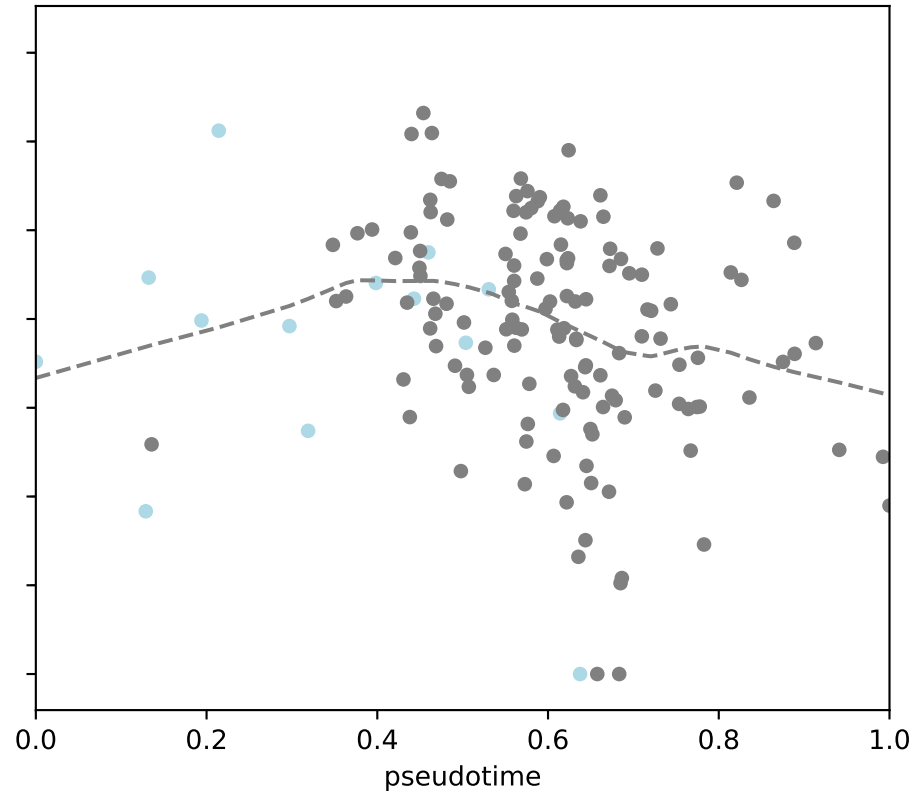
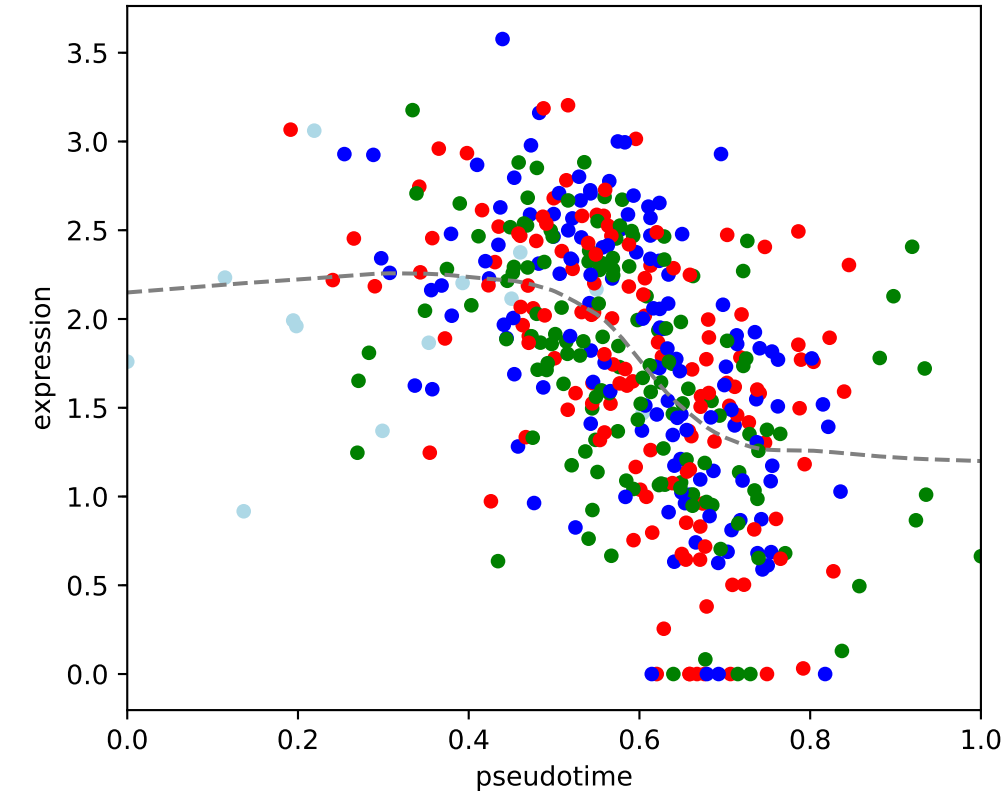
733 1spearman=-0.22



ENST00000532004 RPLP2 (ribosomal protein lateral stalk subunit P2)

733 0spearman=-0.54

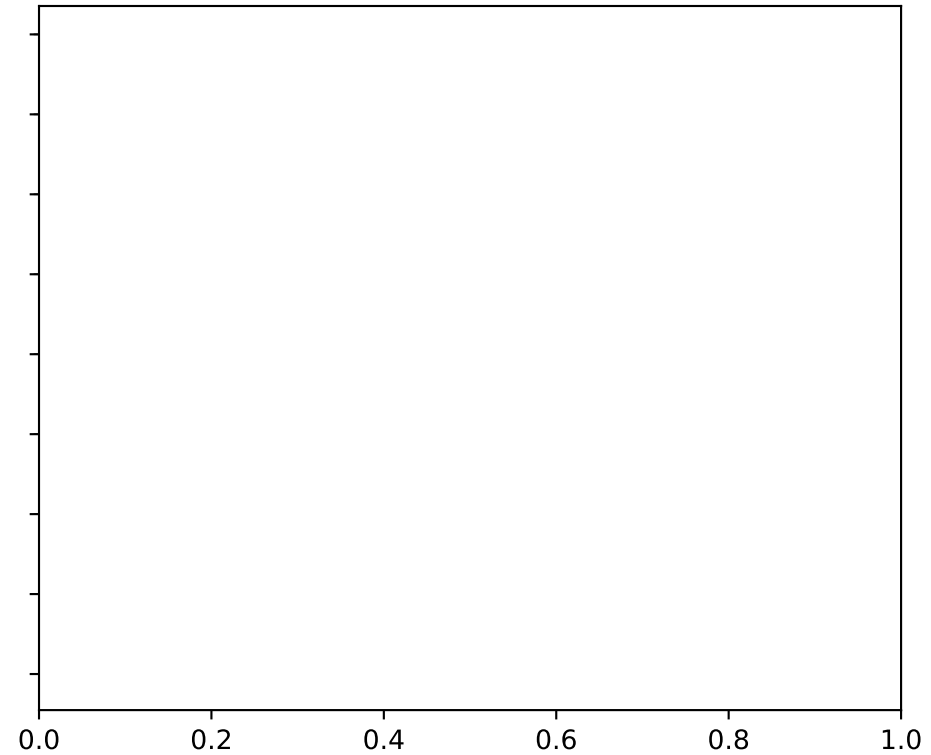
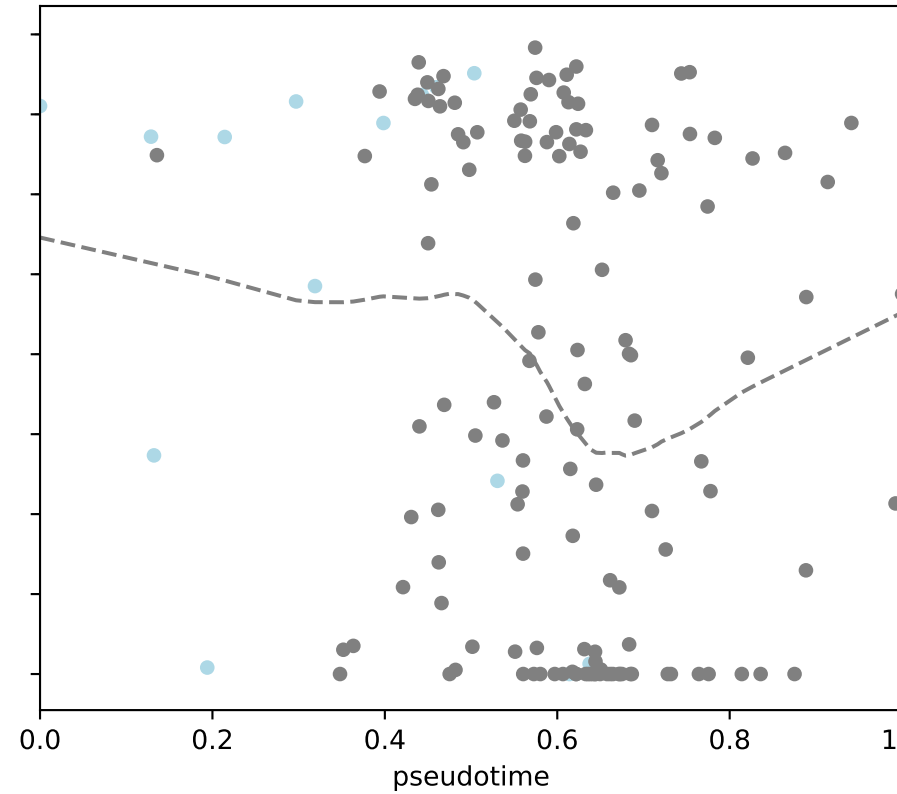
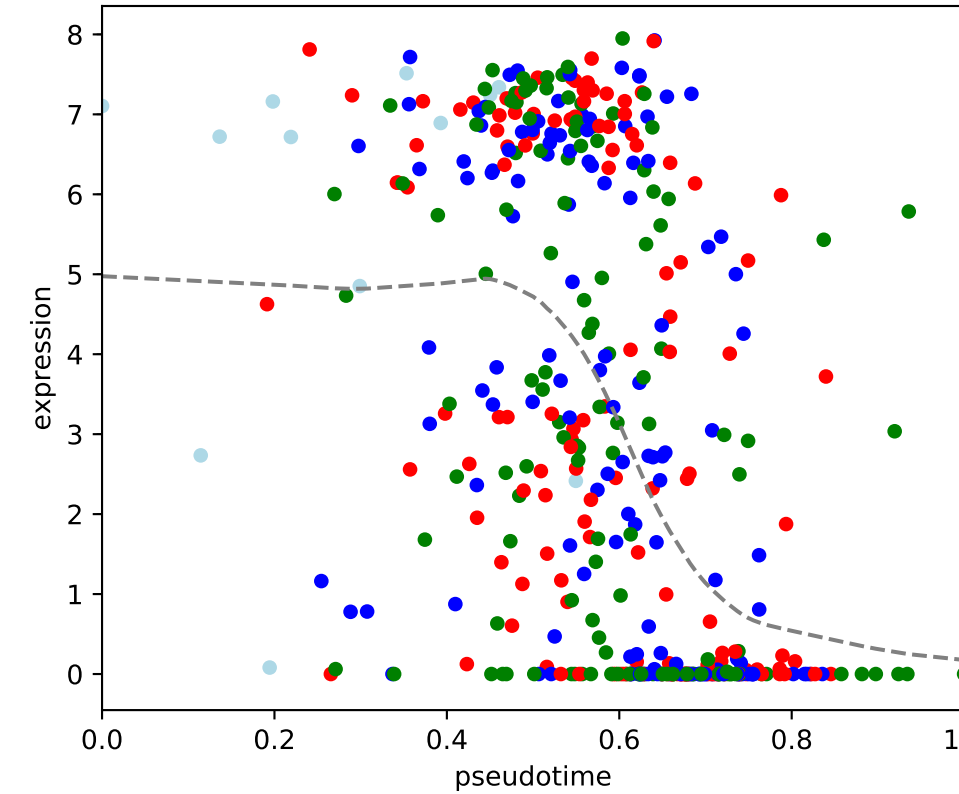
733 1spearman=-0.30



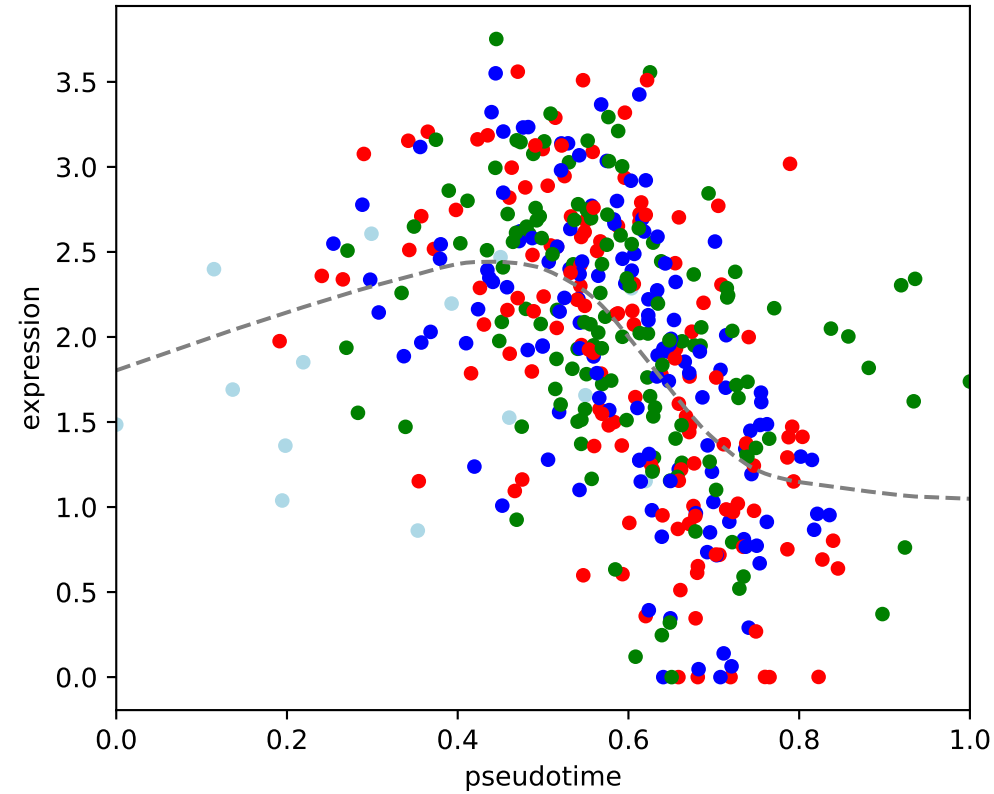
ENST00000619224 RXRG (retinoid X receptor gamma)

733 0spearman=-0.54

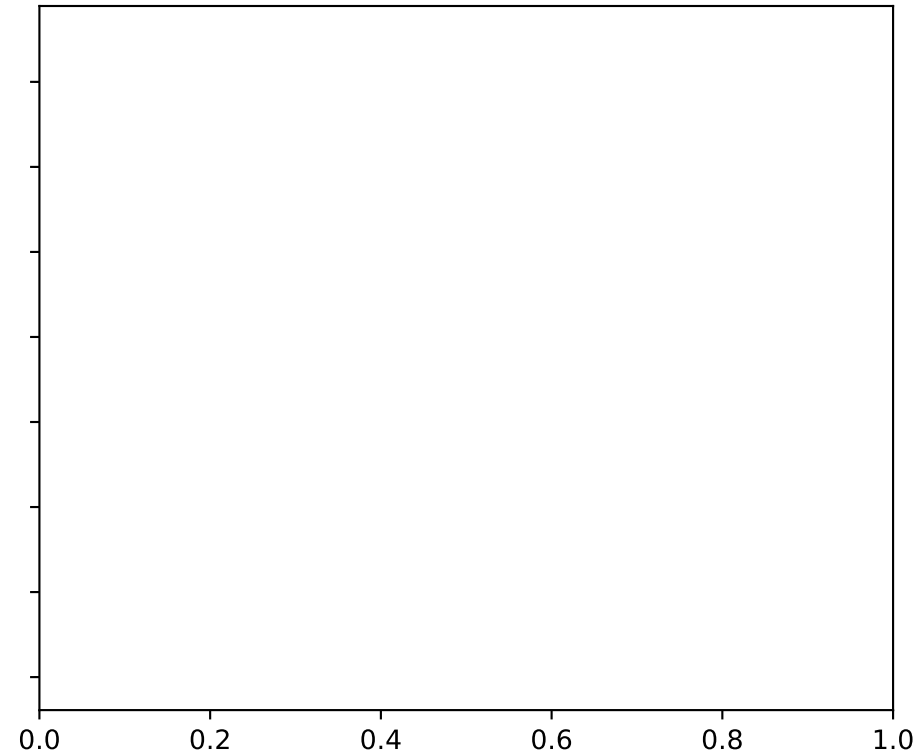
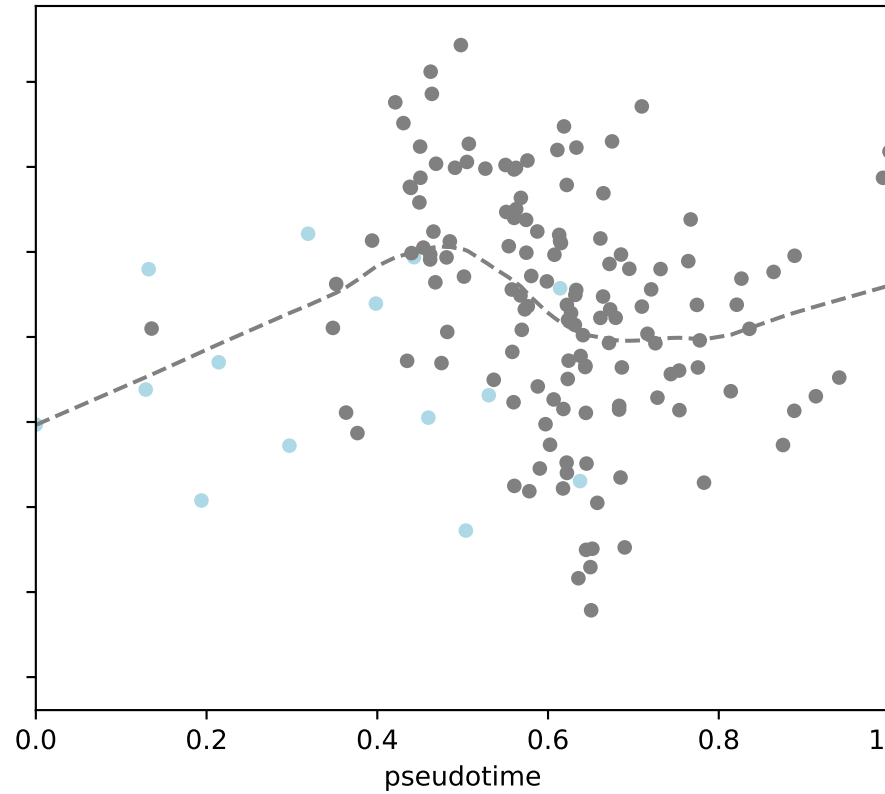
733 1spearman=-0.29



733 0spearman=-0.53



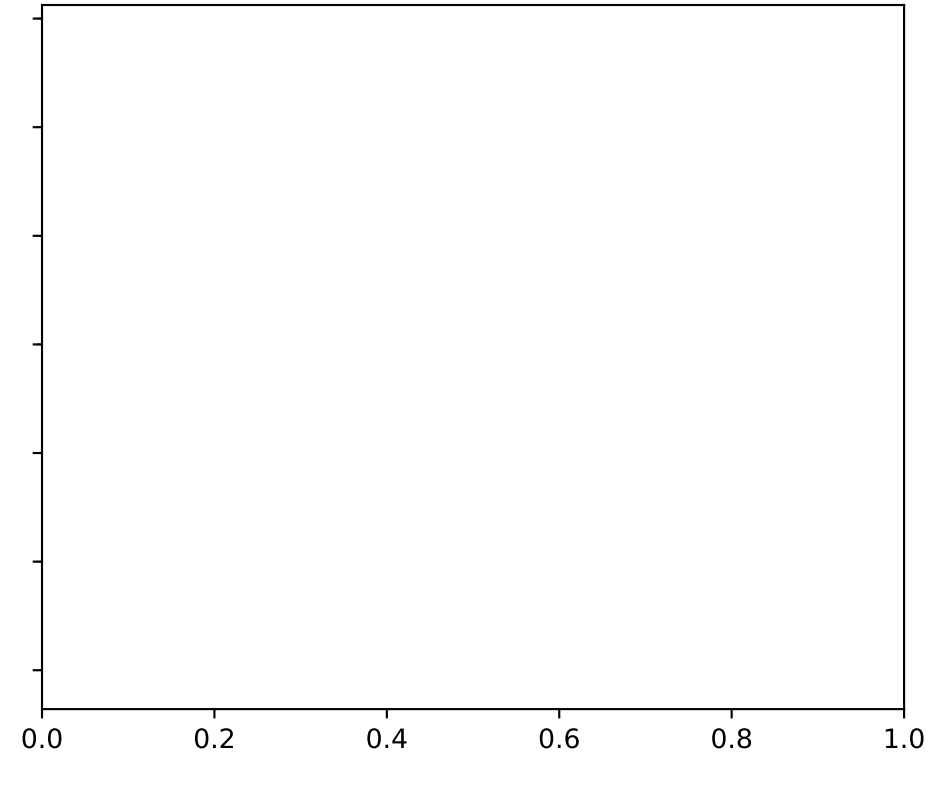
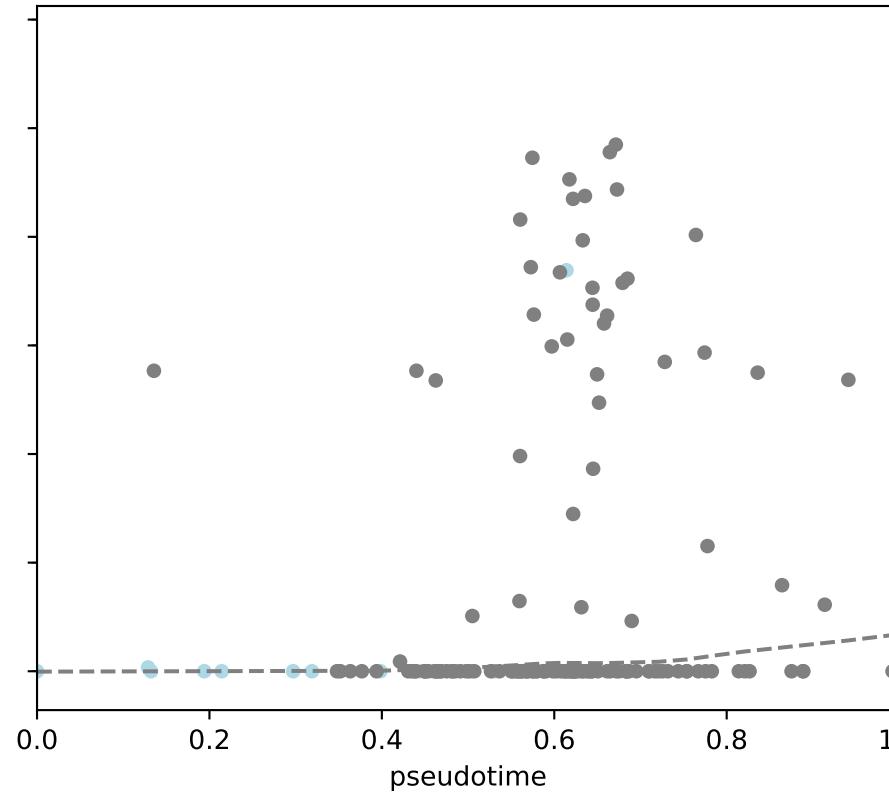
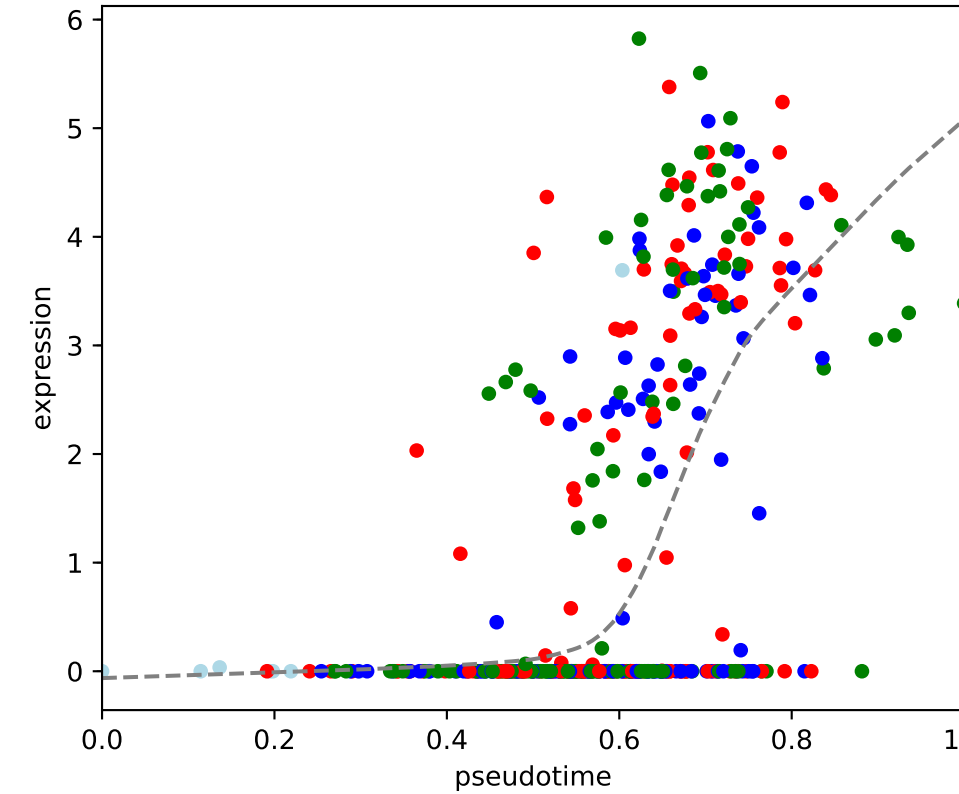
733 1spearman=-0.23



ENST00000394295 ANXA4 (annexin A4)

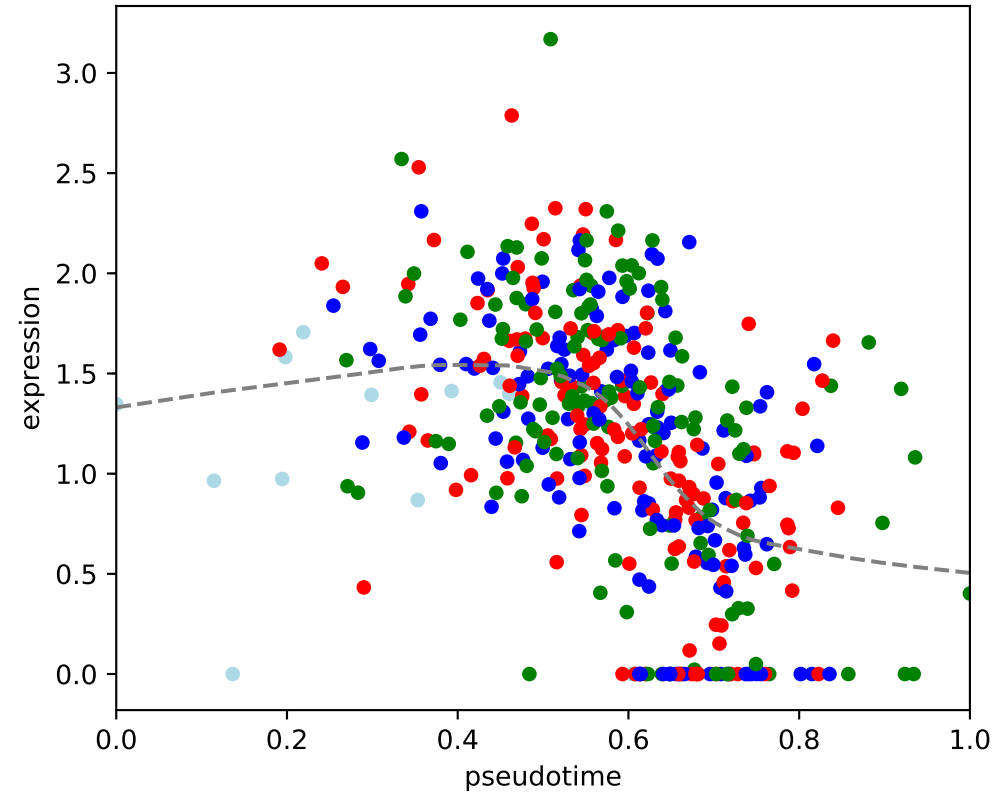
733 0spearman=0.53

733 1spearman=0.22

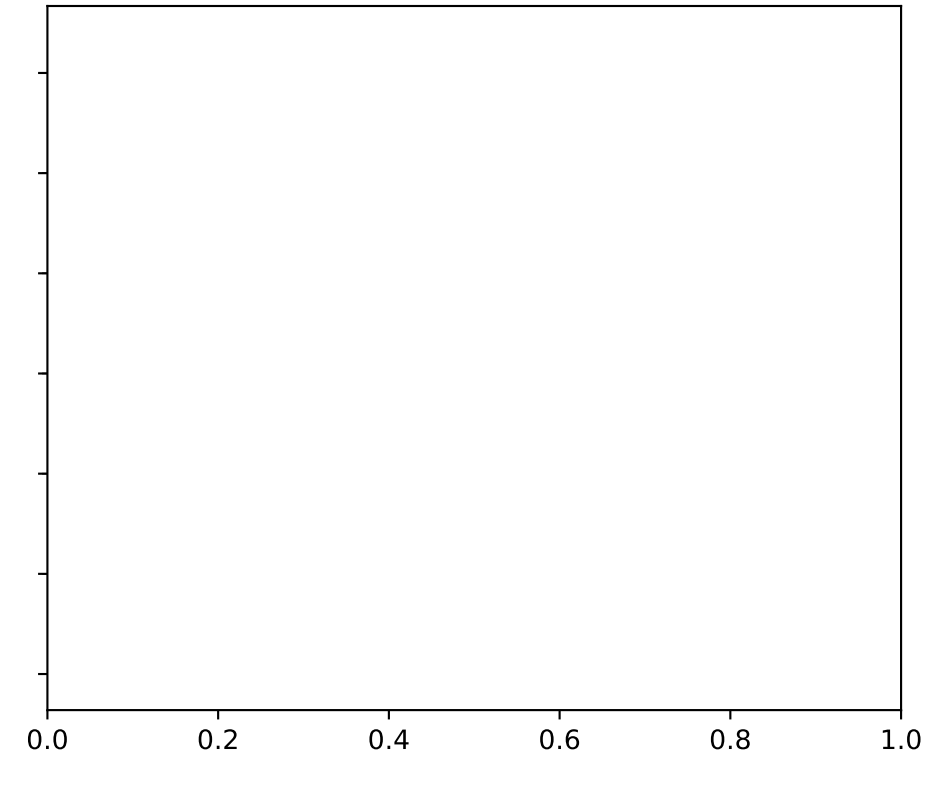
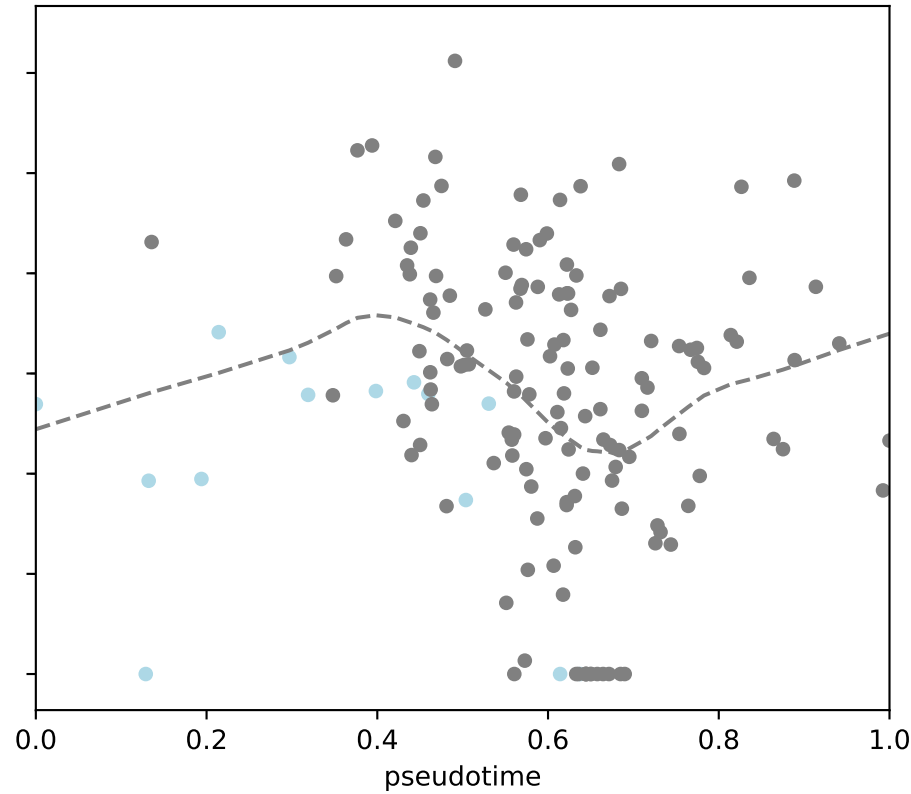


ENST00000597743 CALM3 (calmodulin 3)

733 0spearman=-0.53

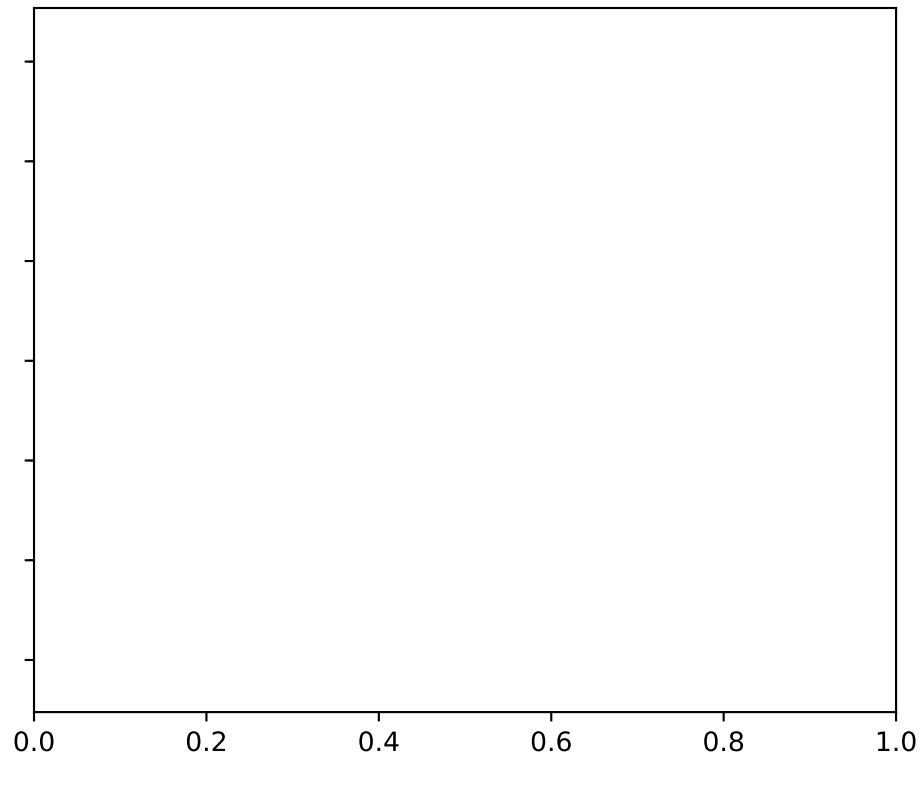
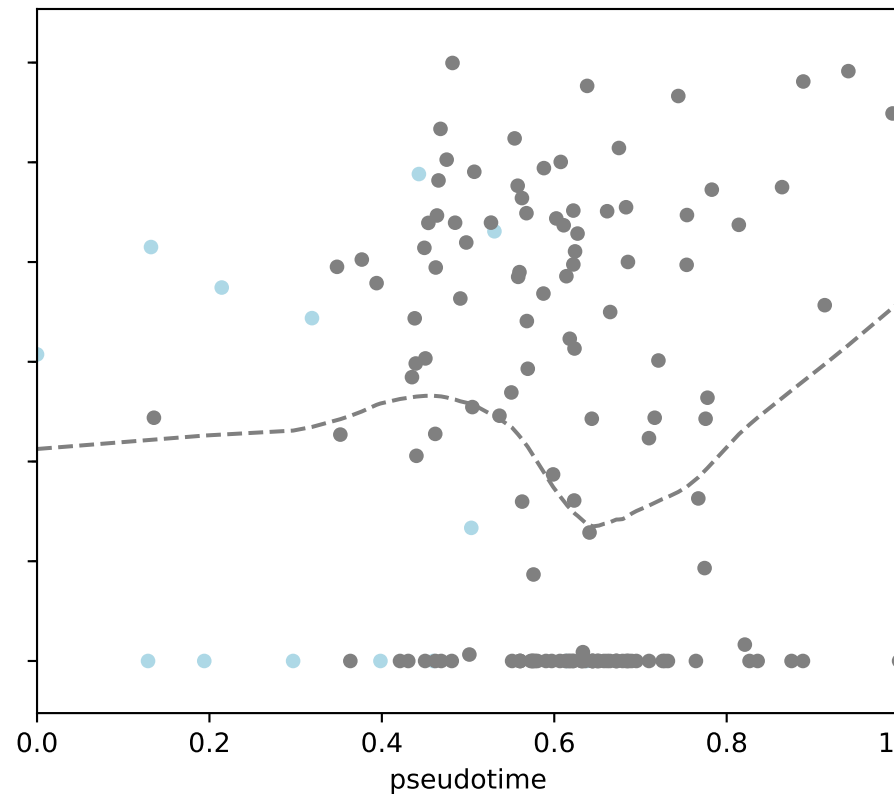
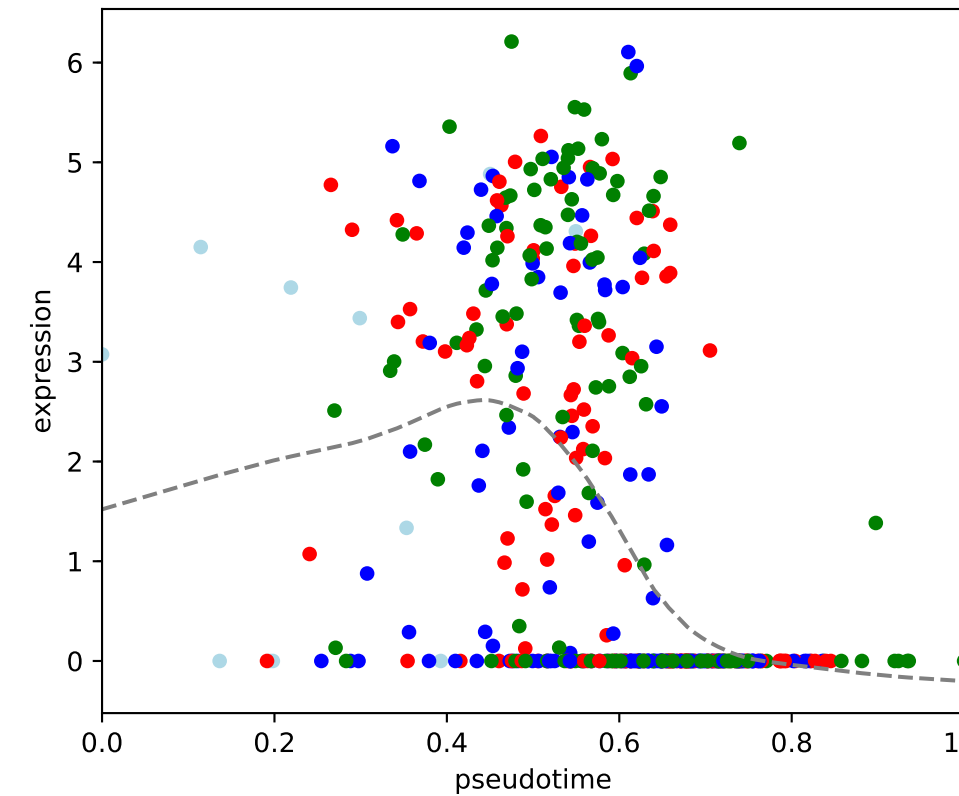


733 1spearman=-0.24



733 0spearman=-0.53

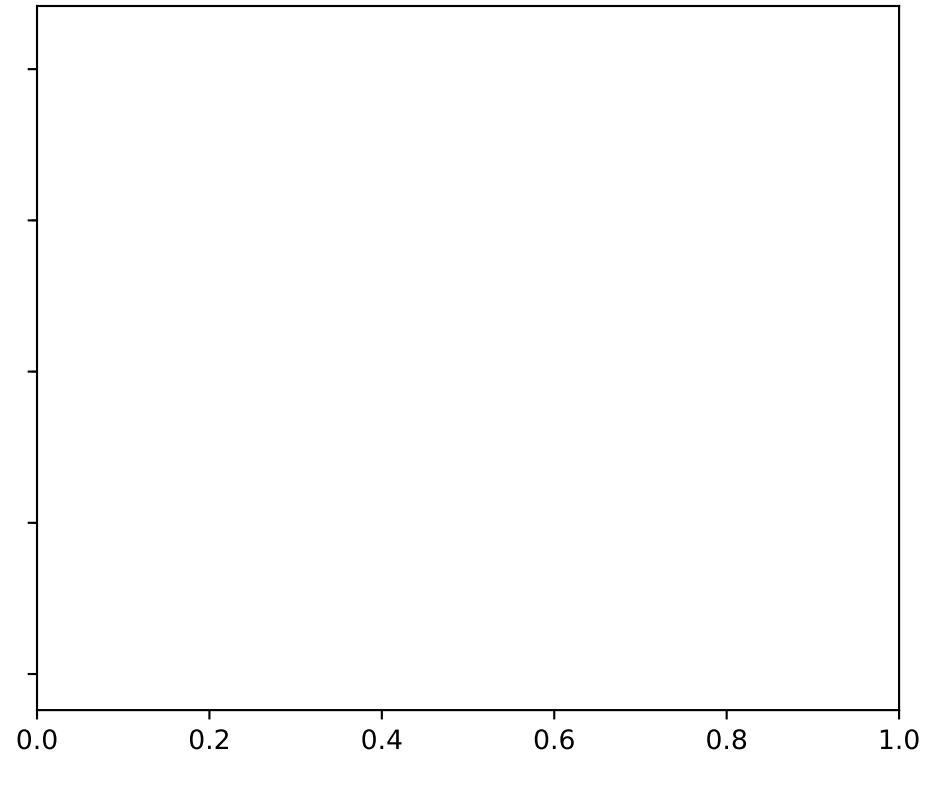
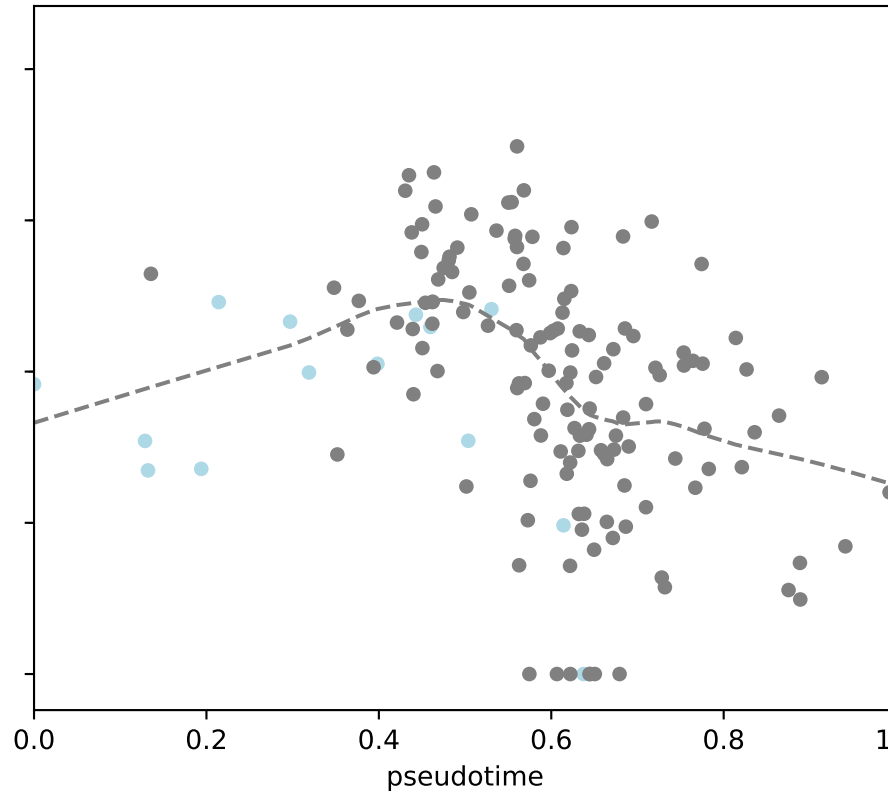
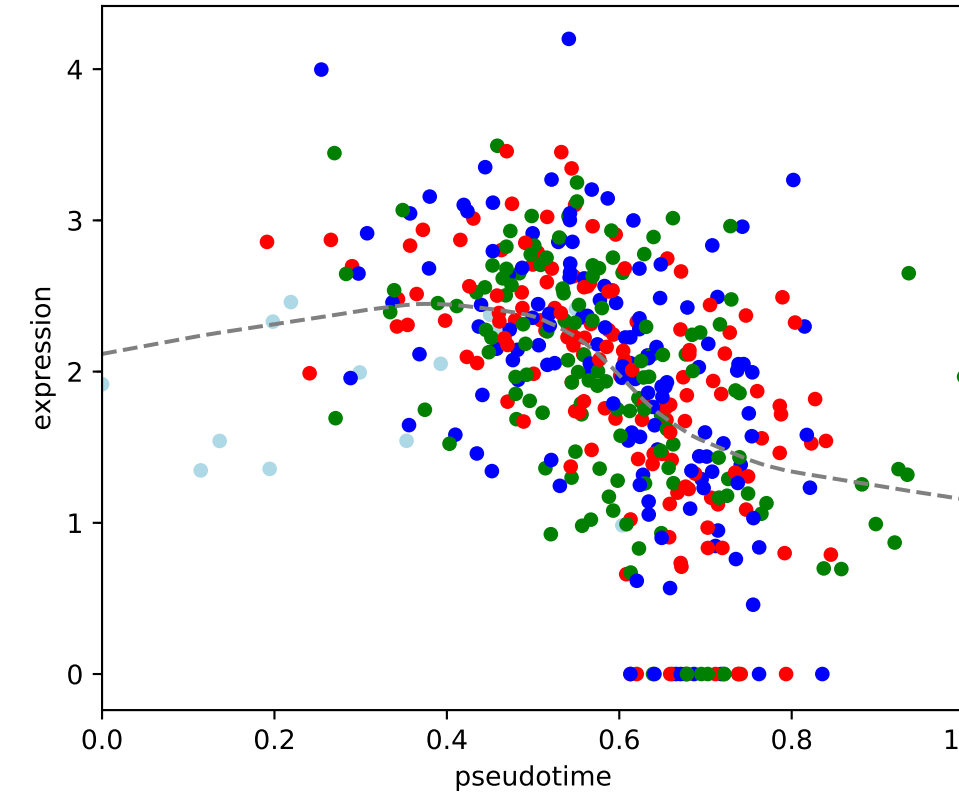
733 1spearman=-0.12



ENST00000515060 ATP5MC1 (ATP synthase membrane subunit c locus 1)

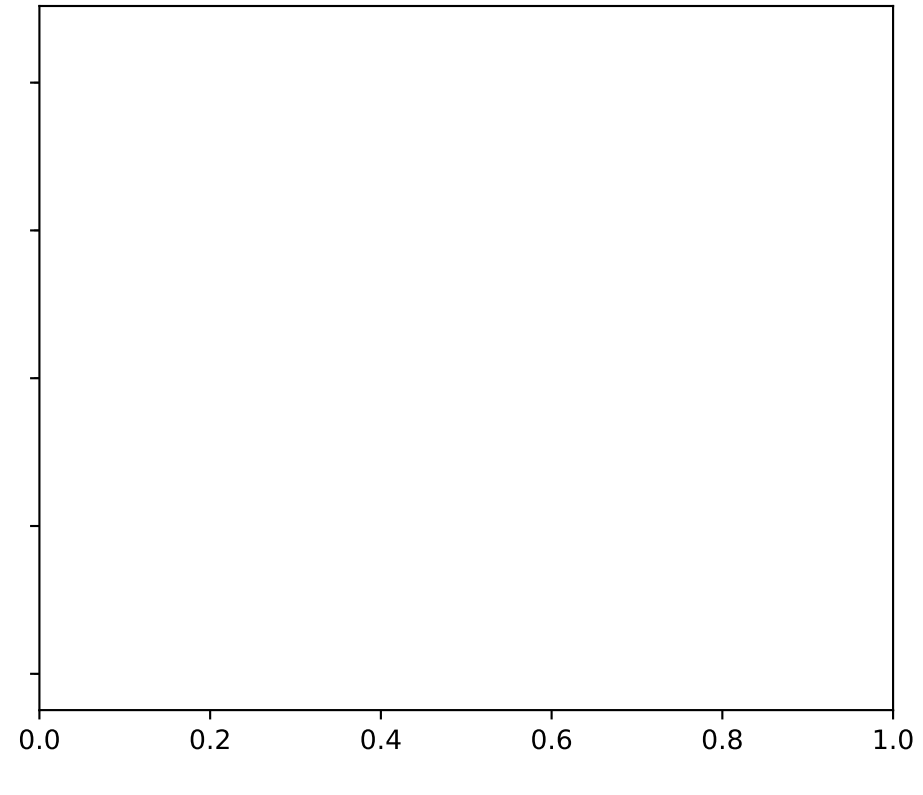
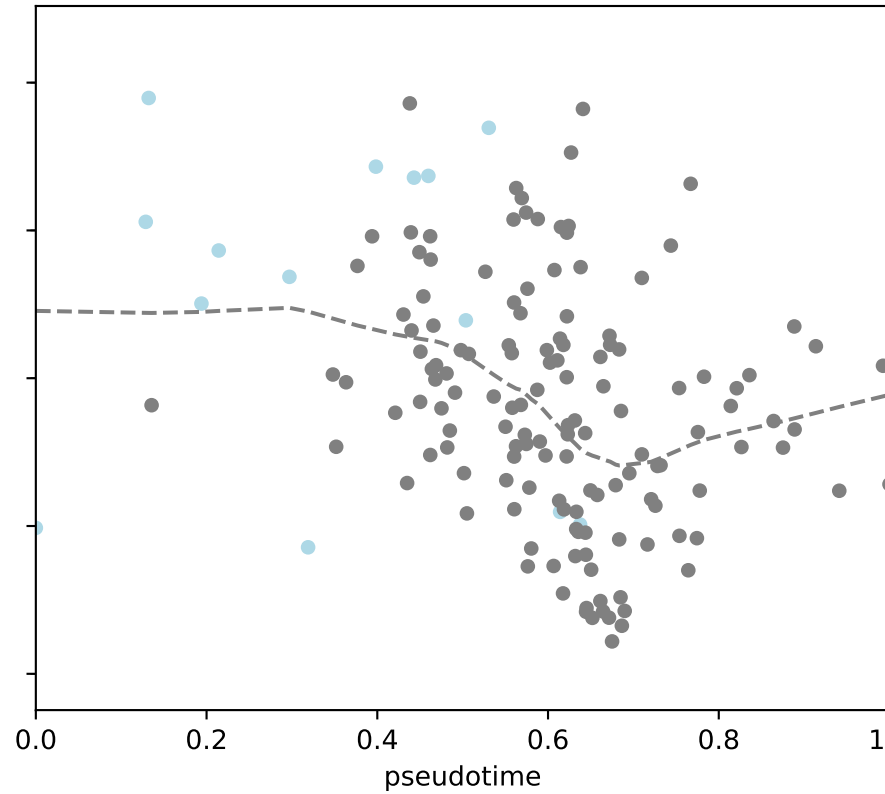
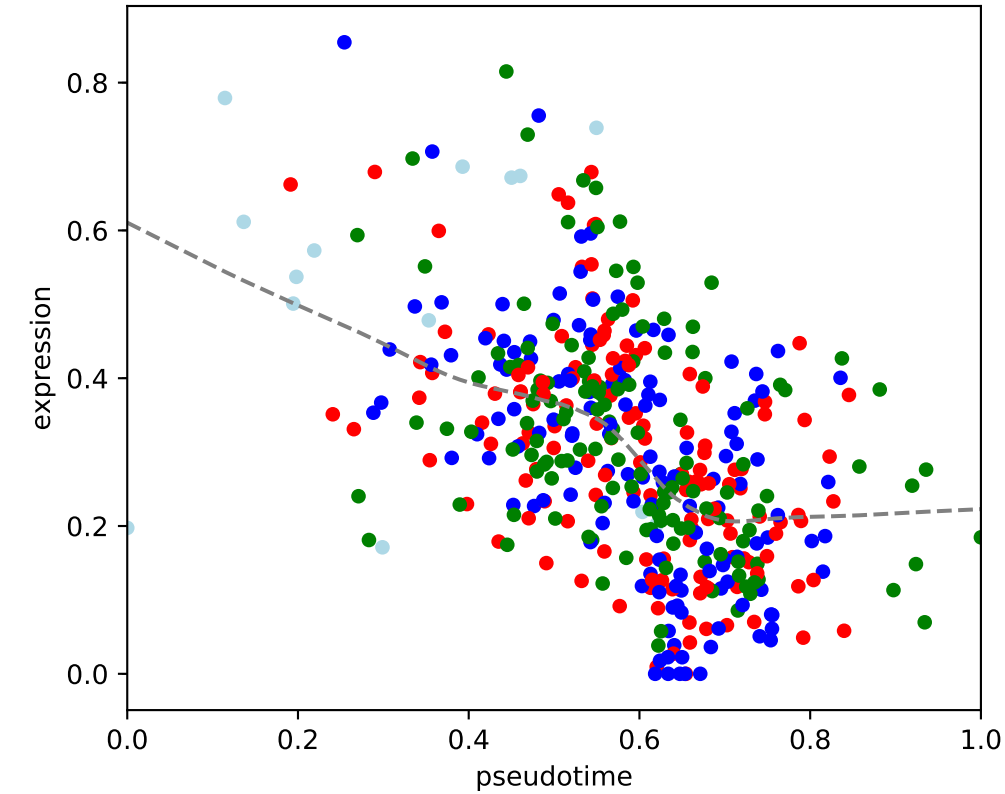
733 0spearman=-0.53

733 1spearman=-0.48



733 0spearman=-0.53

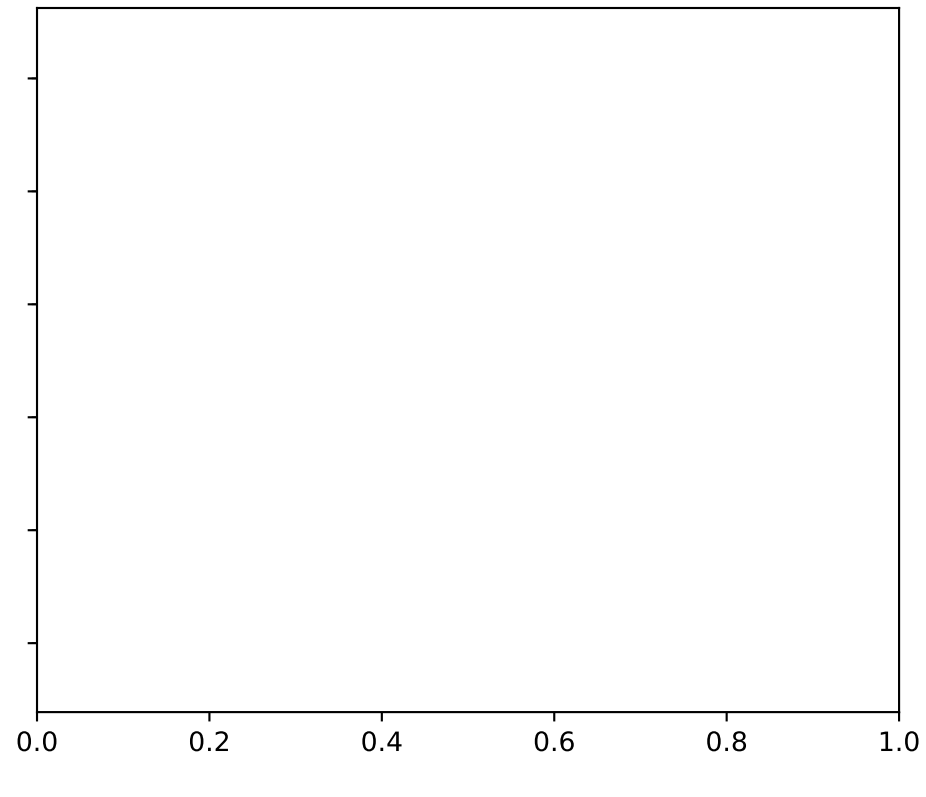
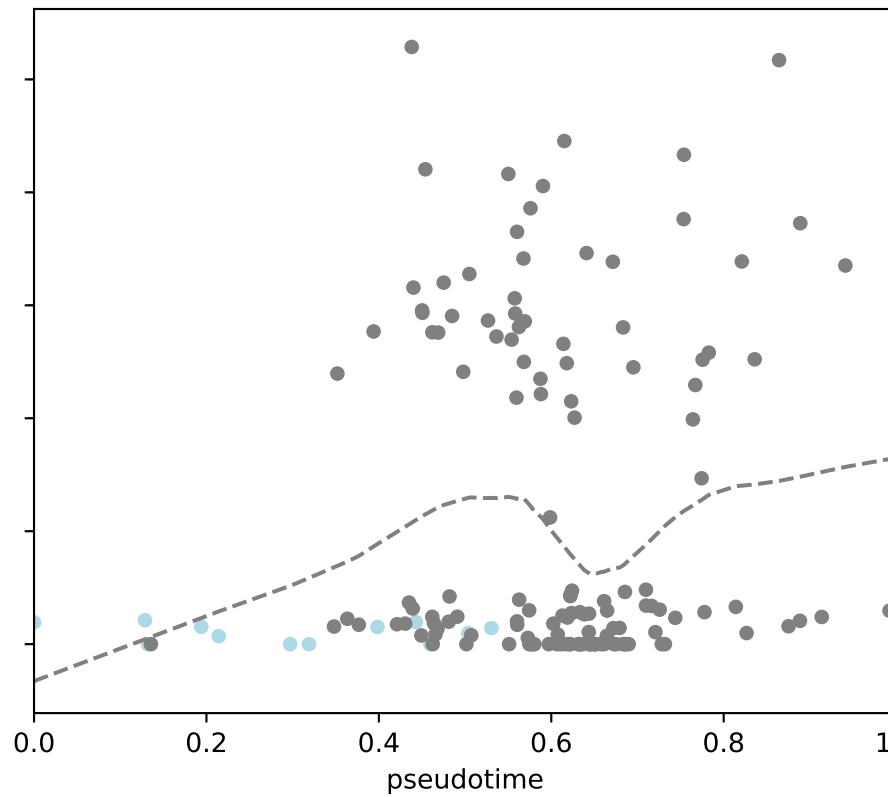
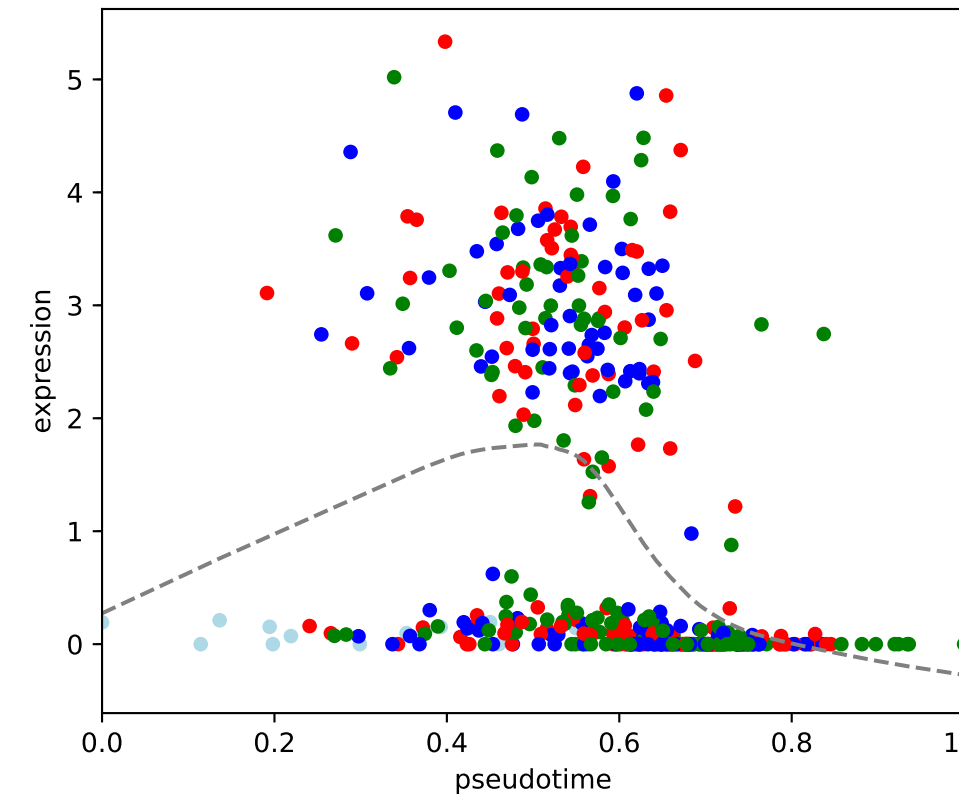
733 1spearman=-0.39



ENST00000371801 OLFM1 (olfactomedin 1)

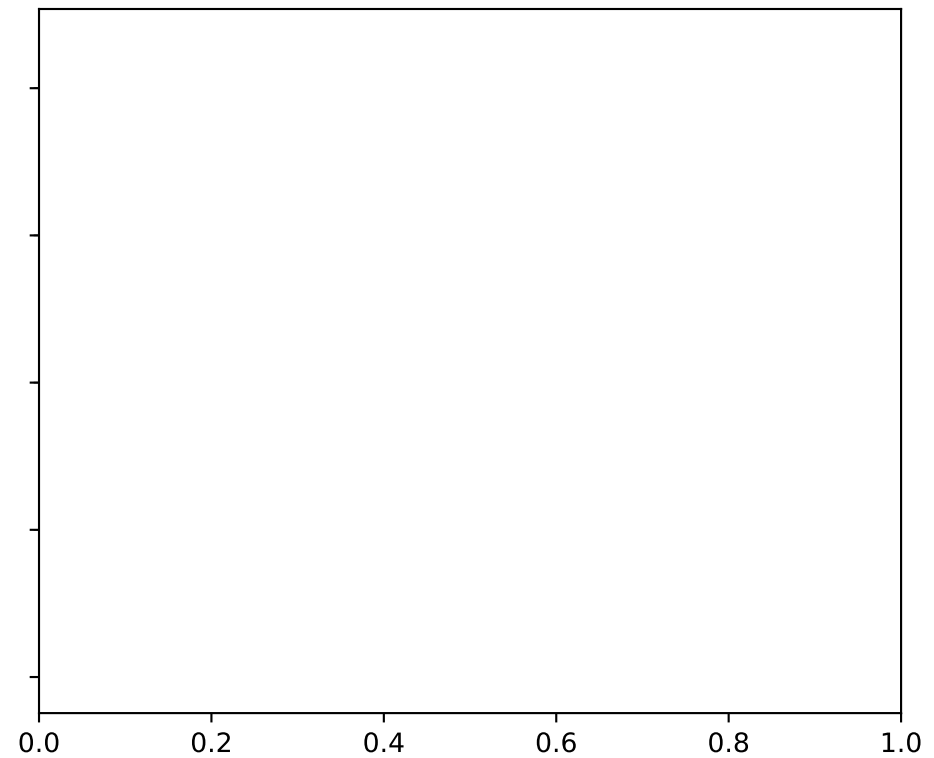
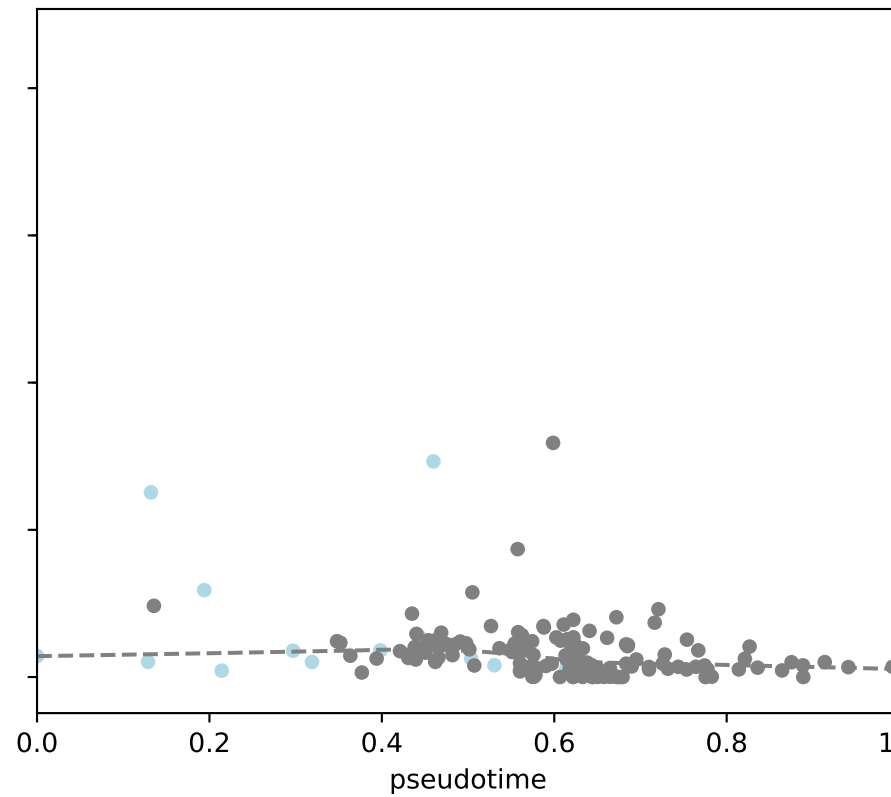
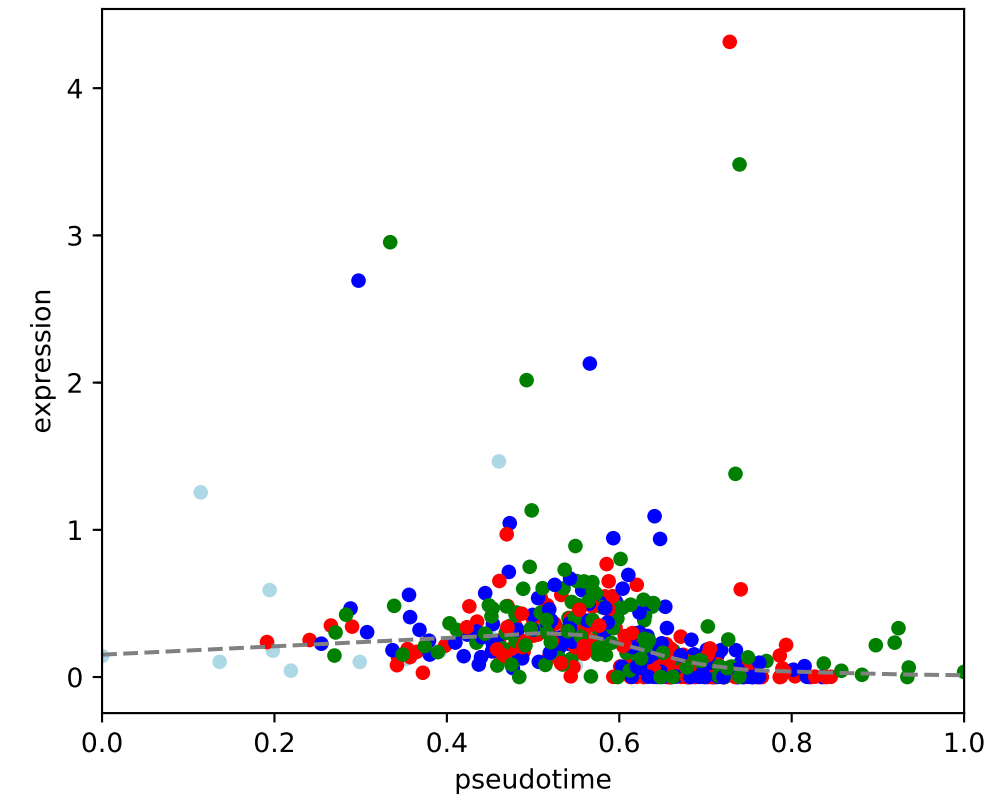
733 0spearman=-0.53

733 1spearman=-0.01



733 0spearman=-0.53

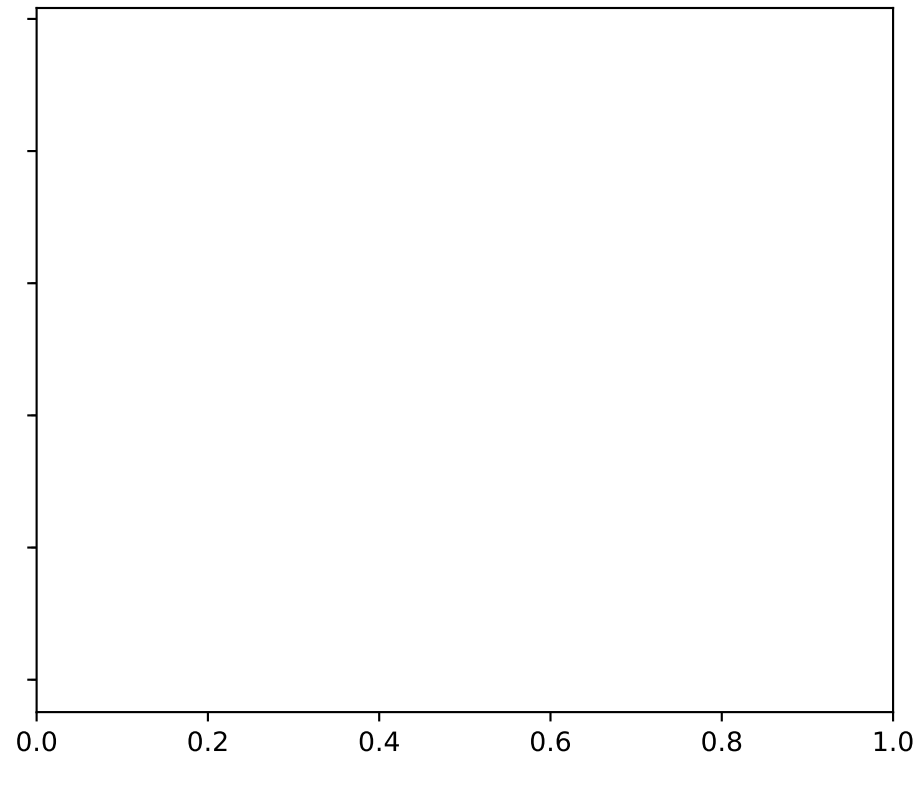
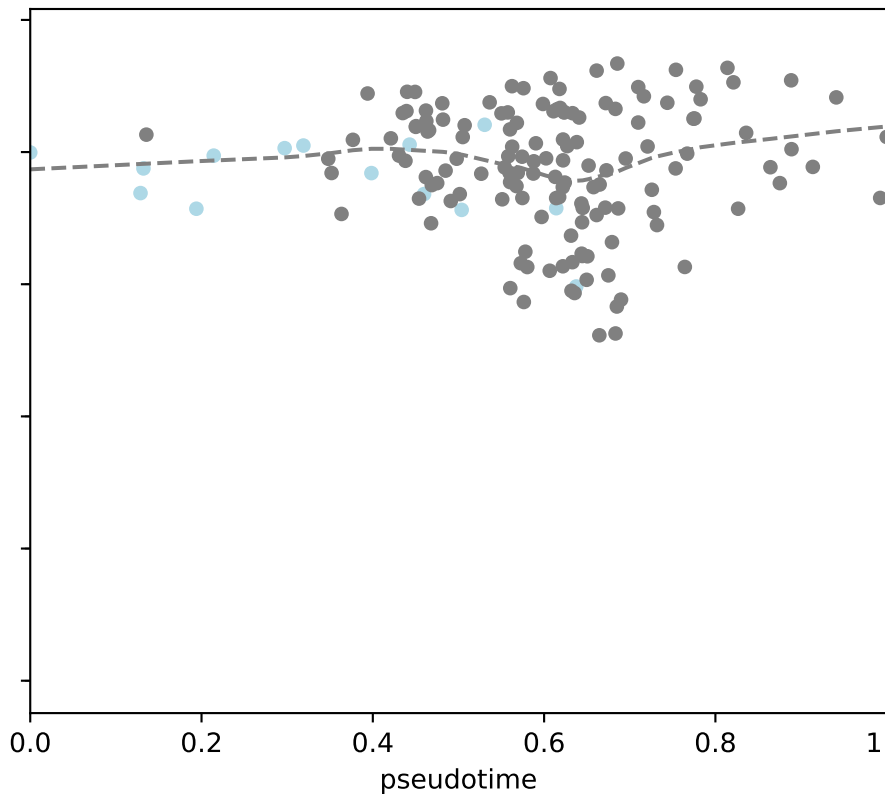
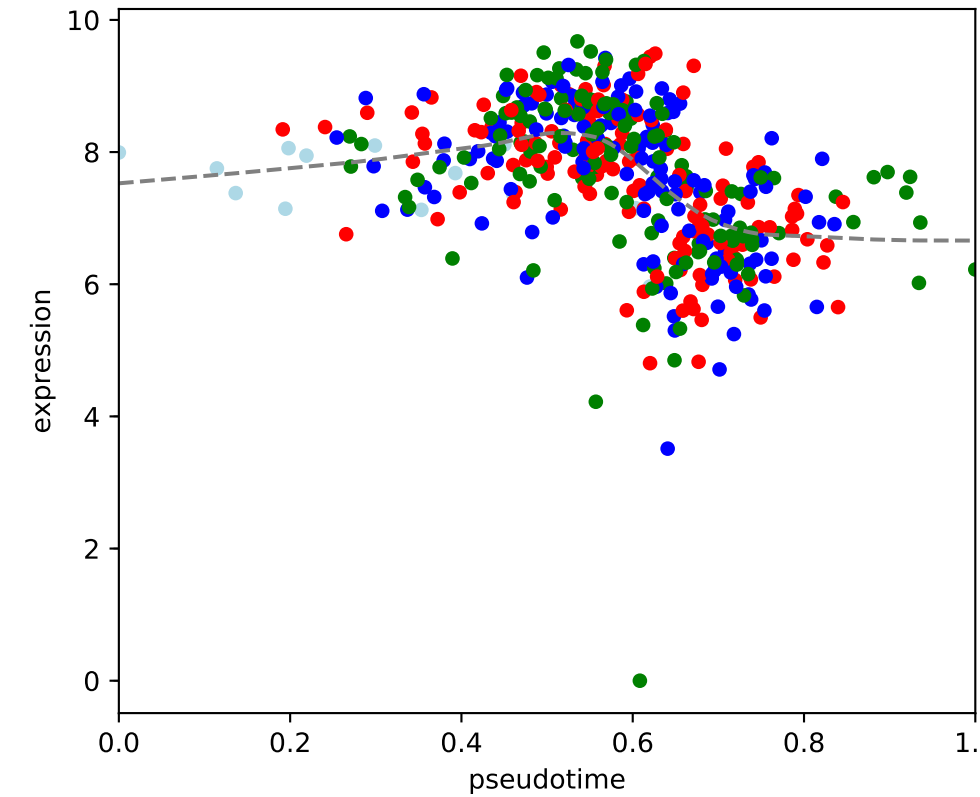
733 1spearman=-0.42



ENST00000357865 STMN1 (stathmin 1)

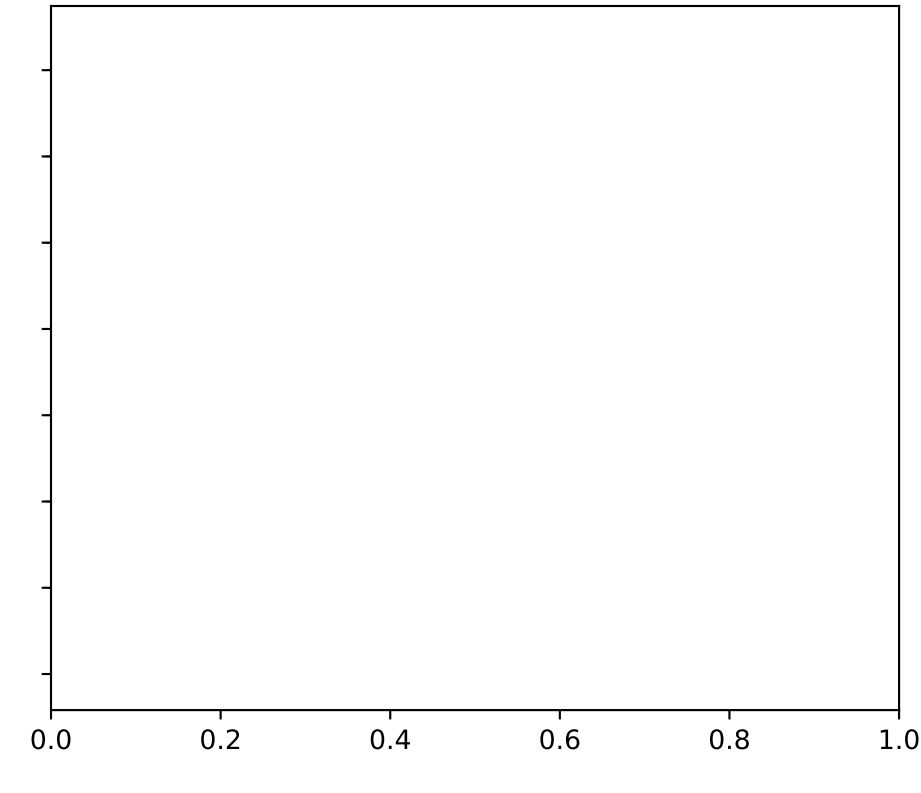
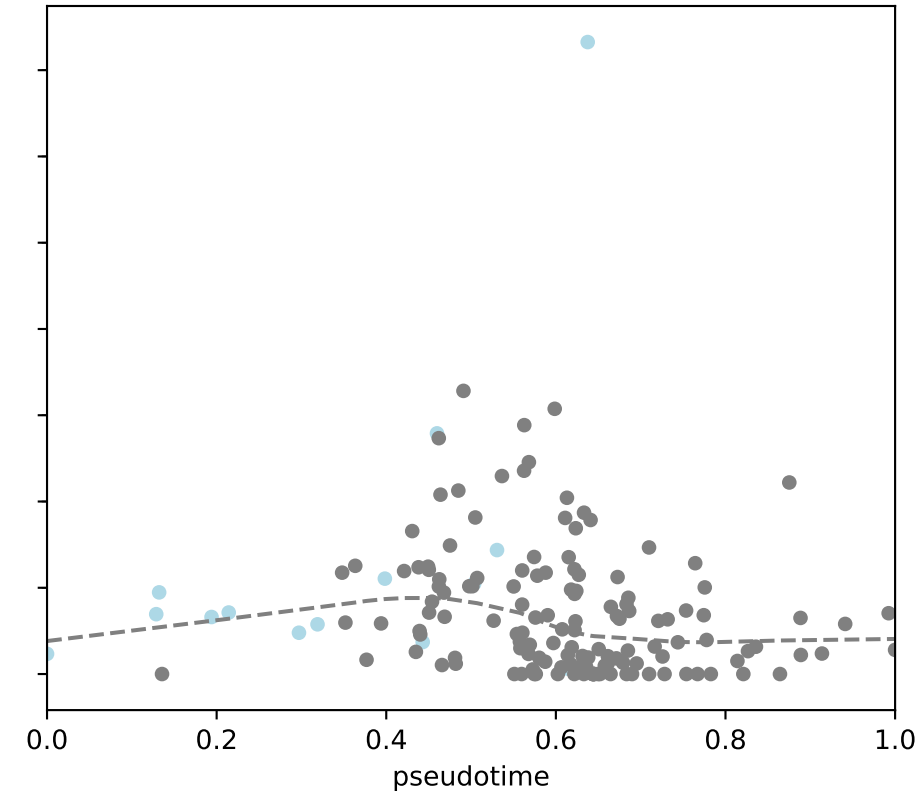
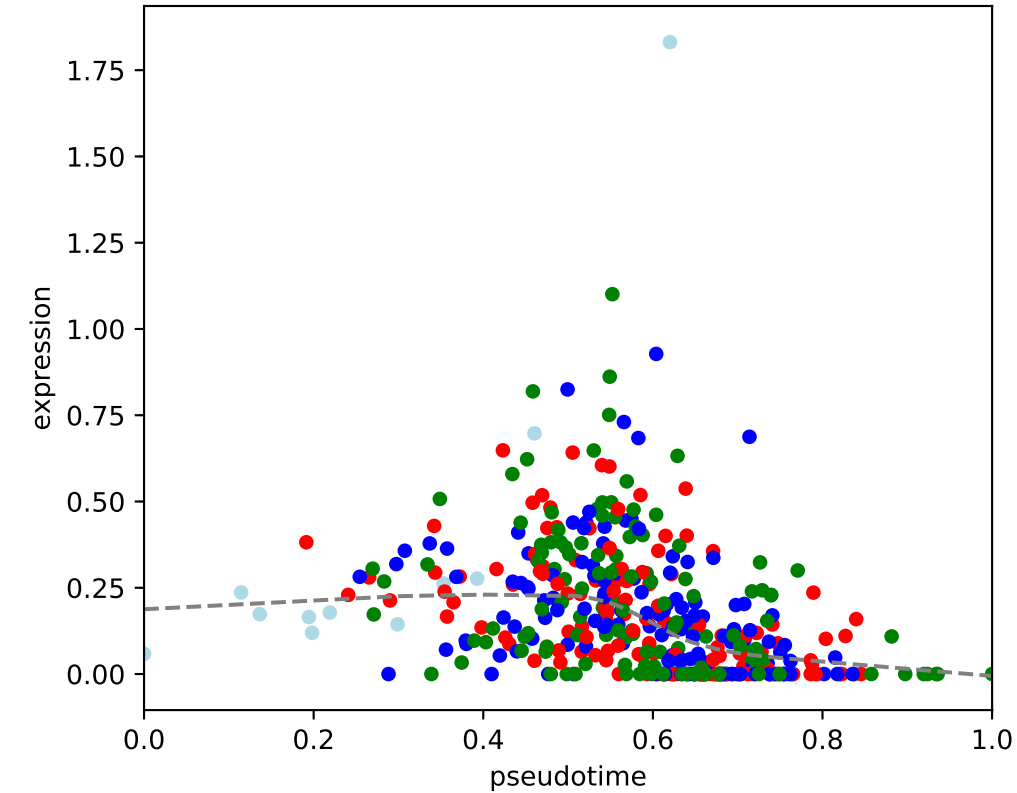
733 0spearman=-0.53

733 1spearman=-0.04



733 0spearman=-0.53

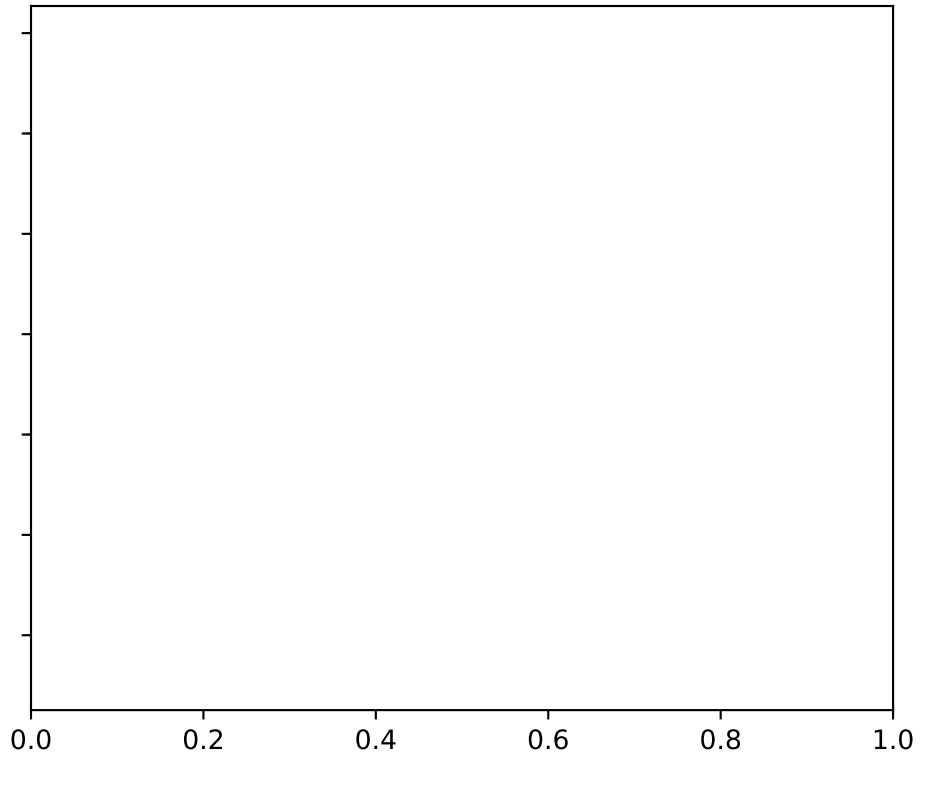
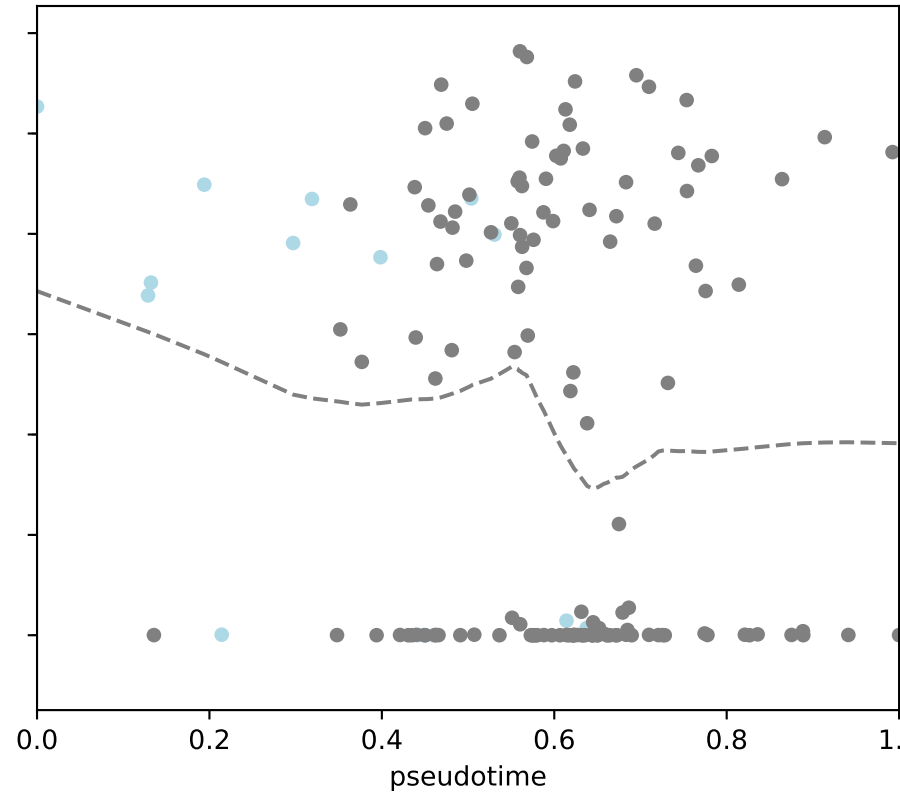
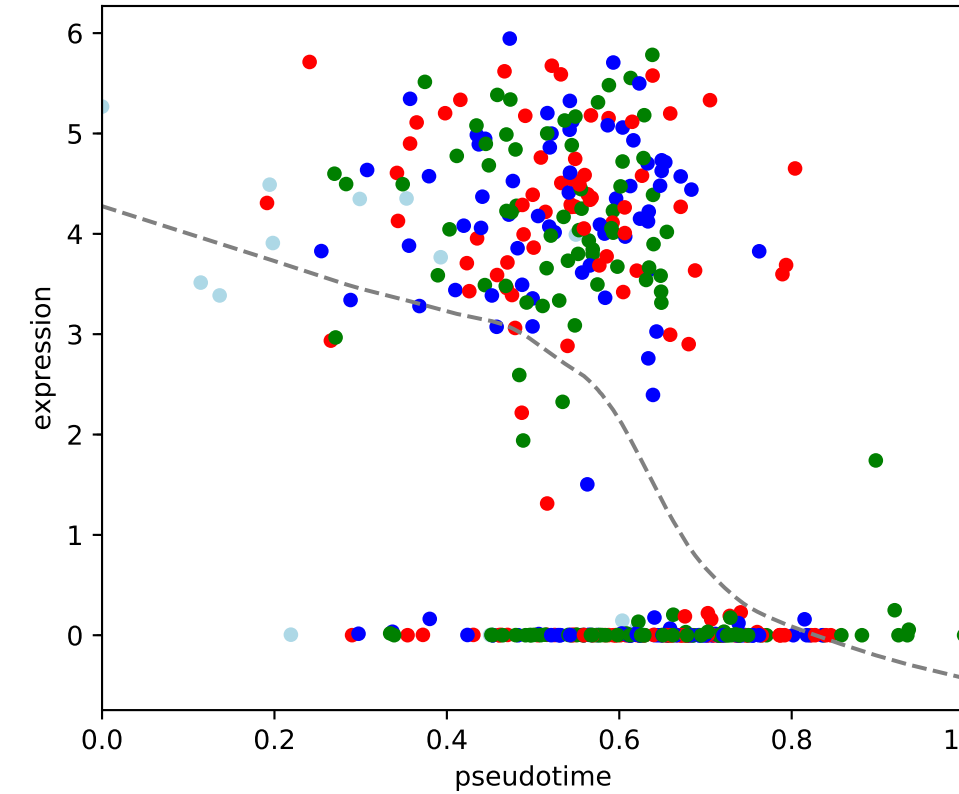
733 1spearman=-0.32



ENST00000633795 DST (dystonin)

733 0spearman=-0.53

733 1spearman=-0.10



ENST00000398153 HDDC2 (HD domain containing 2)

733 0spearman=-0.53

733 1spearman=-0.27

