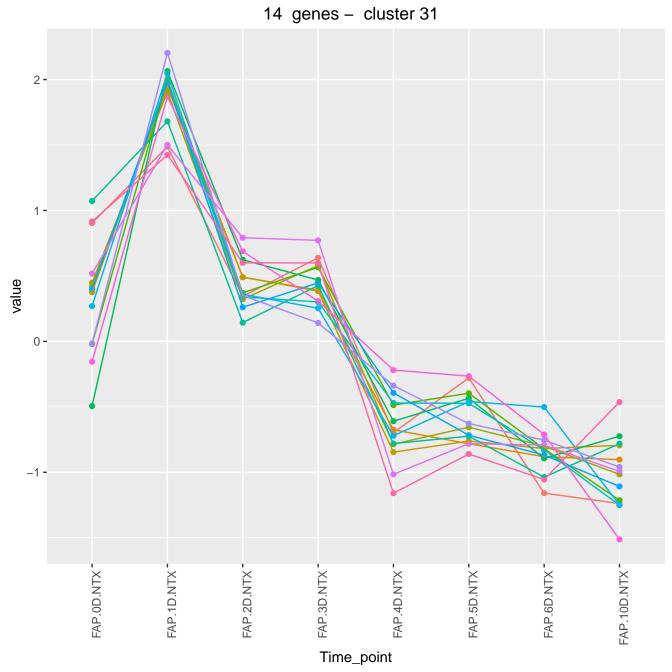
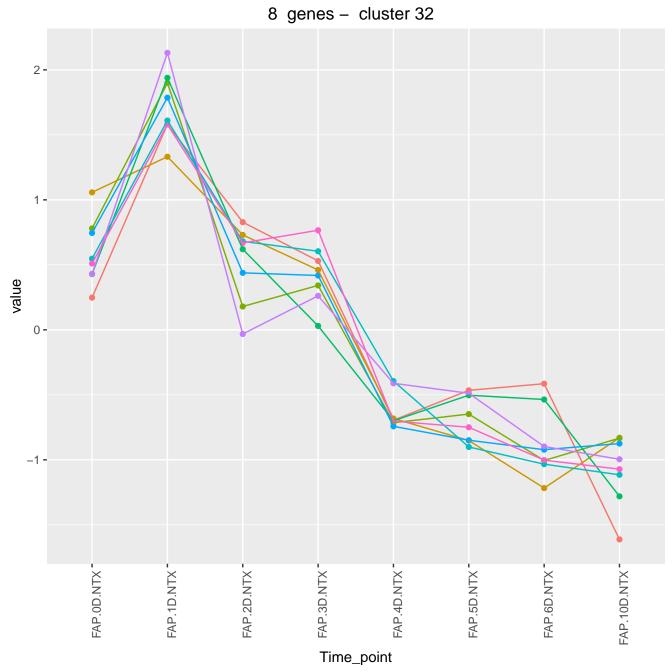
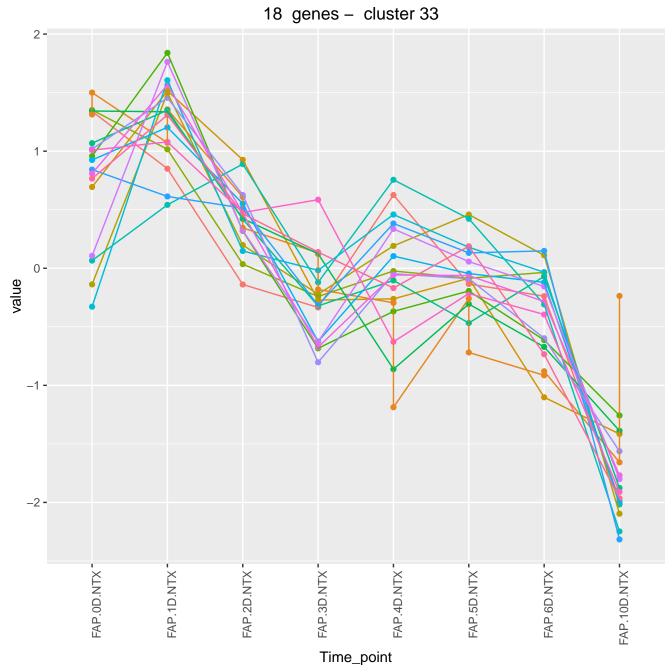
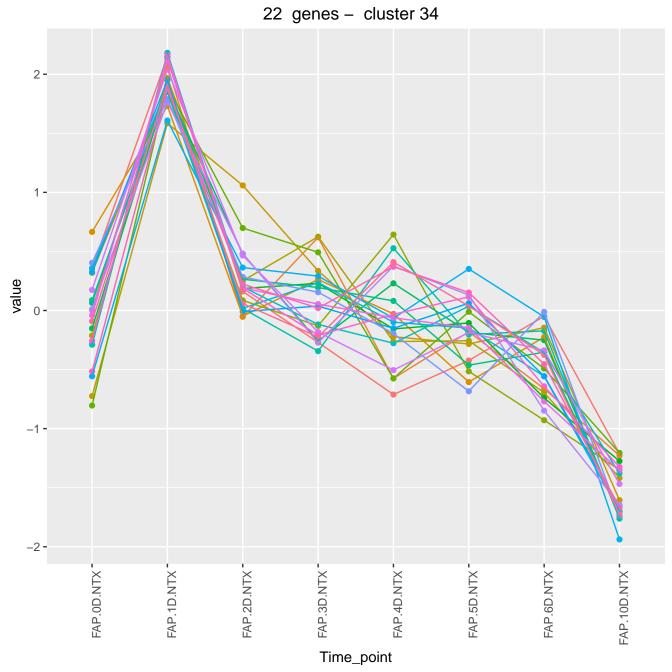


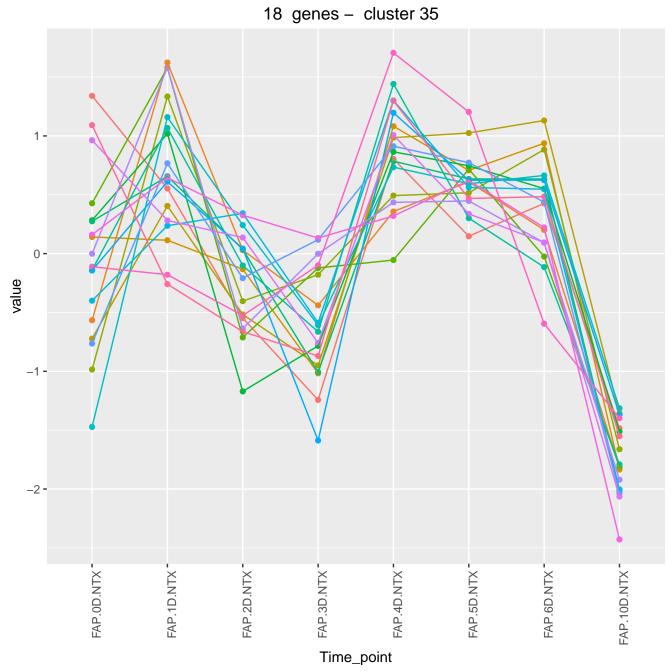
24 genes - cluster 30 2 -1 value 0 --1 **-**-2 **-**FAP.0D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.2D.NTX FAP.3D.NTX FAP.6D.NTX FAP.4D.NTX Time\_point

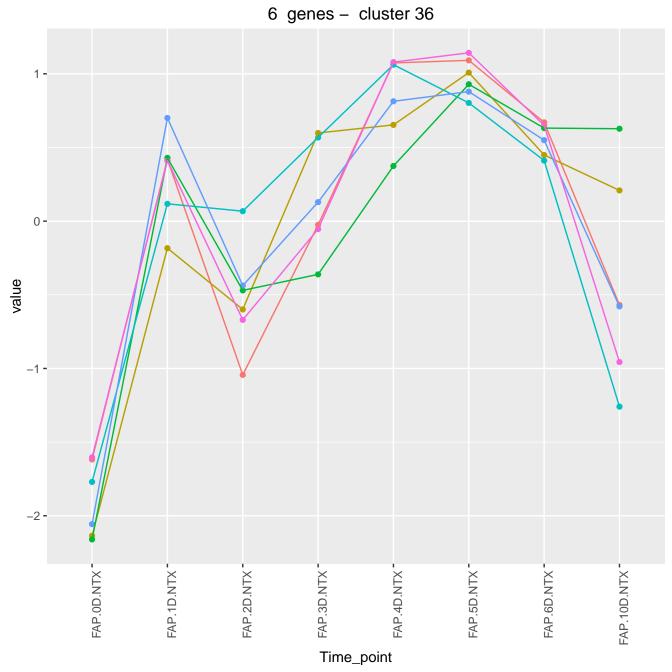


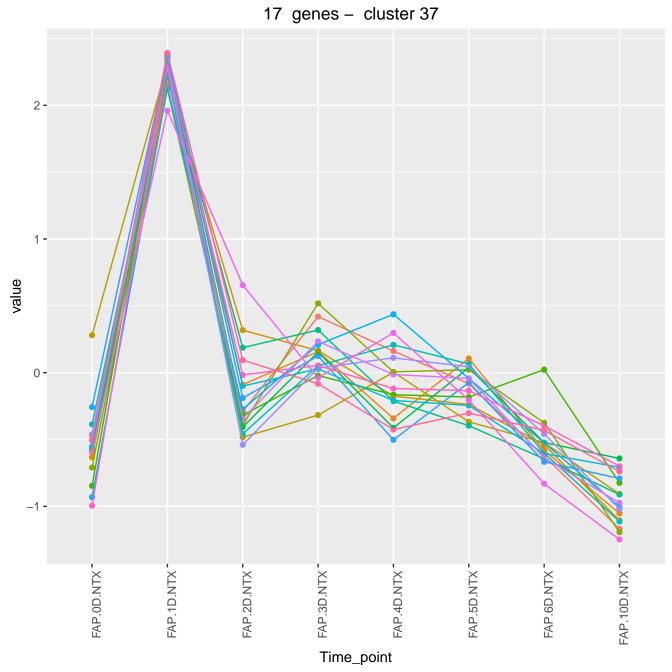




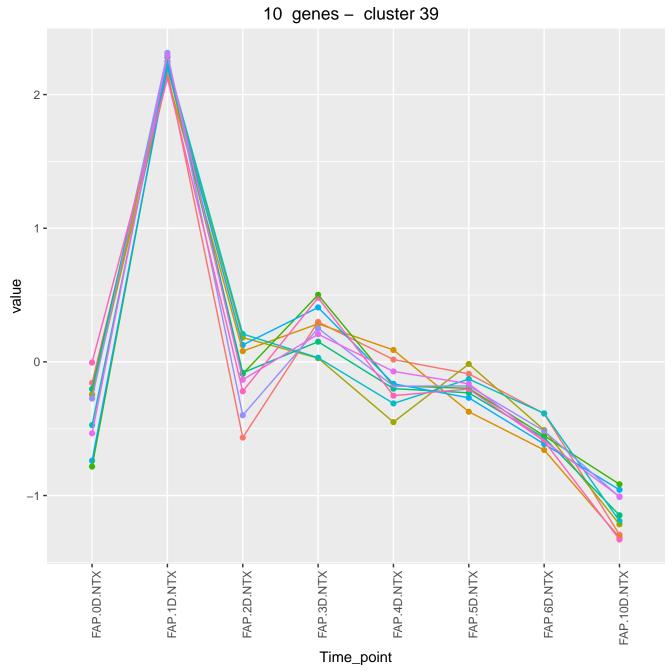


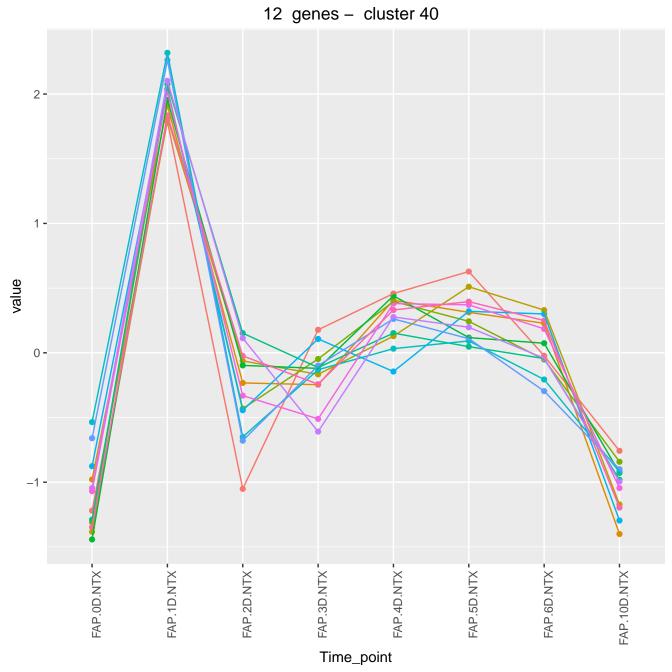


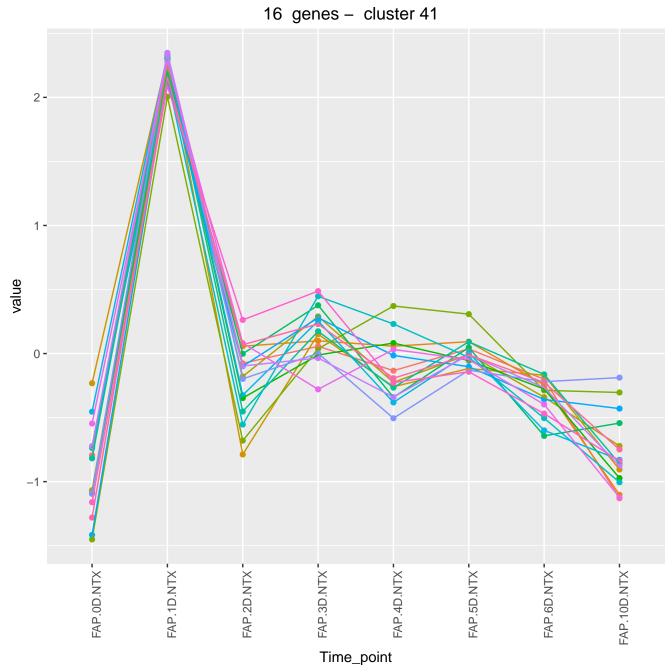


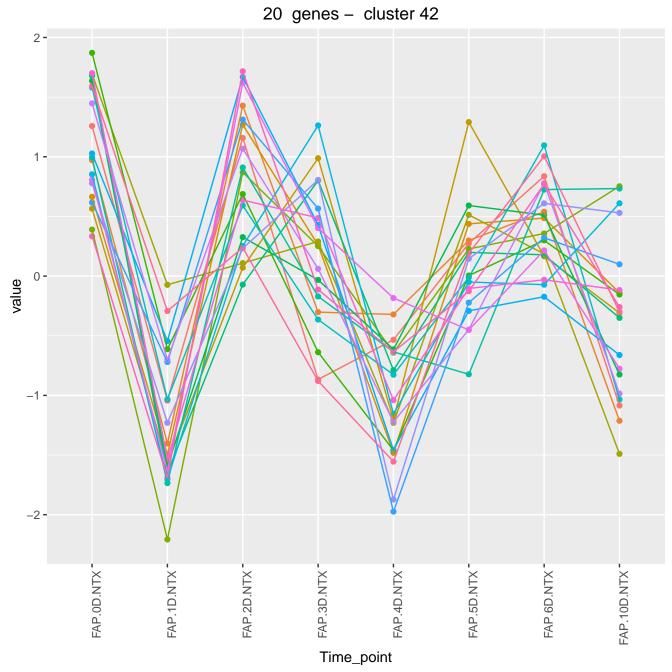


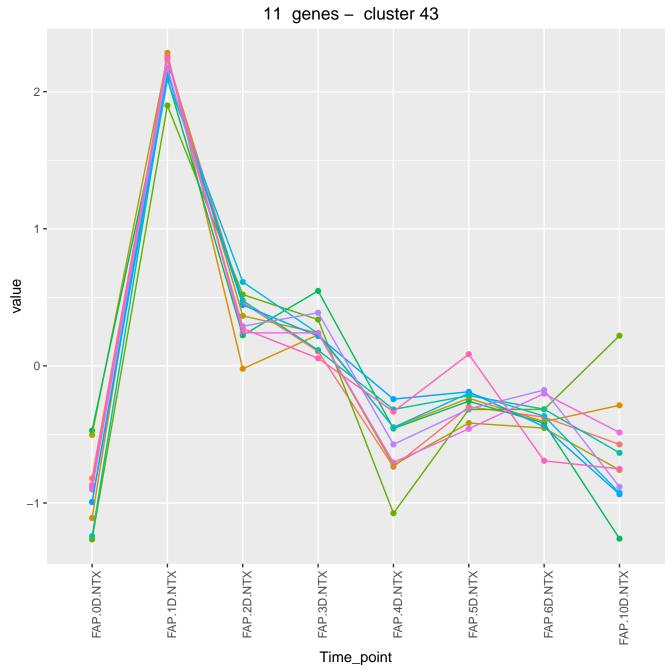
35 genes - cluster 38 2-1 value 0 --1 **-**-2 **-**FAP.0D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.10D.NTX T FAP.2D.NTX FAP.3D.NTX FAP.1D.NTX FAP.6D.NTX FAP.4D.NTX Time\_point

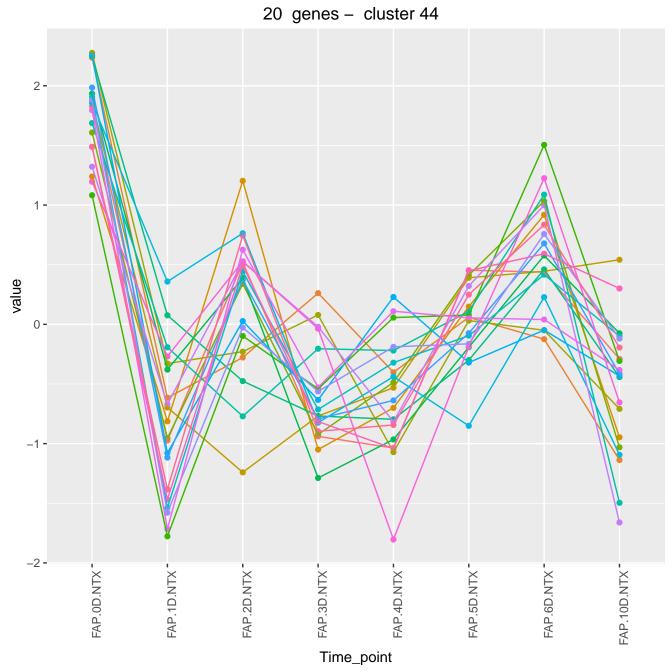


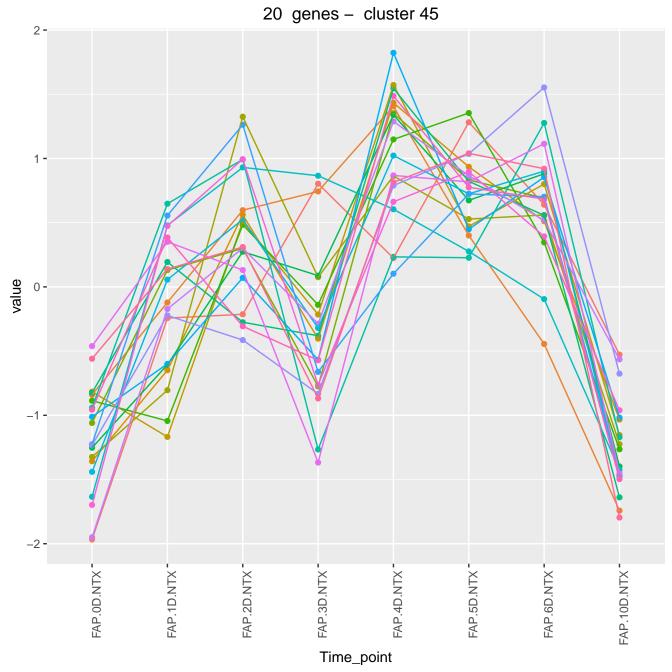


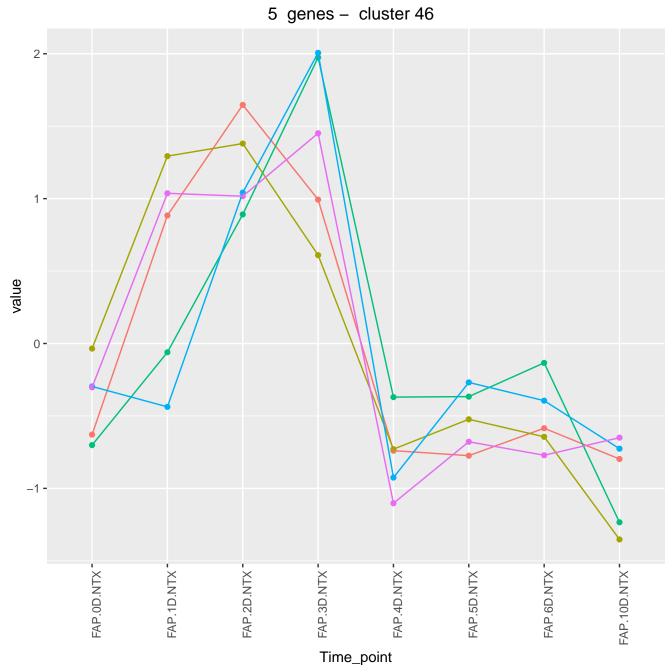


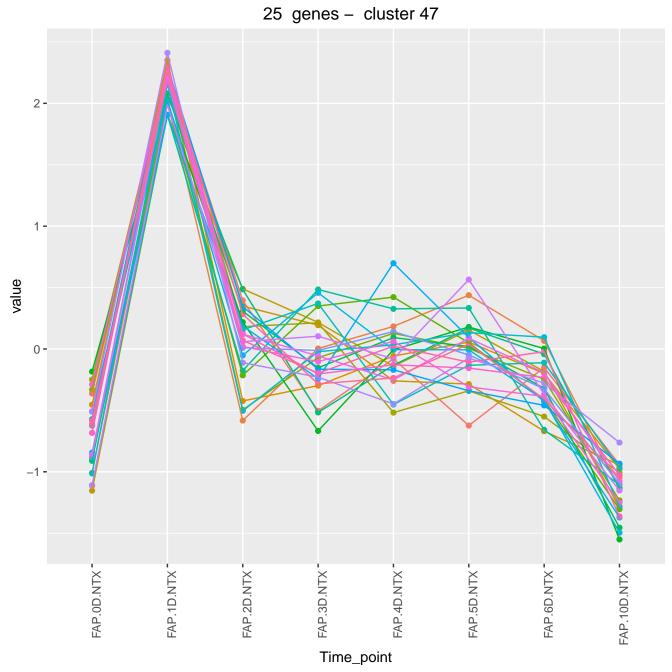


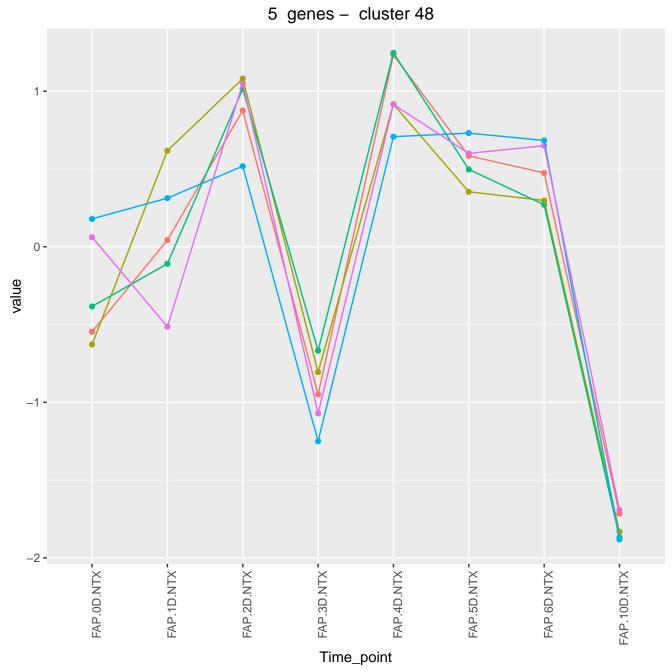


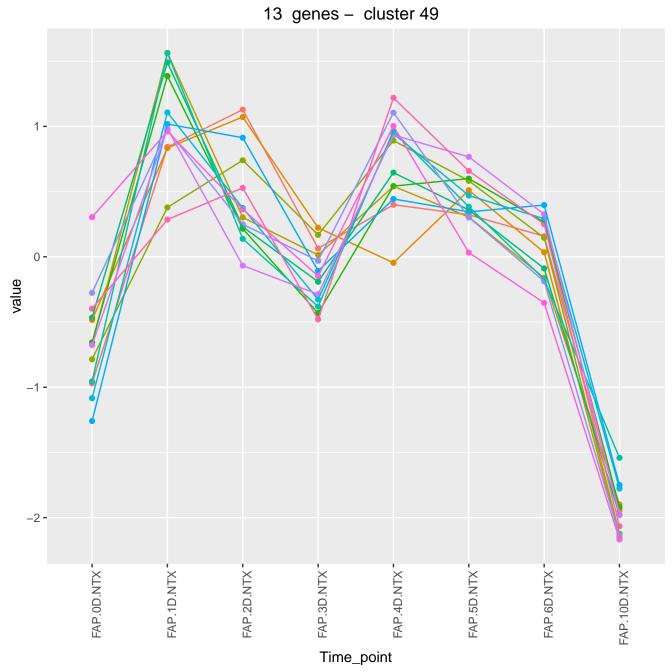


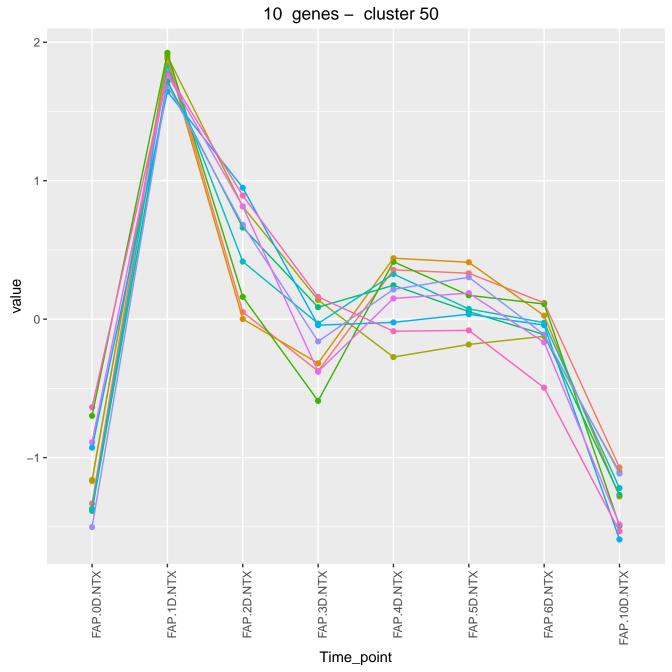


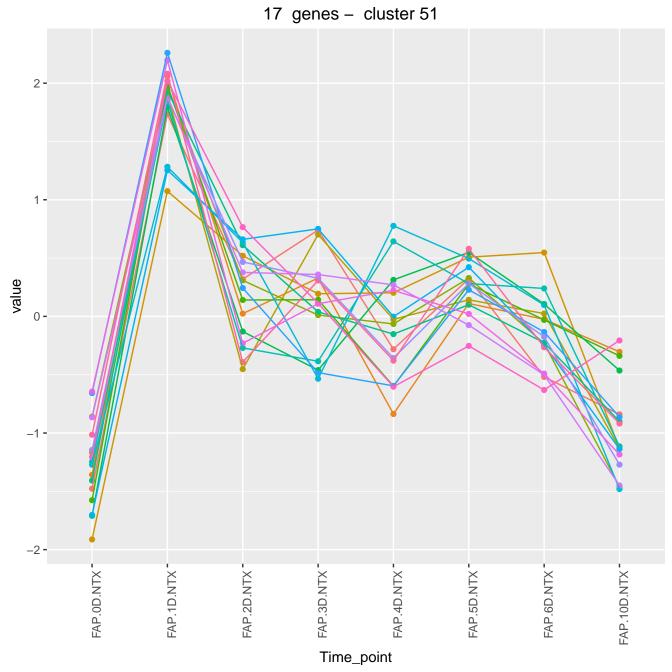


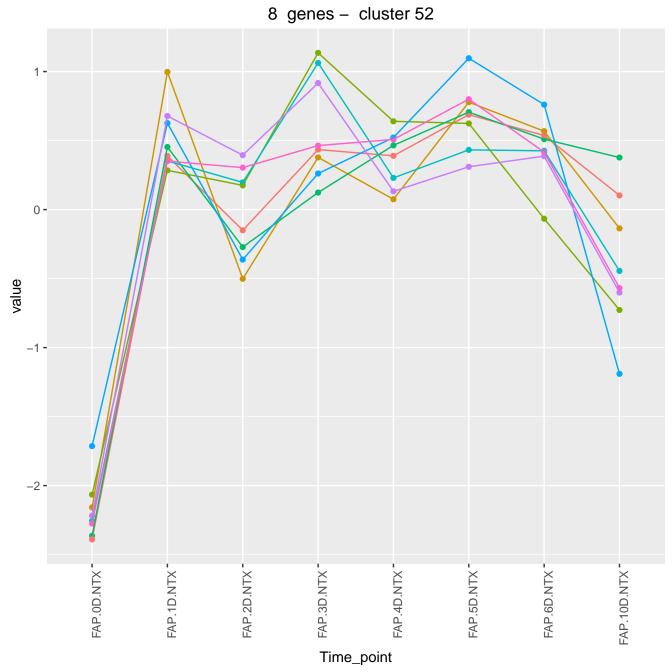


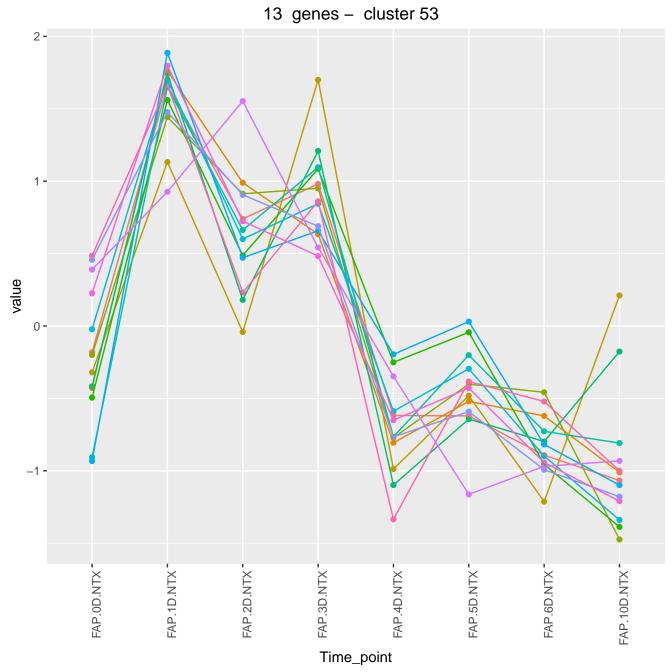


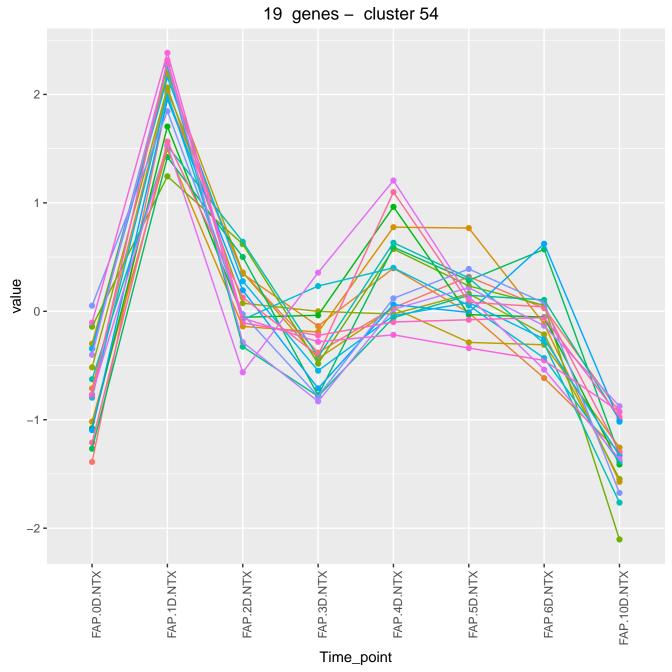


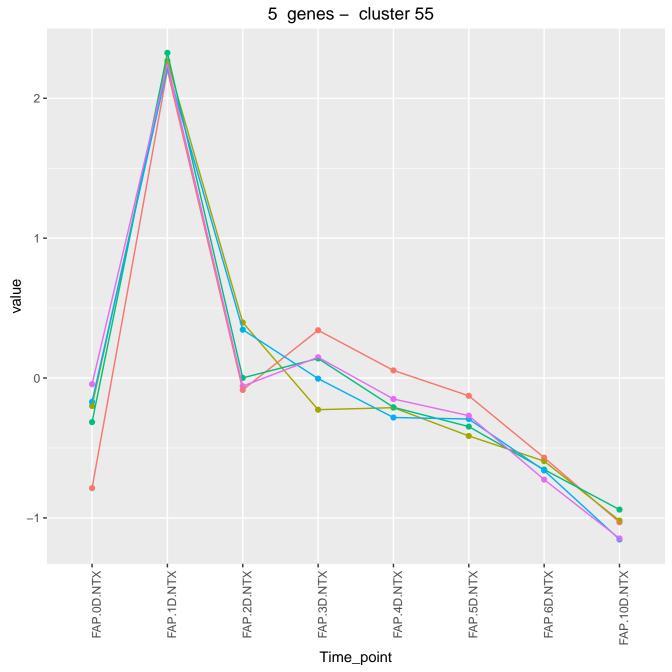


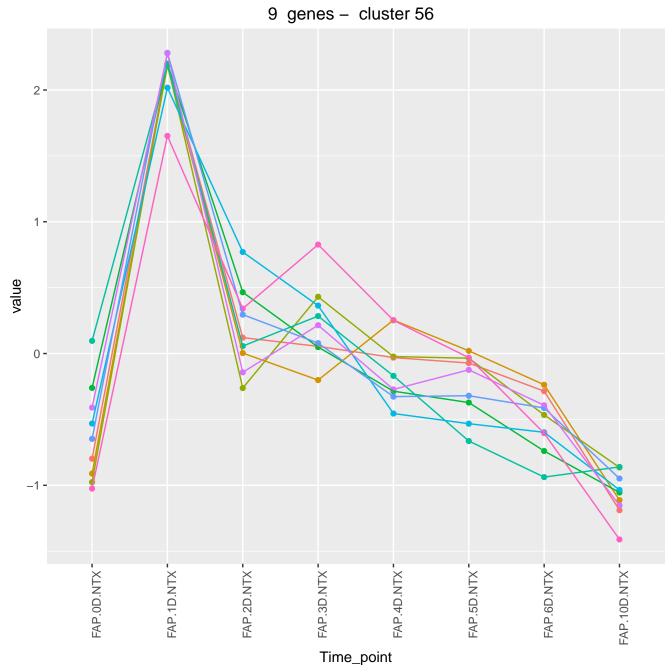


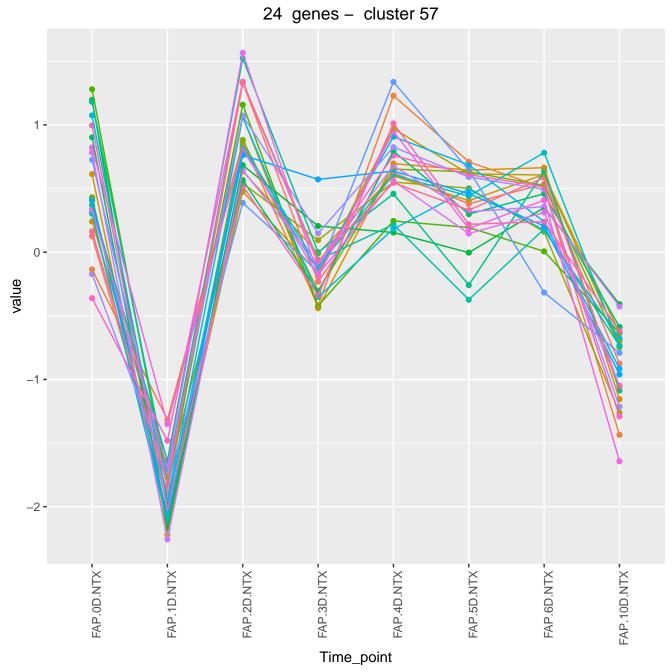


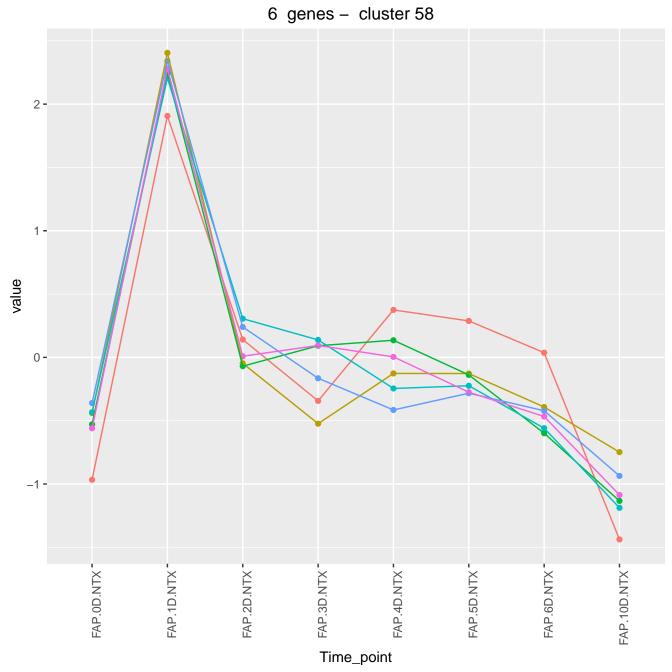


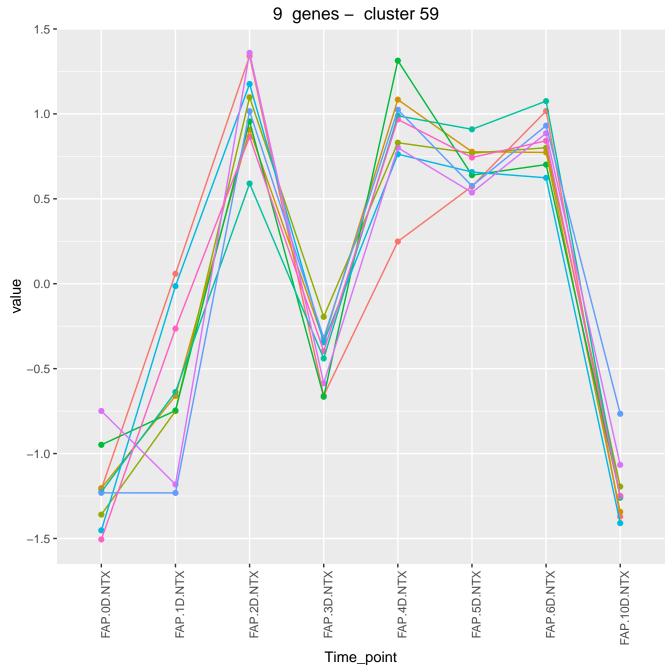


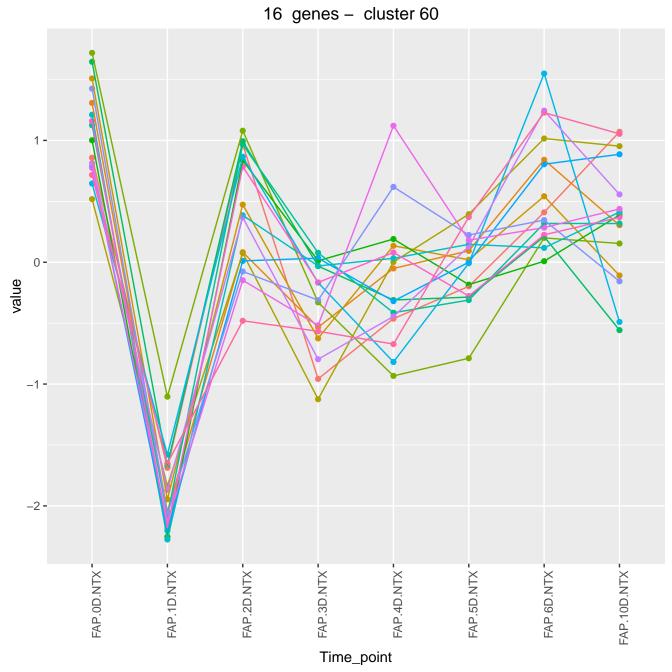


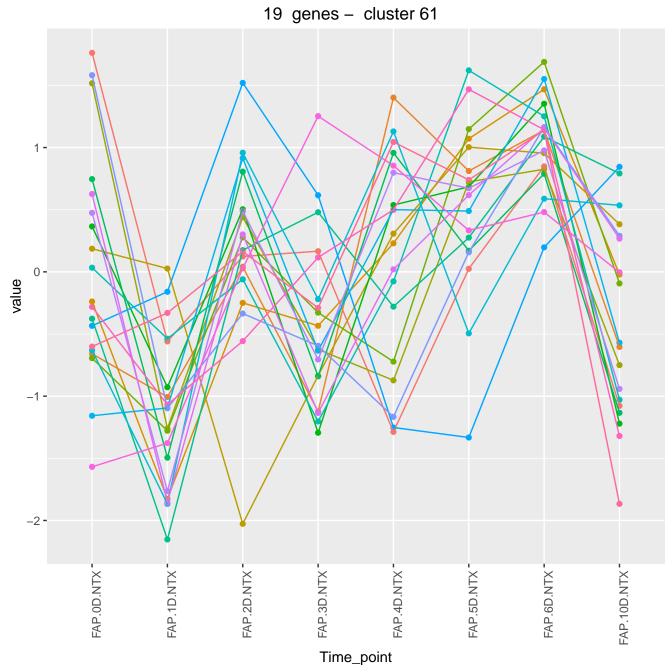


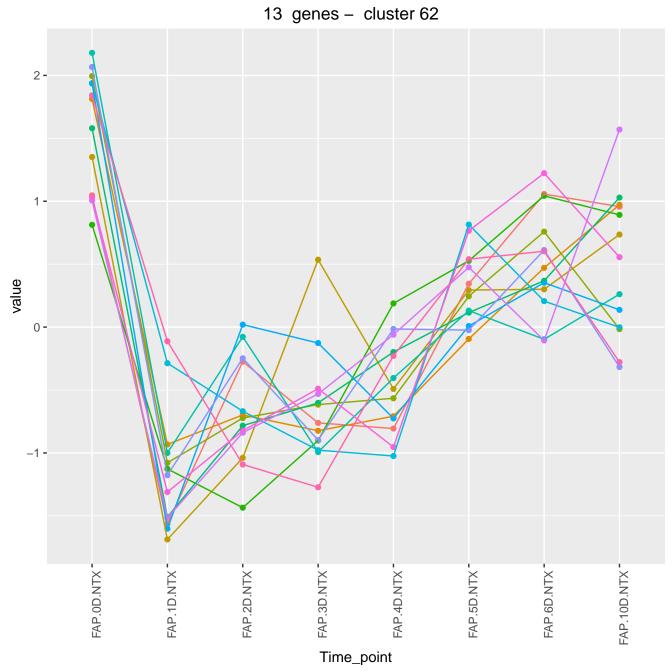




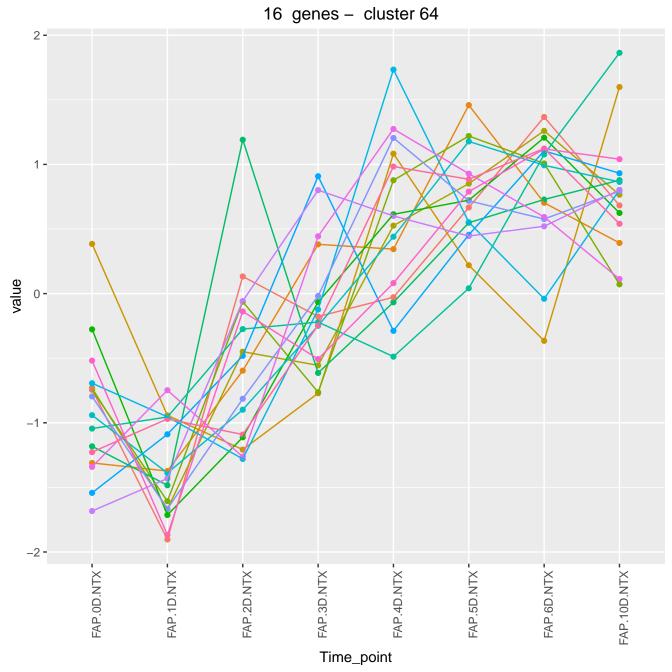


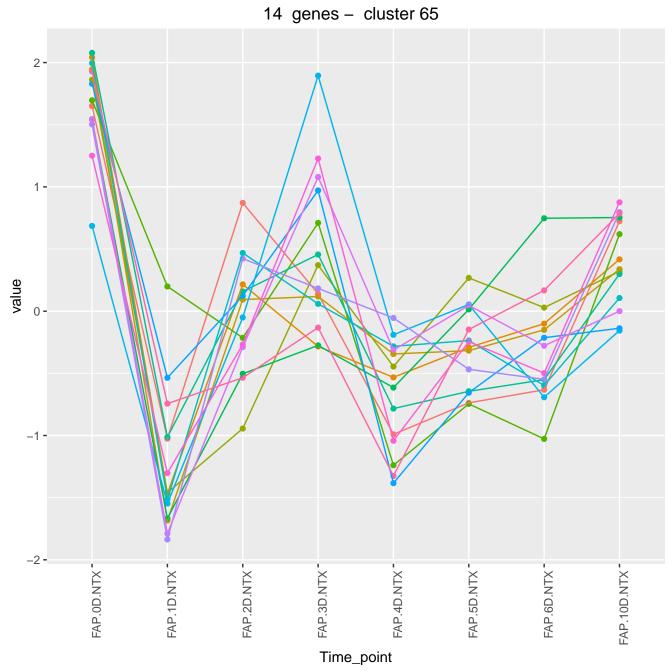


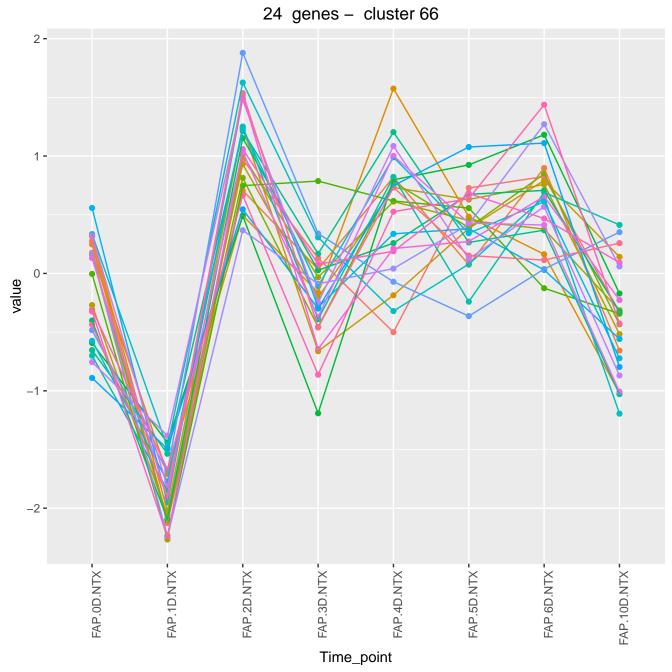


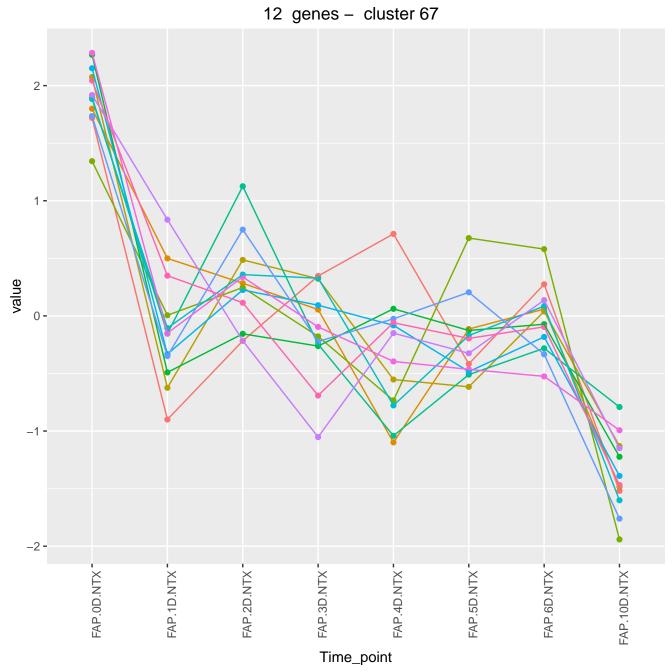


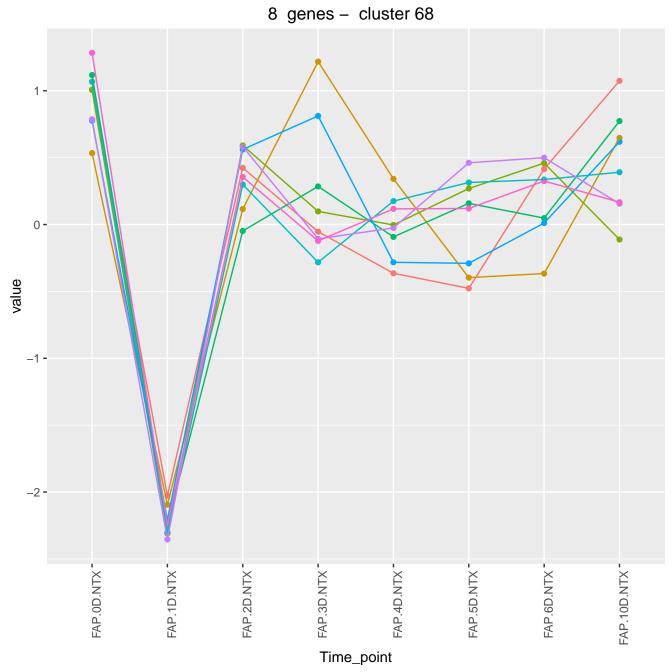
15 genes - cluster 63 2 -1 value **-1** --2 **-**FAP.0D.NTX 7 FAP.2D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.3D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

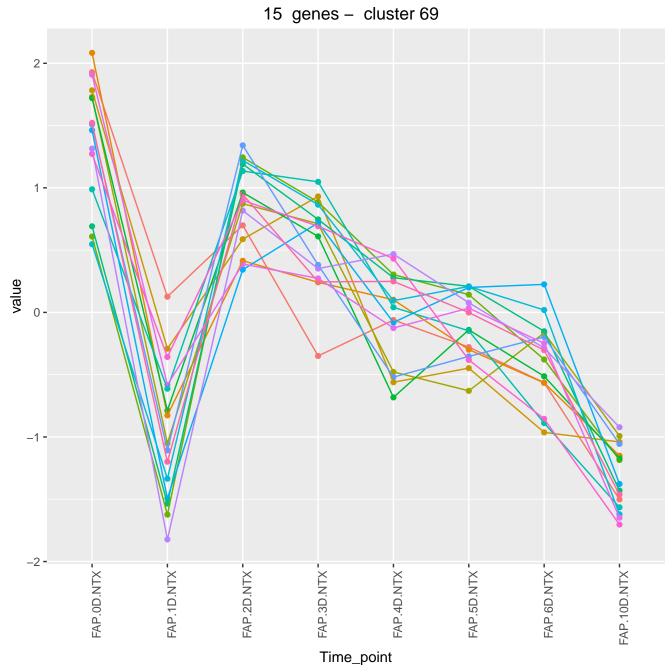


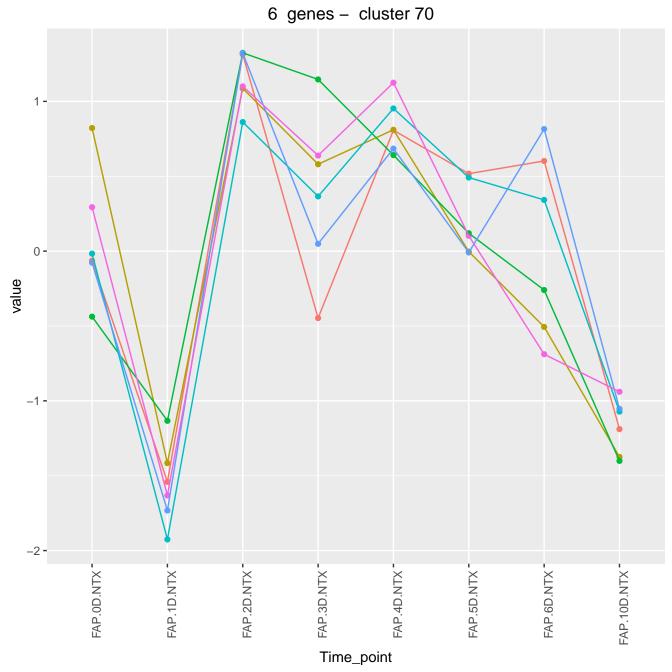


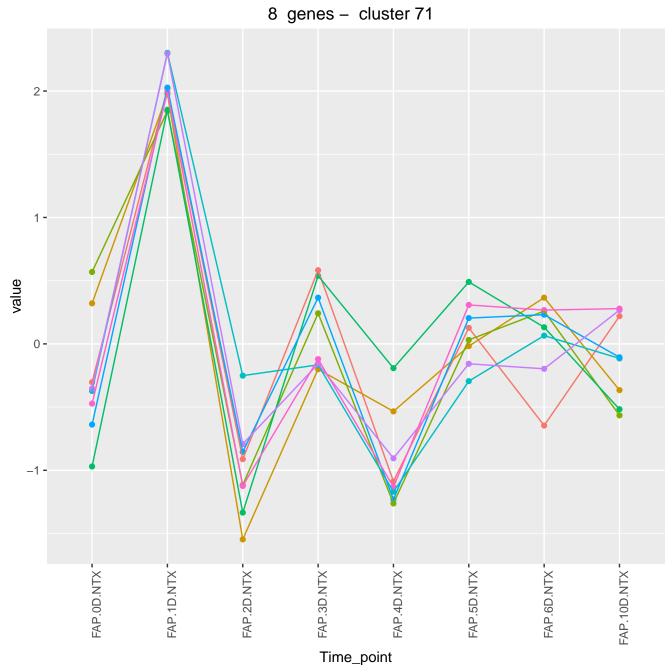




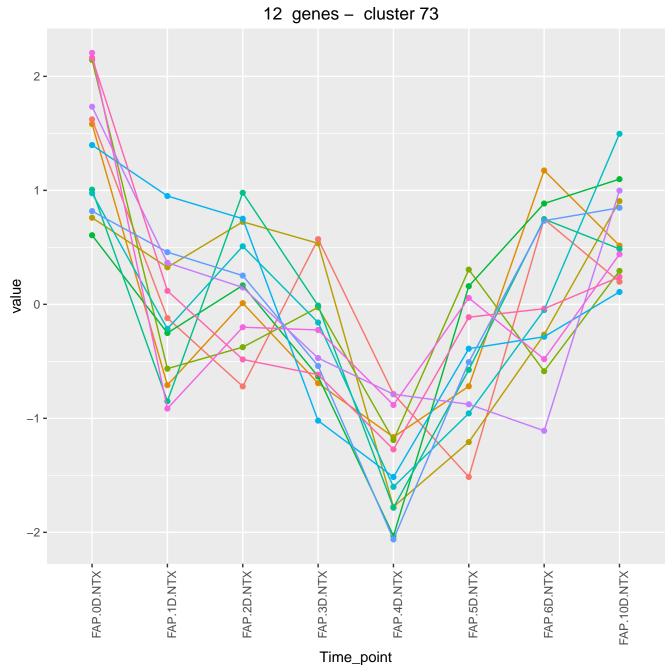


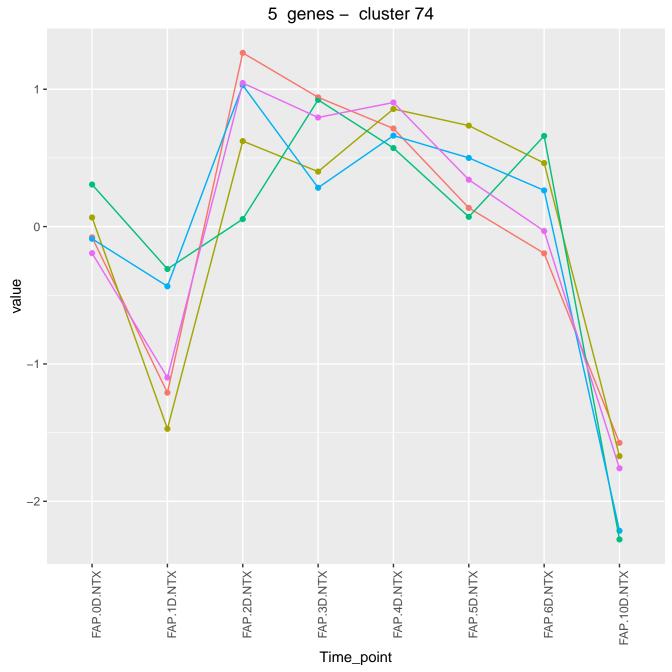




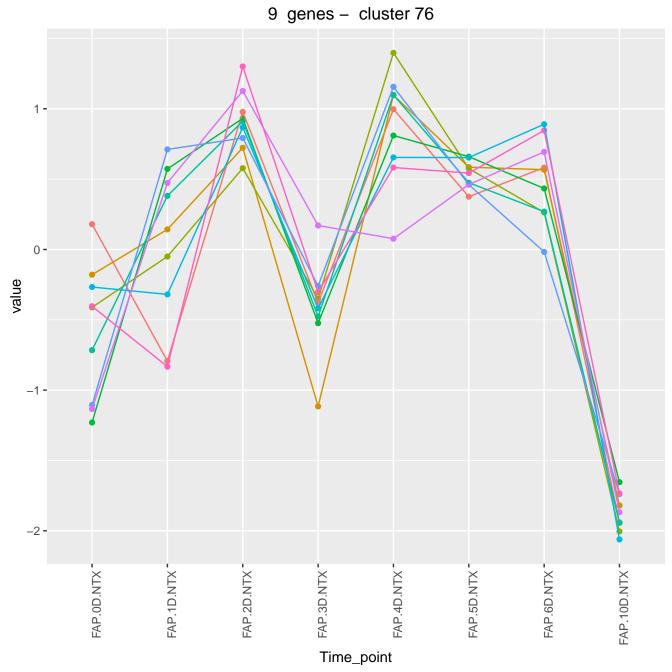


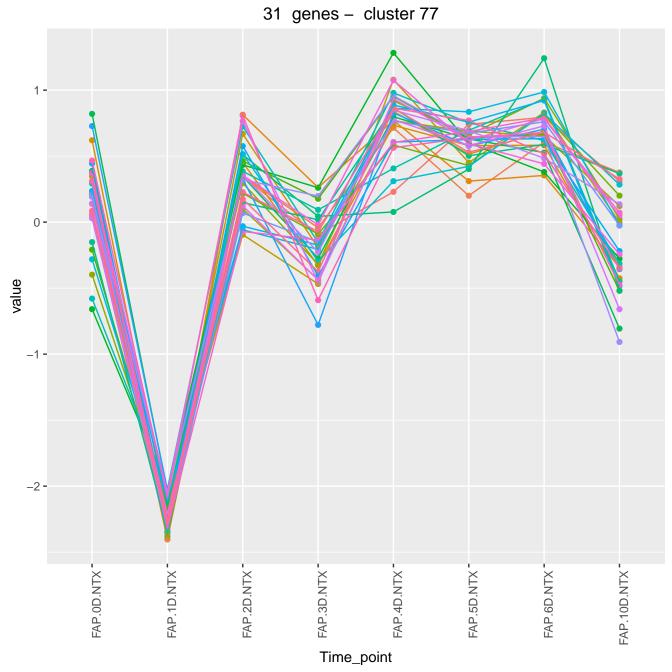
15 genes - cluster 72 2 -1 value 0 --1 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.3D.NTX -FAP.5D.NTX <sup>7</sup> FAP.6D.NTX <sup>7</sup> FAP.10D.NTX -FAP.4D.NTX Time\_point

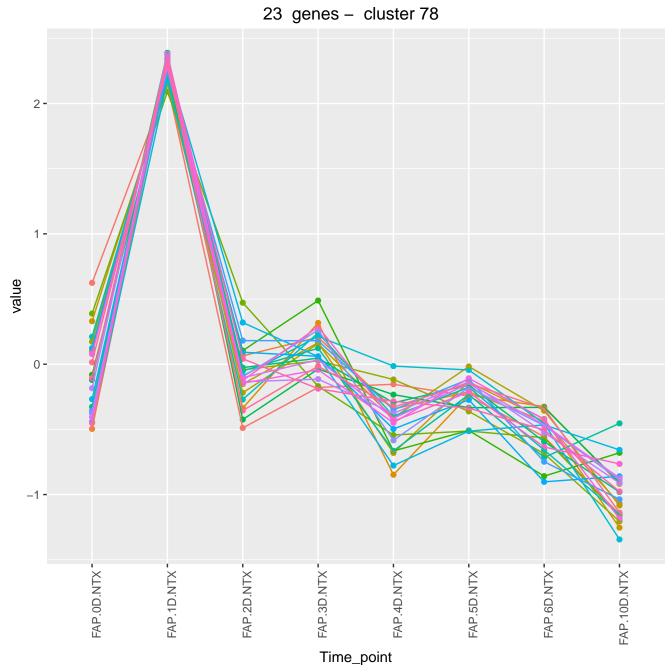


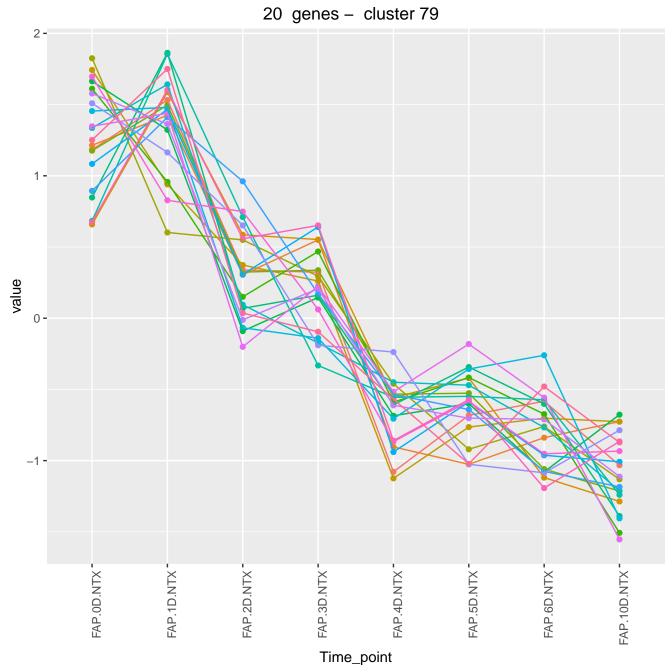


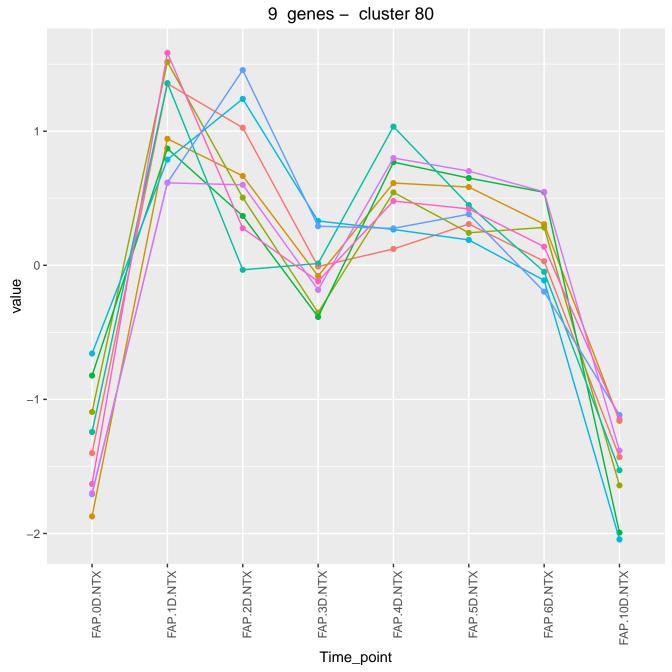
40 genes - cluster 75 2 -1 -0 value \_1 **-**-2 **-**FAP.0D.NTX -FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.4D.NTX FAP.6D.NTX Time\_point



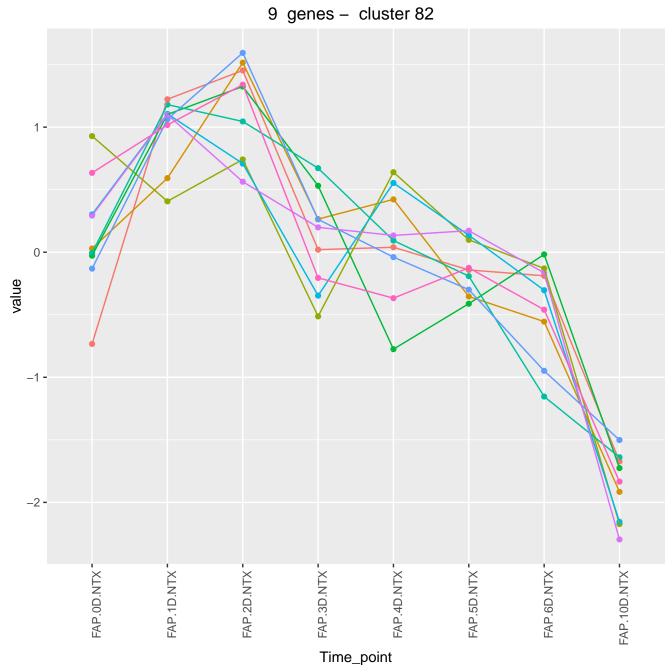


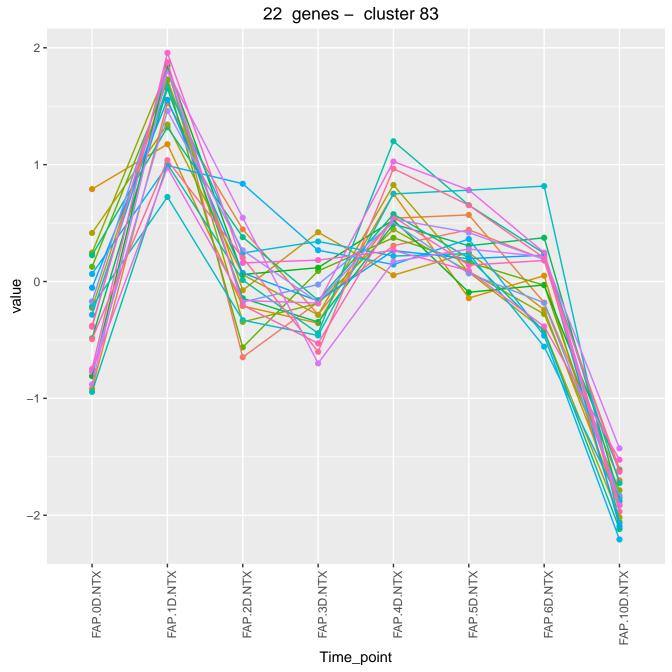




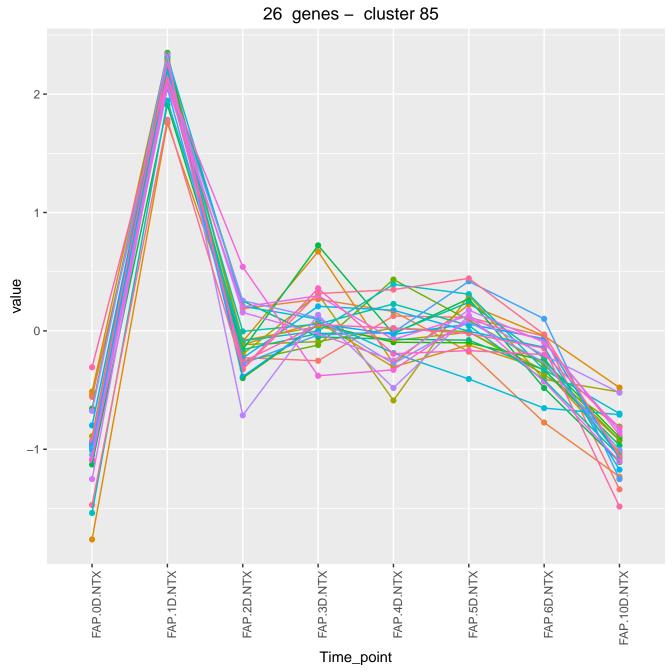


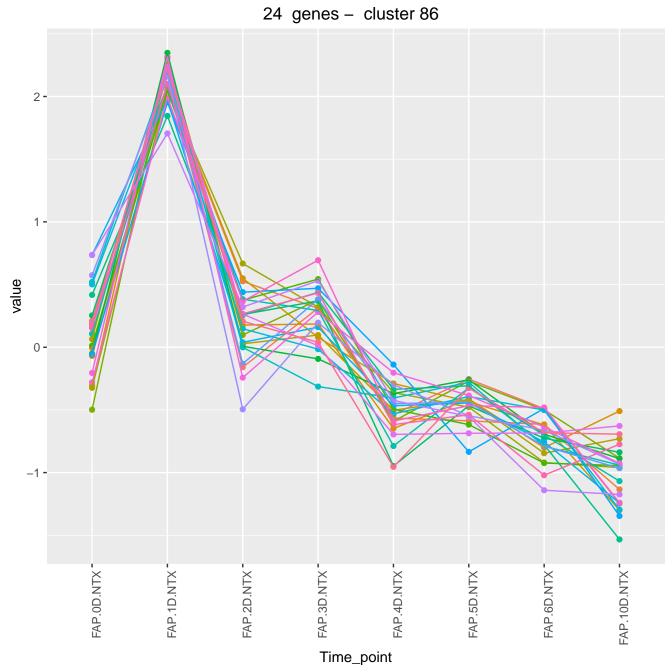
14 genes - cluster 81 2-1 value 0 -**-1-**-2 **-**FAP.0D.NTX 7 FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.6D.NTX -FAP.4D.NTX Time\_point

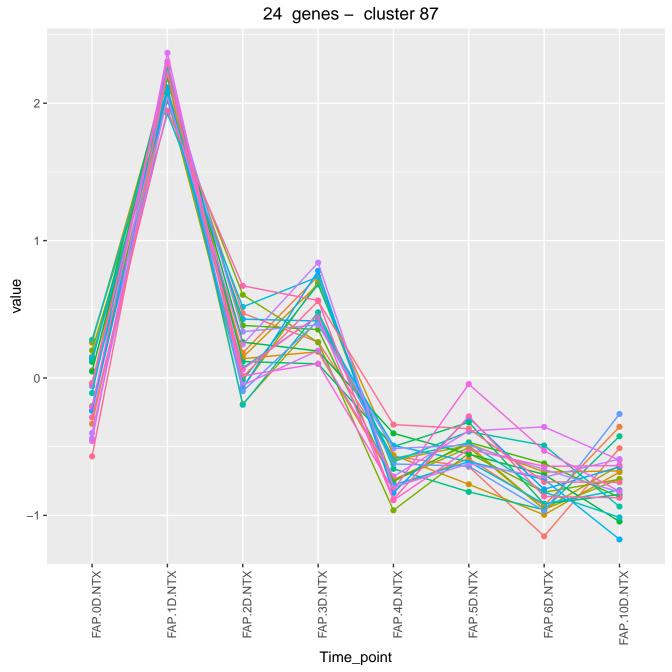


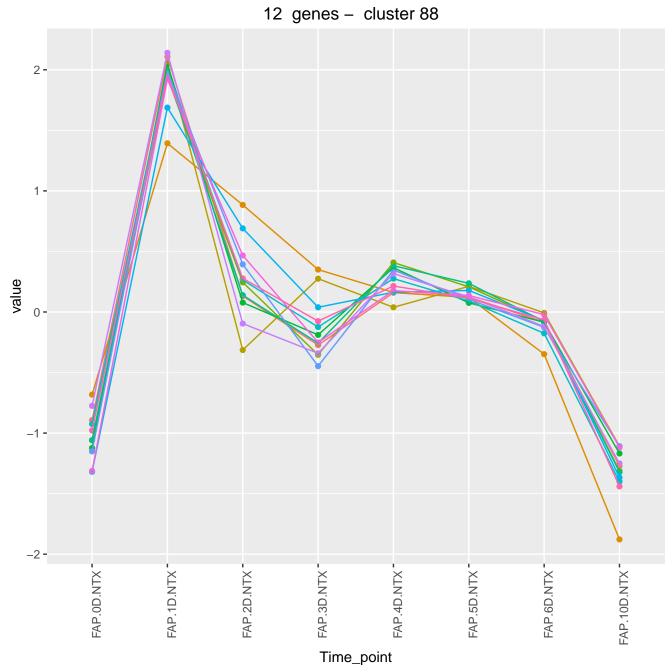


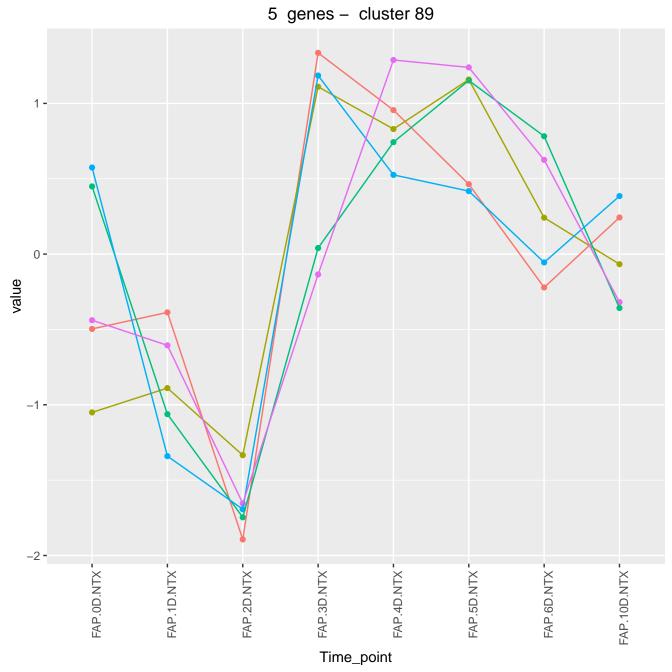
16 genes - cluster 84 2 value 0 -**-1-**FAP.0D.NTX 7 FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

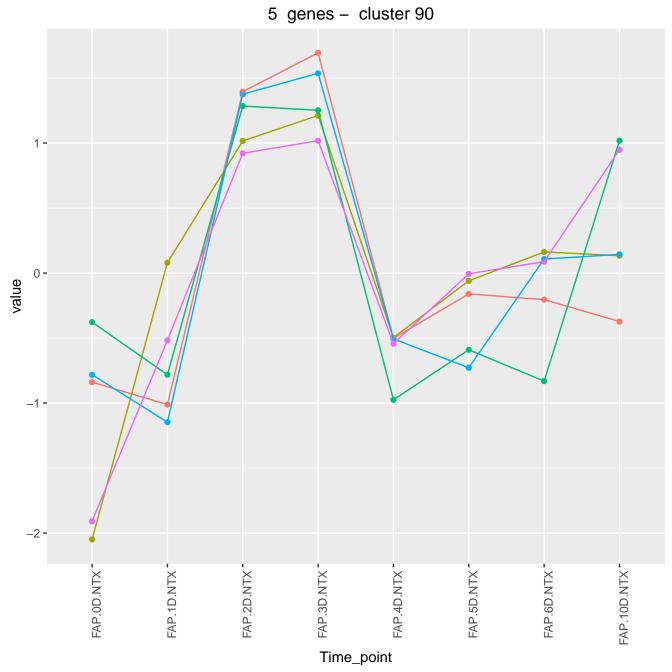


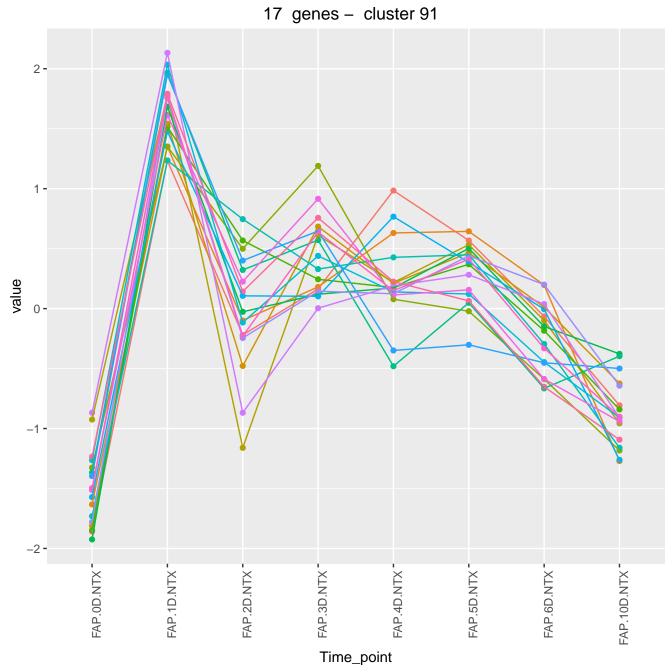


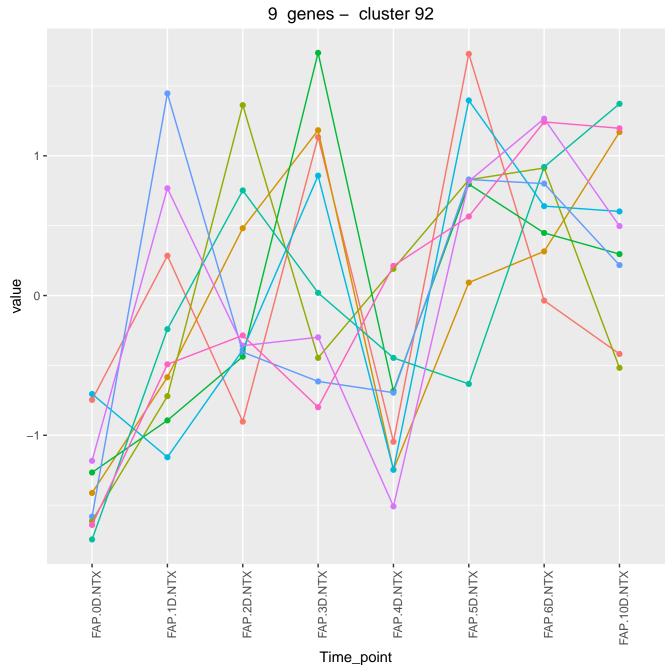


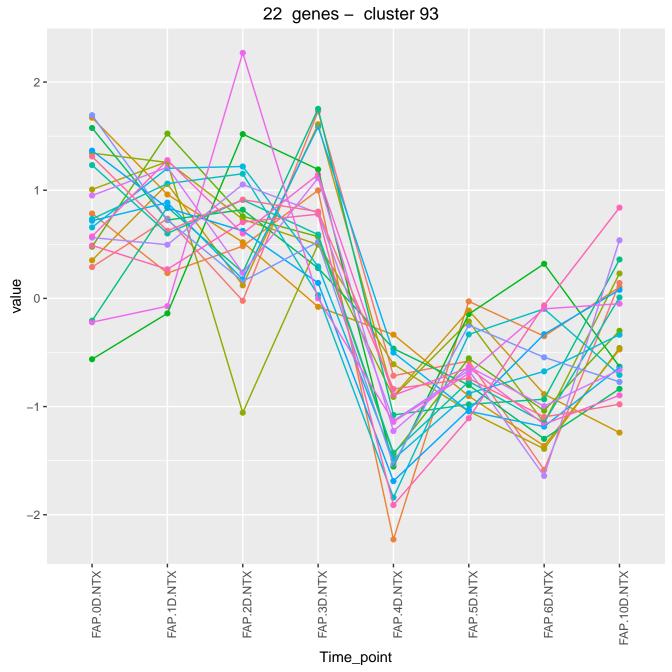


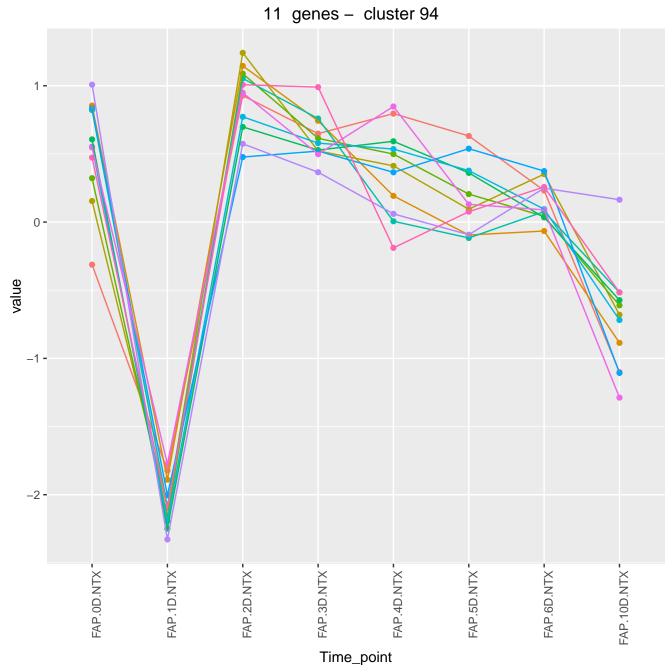


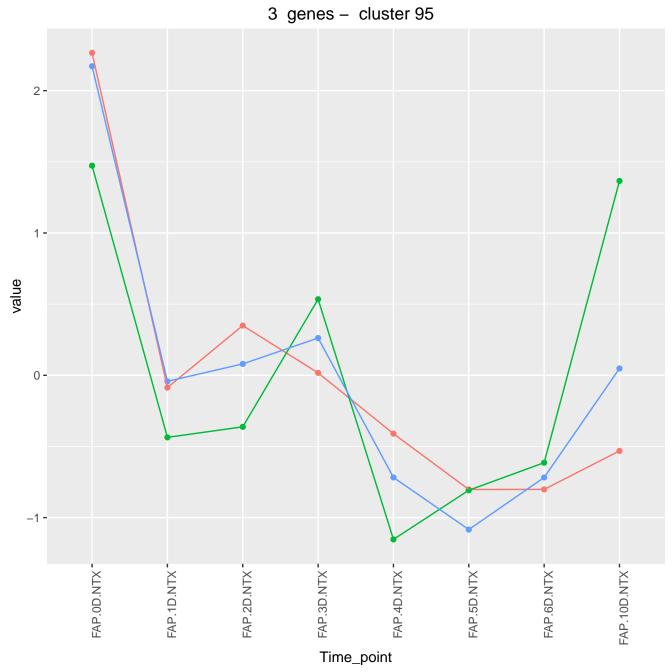


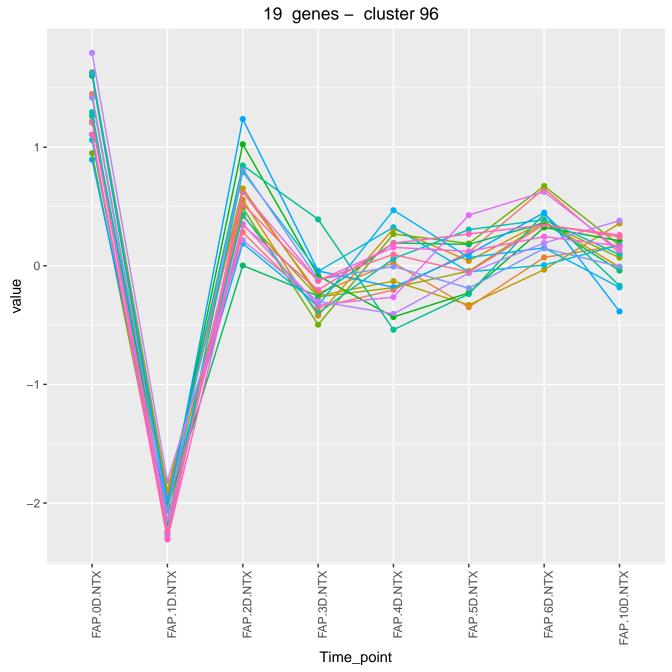


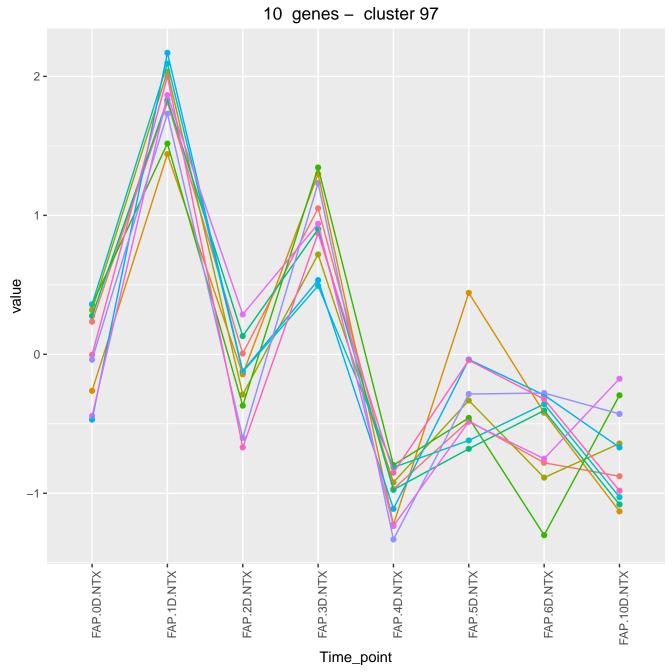


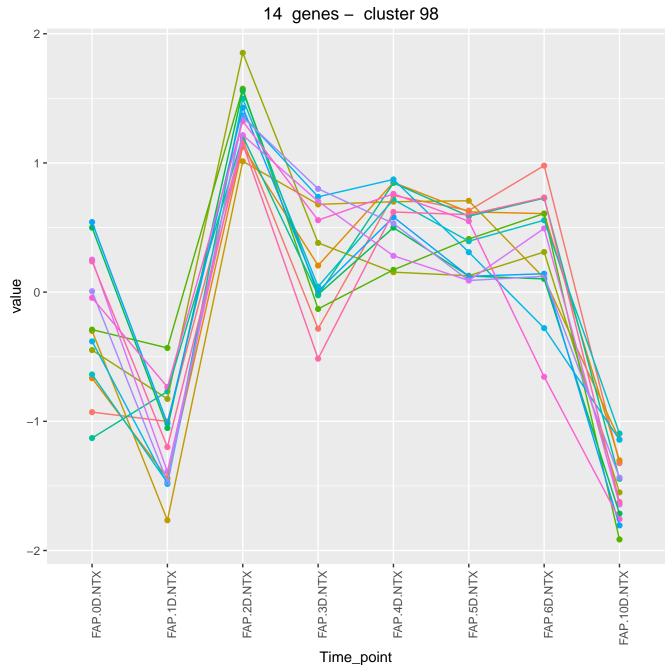


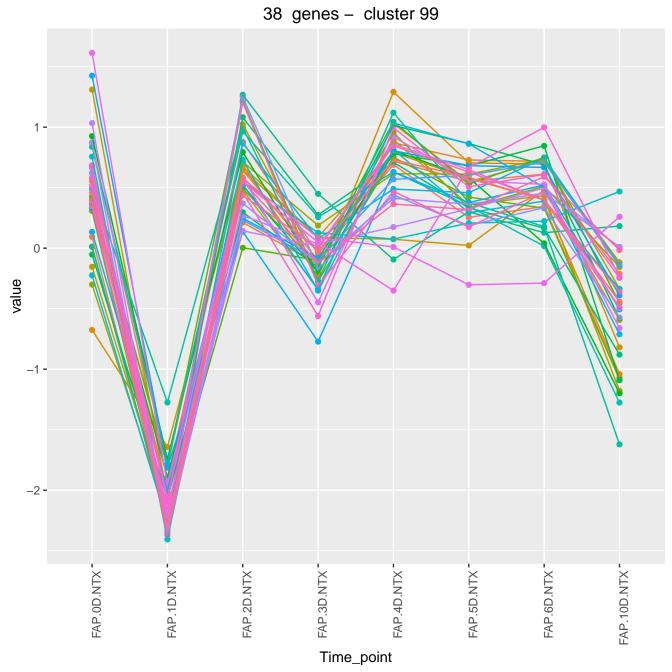


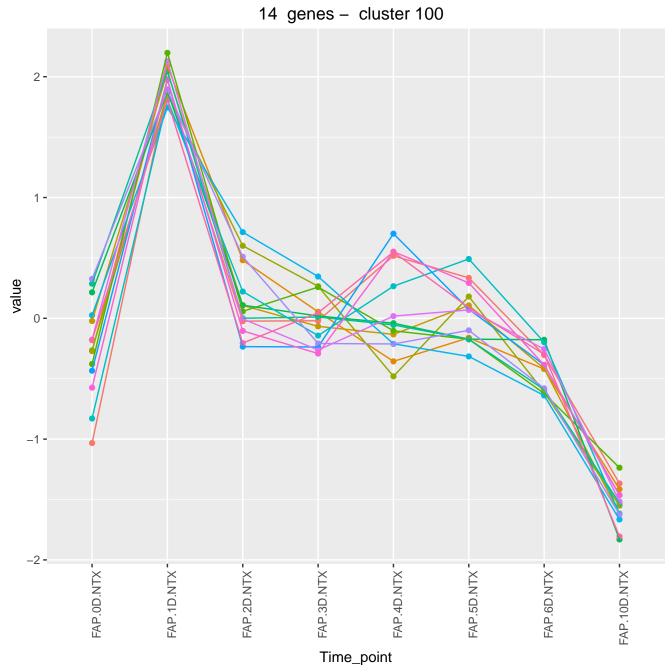


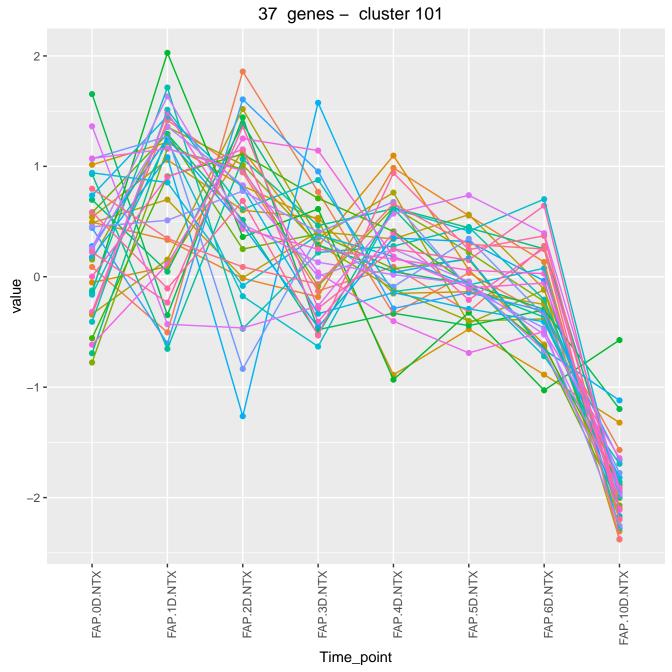


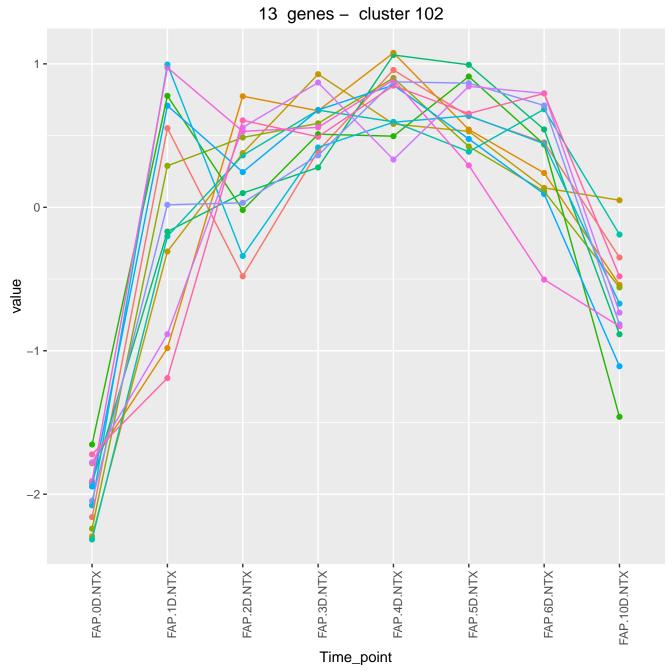


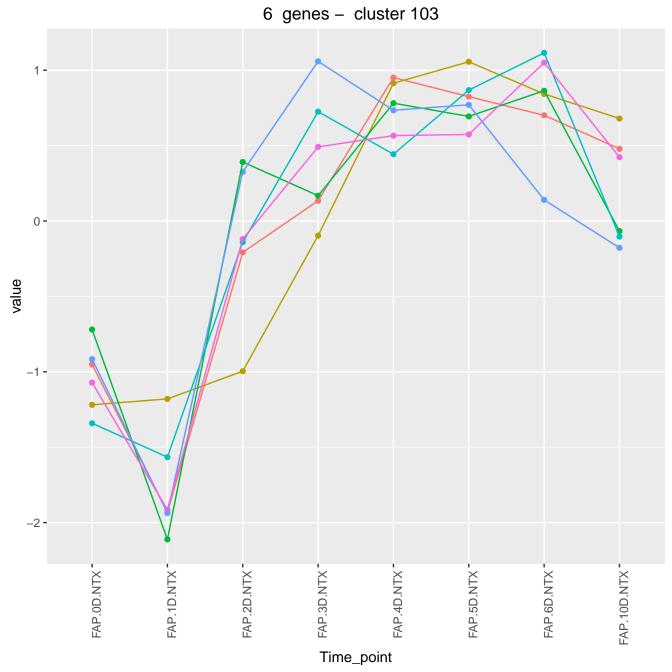


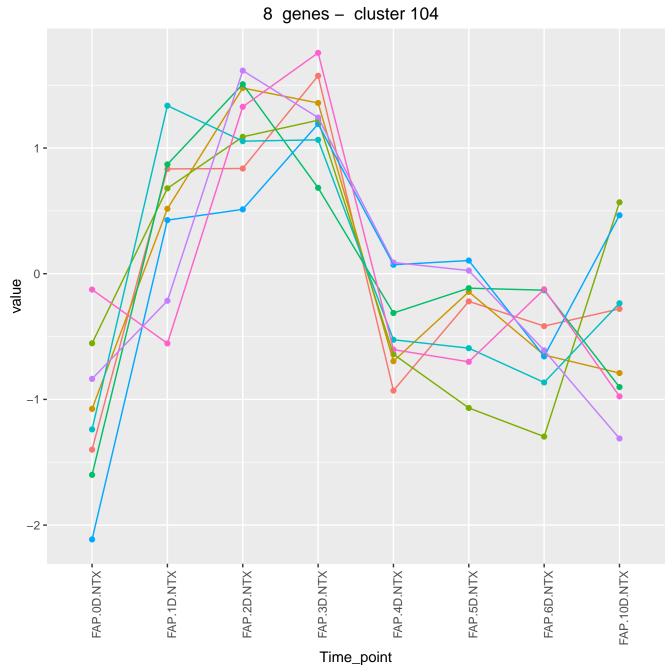


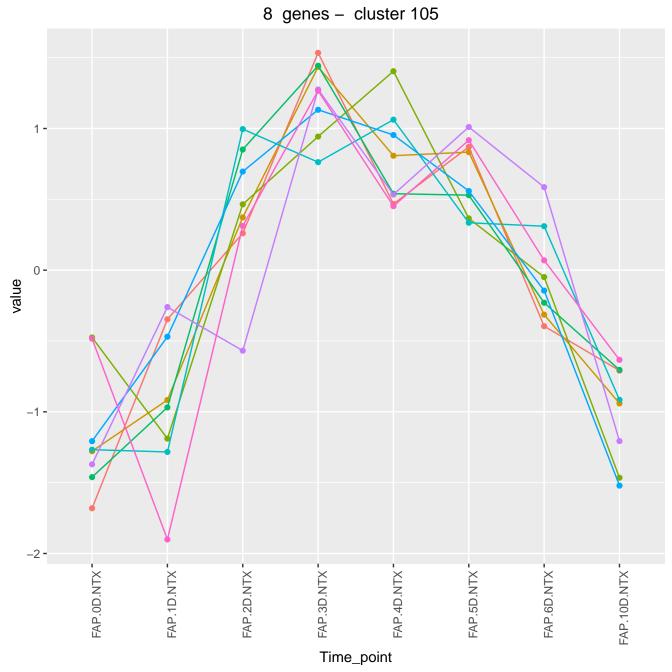


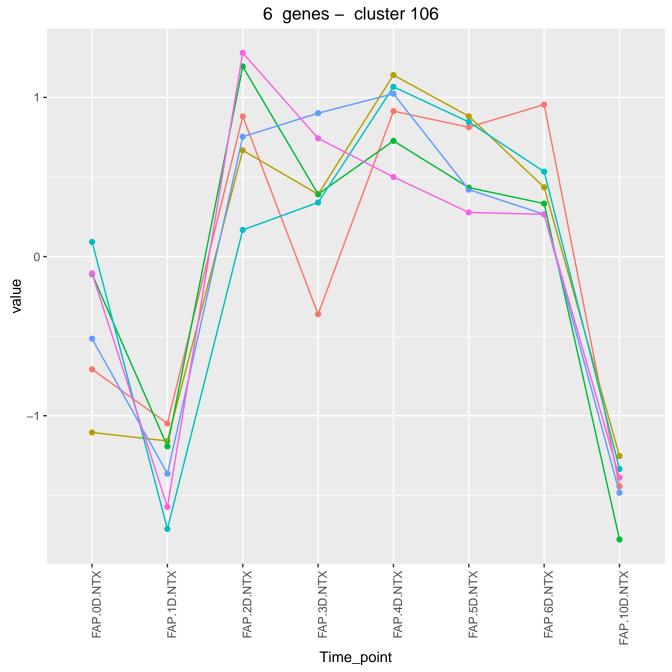


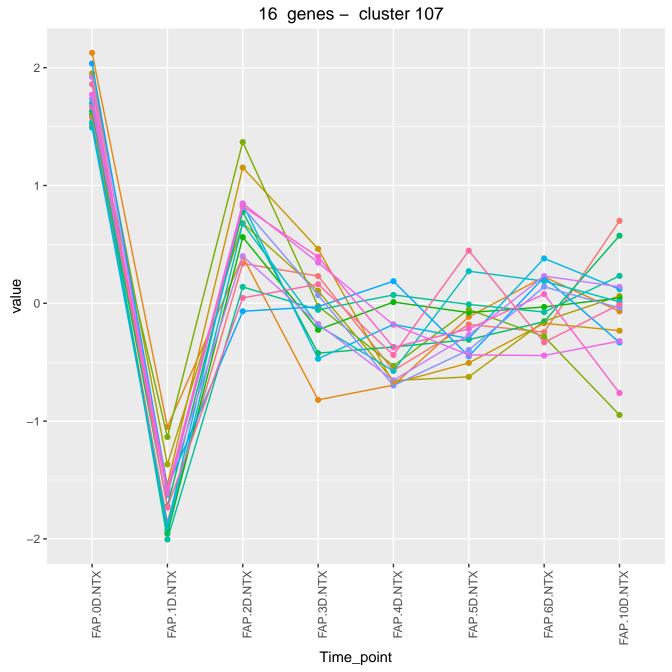


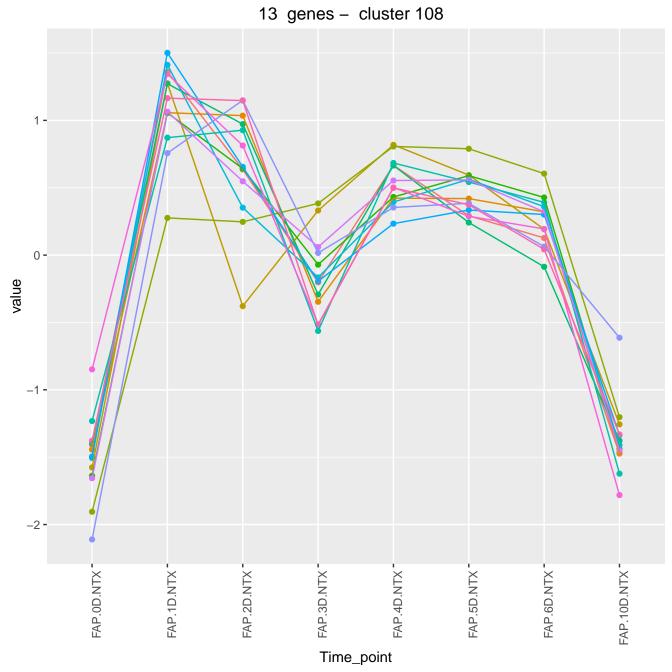


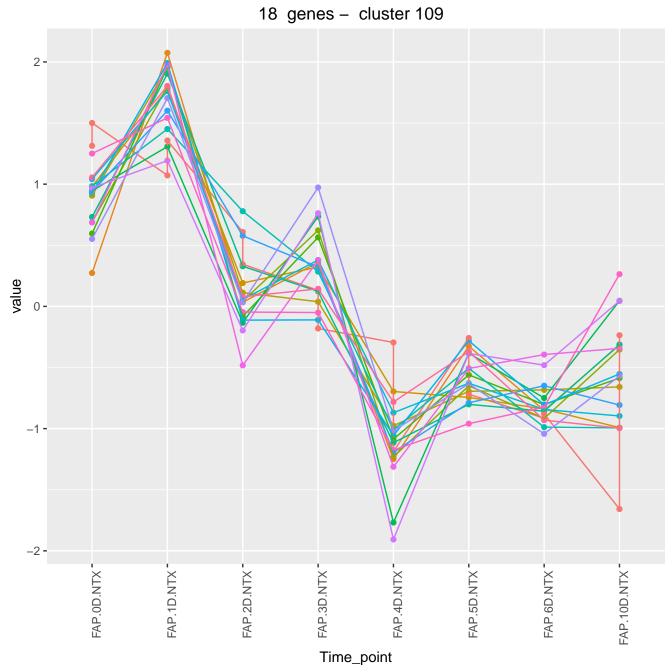


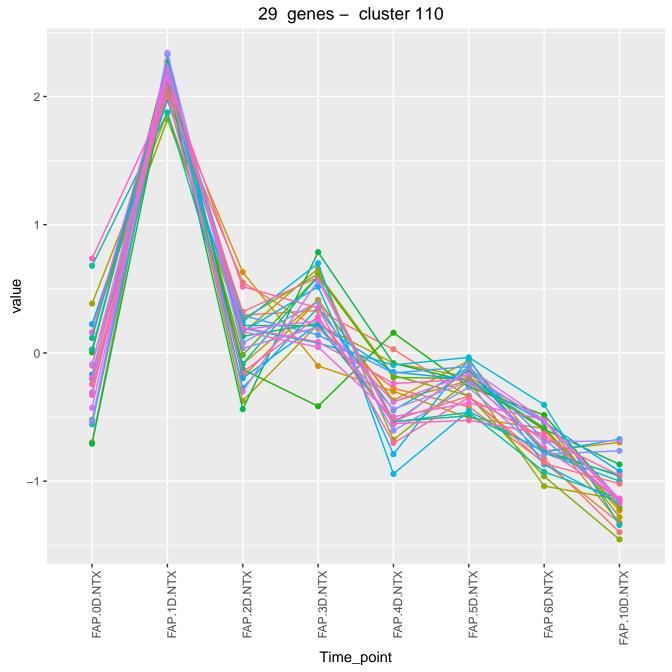


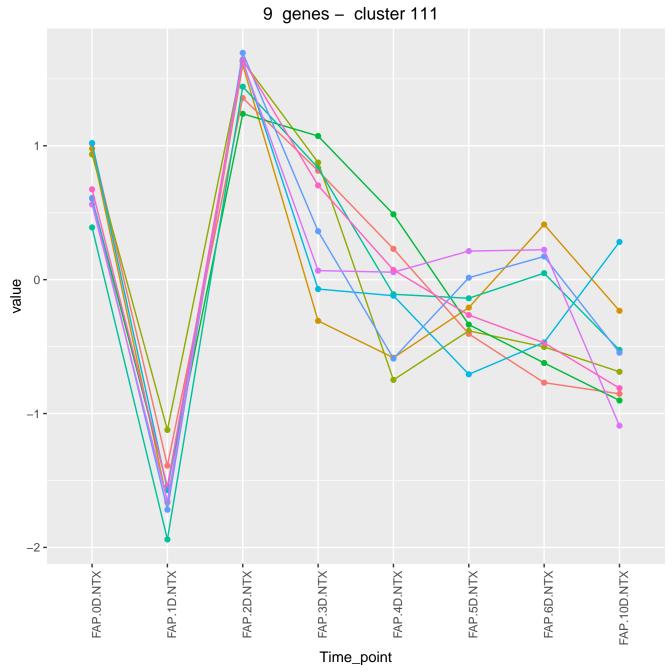


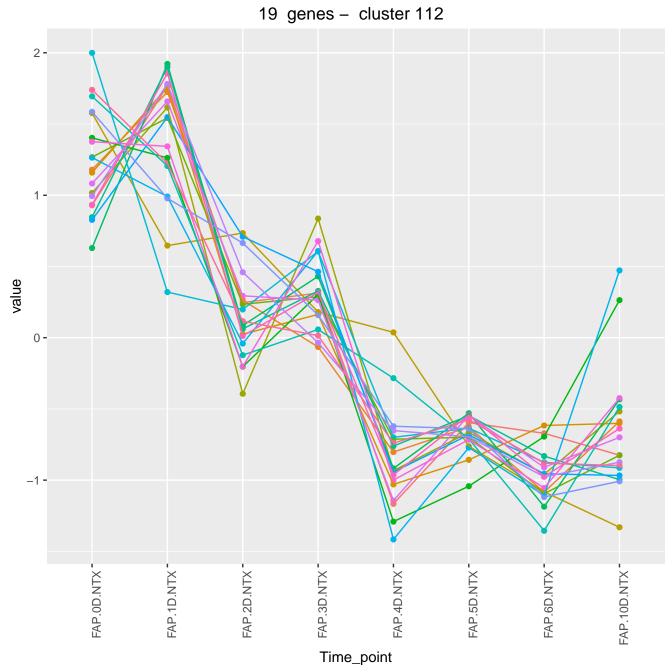


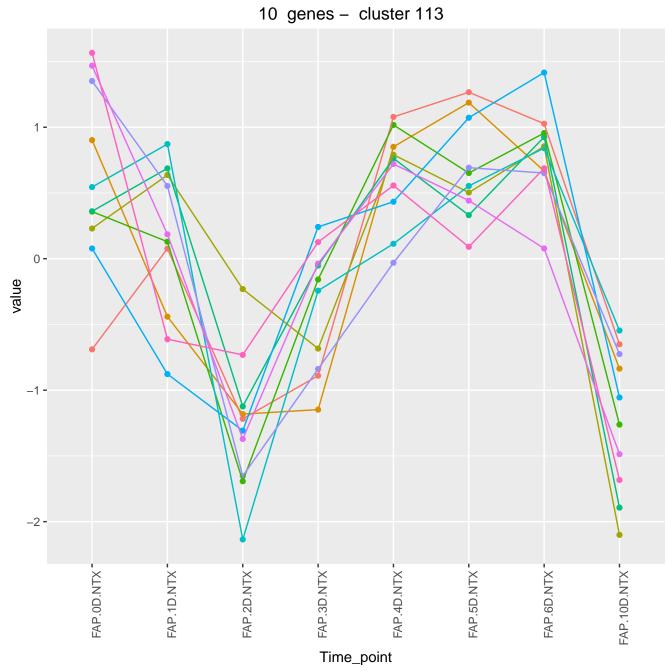


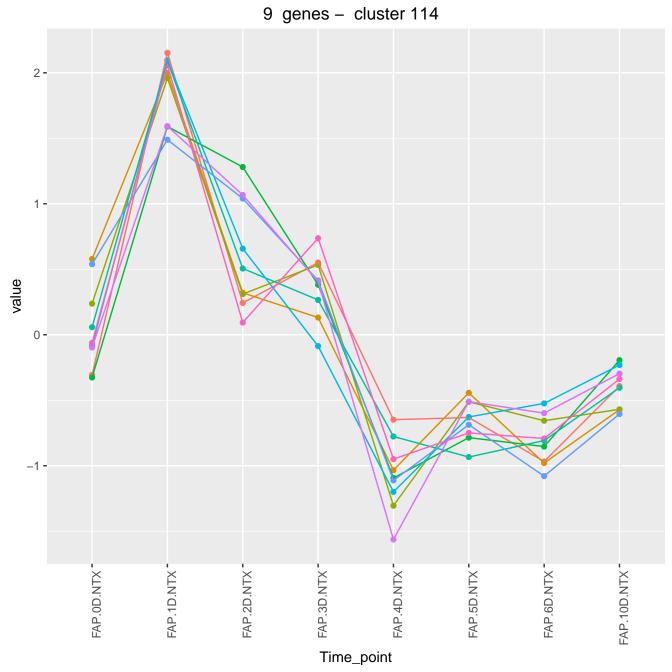


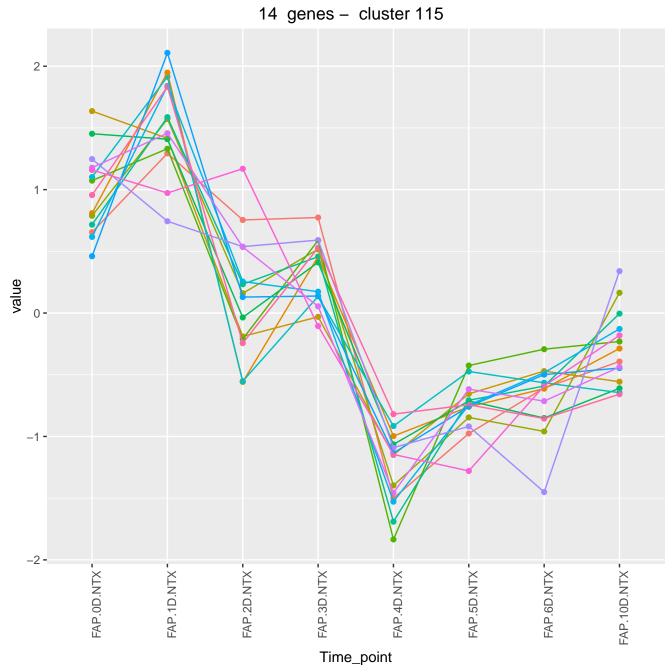


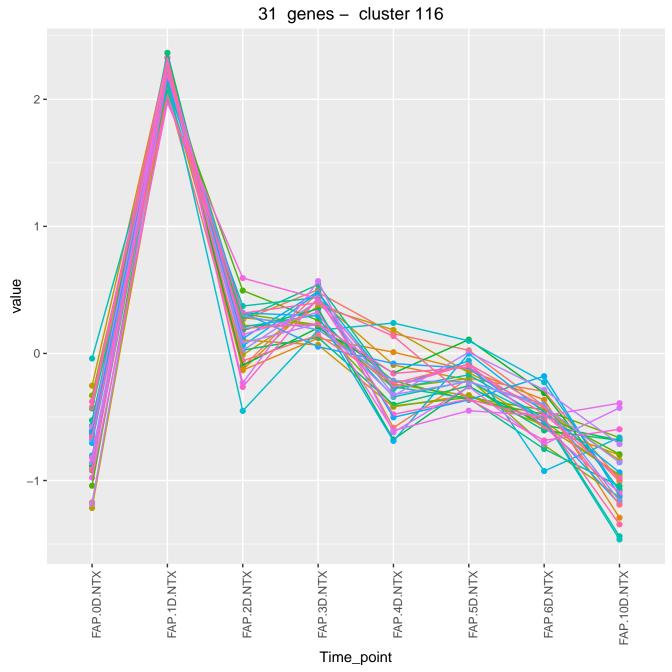


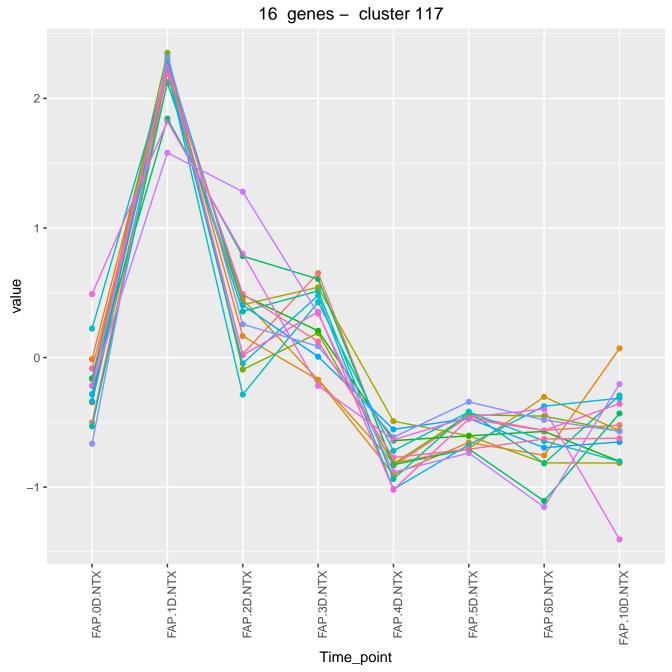


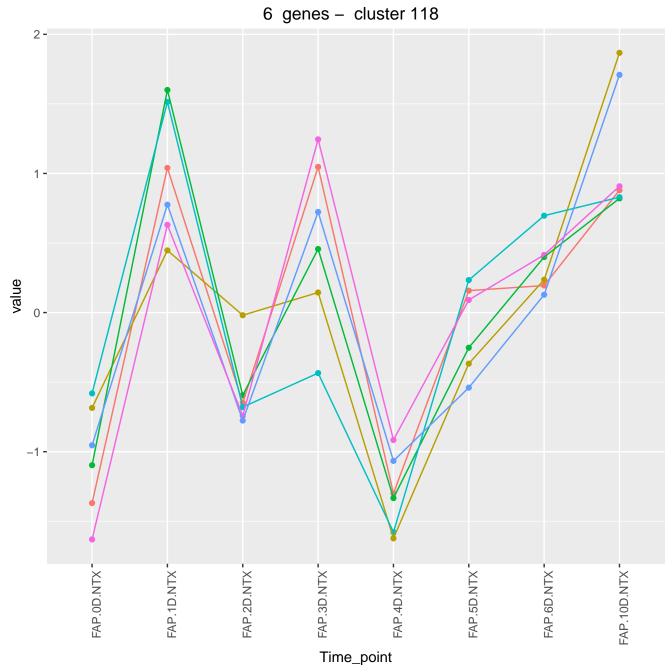


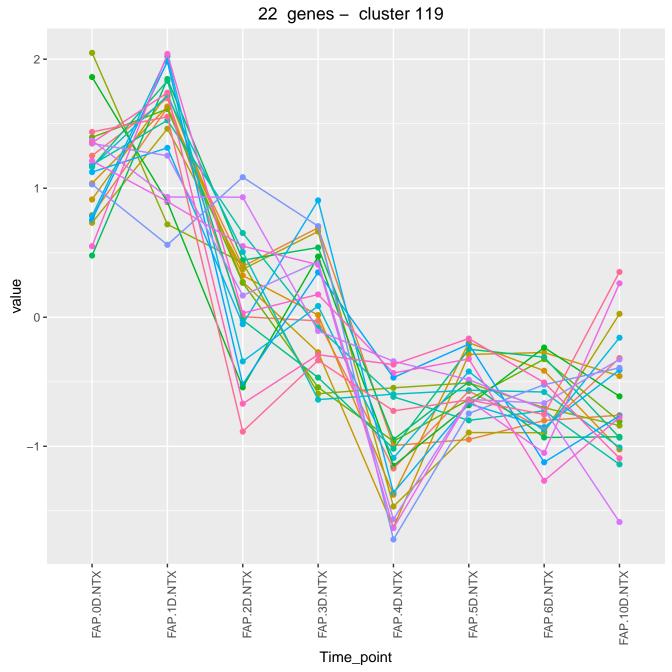


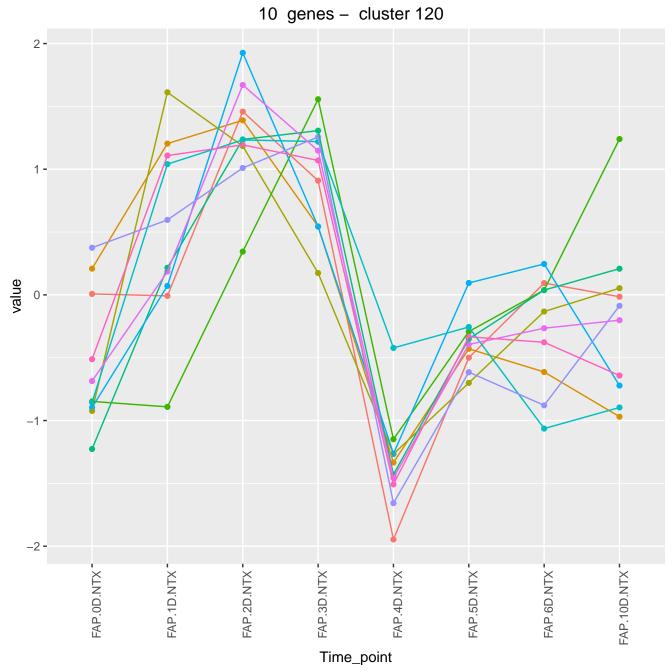


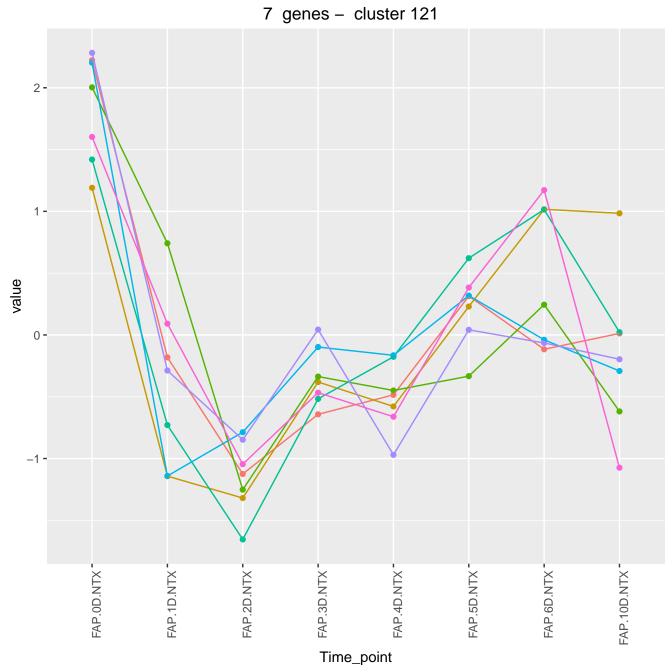


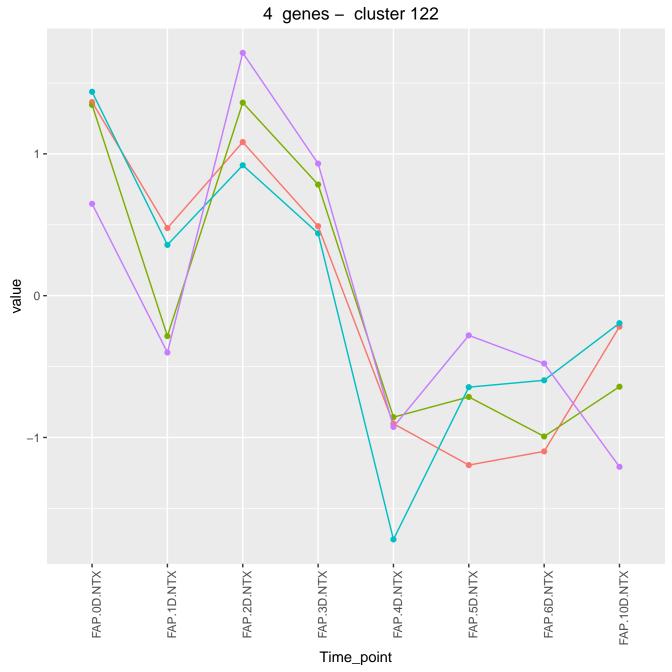


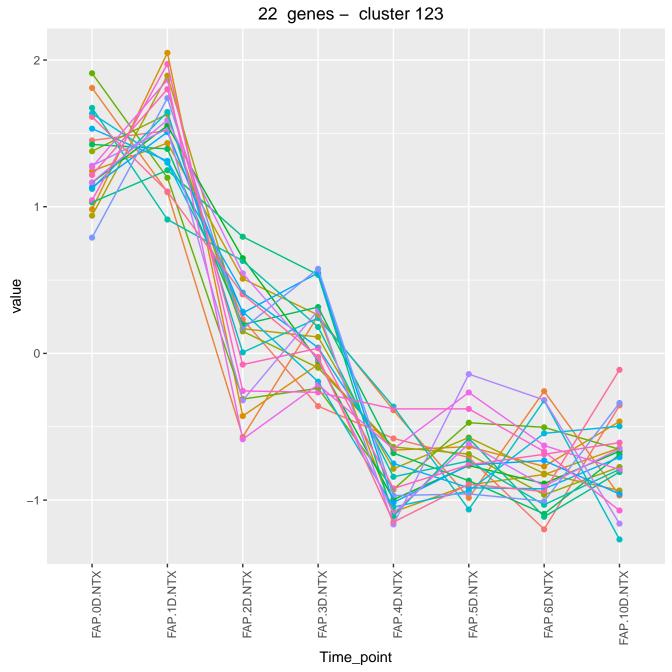


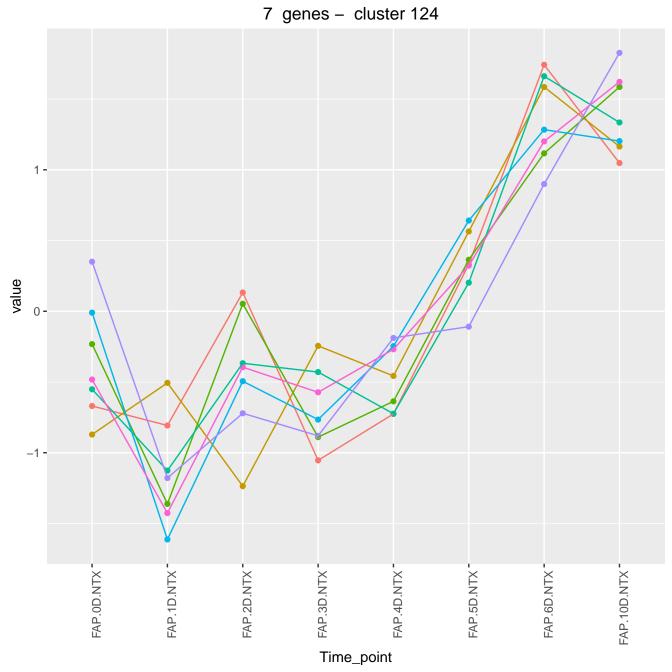


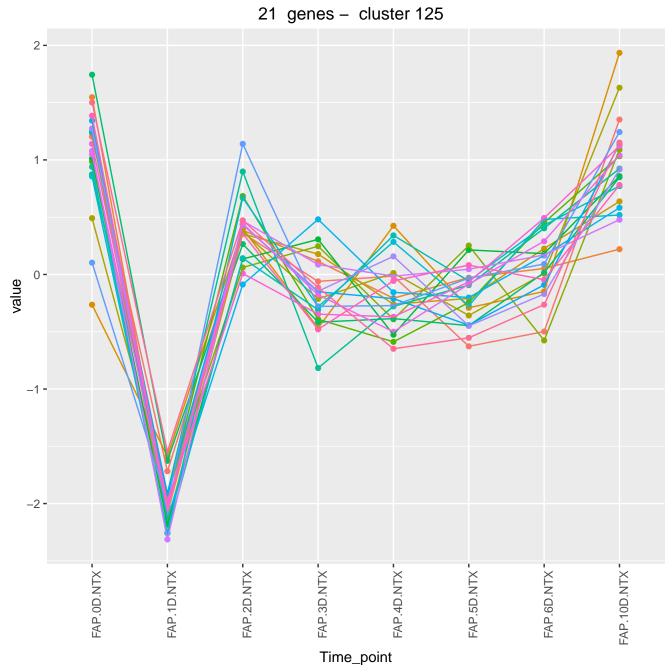


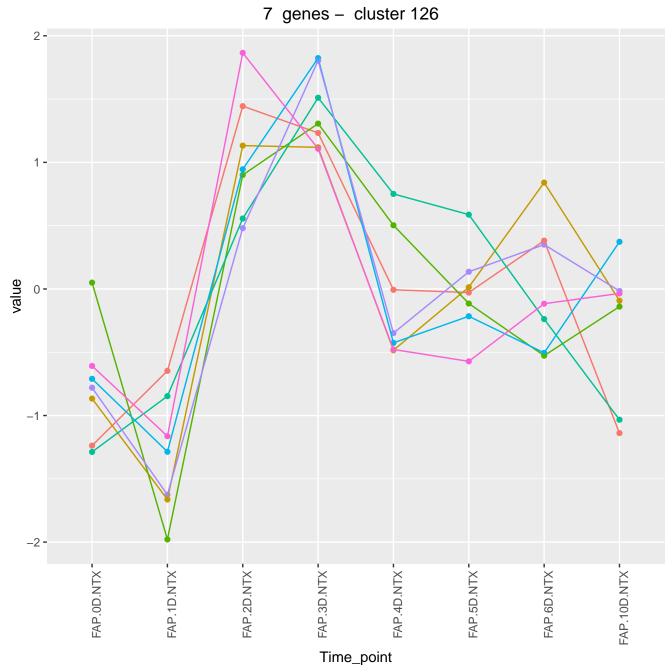




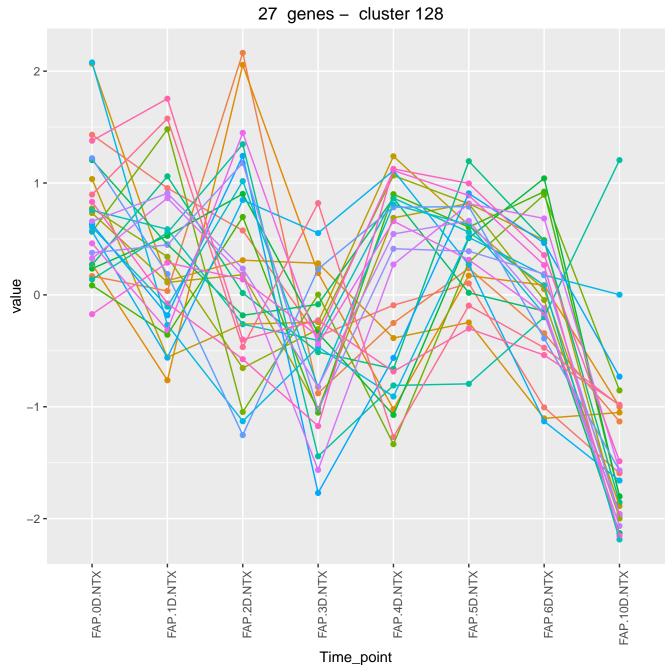


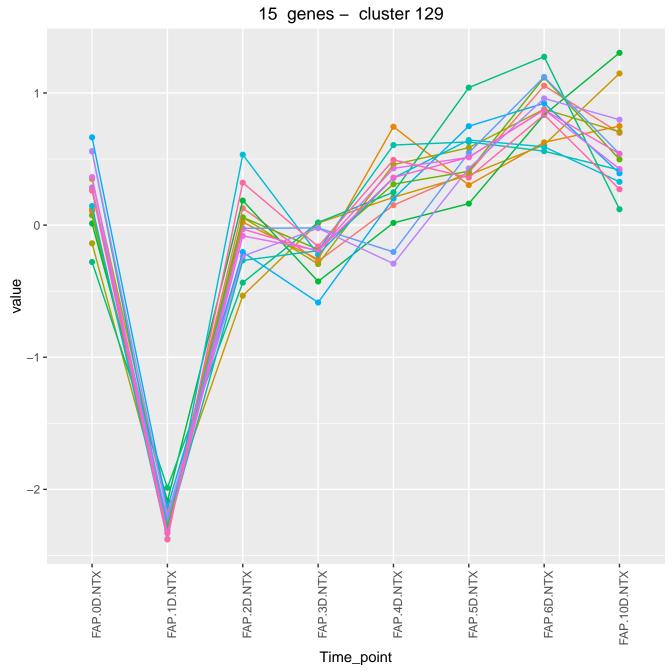


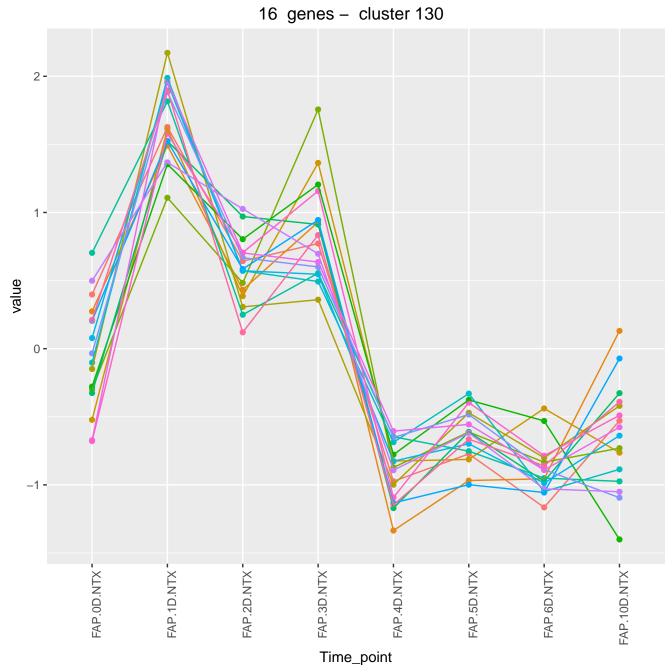


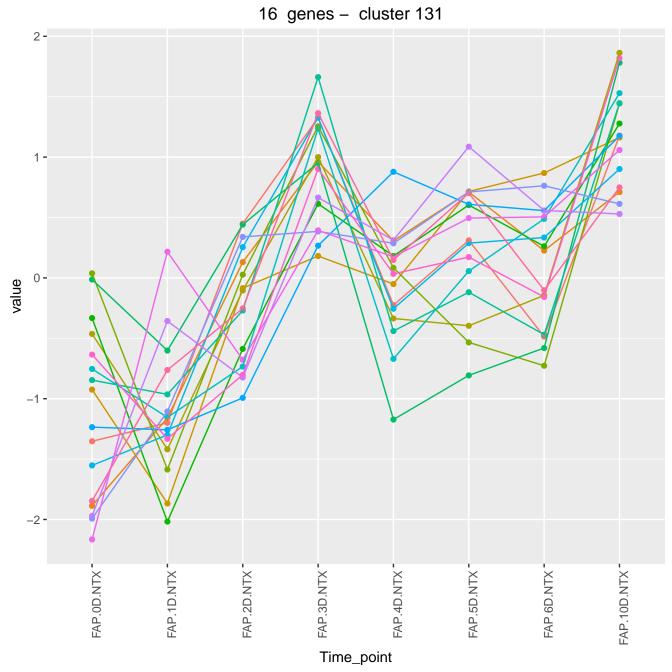


34 genes - cluster 127 2 -1 value 0 -\_1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.3D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

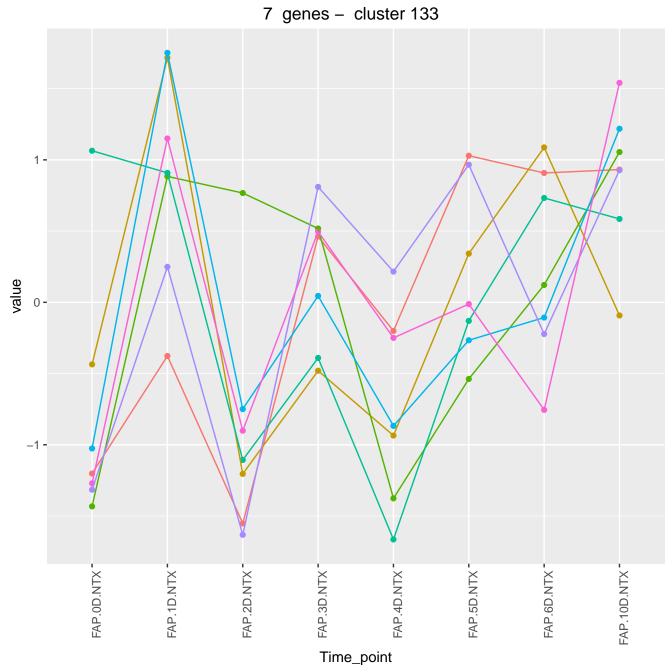


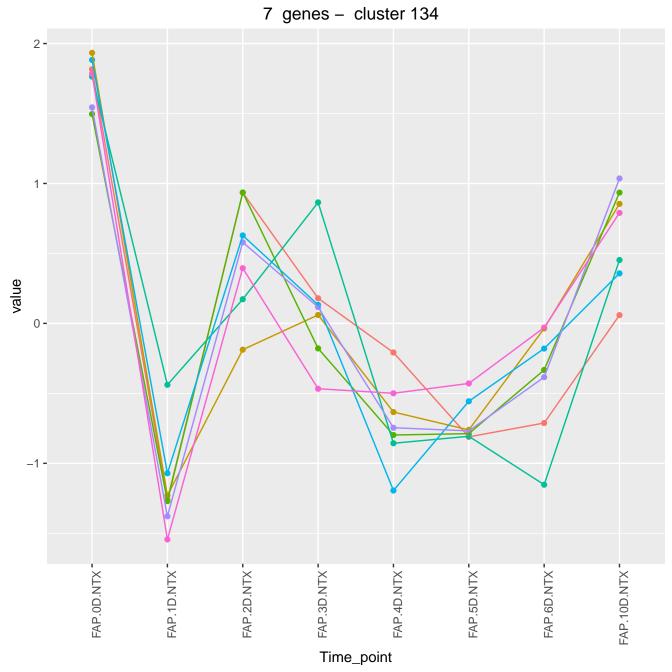


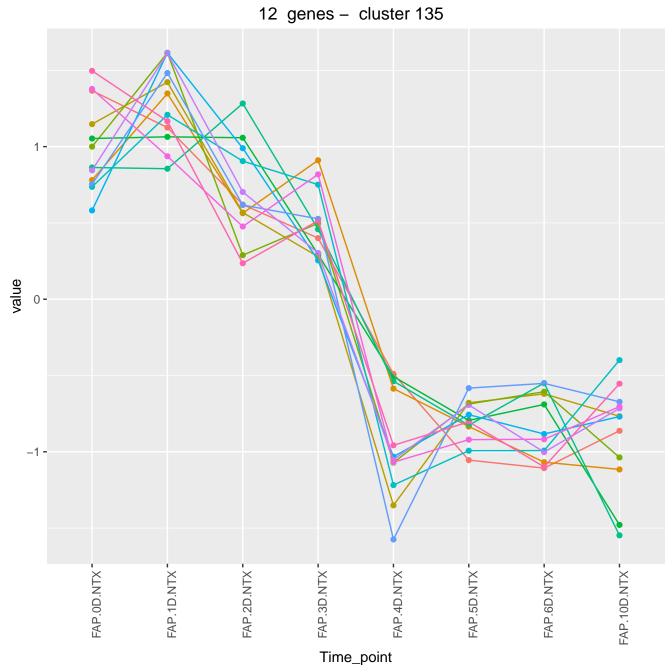


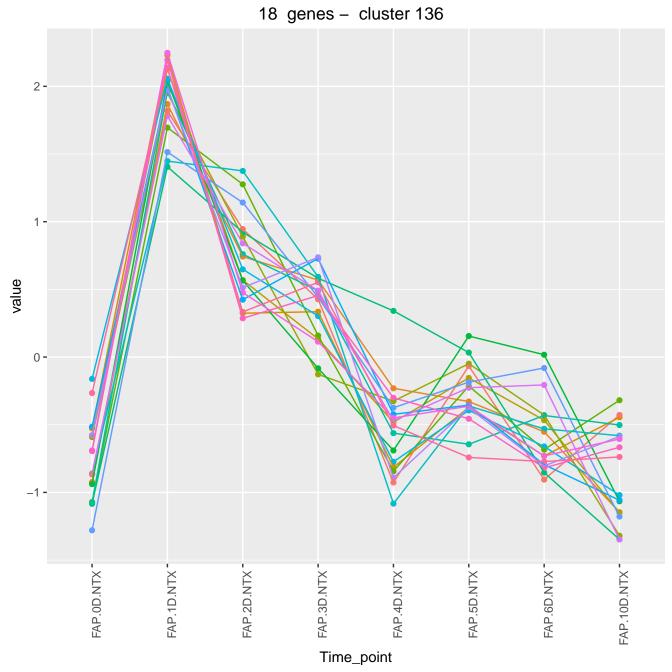


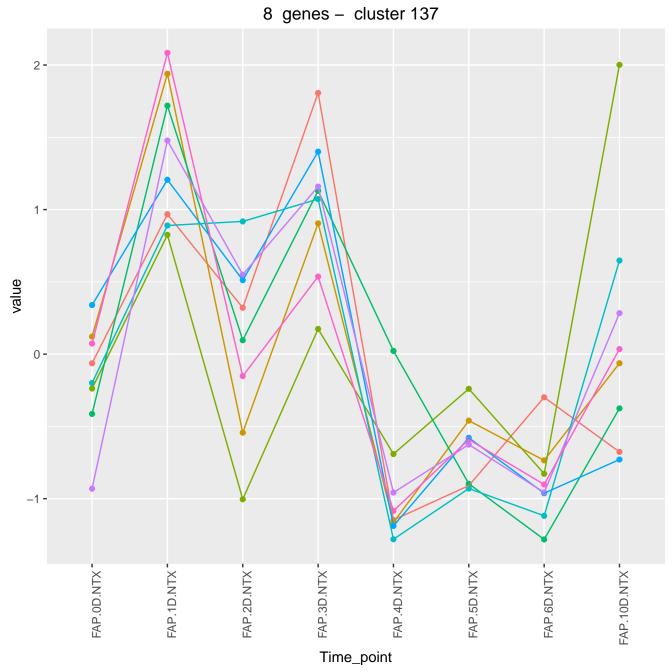
26 genes - cluster 132 2-1 value 0 -\_1 **-**-2 **-**FAP.0D.NTX 7 FAP.2D.NTX -FAP.3D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.4D.NTX FAP.6D.NTX -Time\_point

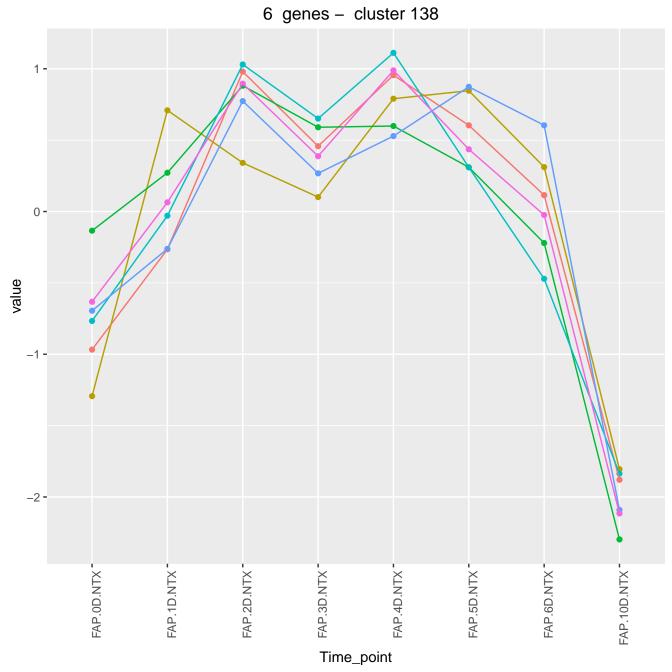


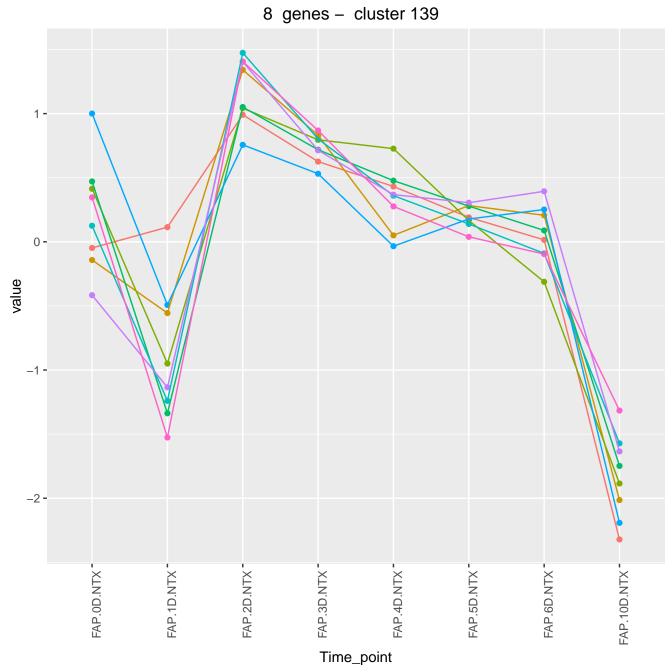


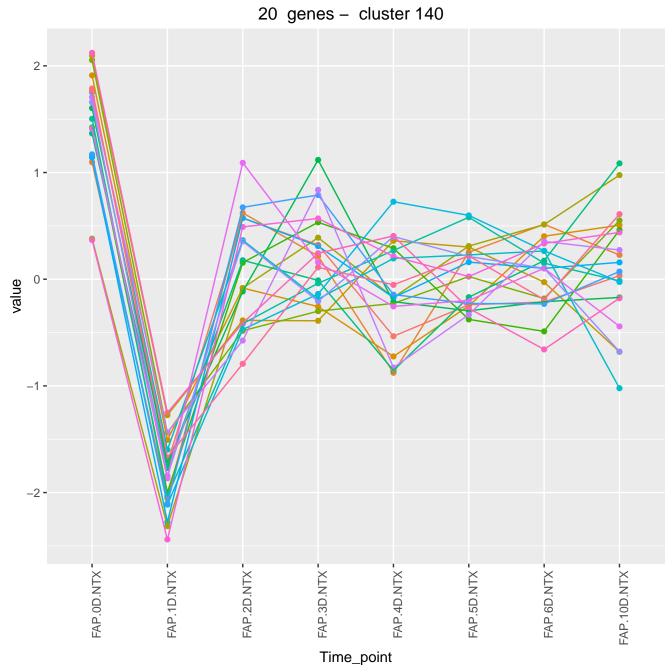


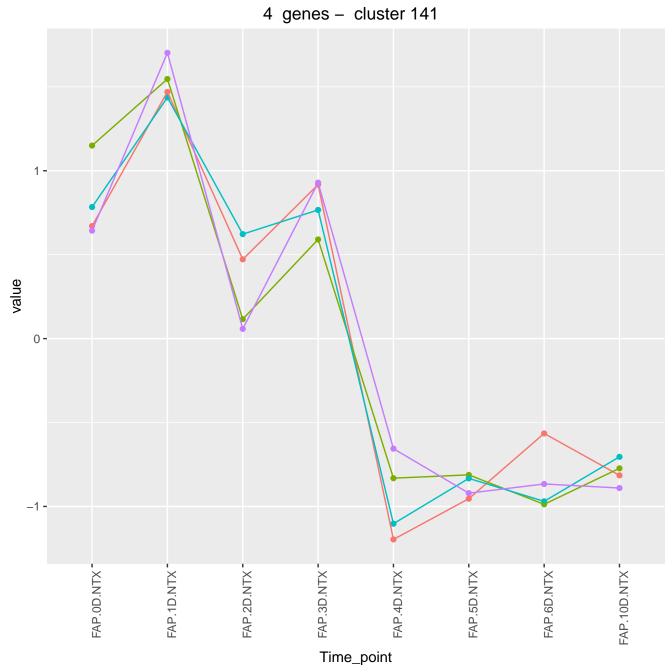


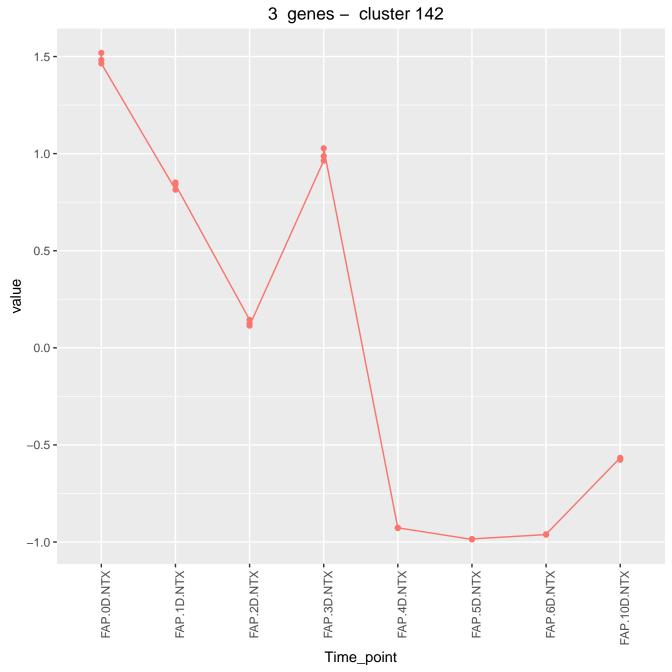


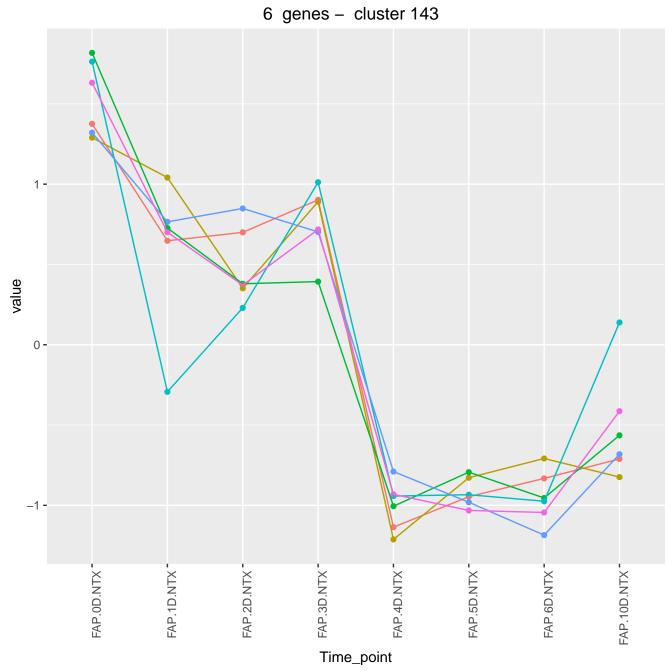


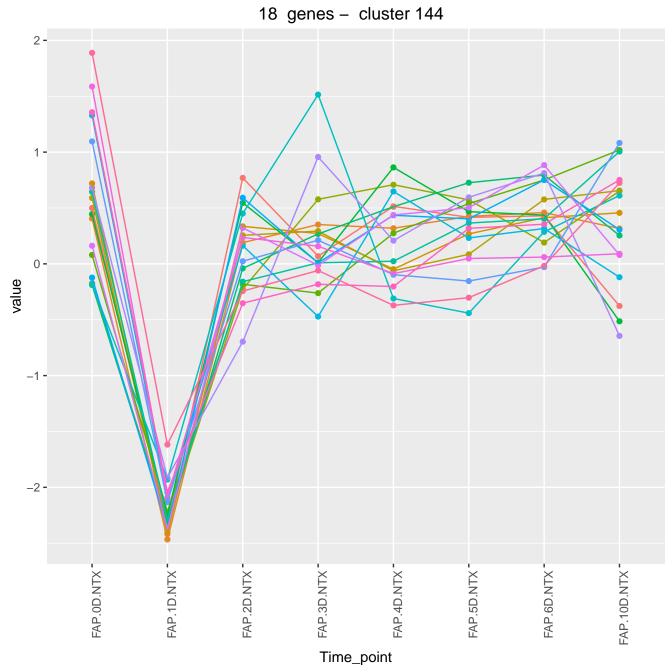


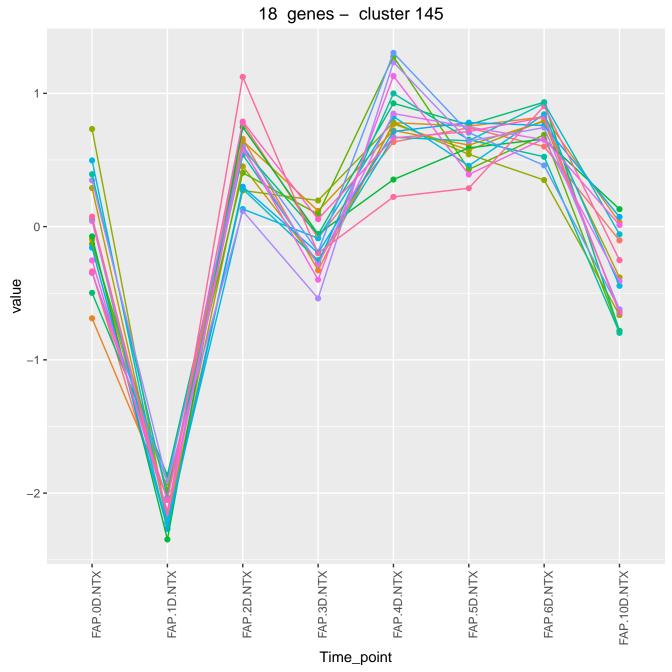


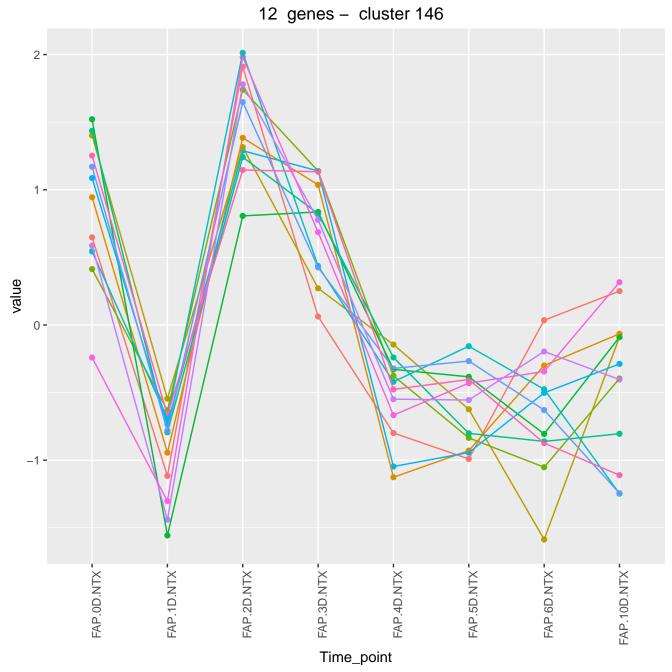


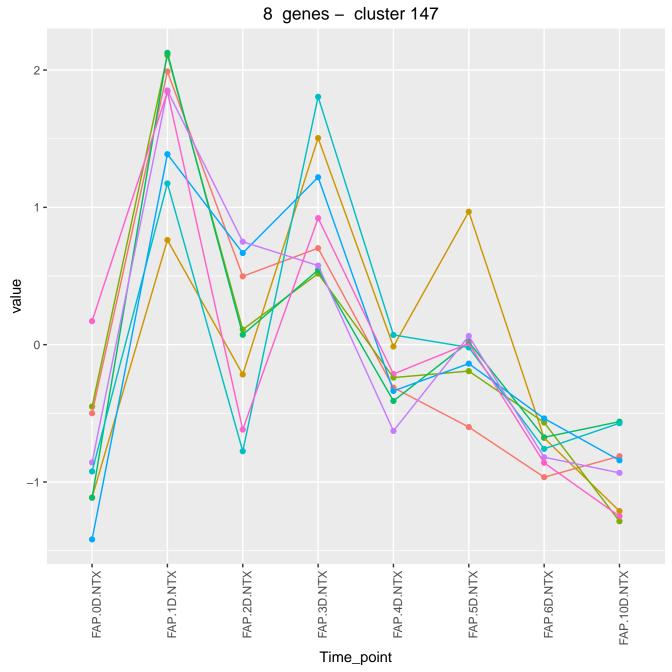


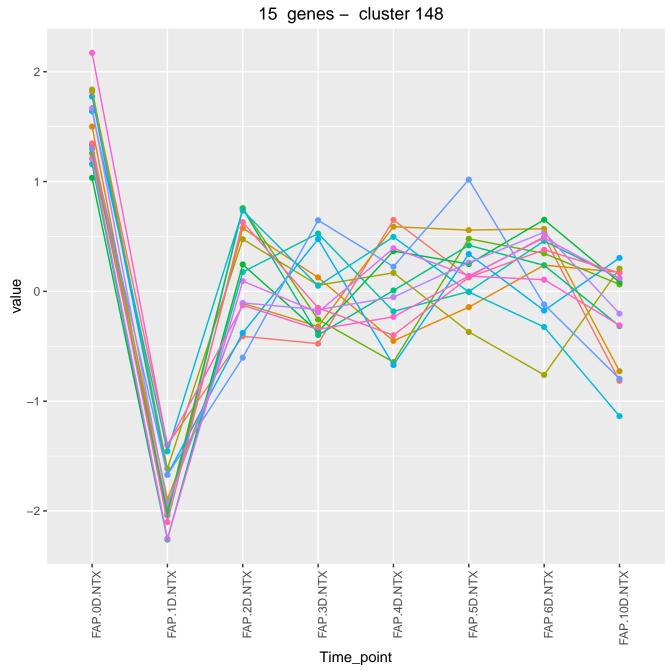


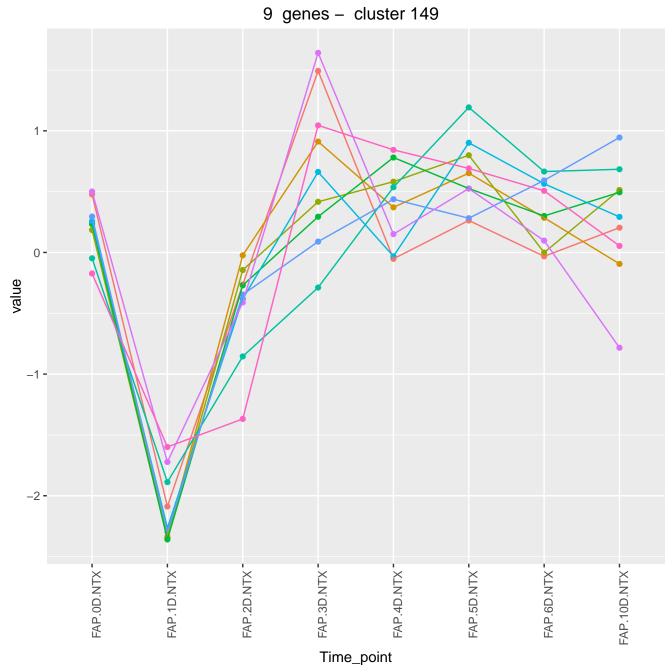


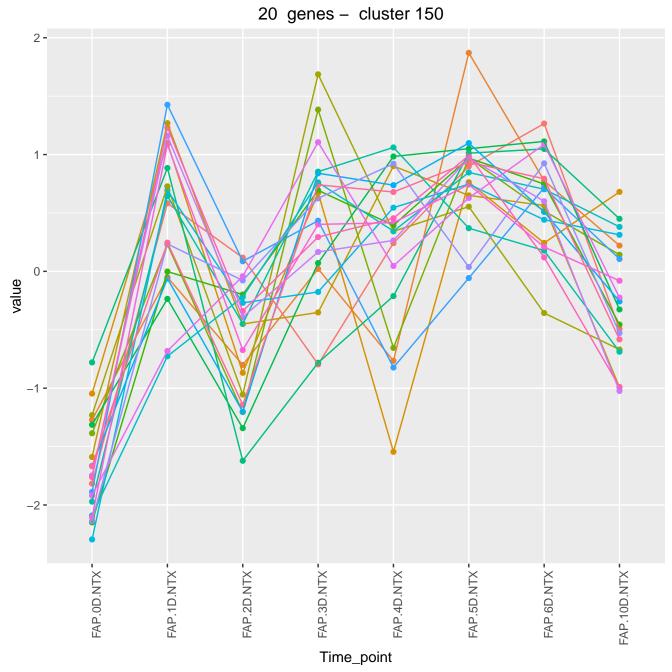


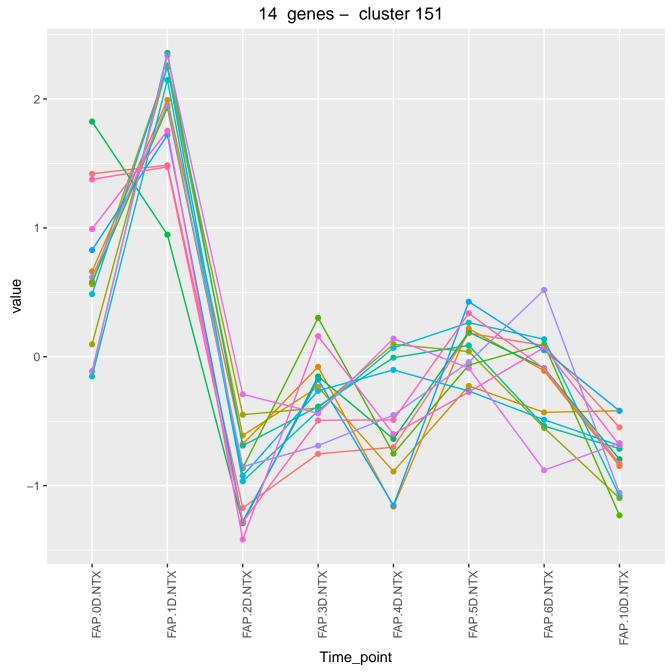


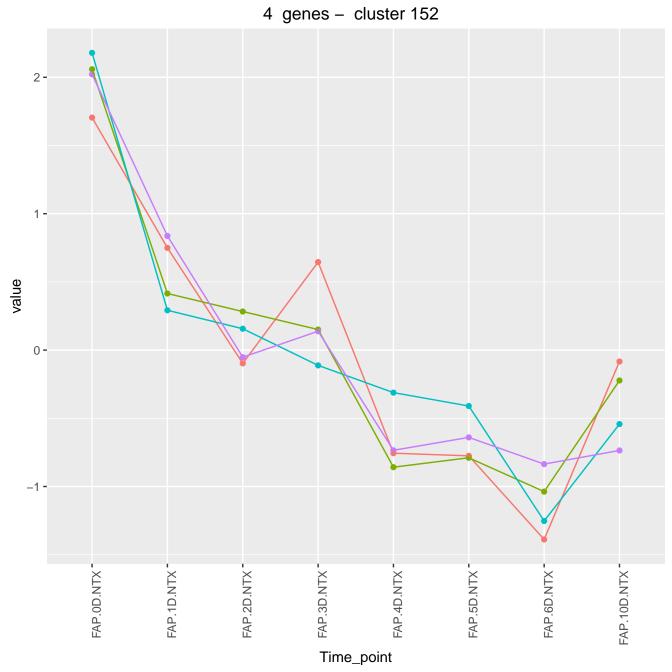


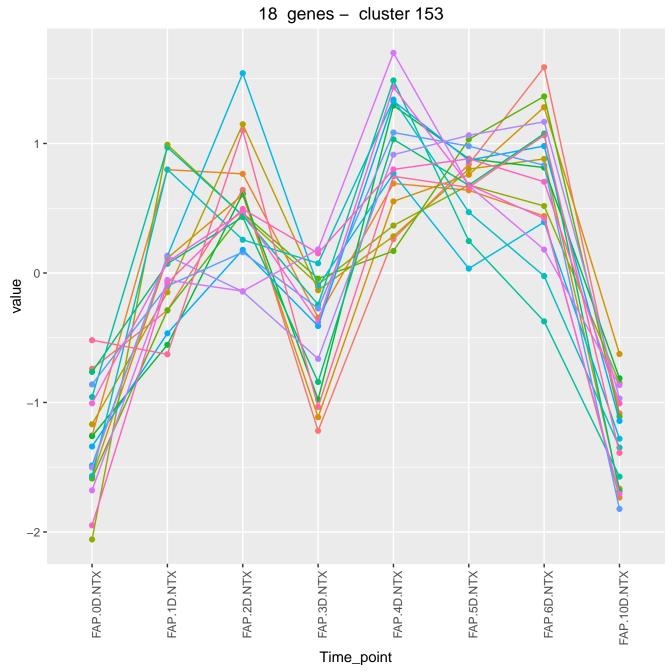


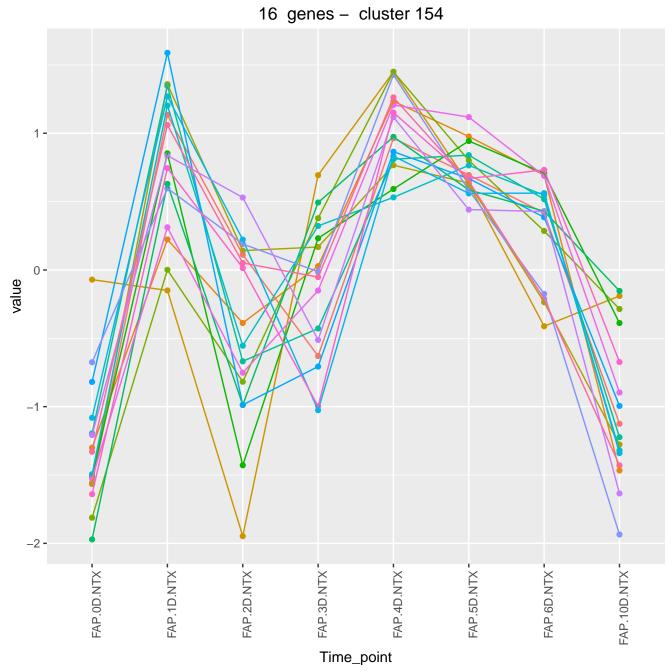


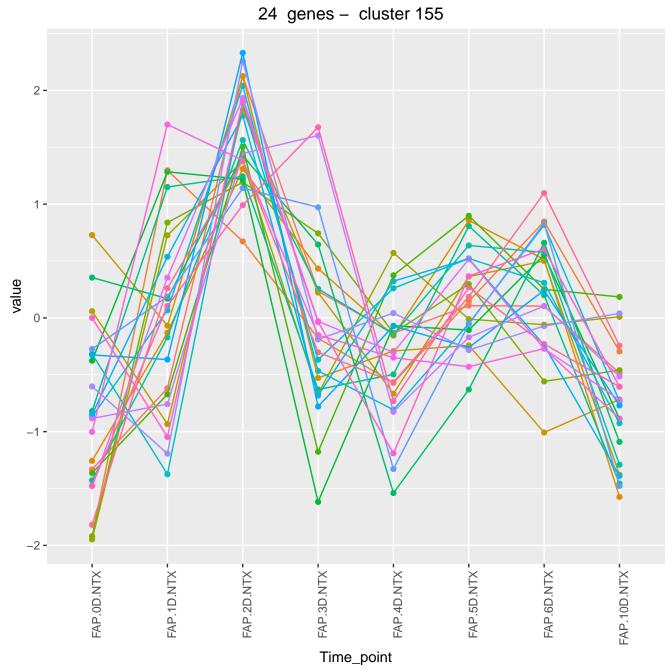


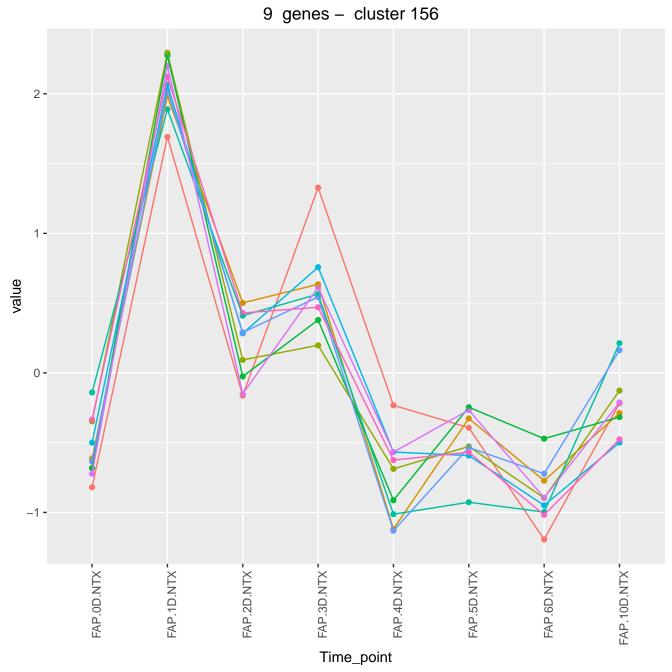


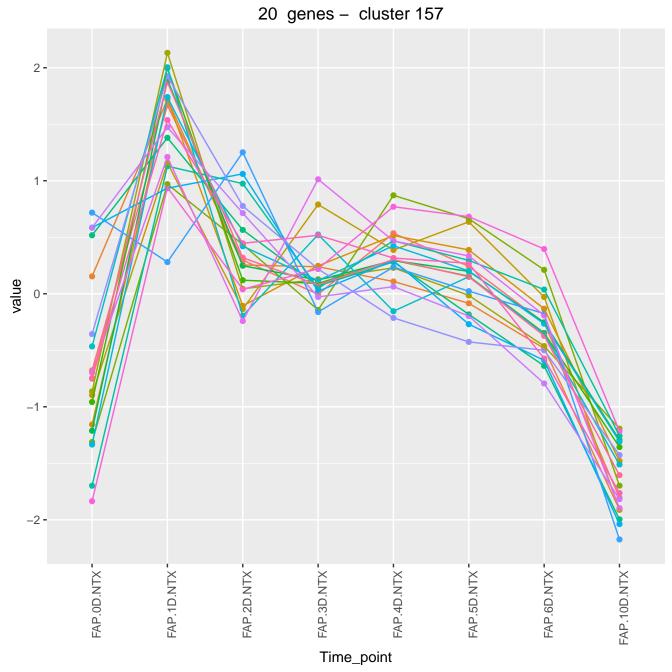


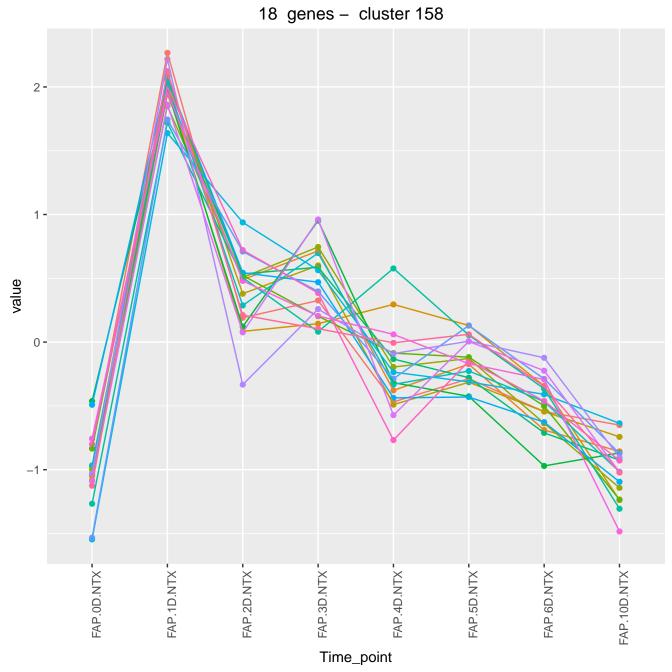


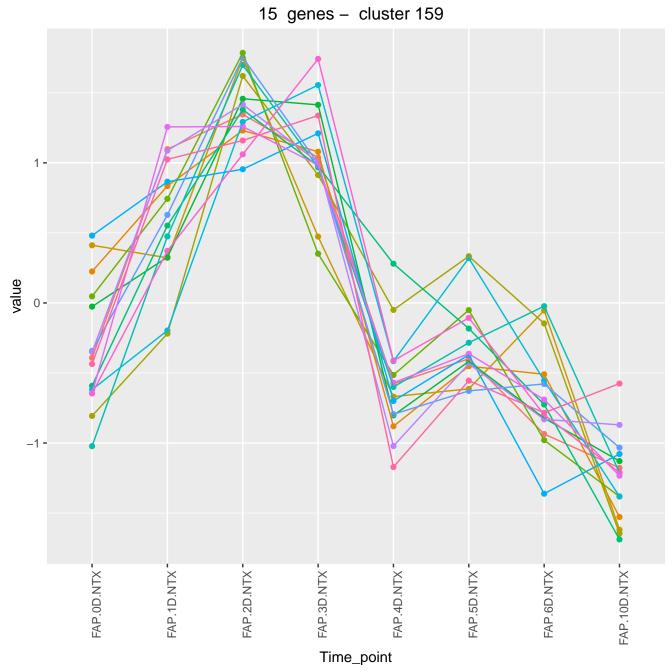


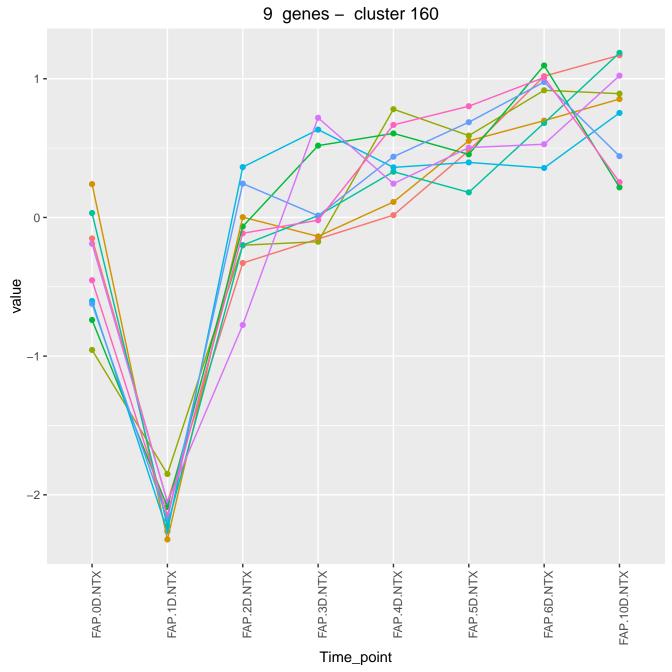


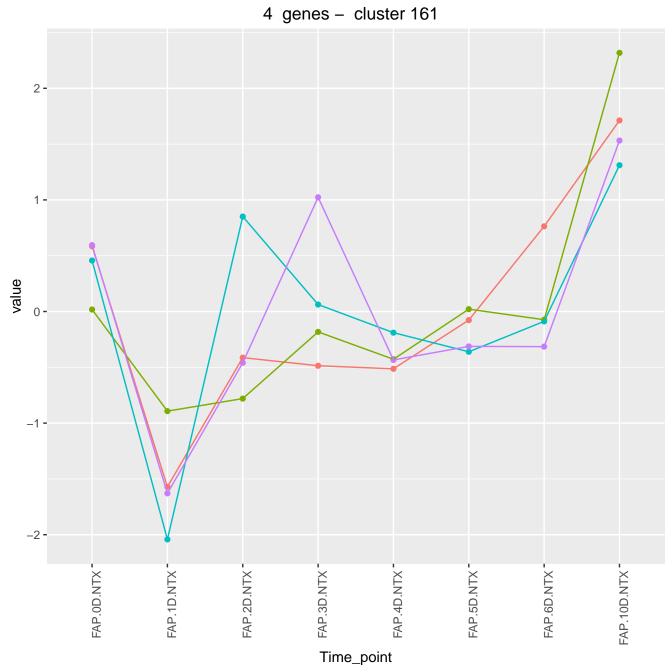


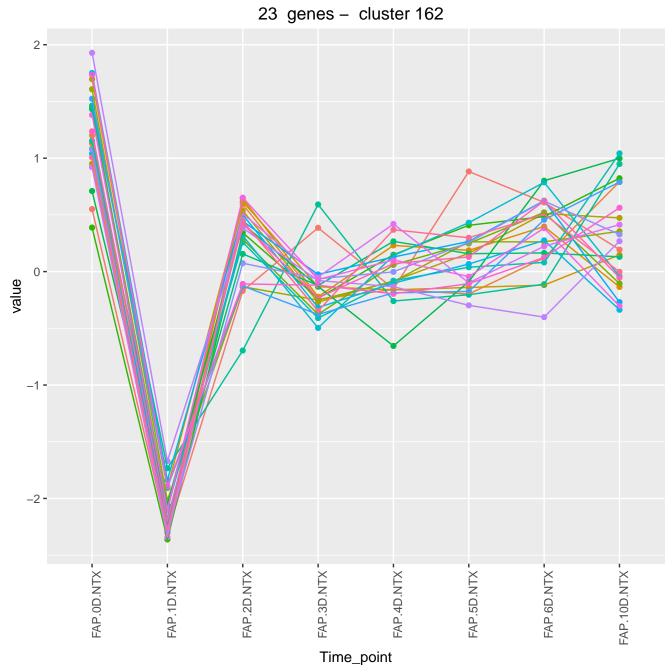


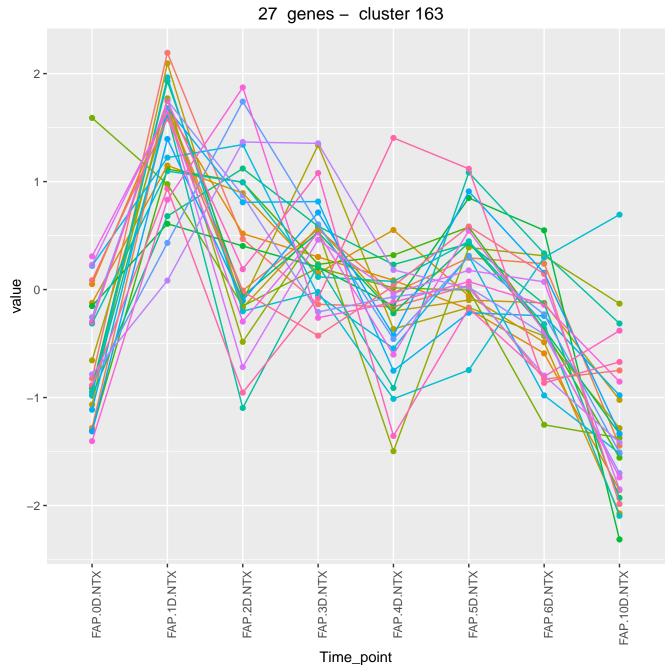


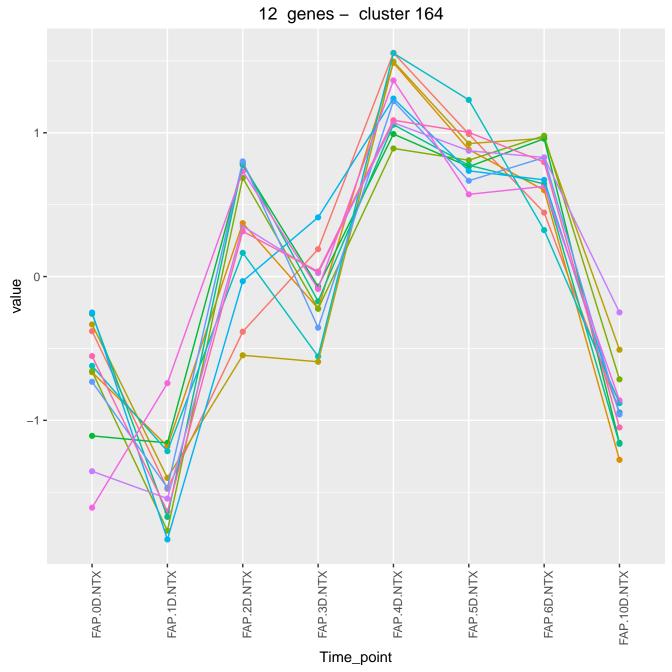


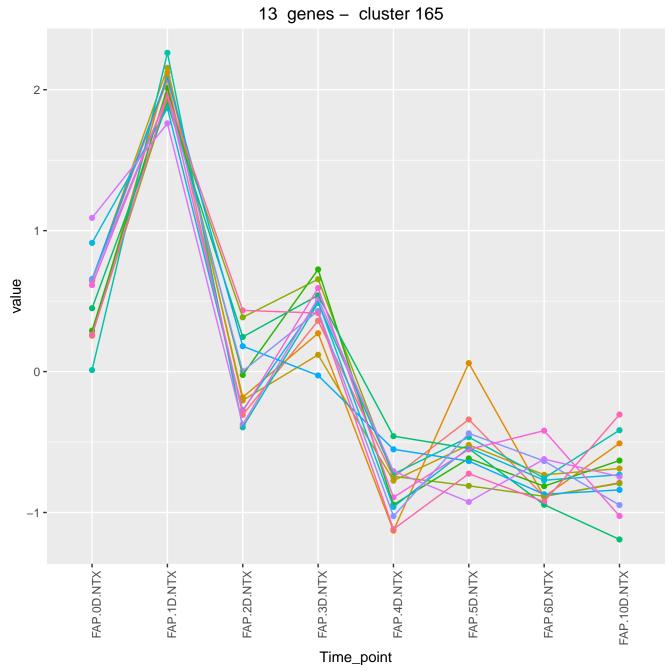


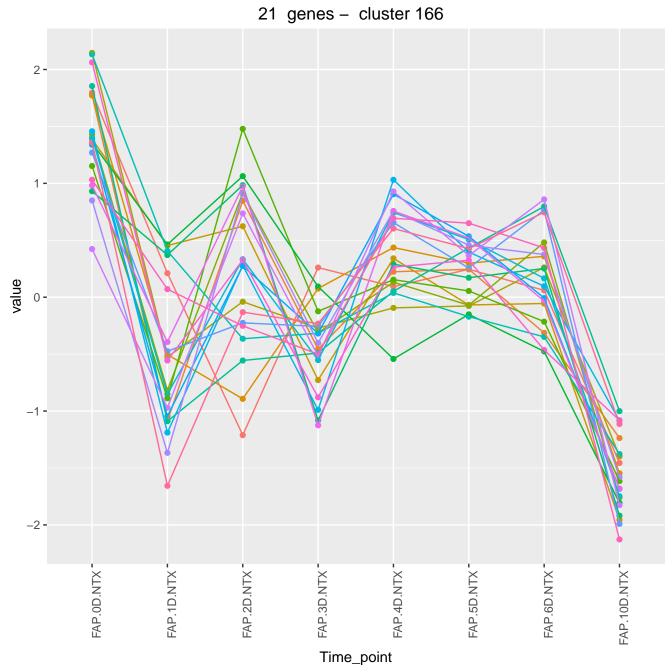


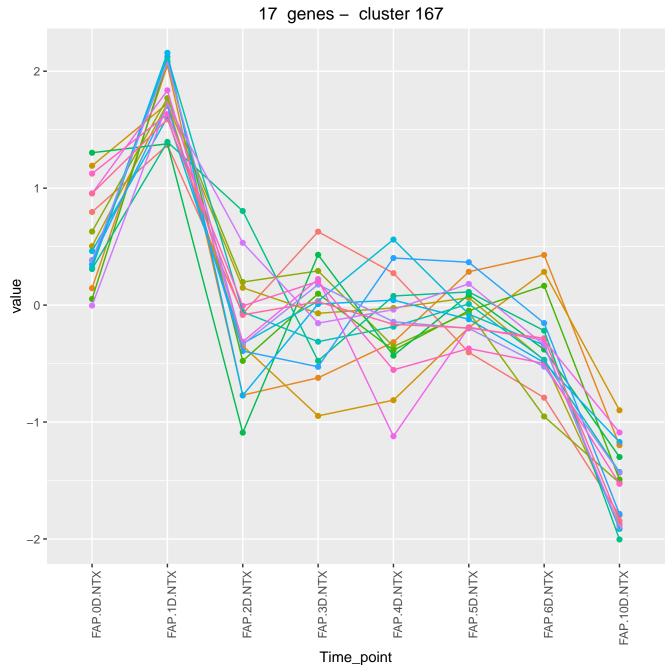


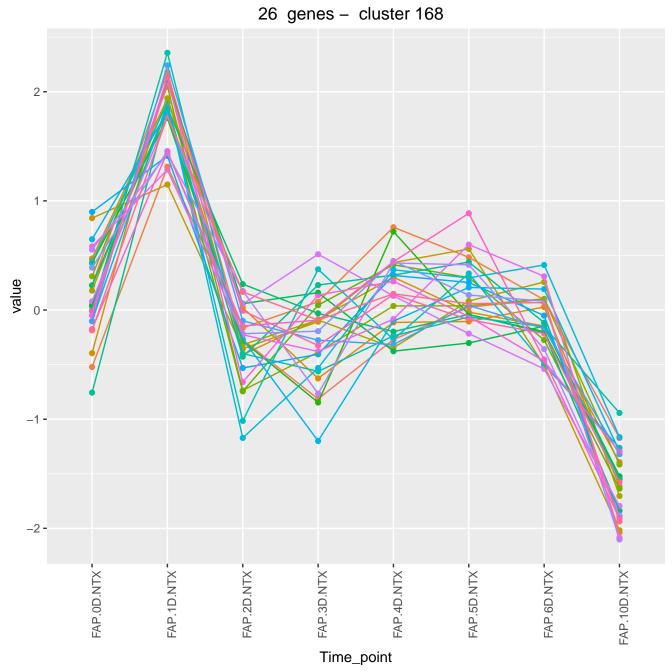


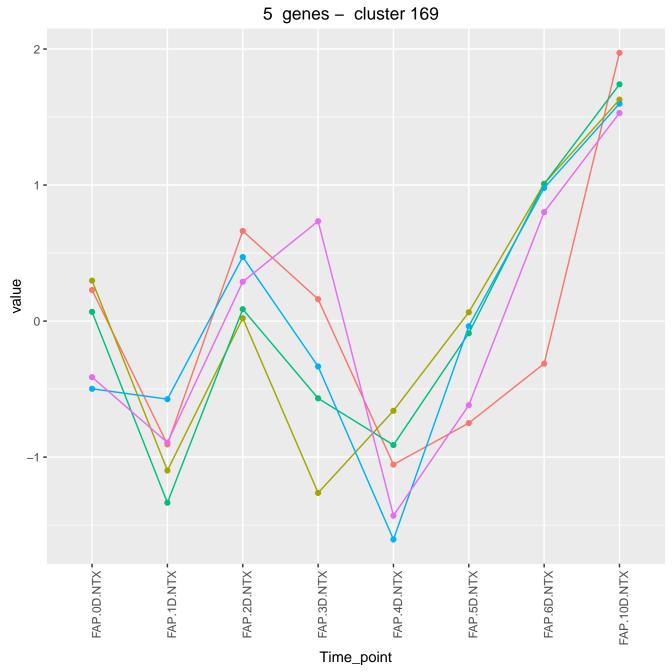


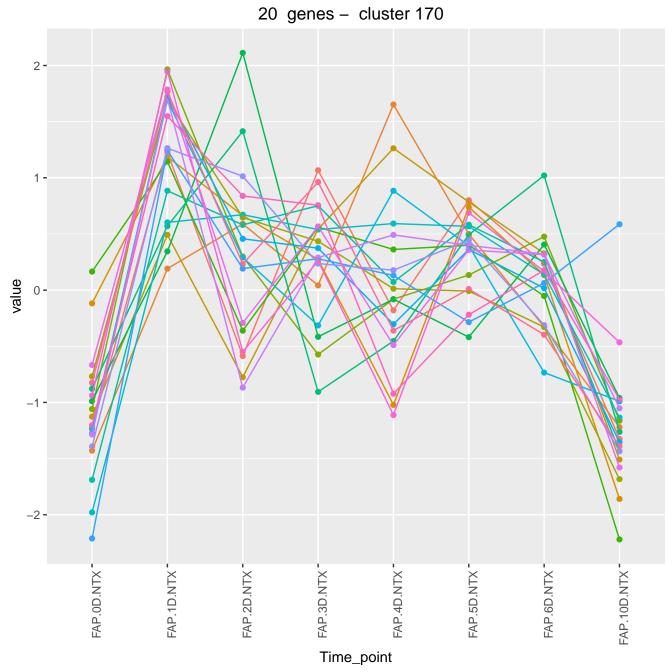


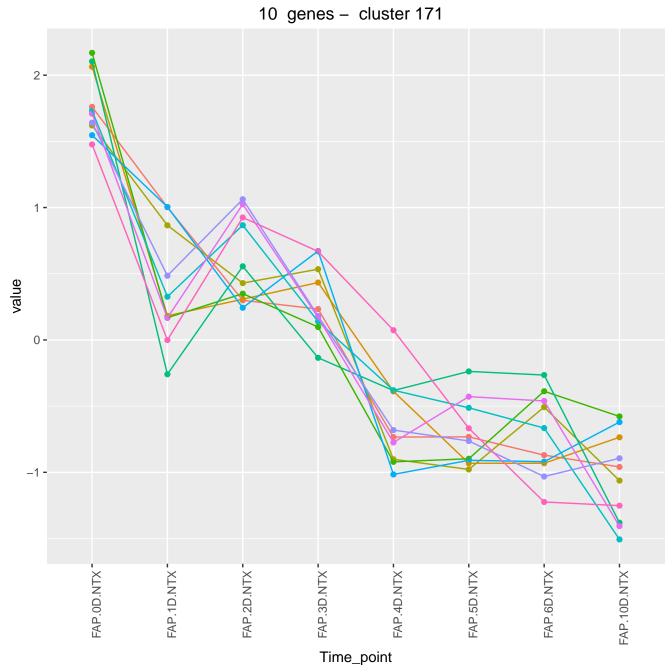


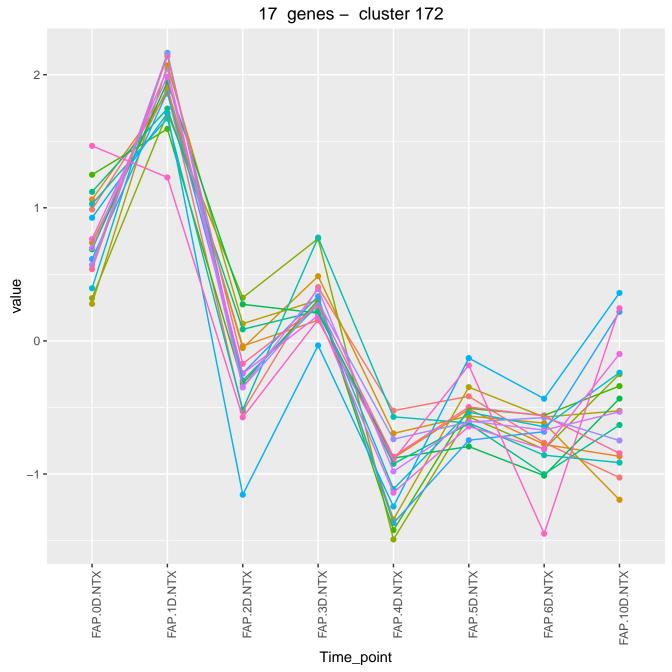


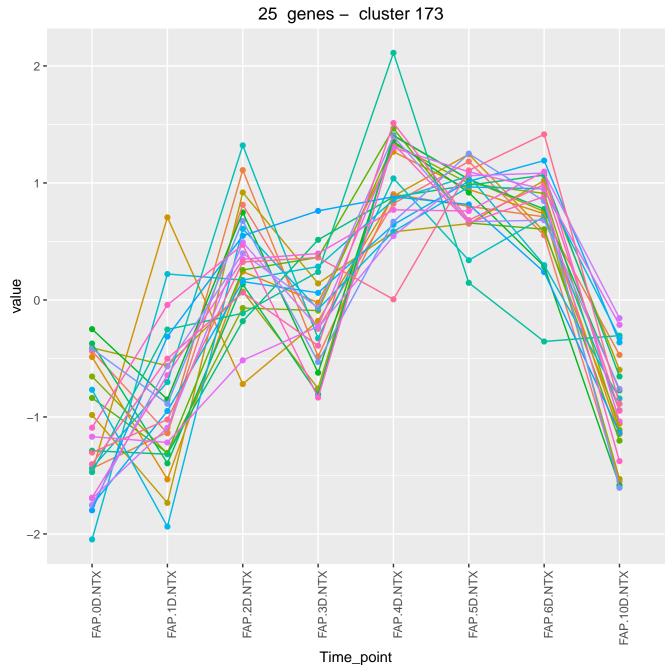


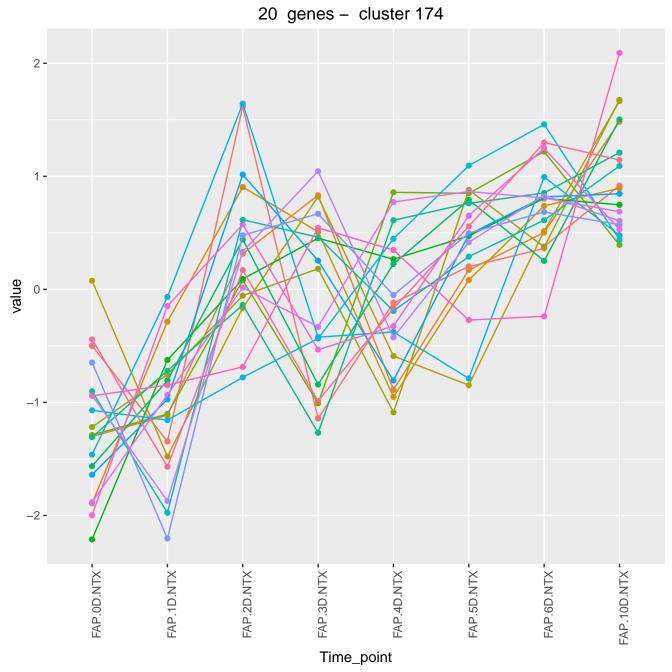


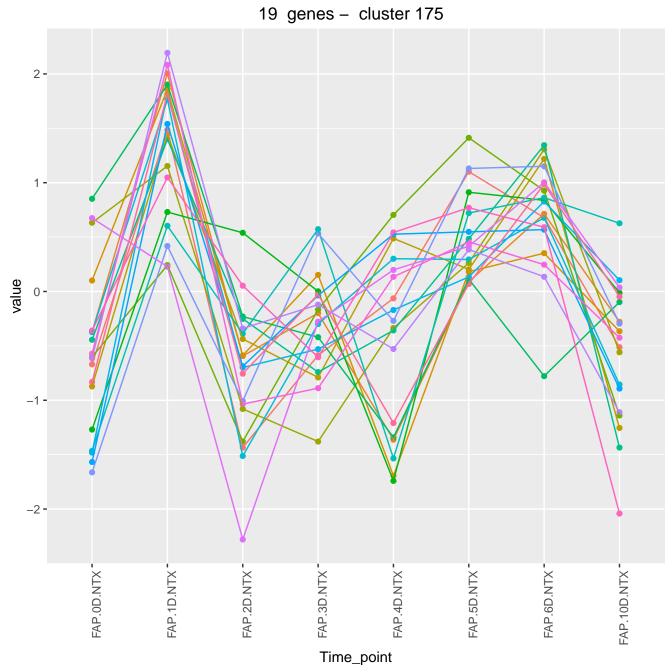


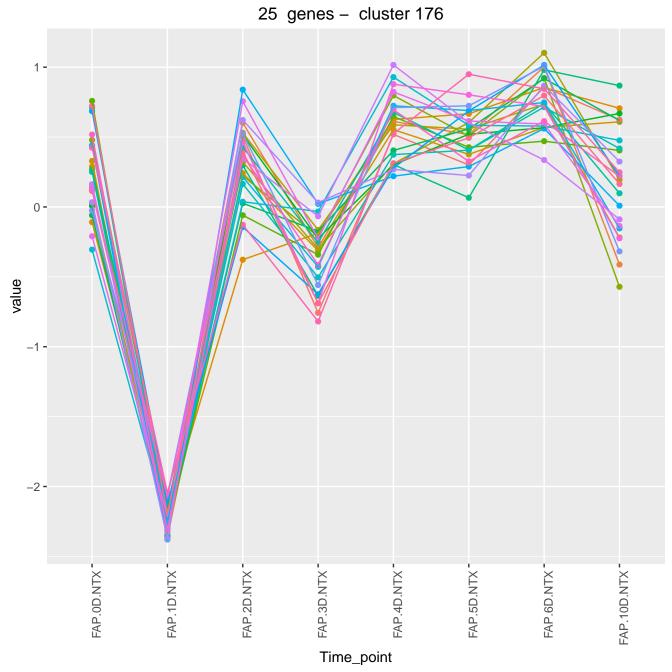


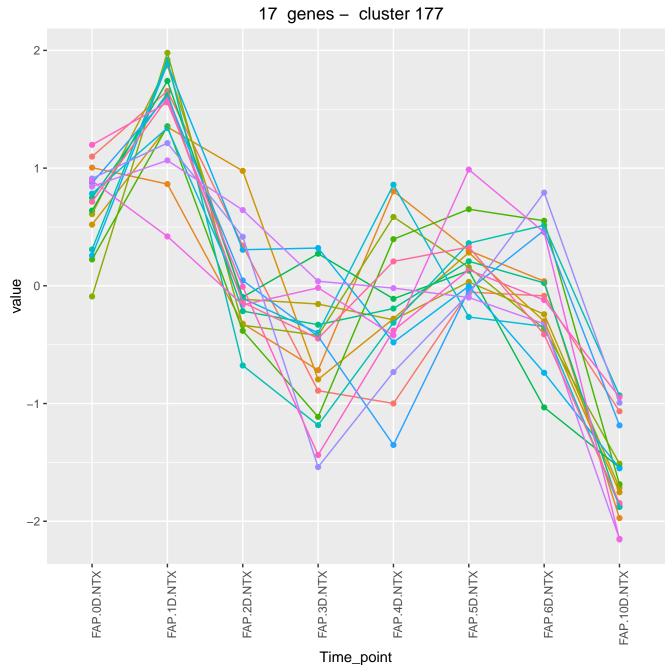


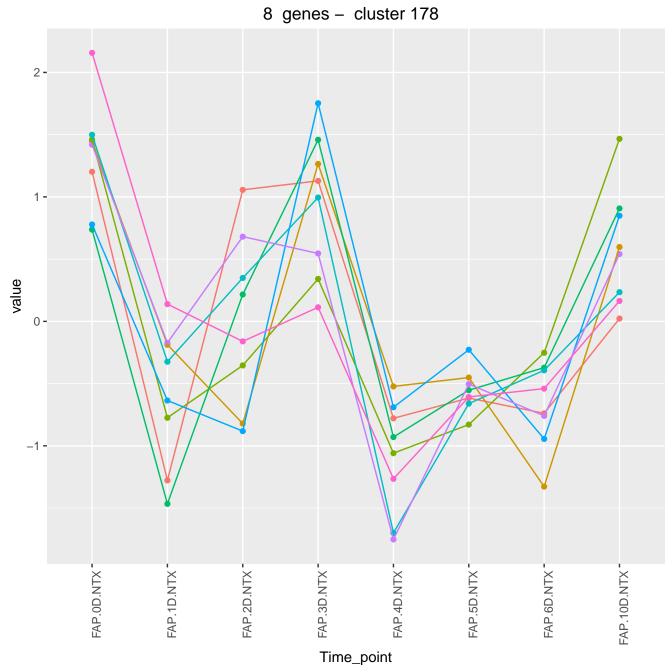


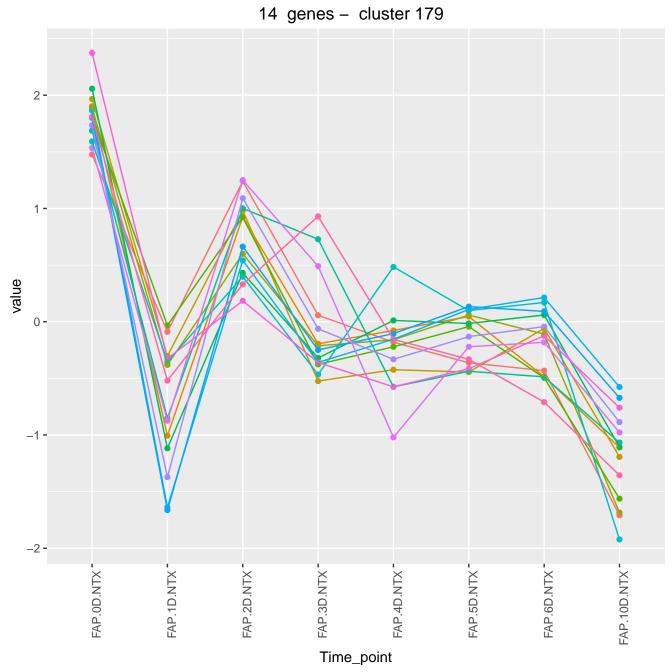


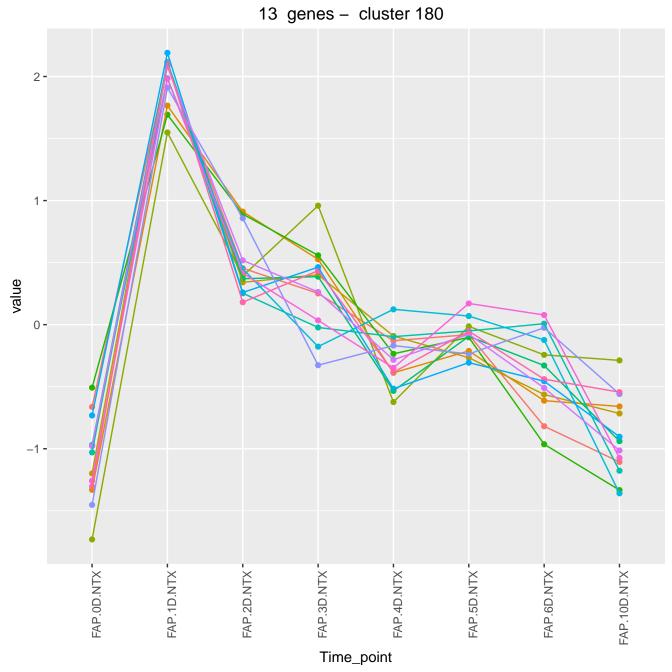


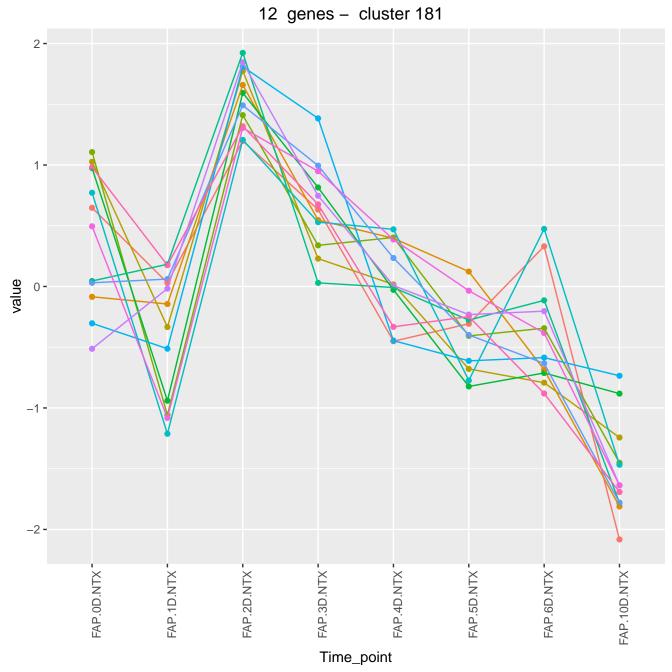


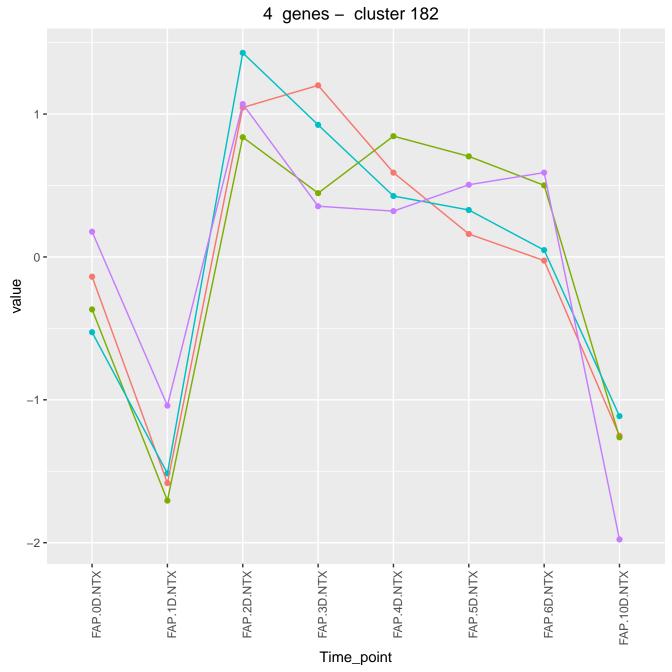


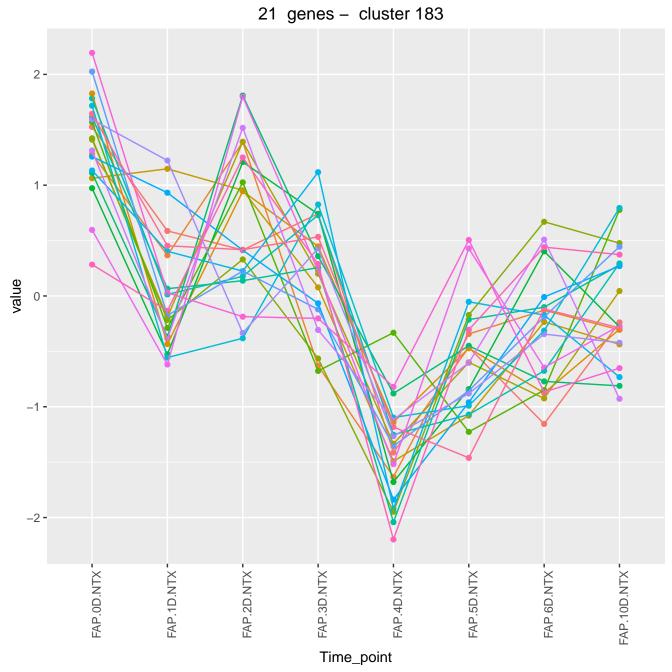


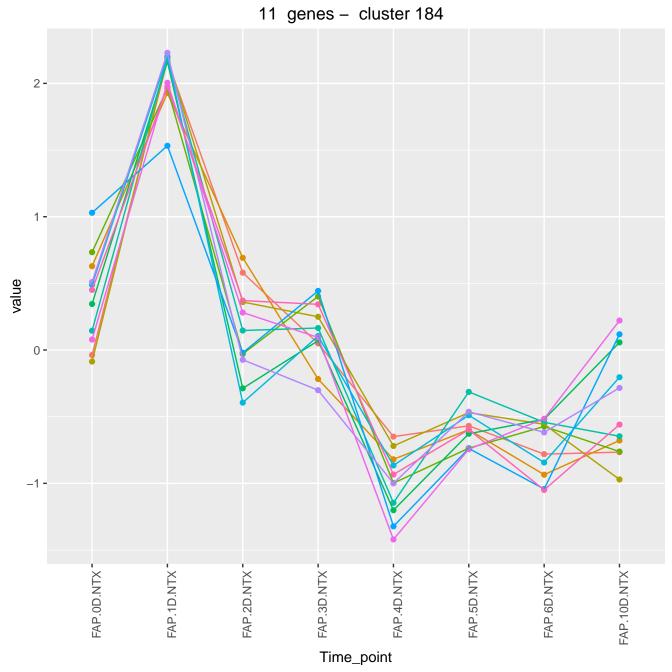


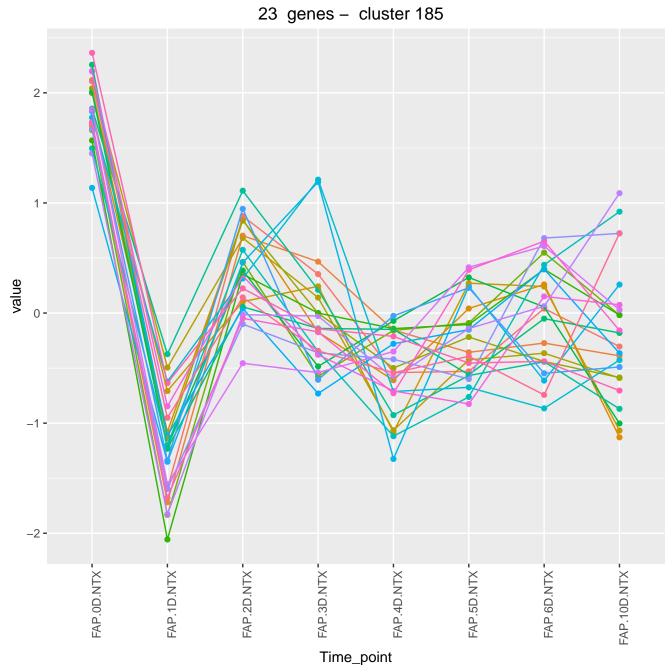


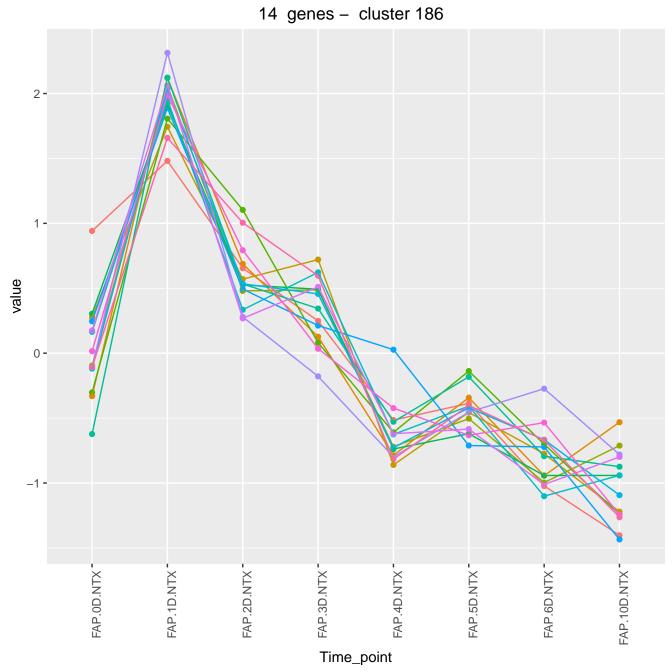


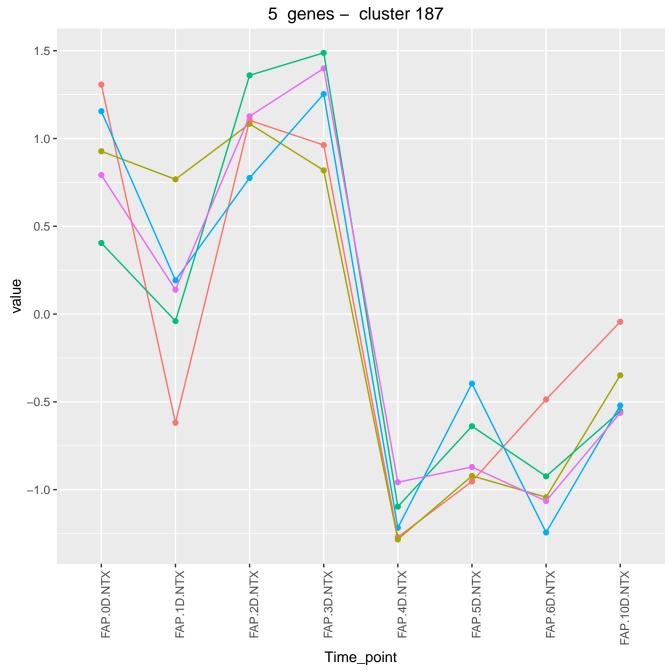


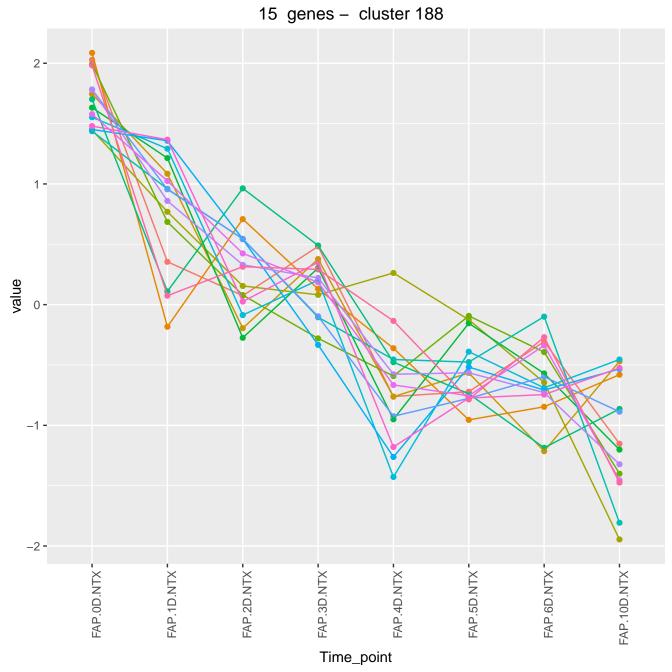


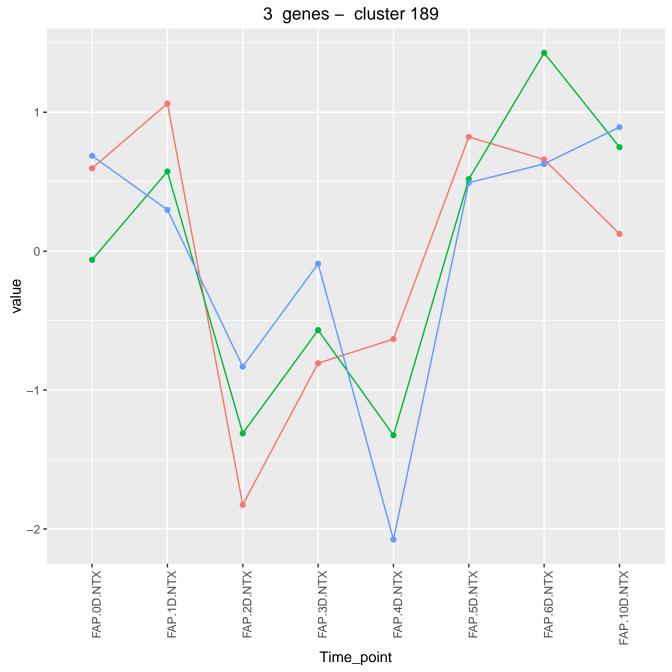


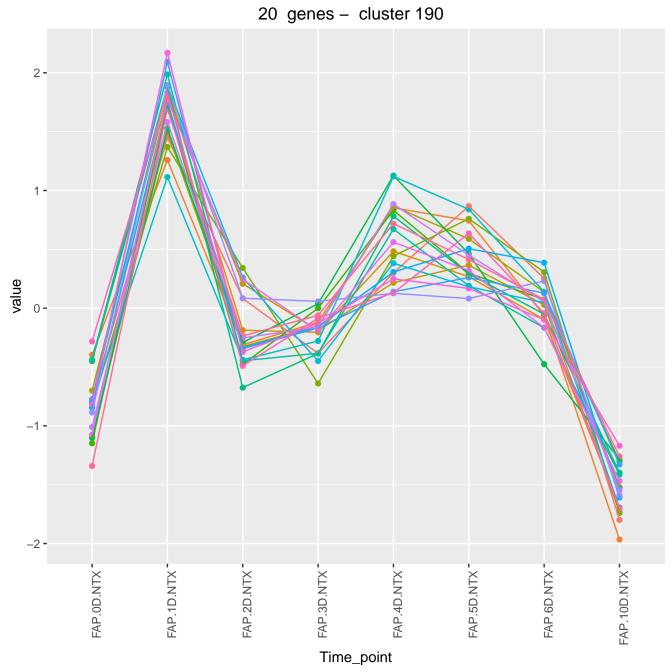


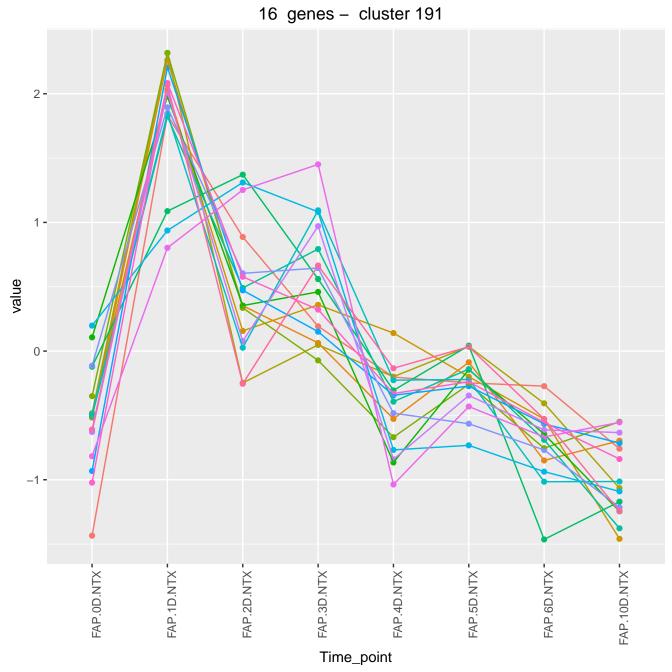


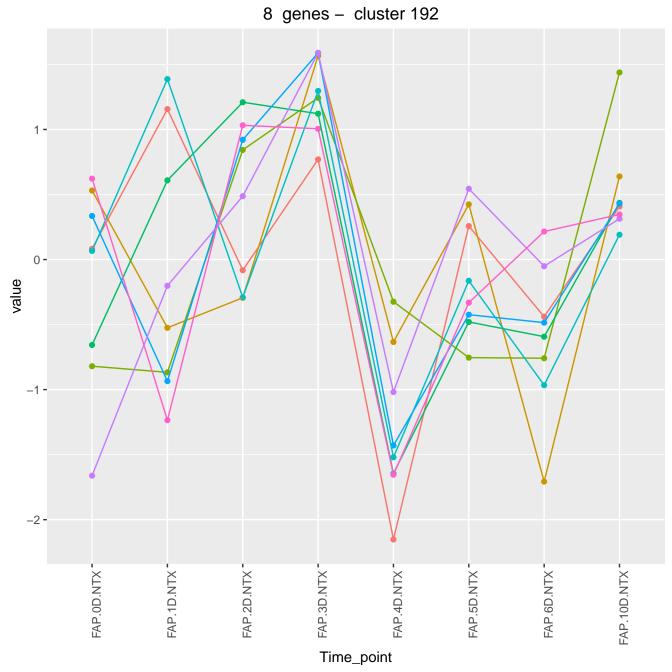


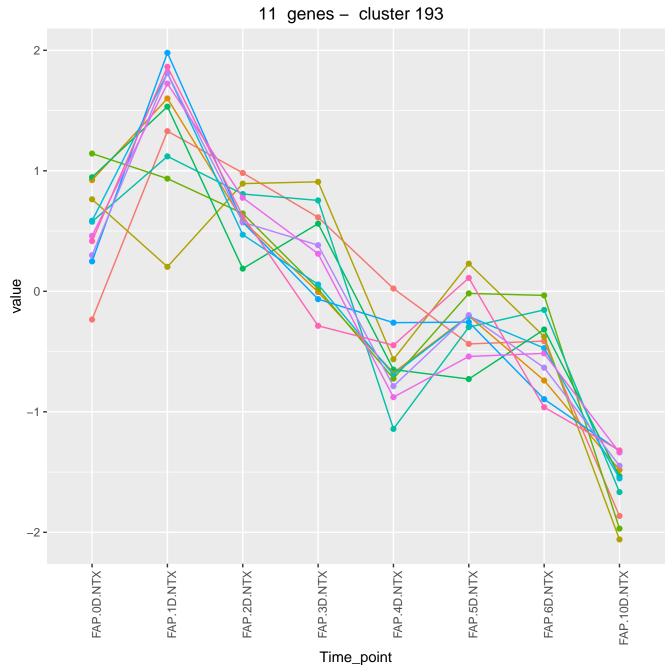


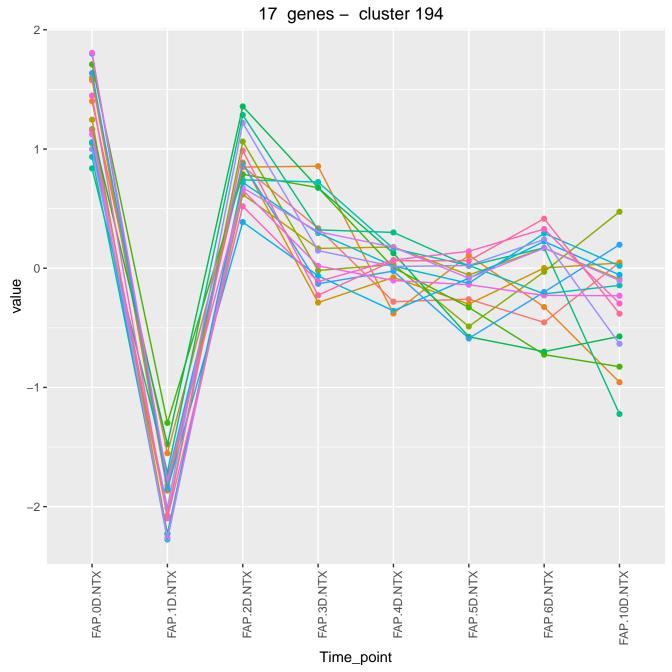




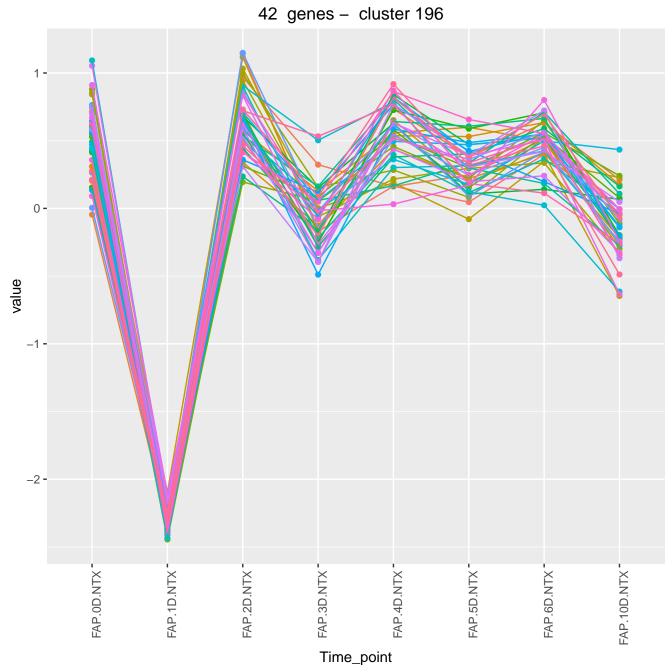


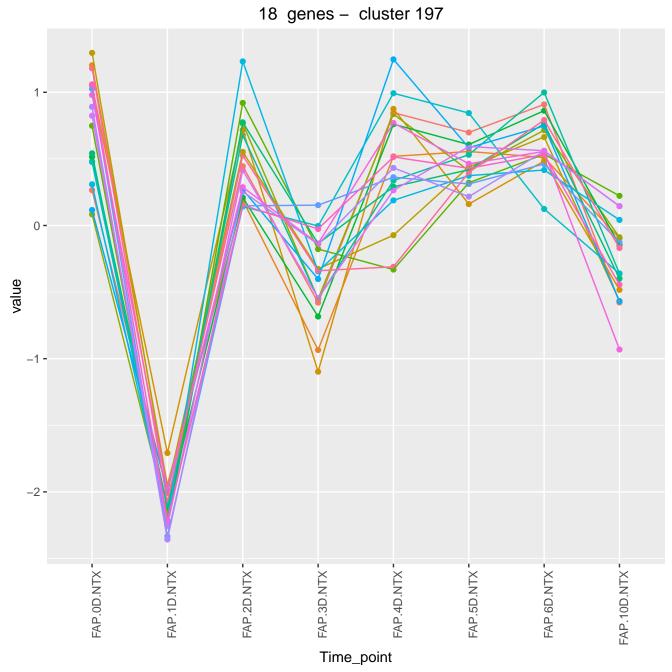


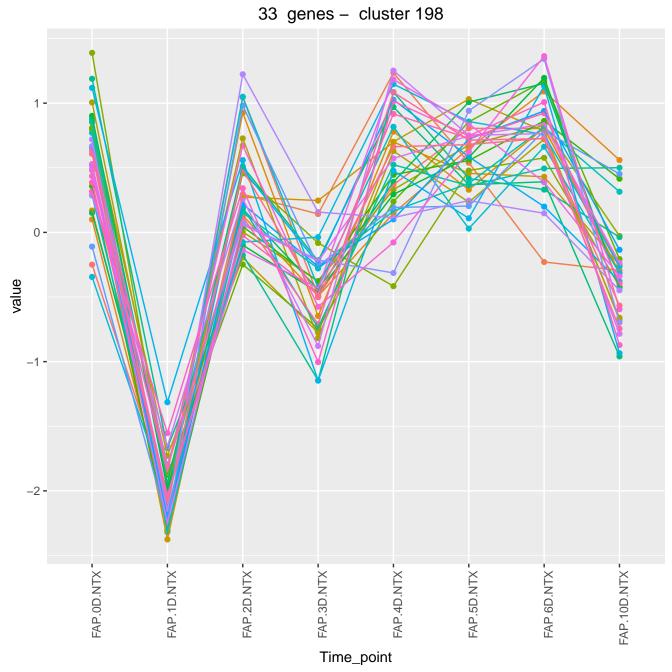


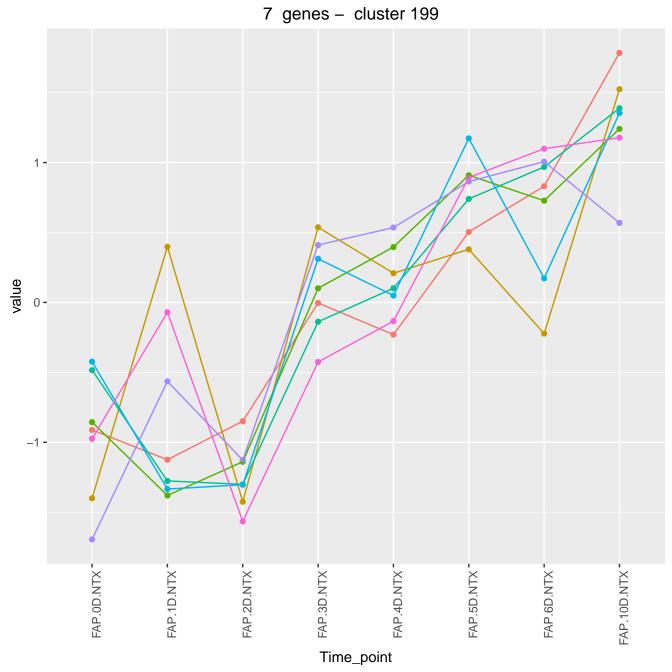


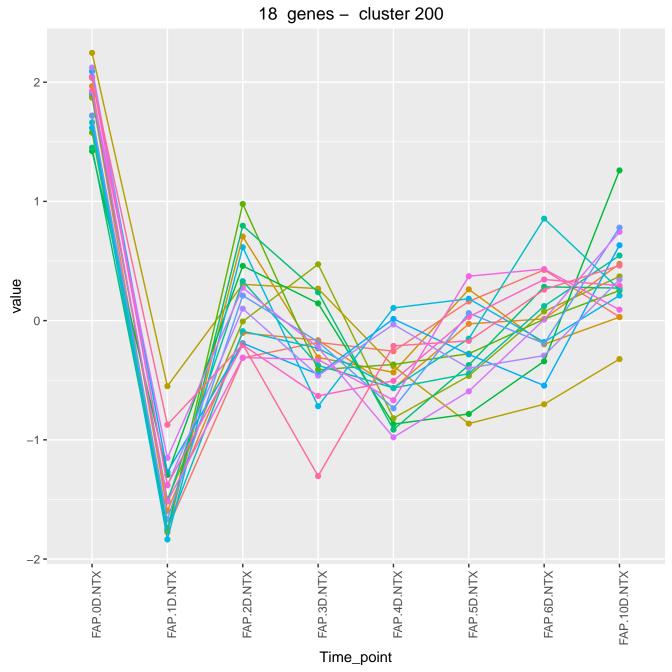
17 genes - cluster 195 2-1 value 0 -**-1-**-2 **-**FAP.0D.NTX 7 FAP.2D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.3D.NTX FAP.6D.NTX -FAP.4D.NTX Time\_point

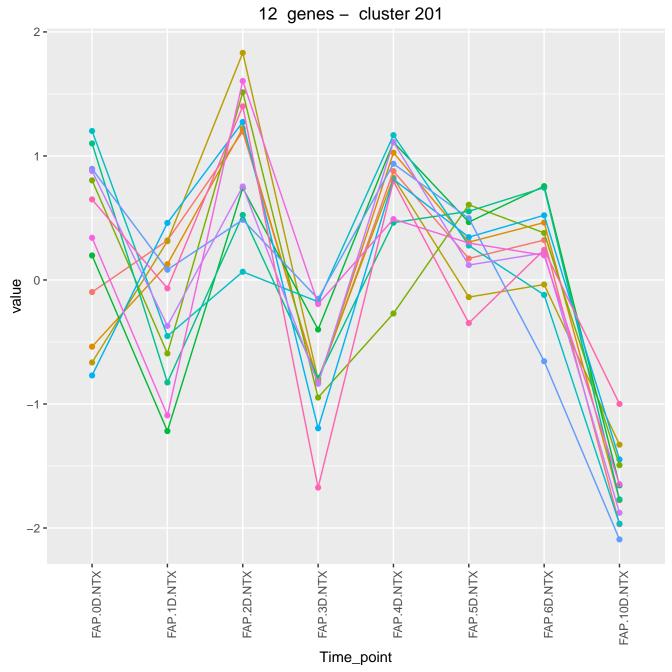


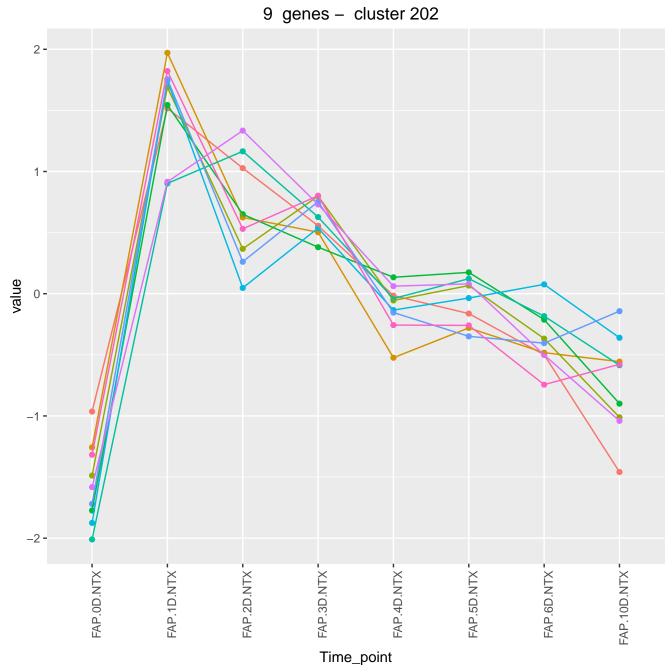


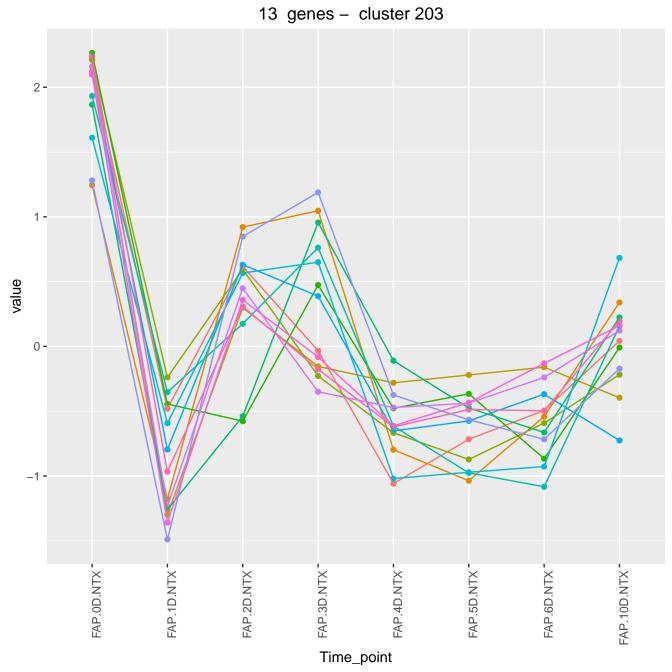


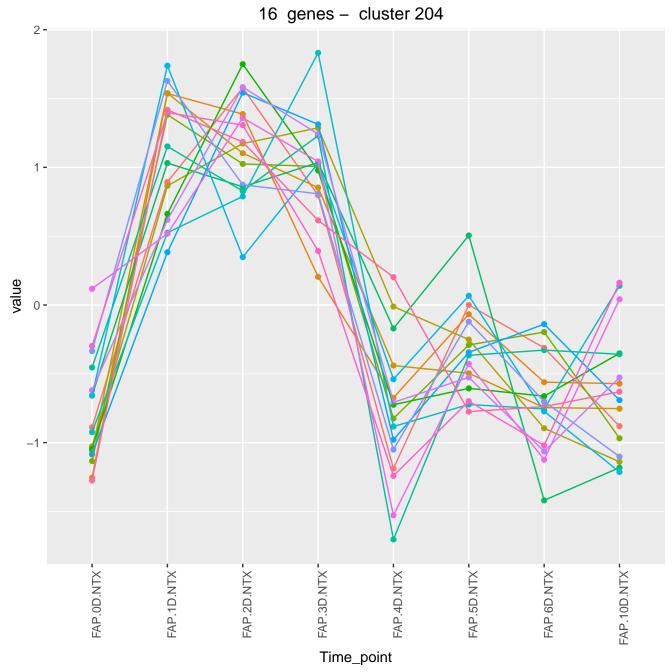


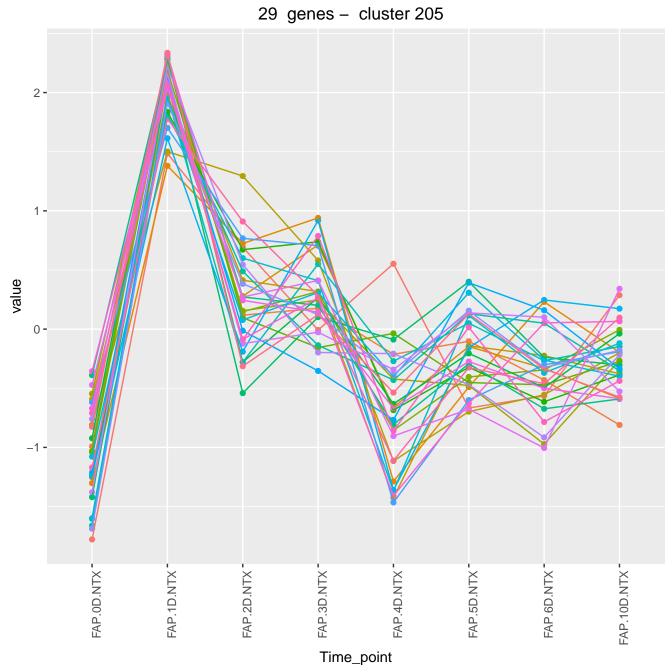


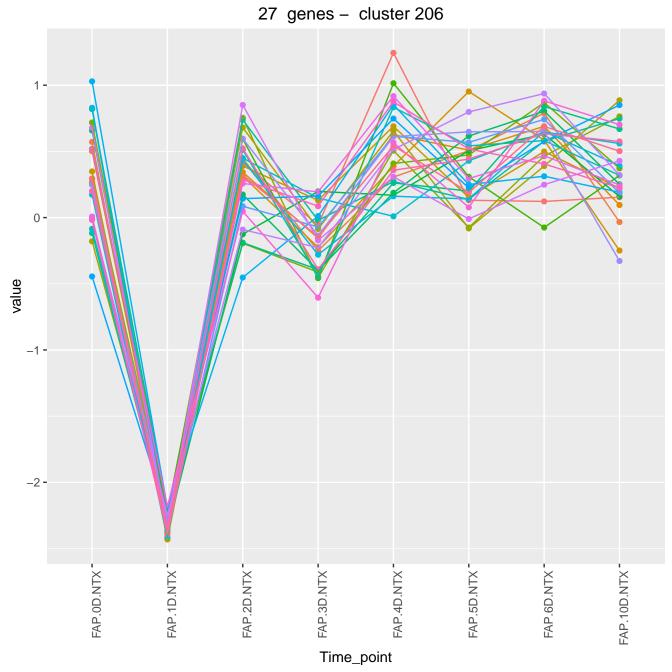


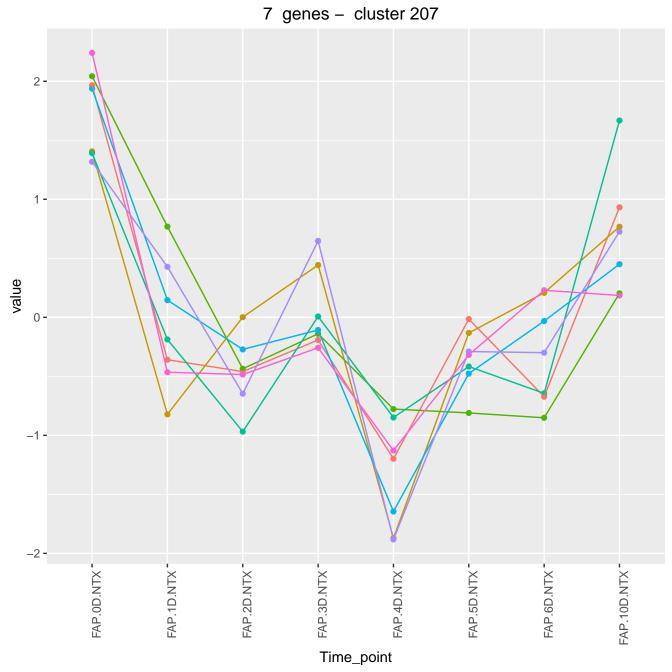


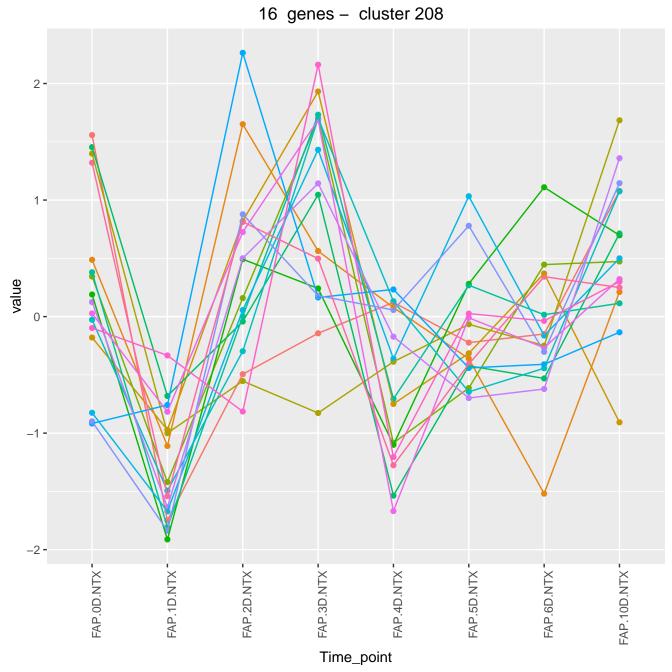


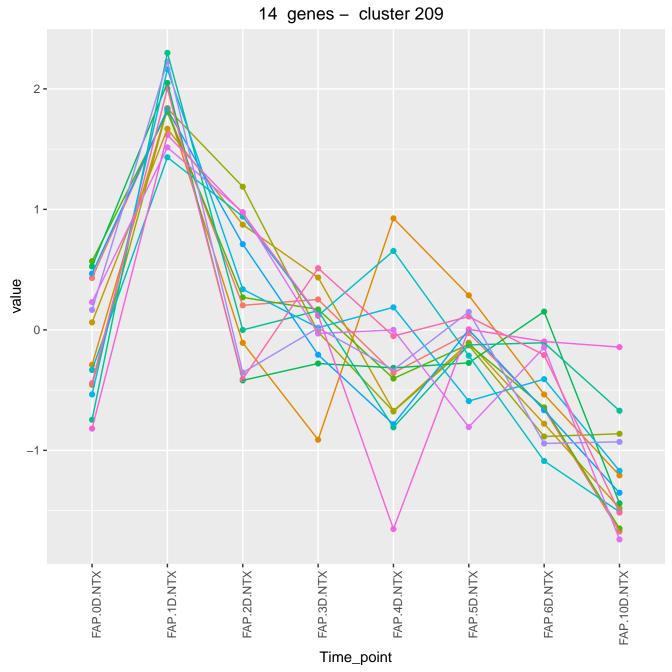


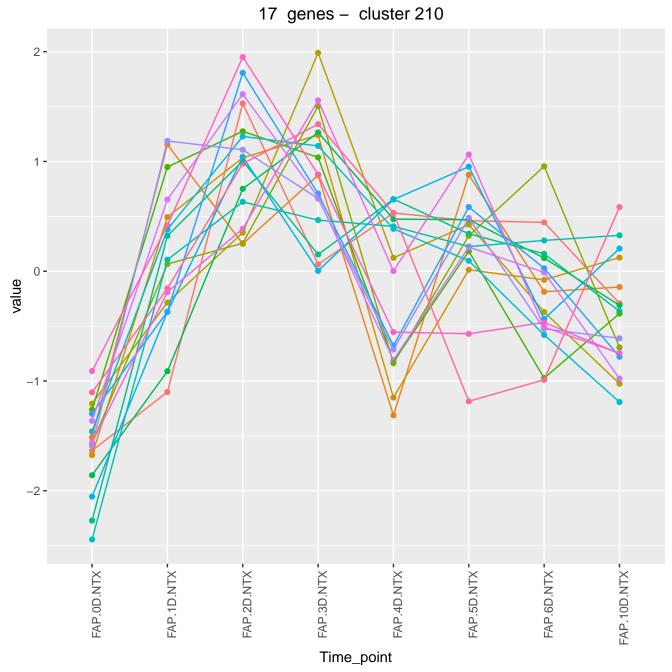


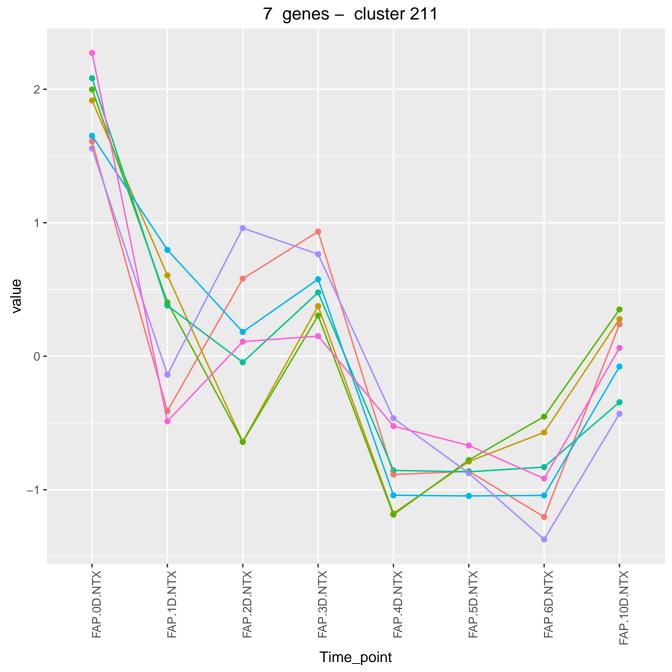


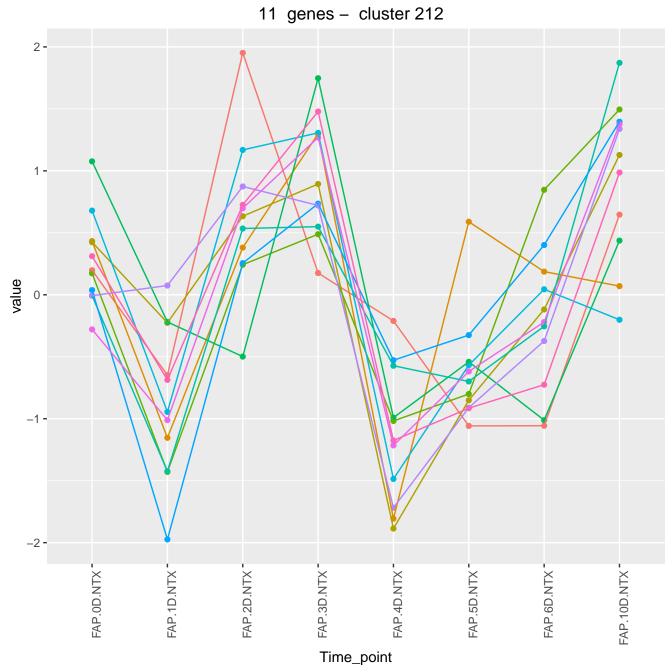


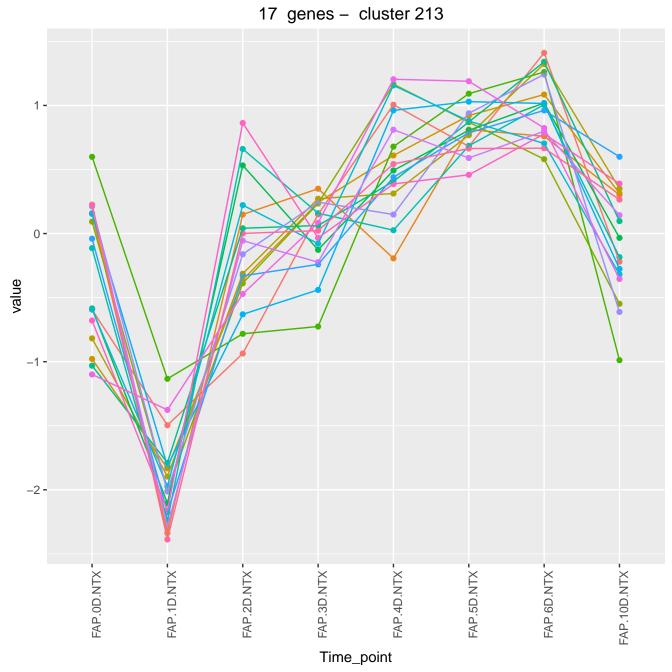


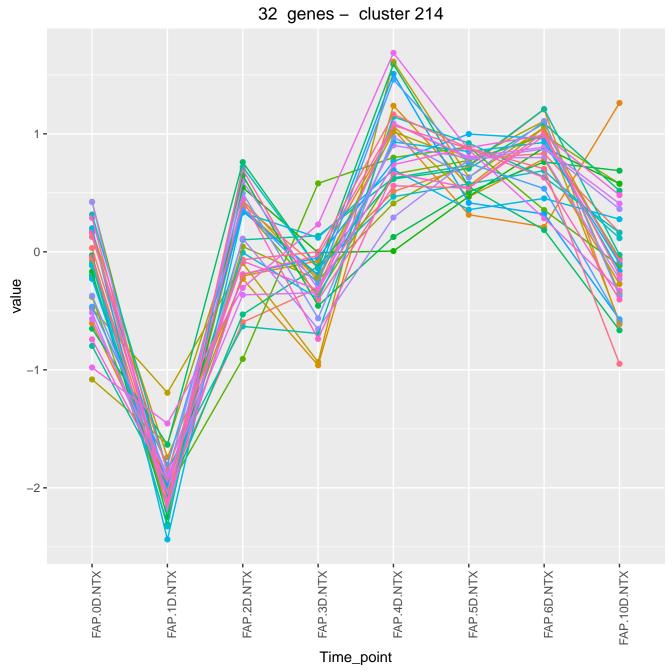


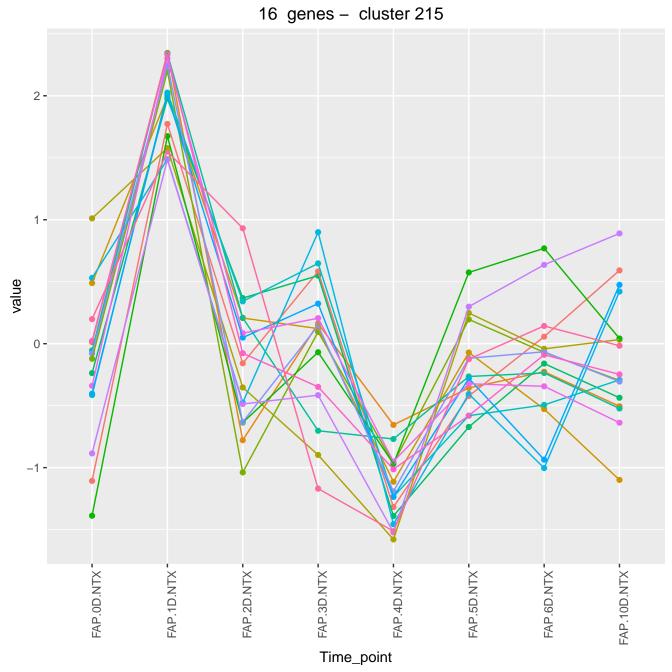




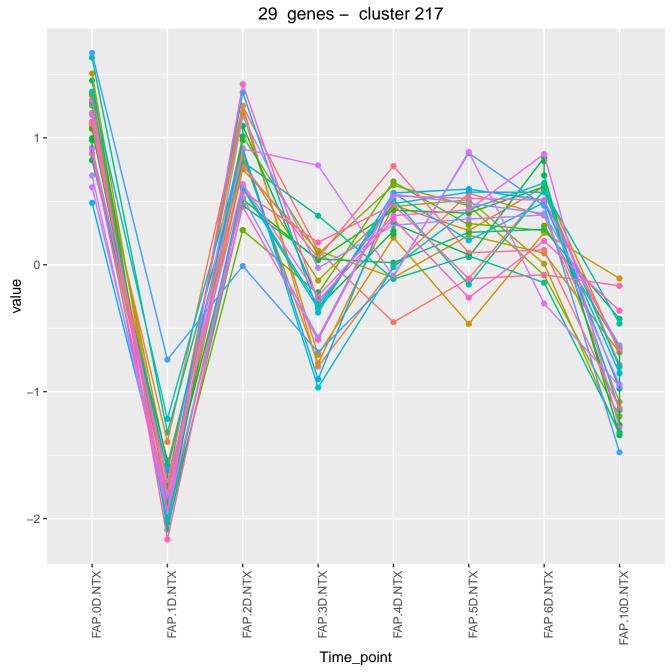


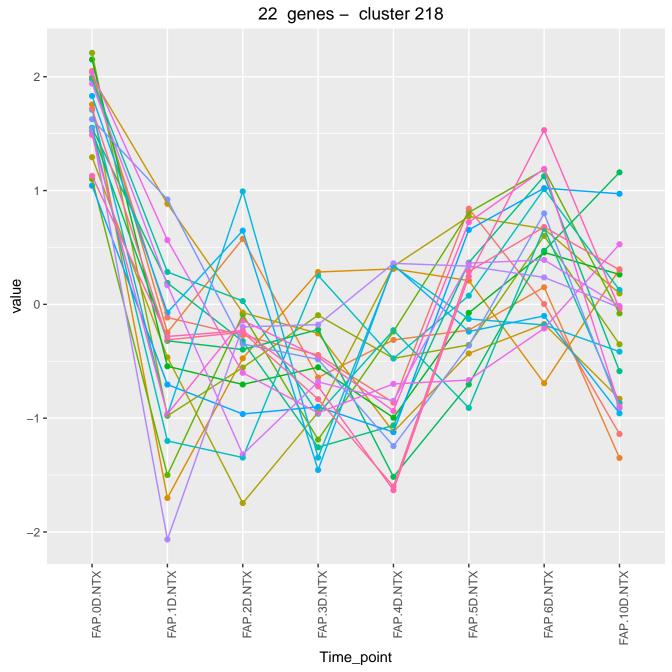


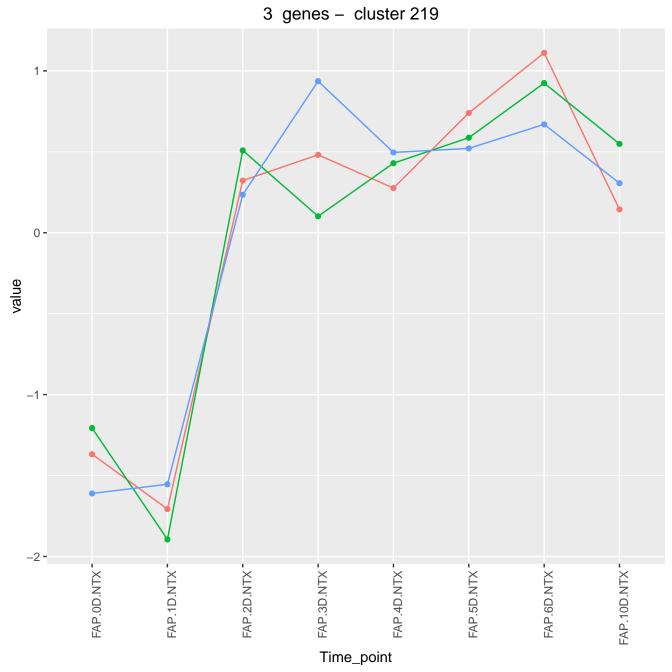


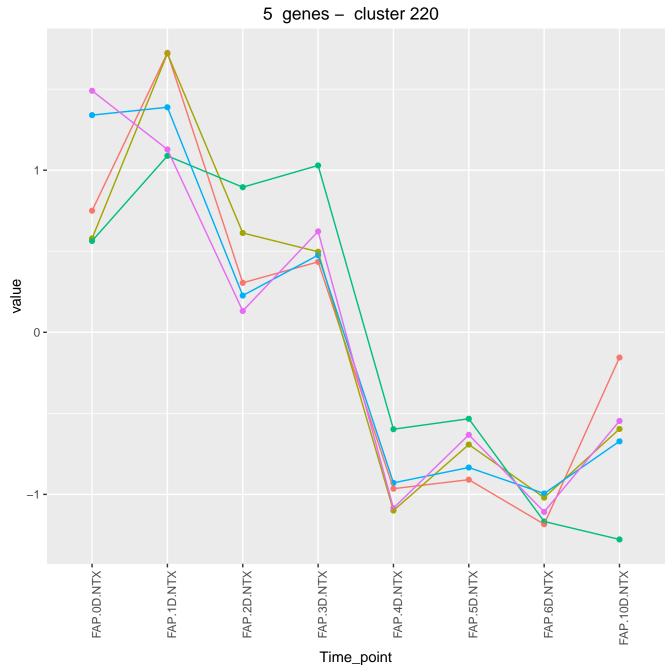


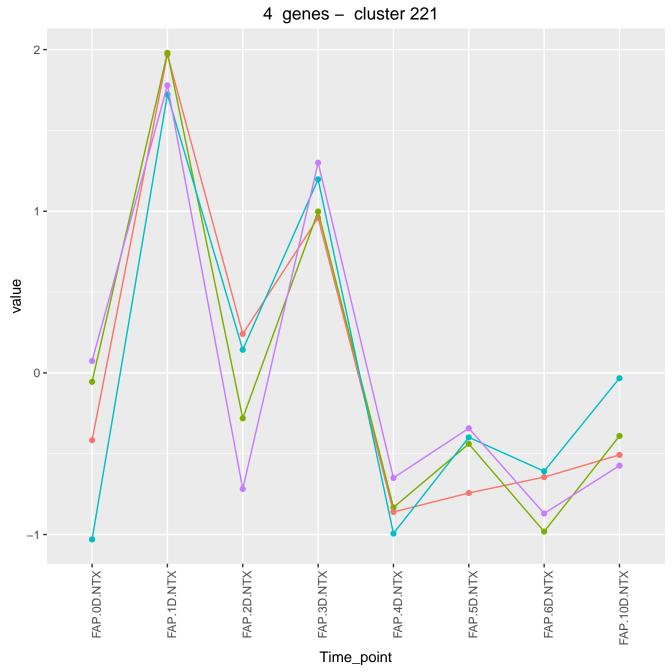
31 genes - cluster 216 2 -1 value -1 **-**-2 **-**FAP.0D.NTX 7 FAP.3D.NTX -FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX -FAP.2D.NTX FAP.6D.NTX -FAP.4D.NTX Time\_point

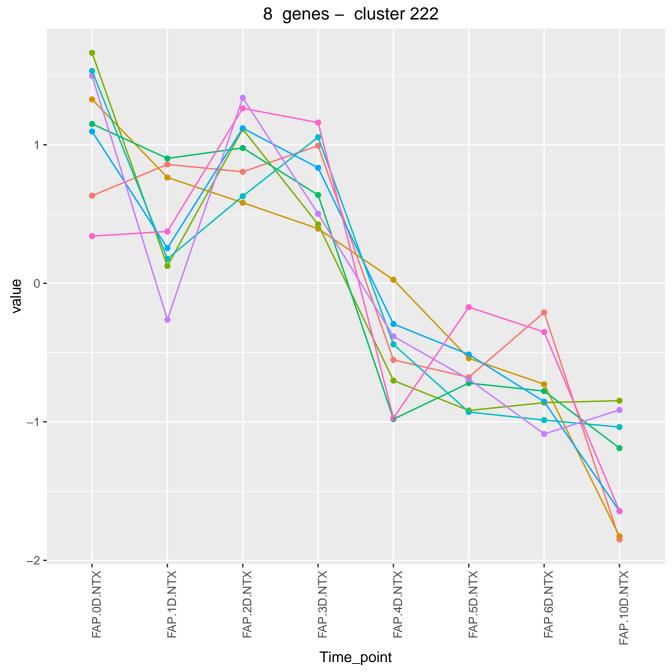


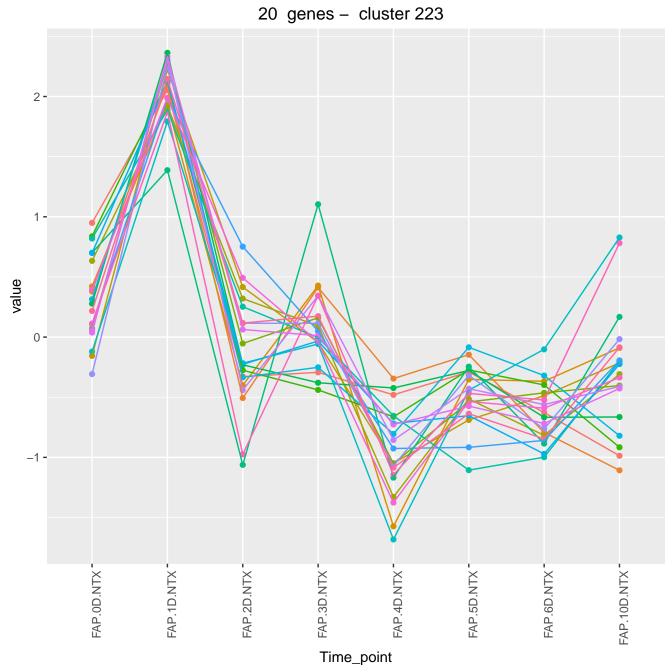


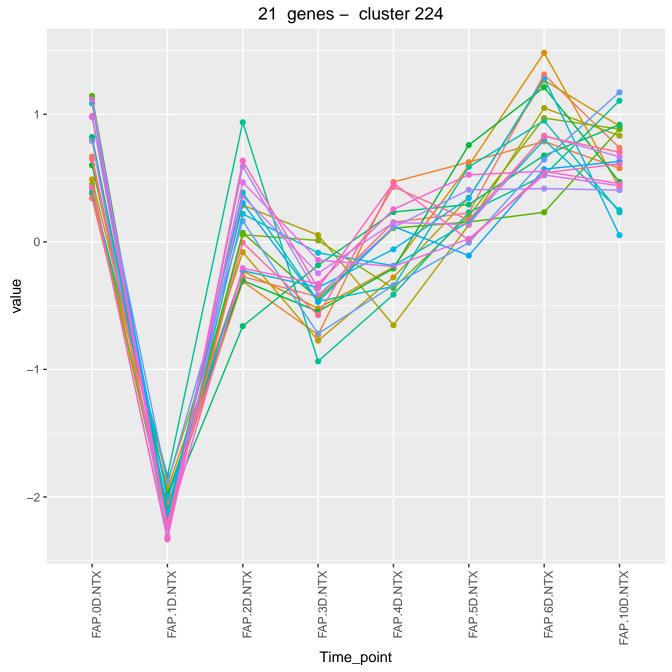


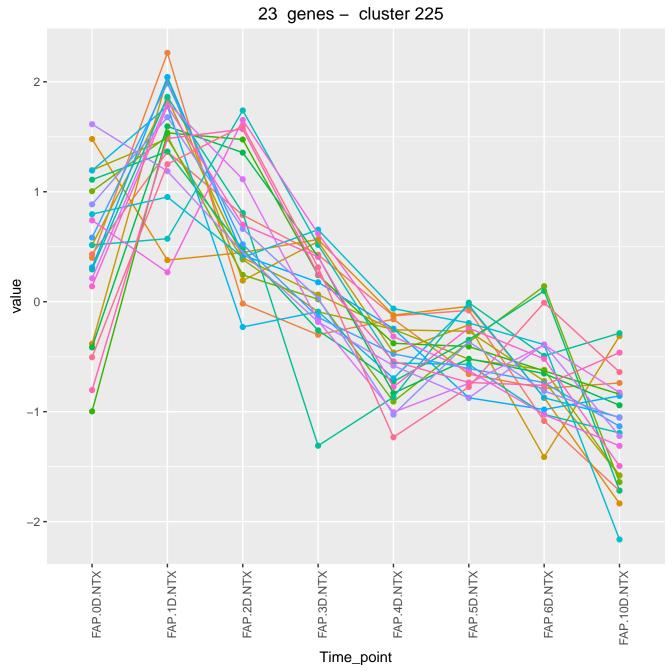


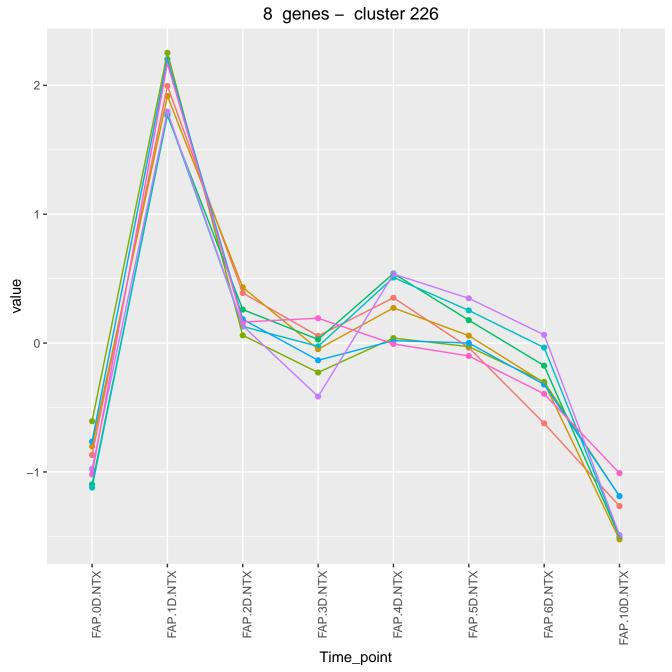


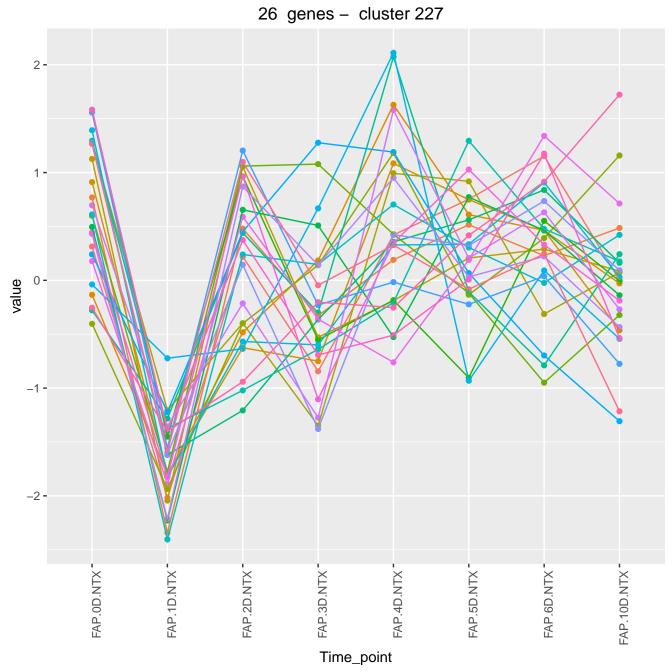


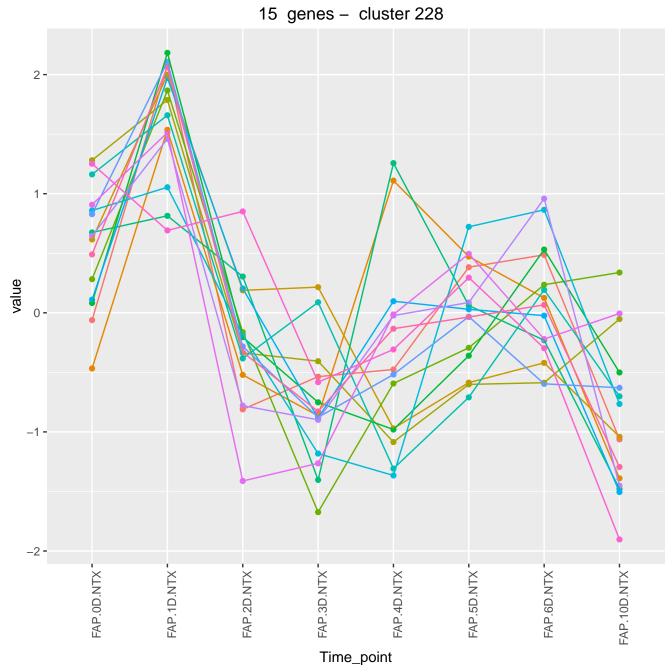


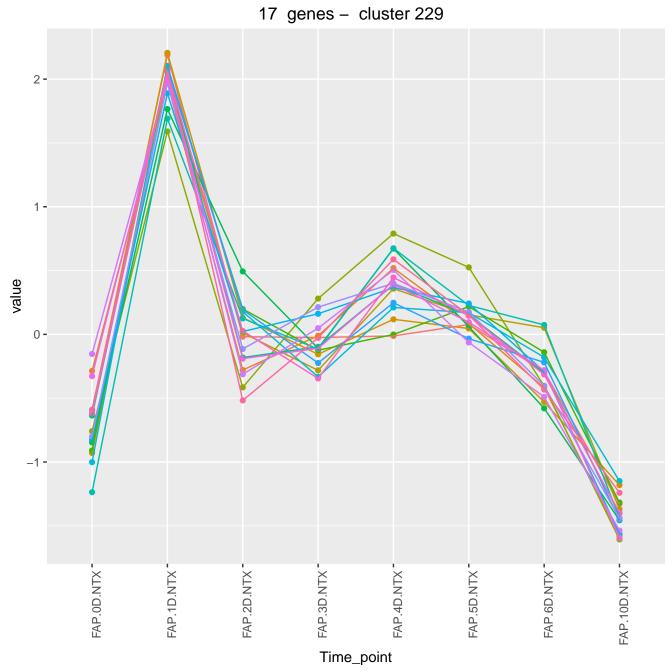


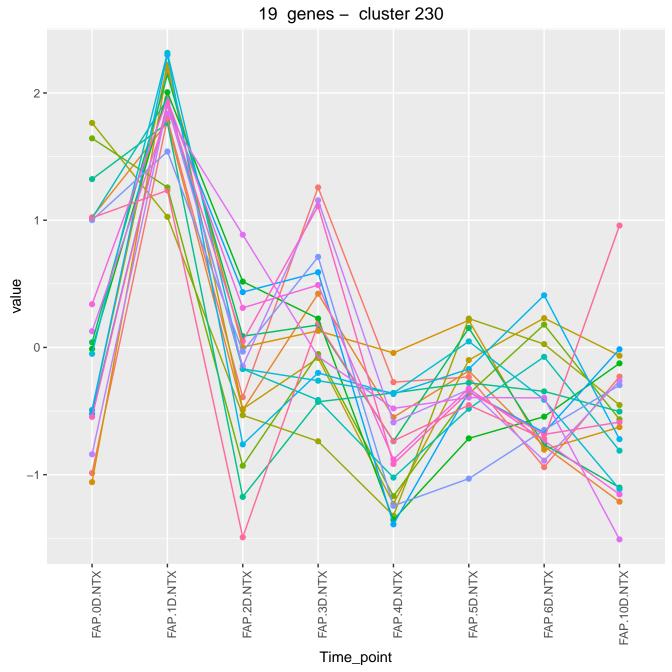


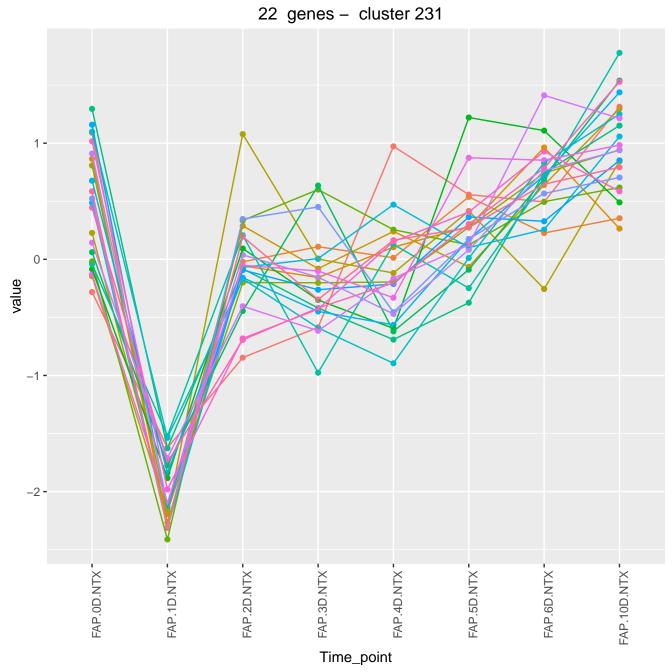


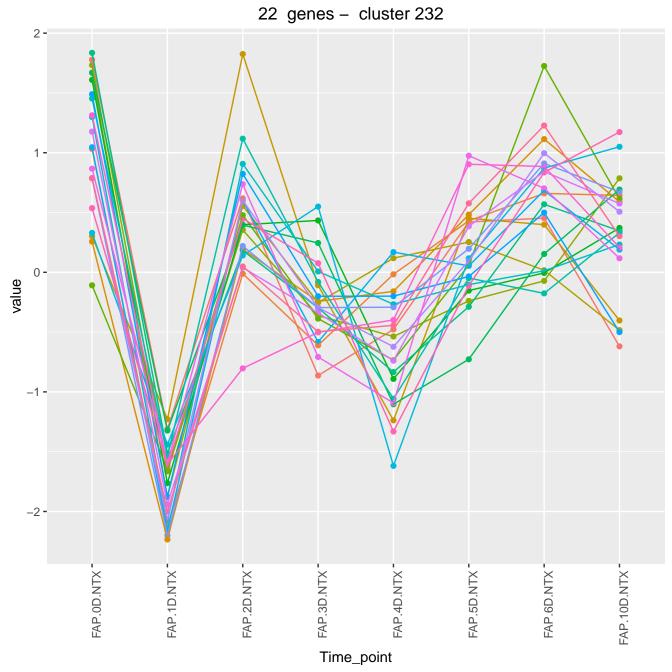




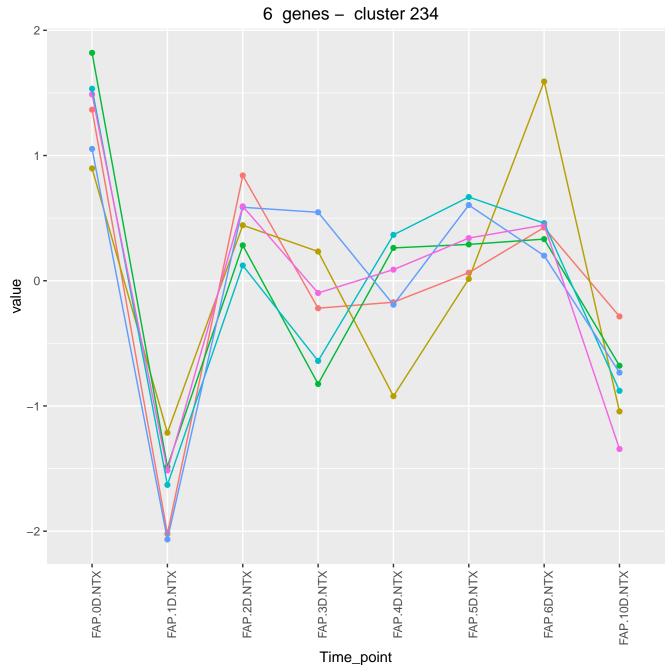




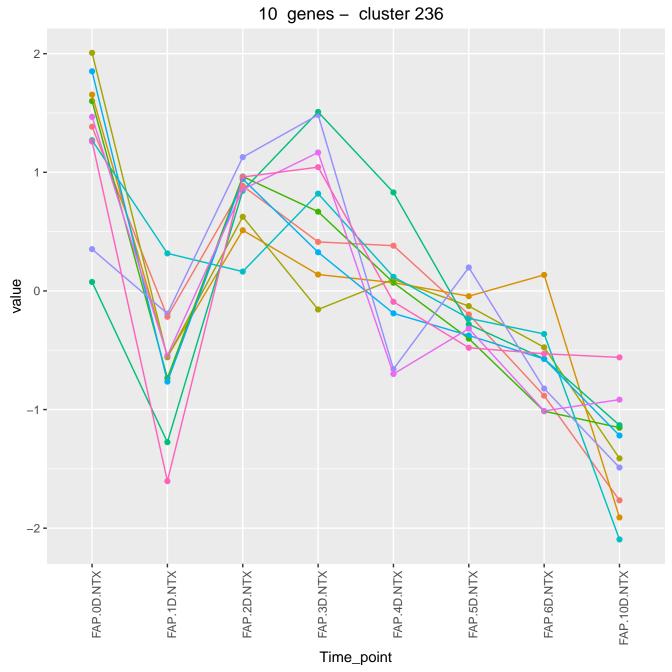


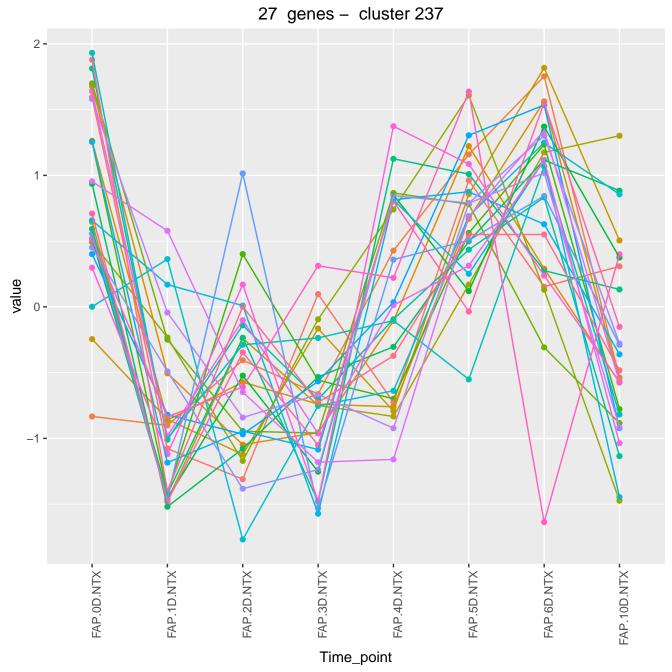


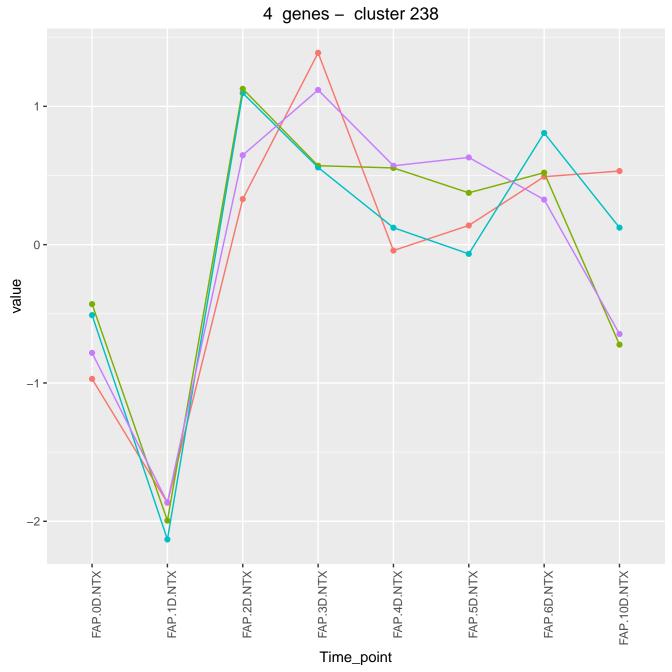
27 genes - cluster 233 2 -1 value **-1** --2 **-**FAP.0D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.4D.NTX FAP.6D.NTX -Time\_point

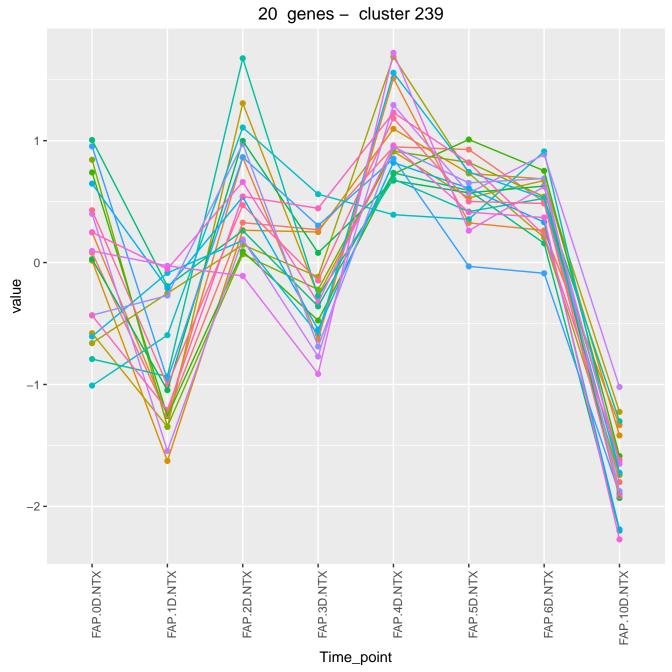


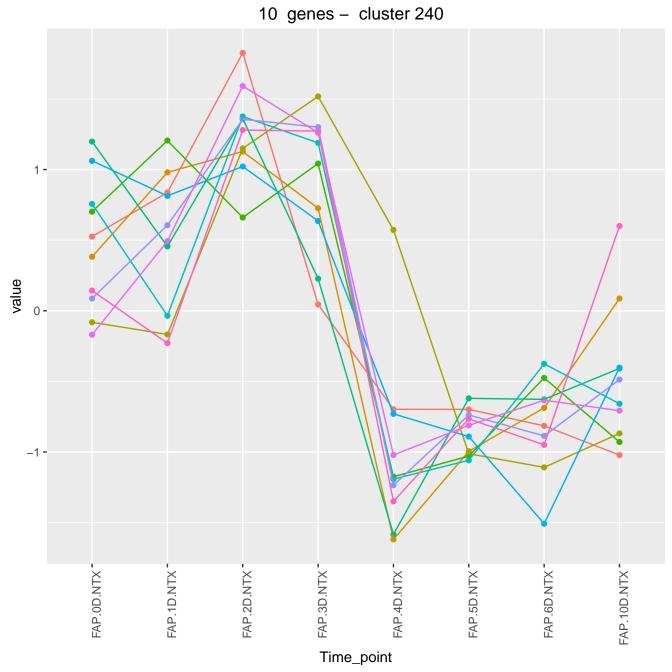
14 genes - cluster 235 2 -1 value 0 --1 **-**-2 **-**FAP.0D.NTX 7 FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.4D.NTX <sup>7</sup> FAP.10D.NTX -FAP.5D.NTX -FAP.1D.NTX FAP.6D.NTX Time\_point

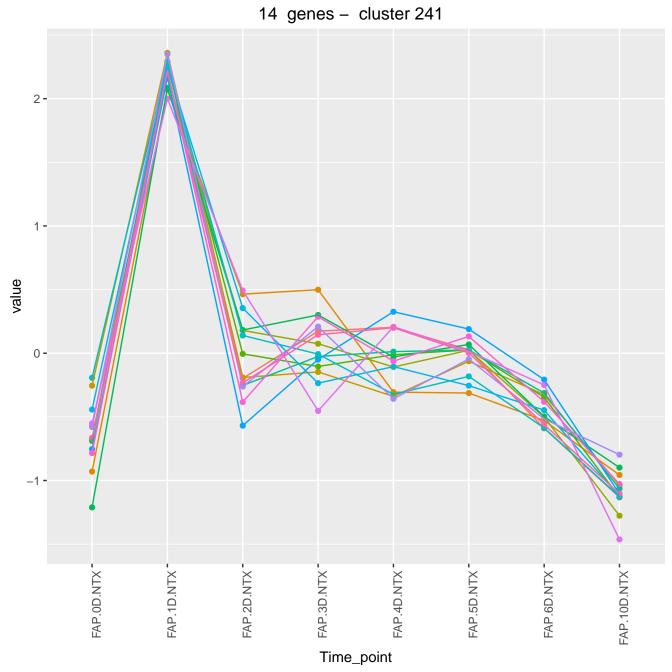


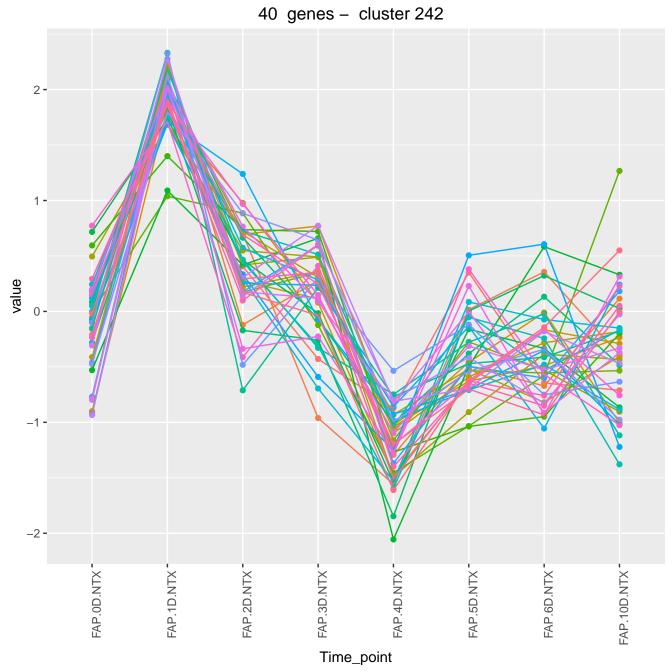


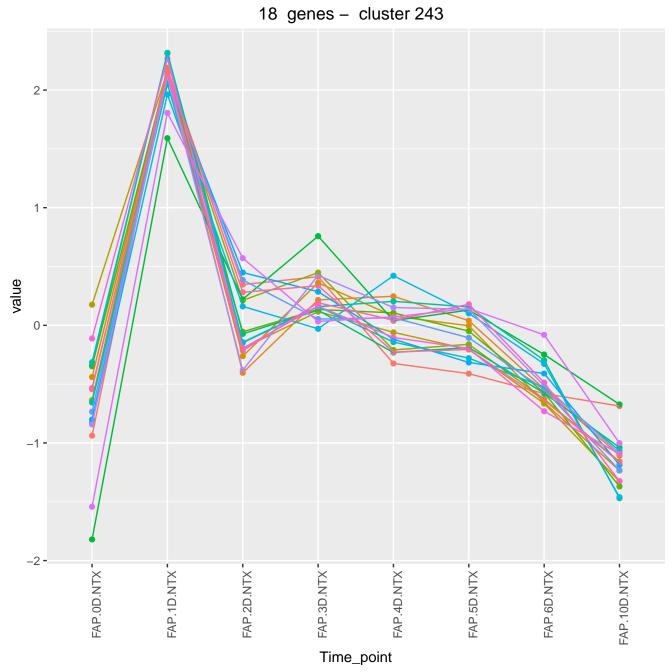


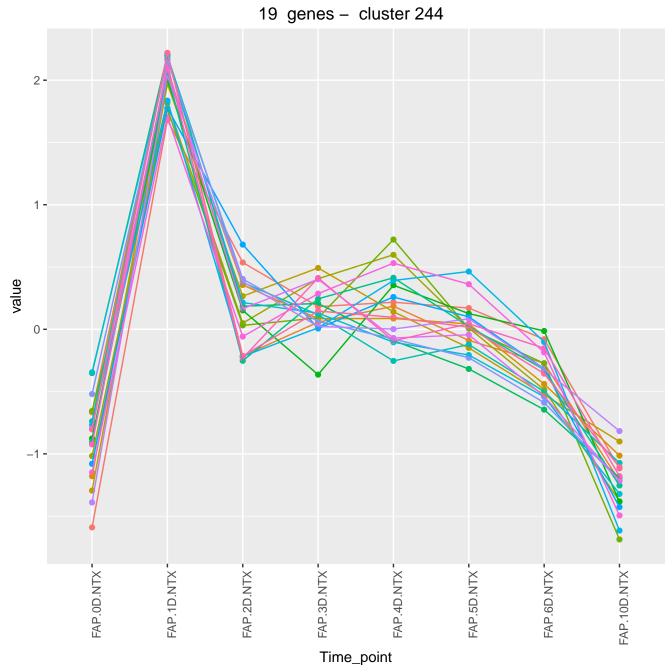


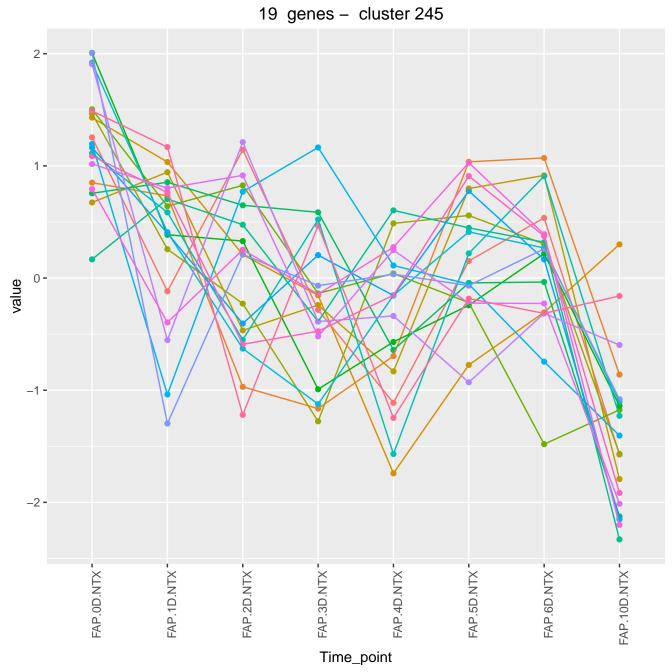


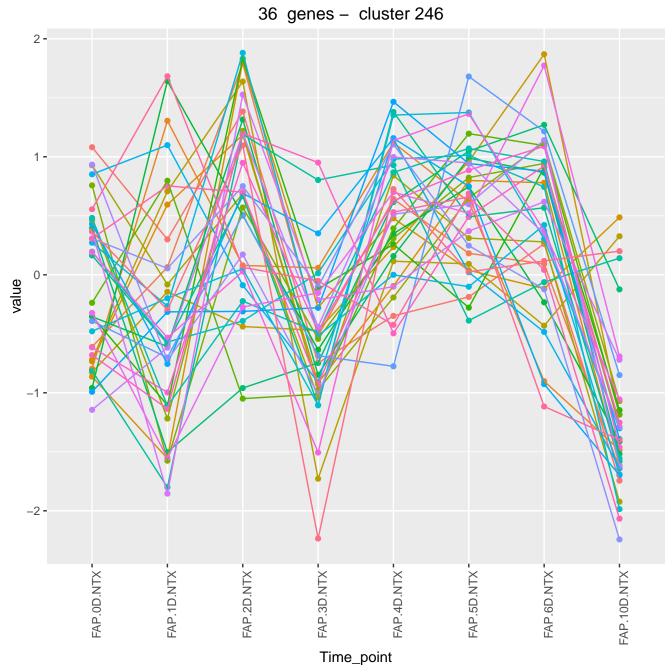


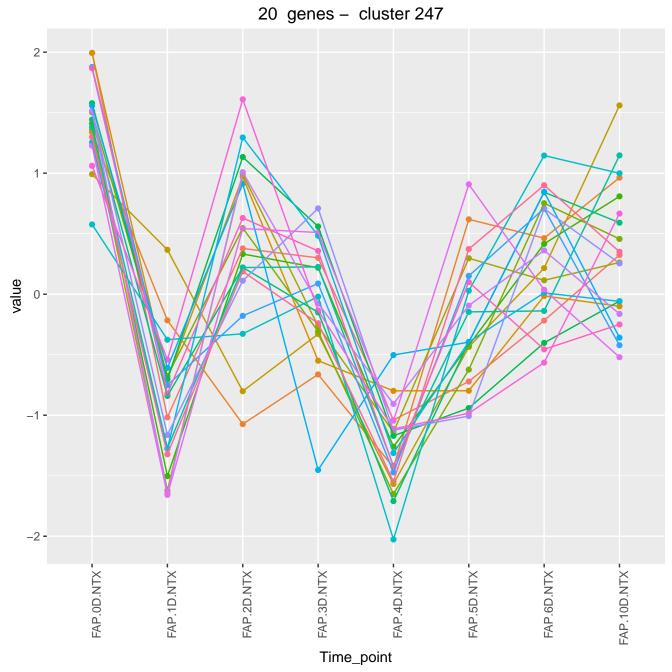


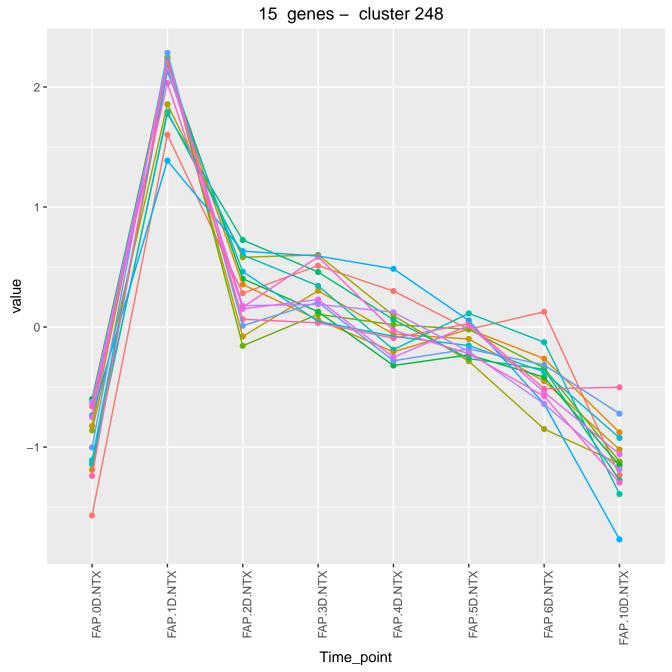


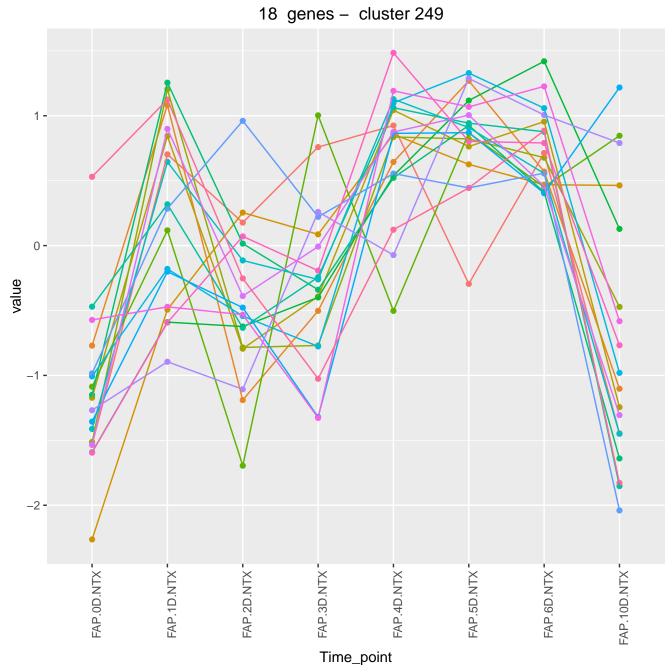


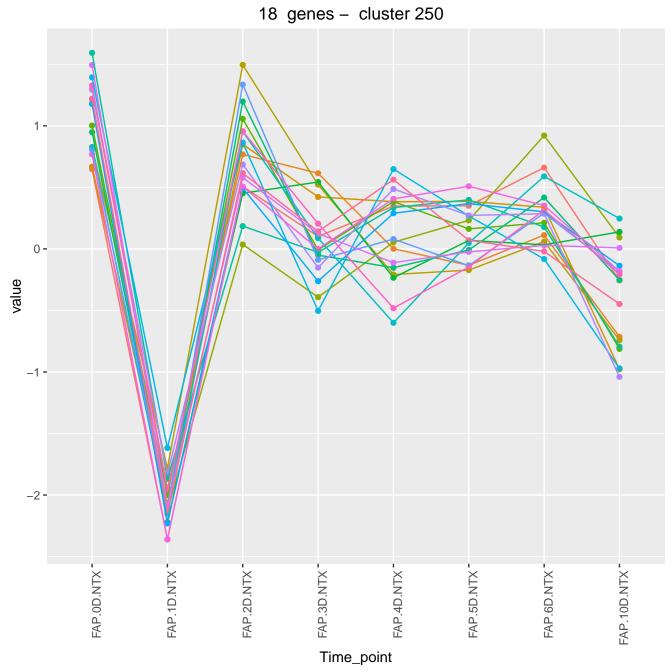


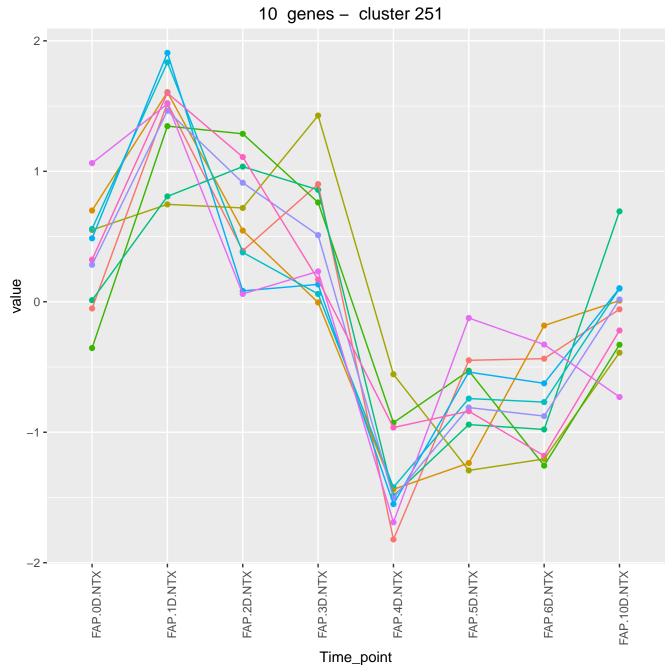


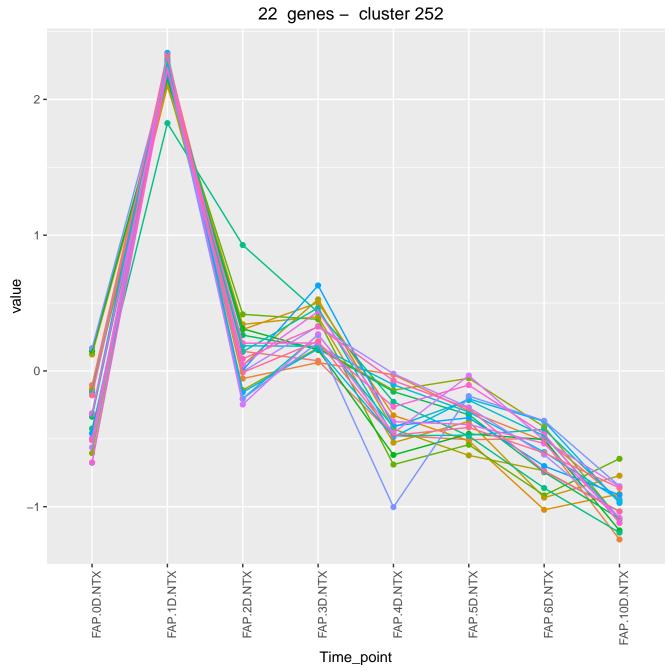


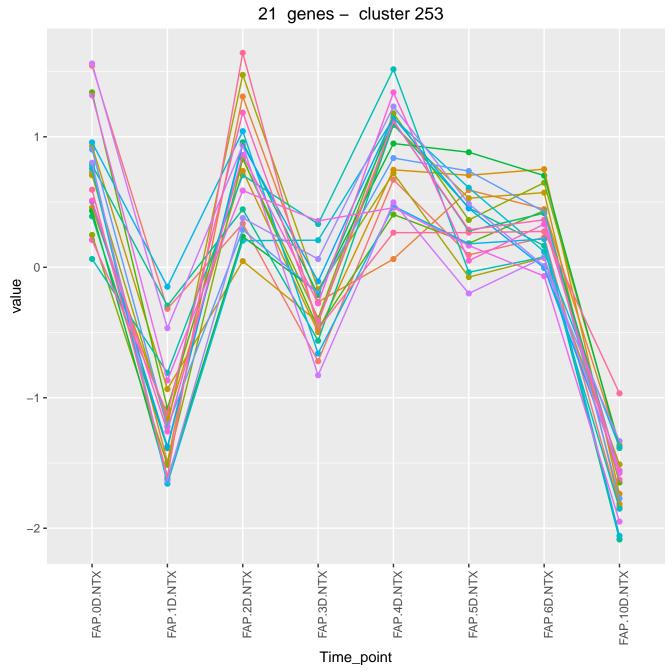




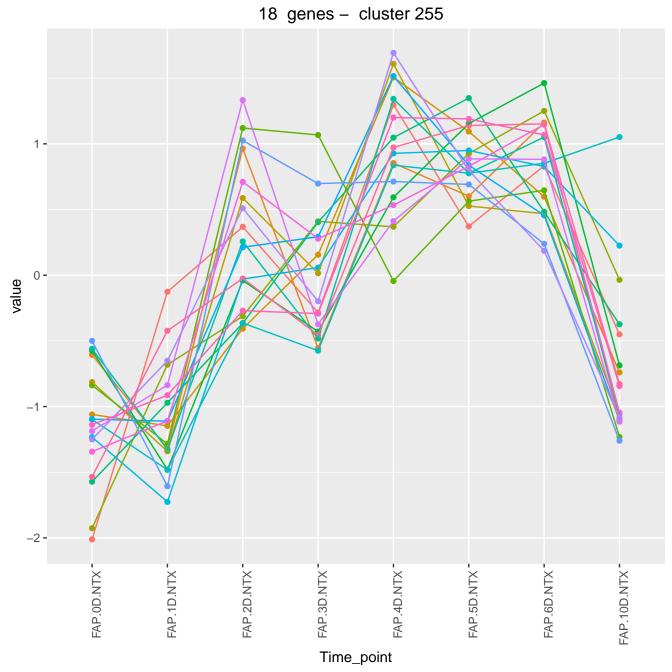


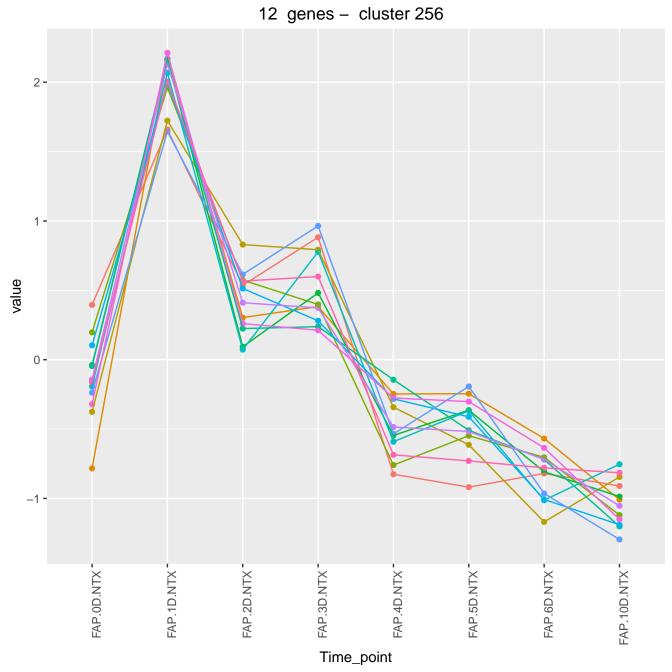


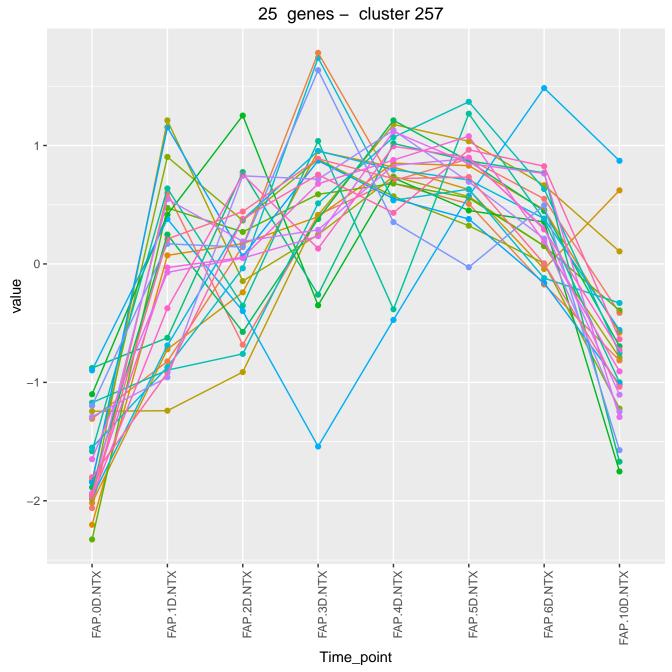


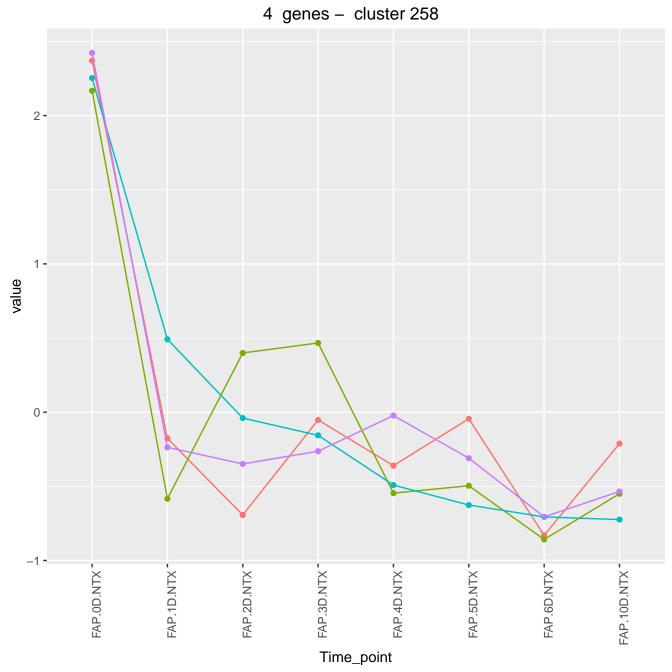


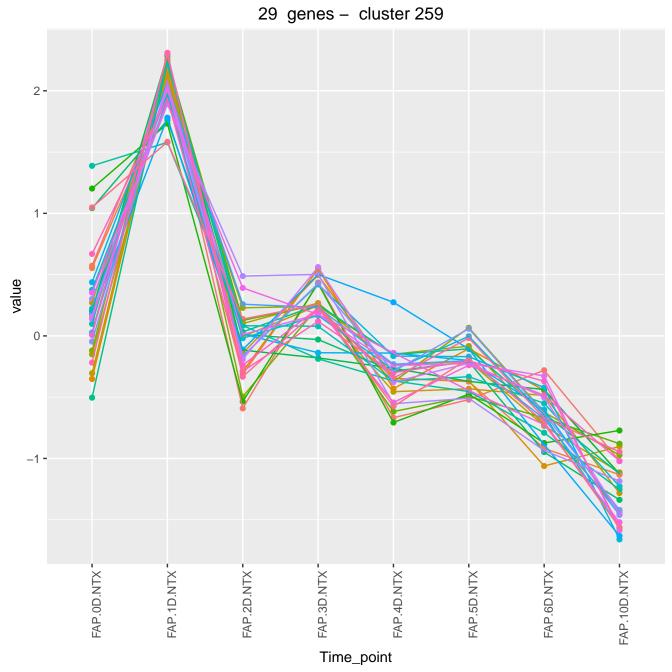
24 genes - cluster 254 2 -1 value 0 --1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.6D.NTX <sup>7</sup> FAP.4D.NTX Time\_point

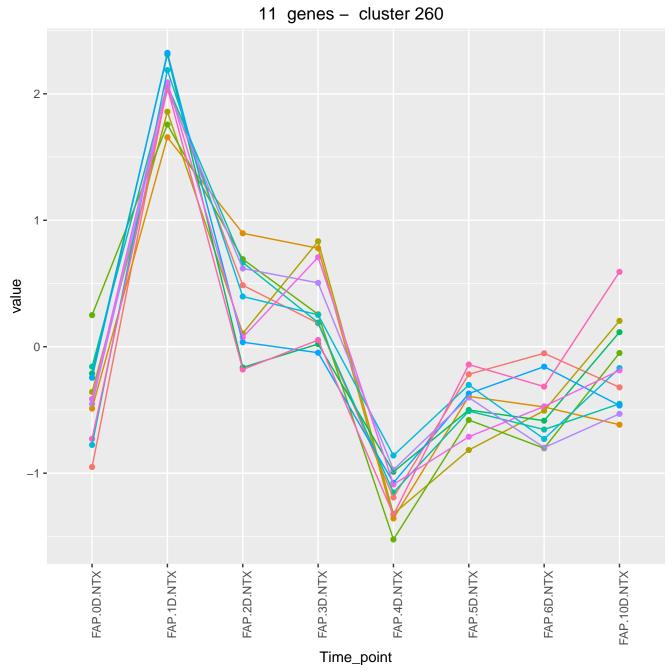


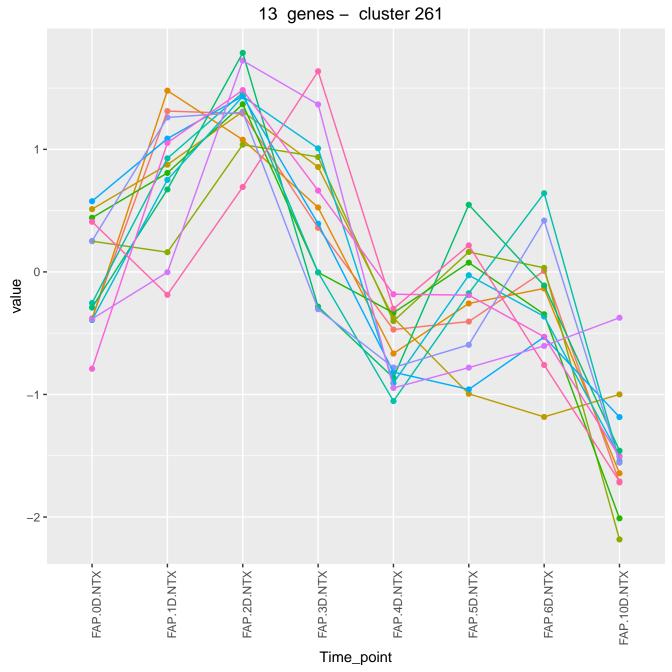


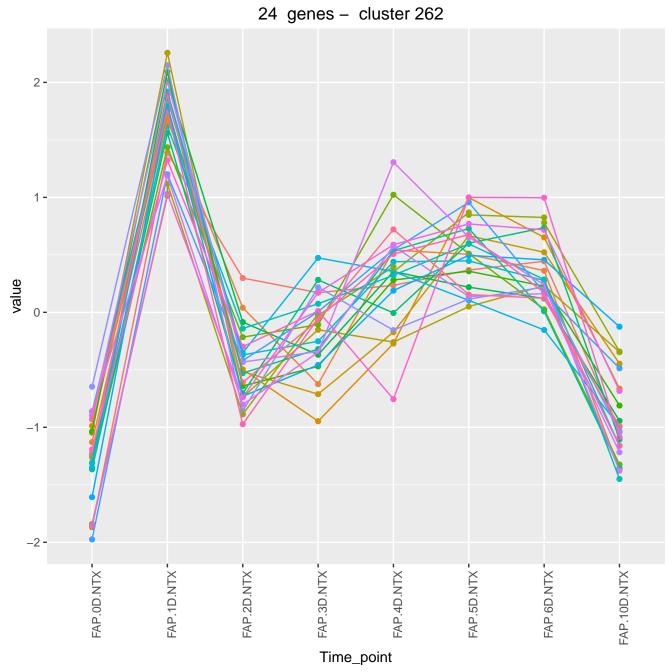


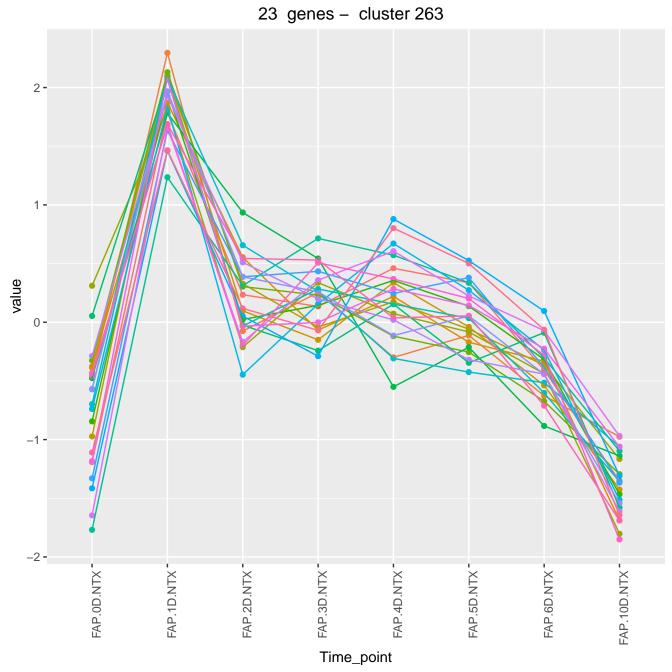


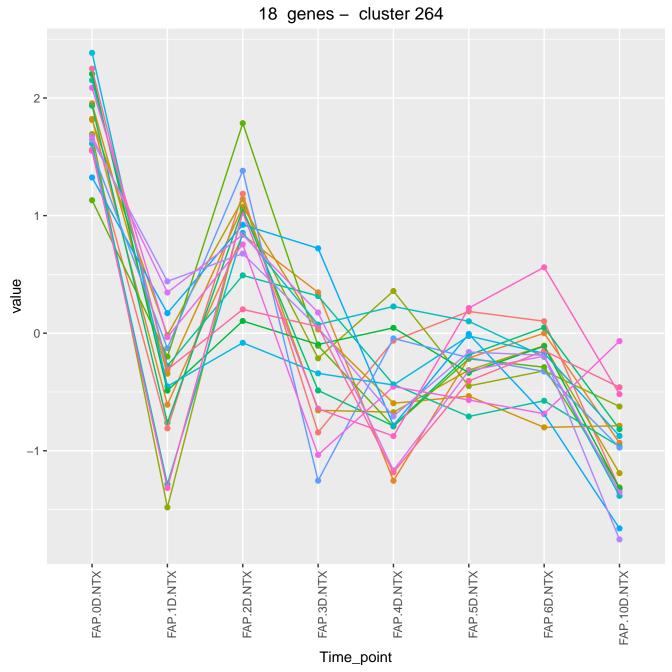


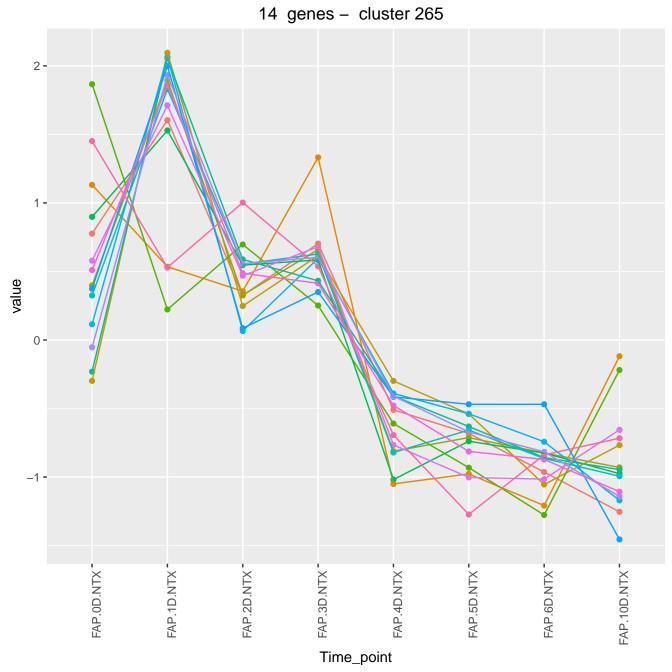


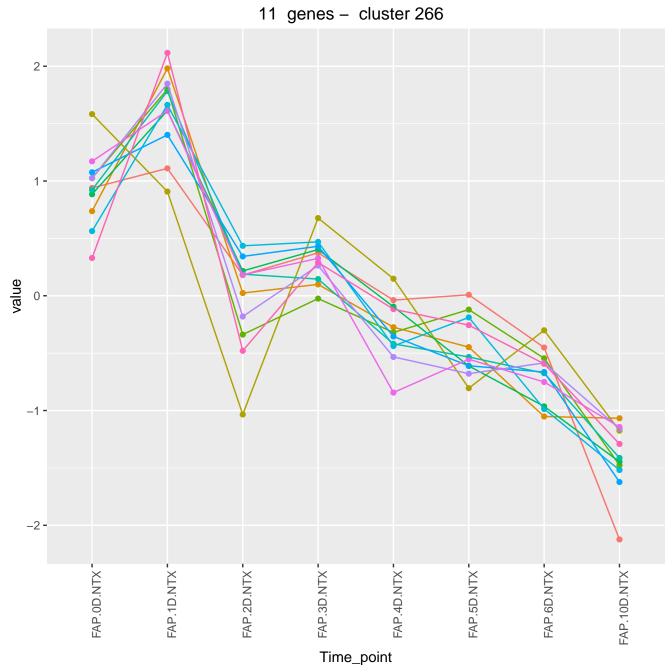


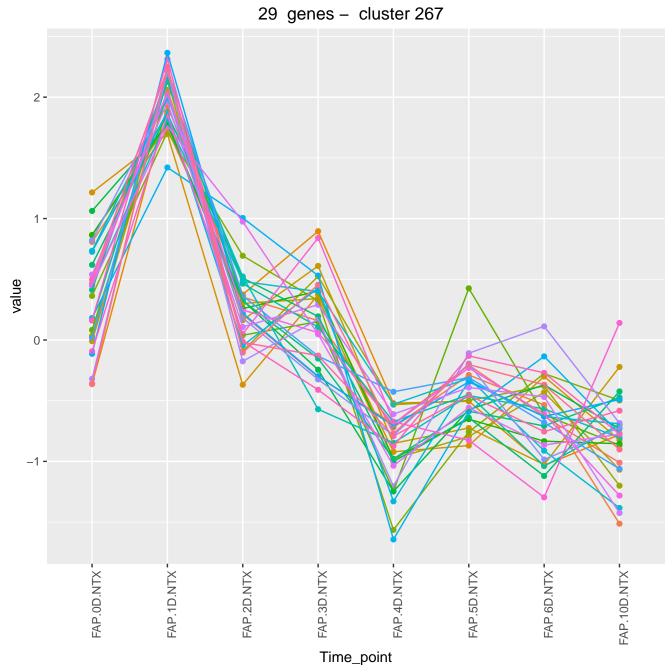


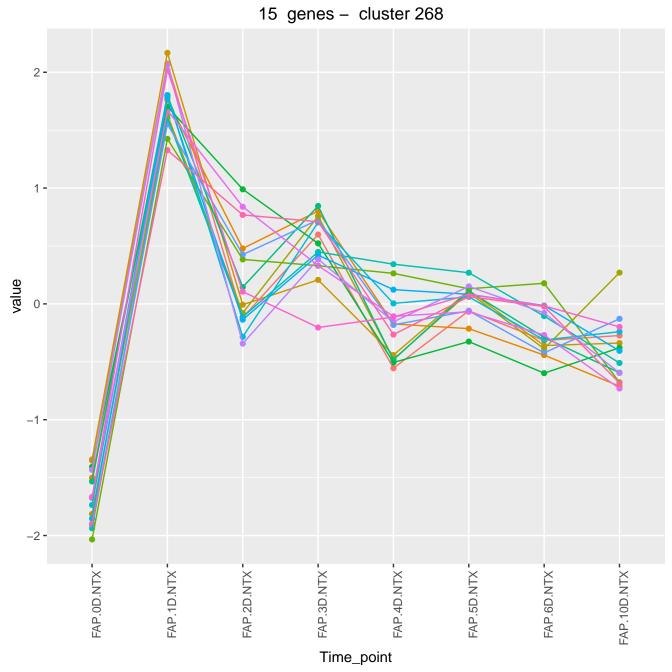


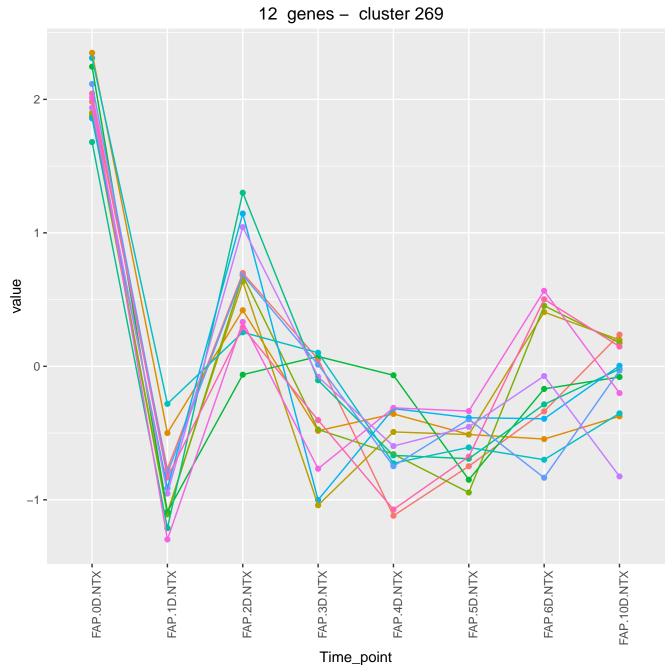


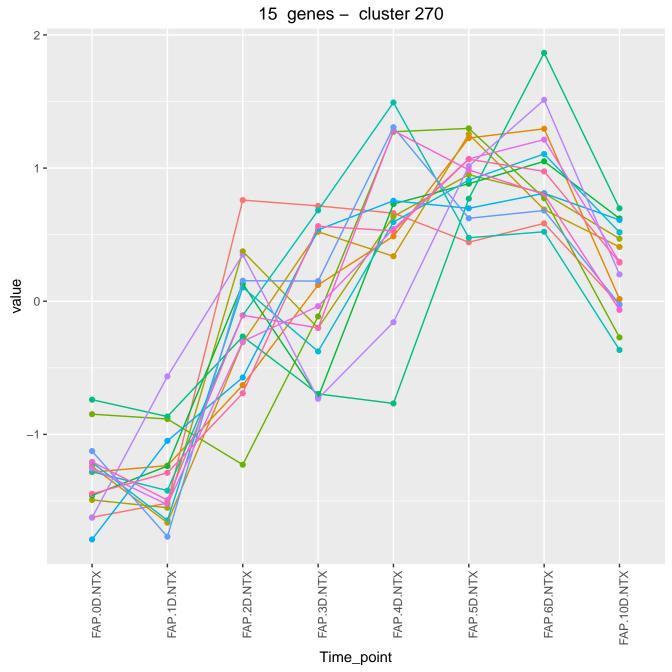


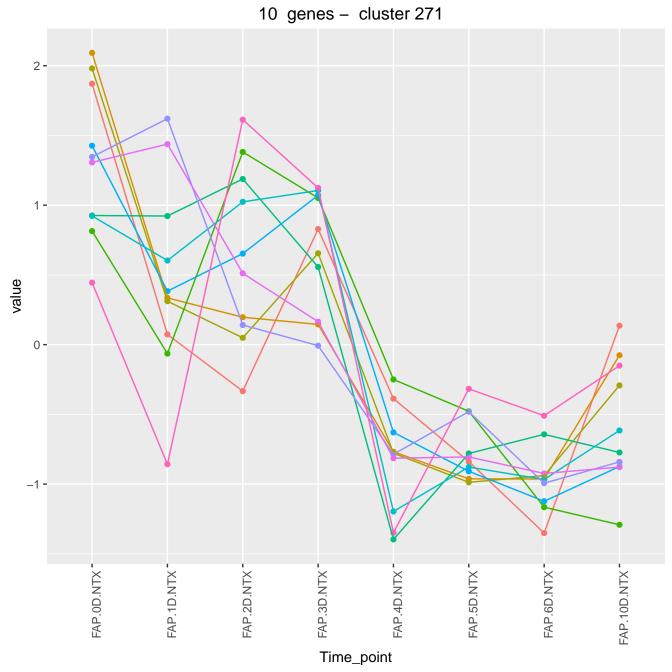




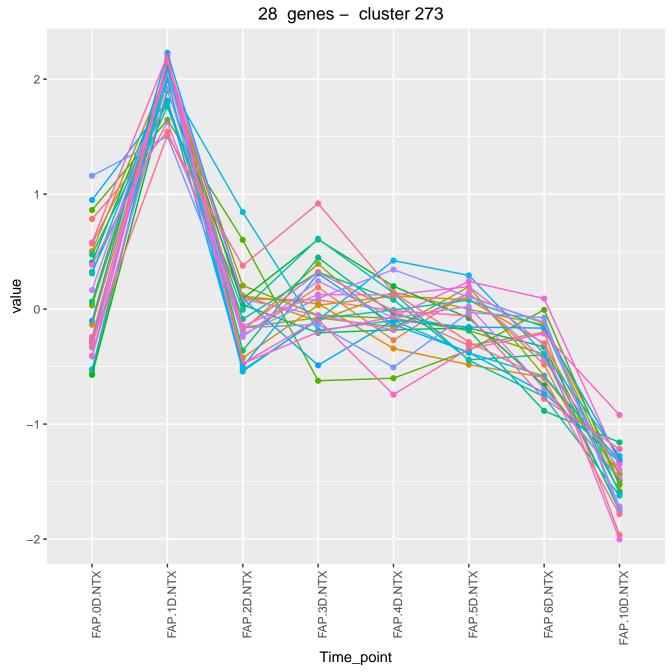




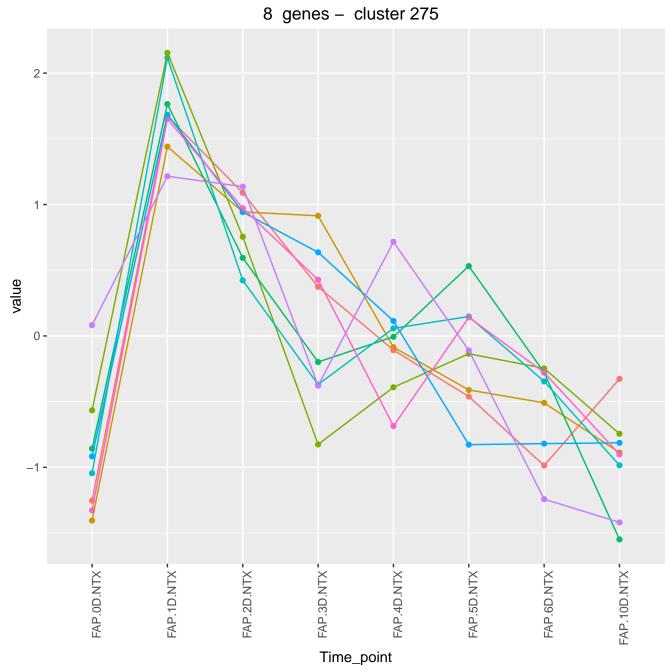


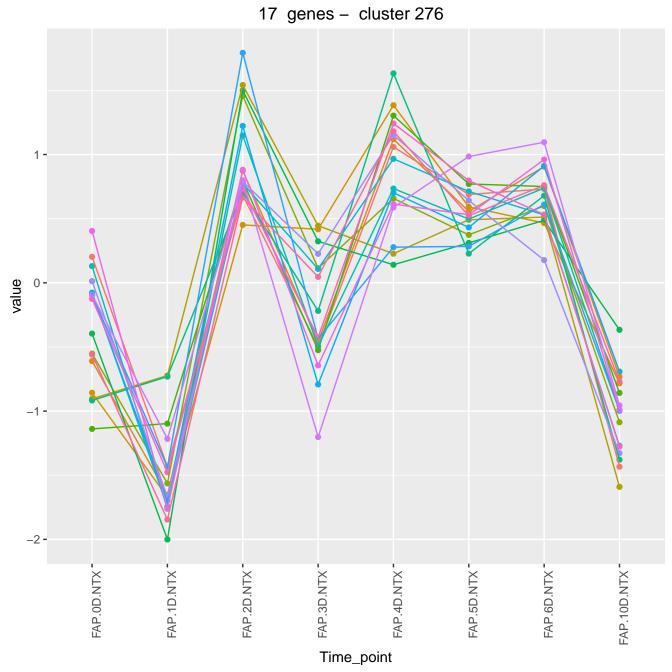


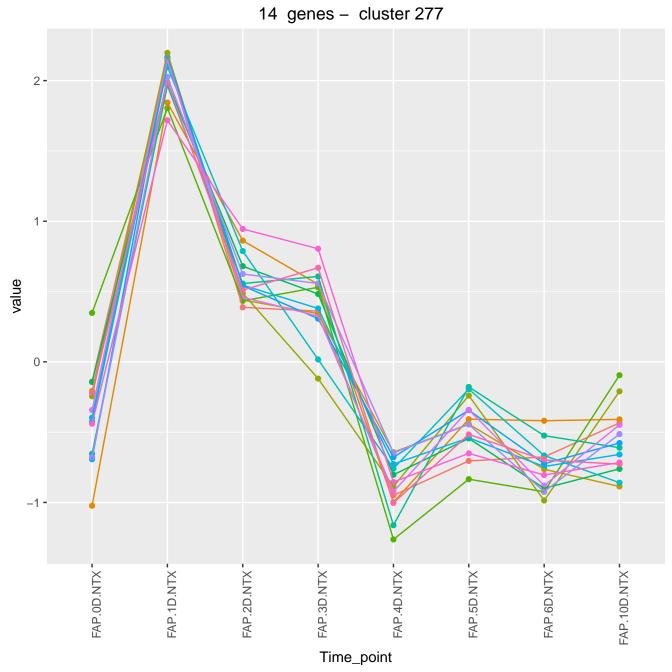
24 genes - cluster 272 1 -0 value \_1 **-**-2 **-**FAP.0D.NTX FAP.1D.NTX FAP.5D.NTX FAP.10D.NTX FAP.2D.NTX FAP.3D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

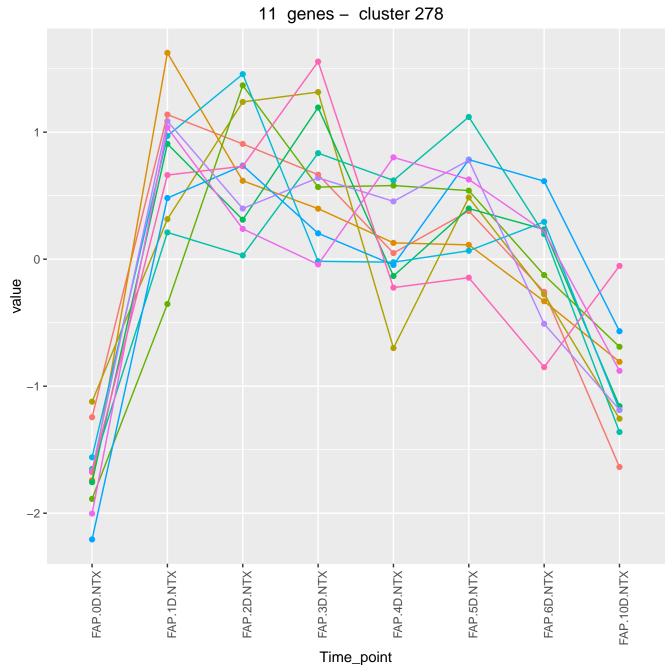


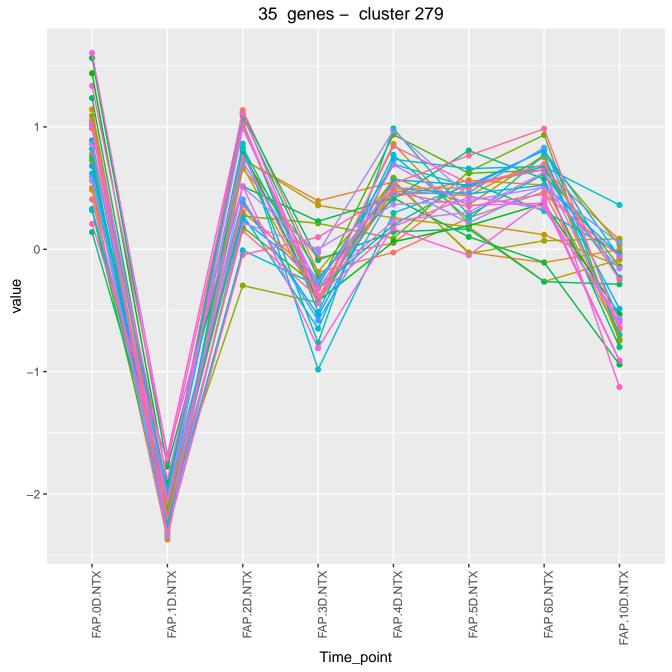
18 genes - cluster 274 1 -0 value \_1 **-**-2 **-**FAP.0D.NTX 7 FAP.10D.NTX -FAP.1D.NTX FAP.3D.NTX -FAP.5D.NTX FAP.2D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

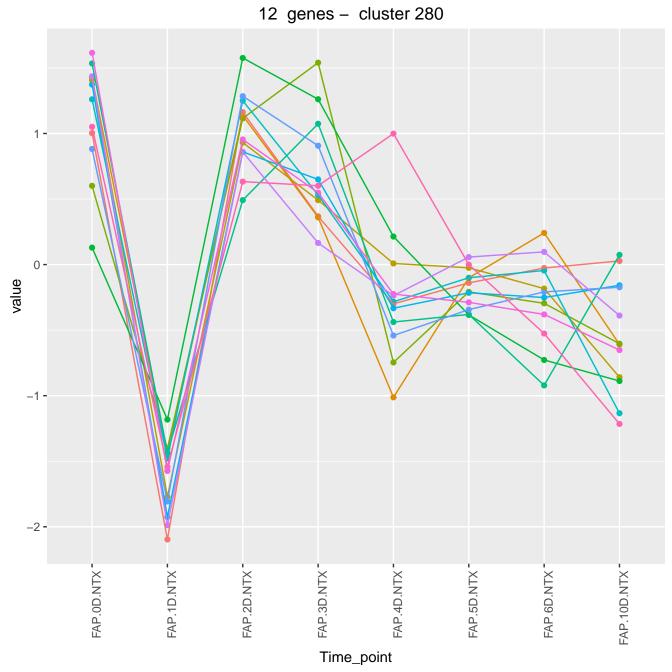


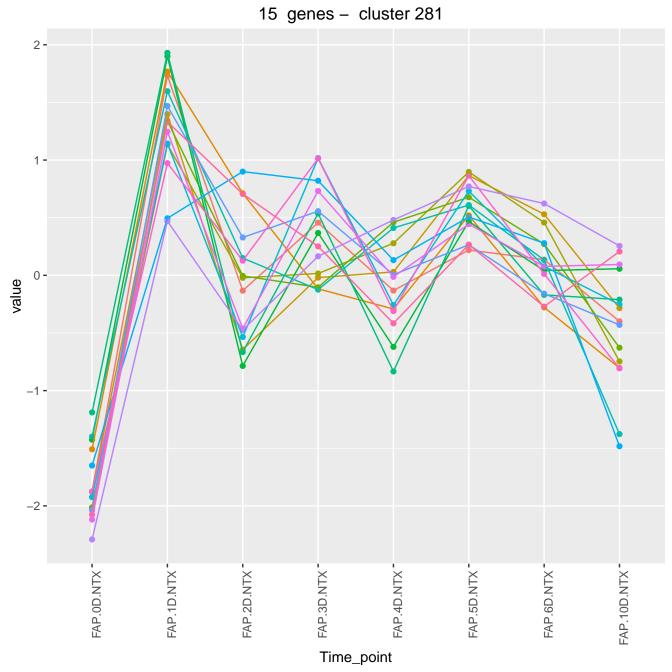


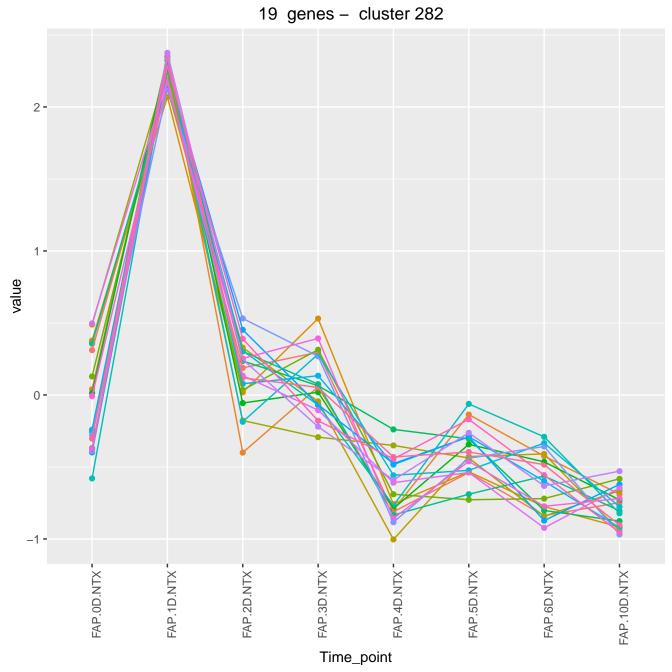


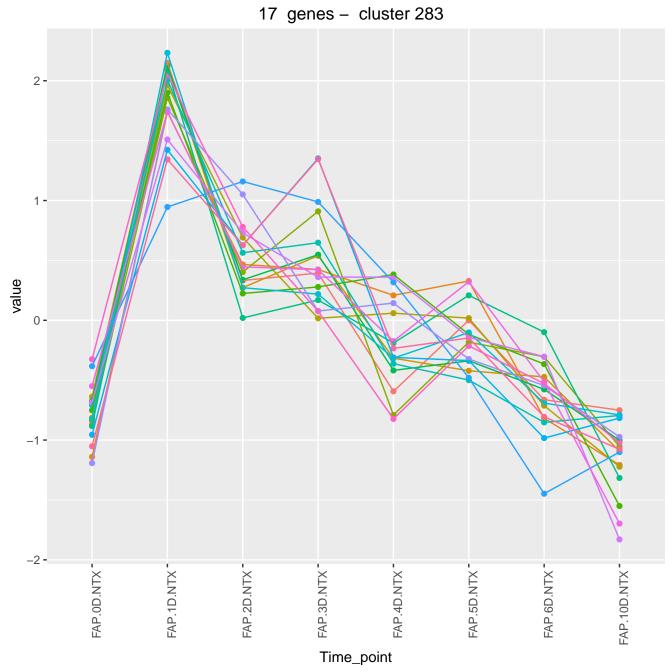


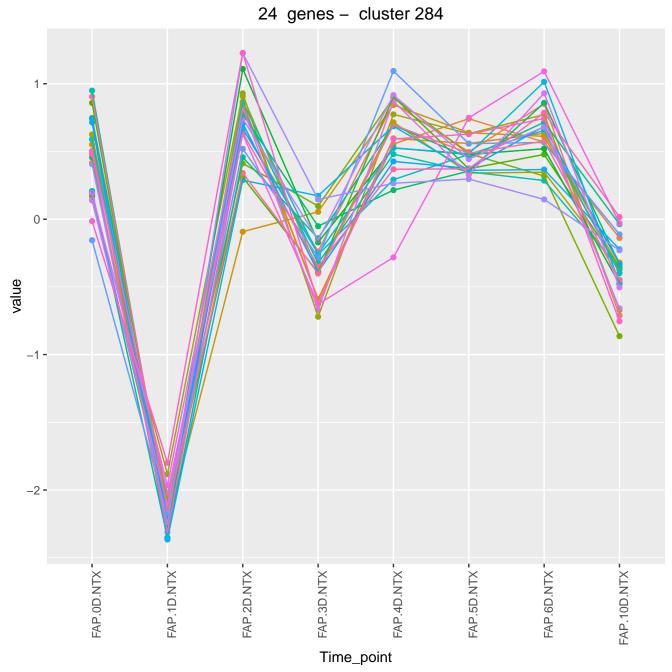


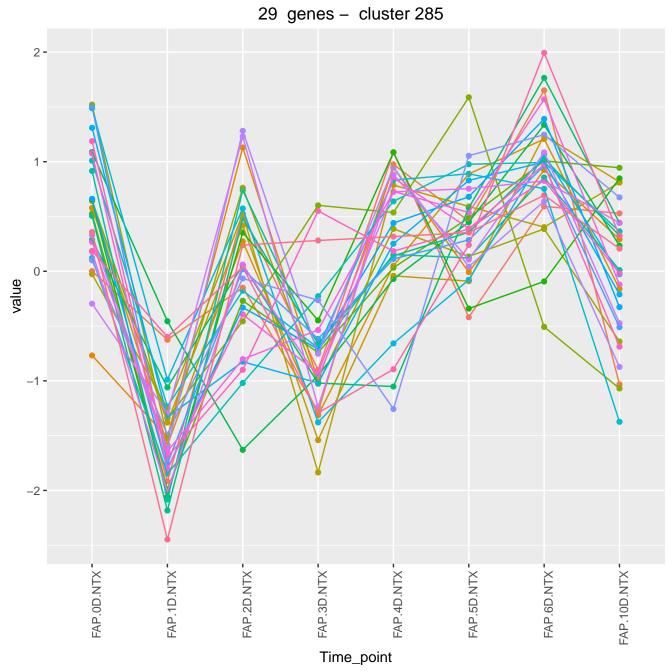


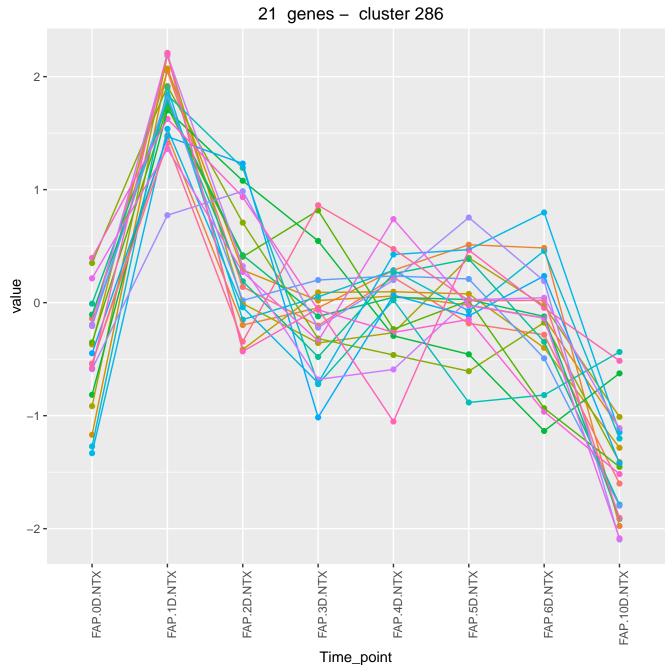


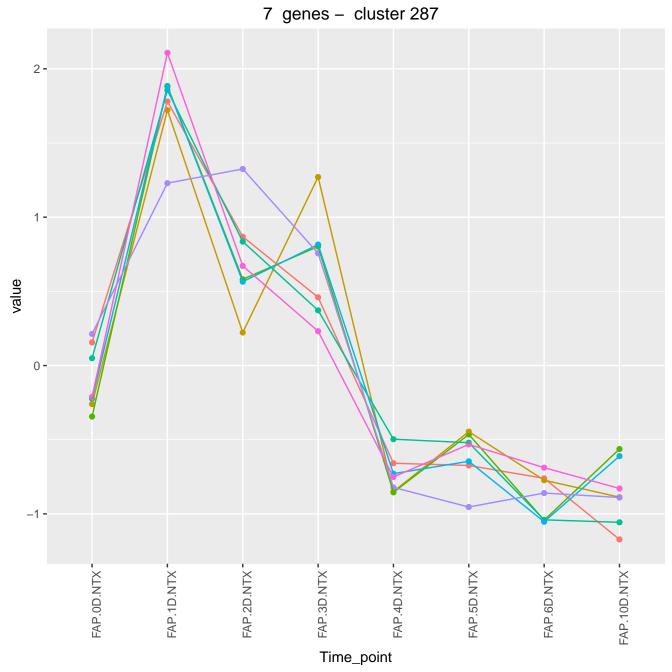


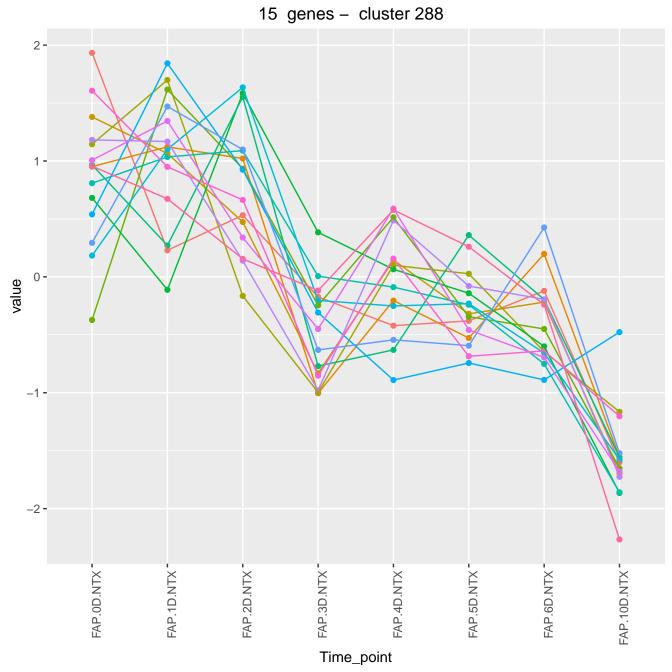


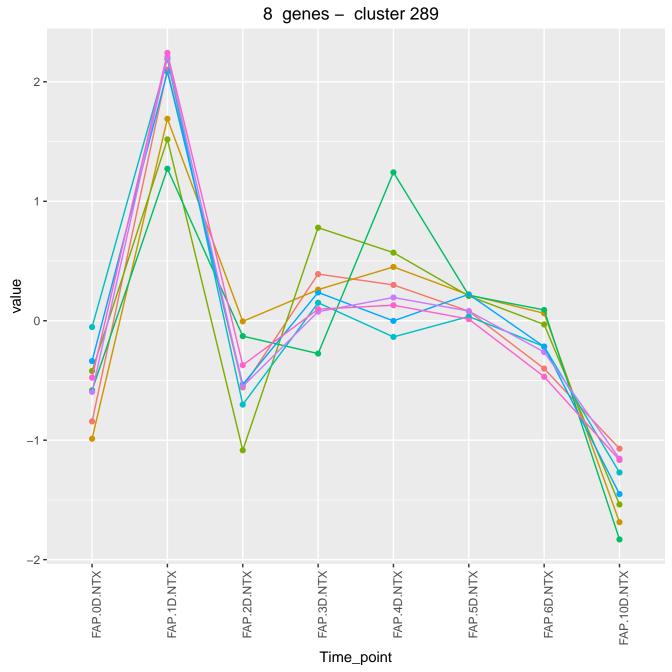


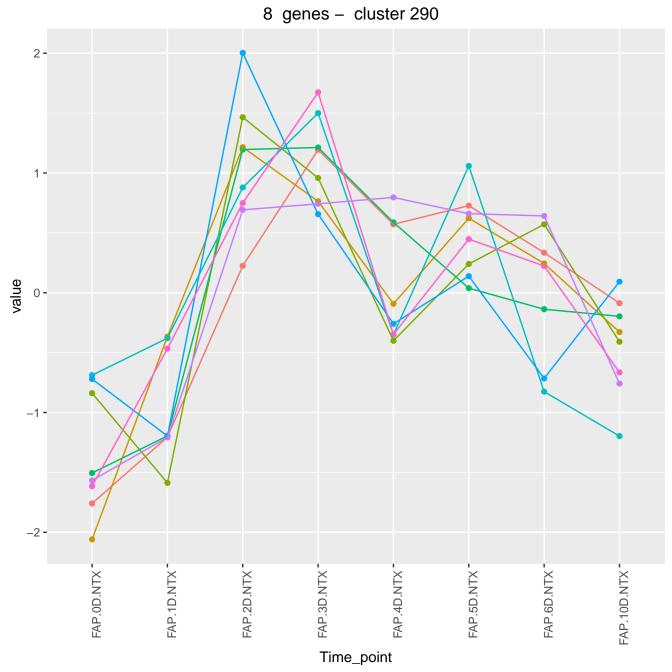


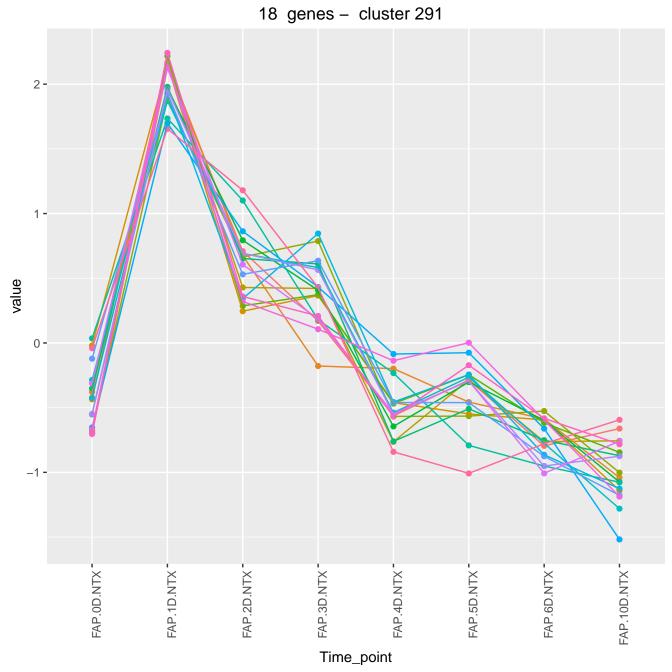


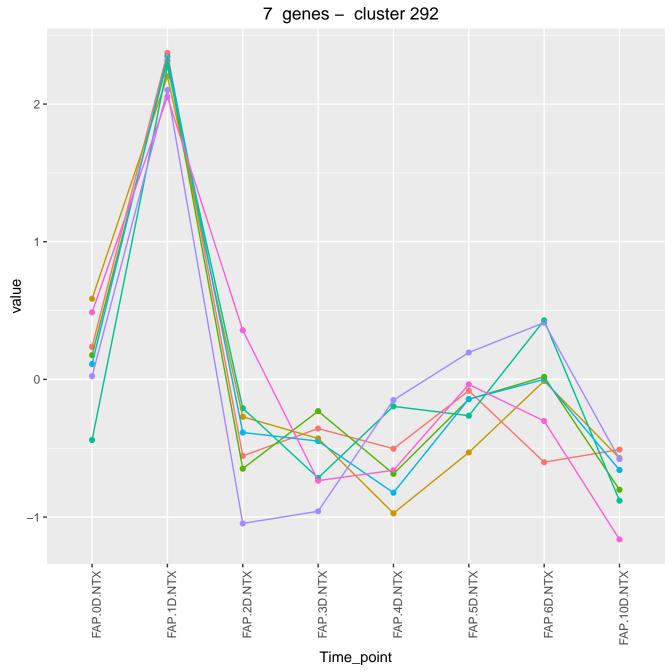


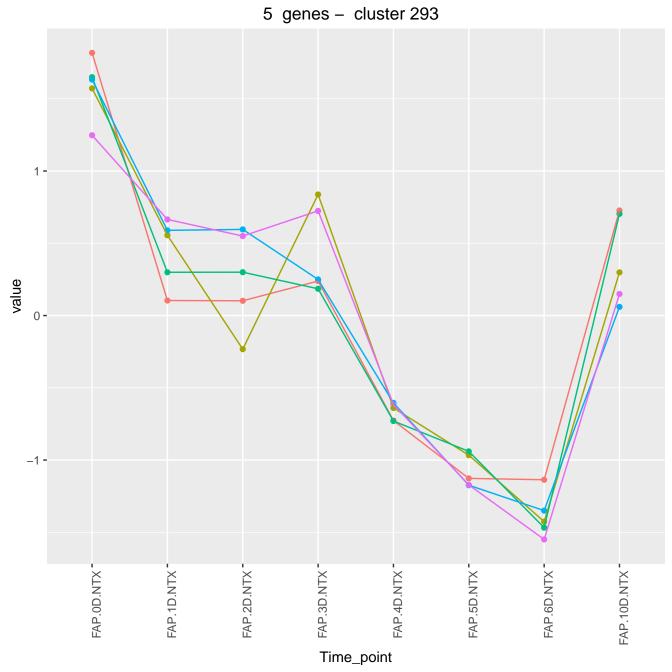


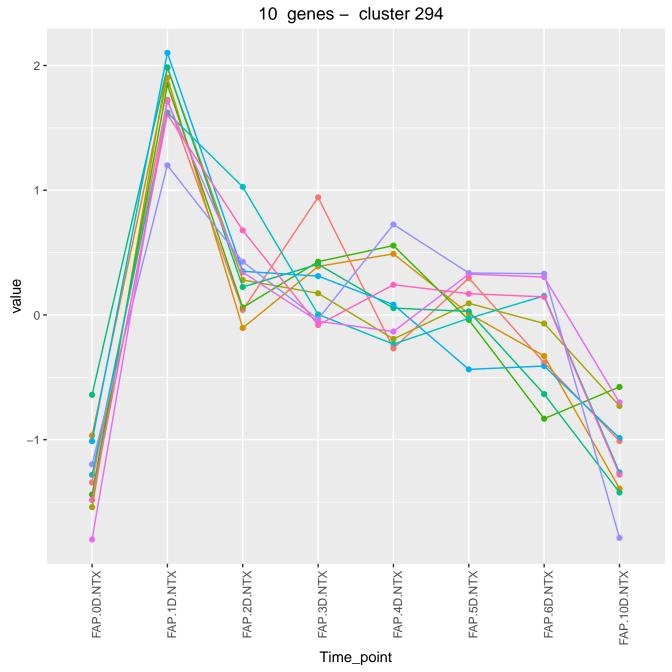


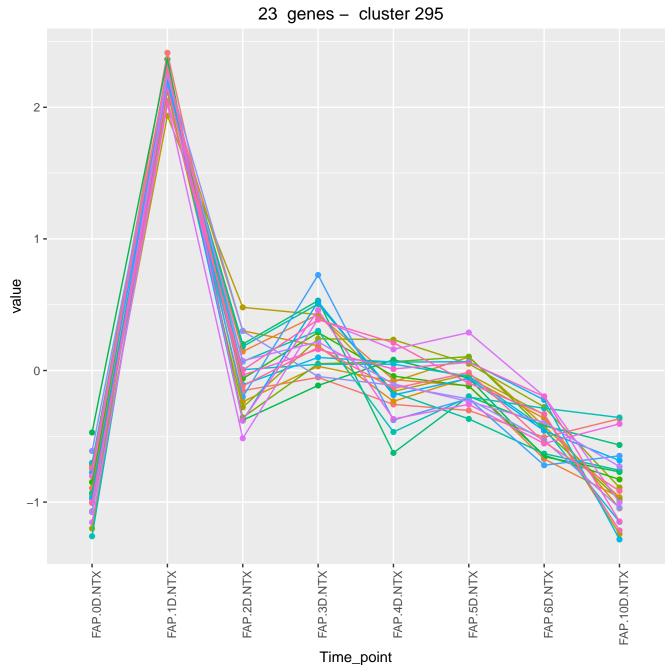


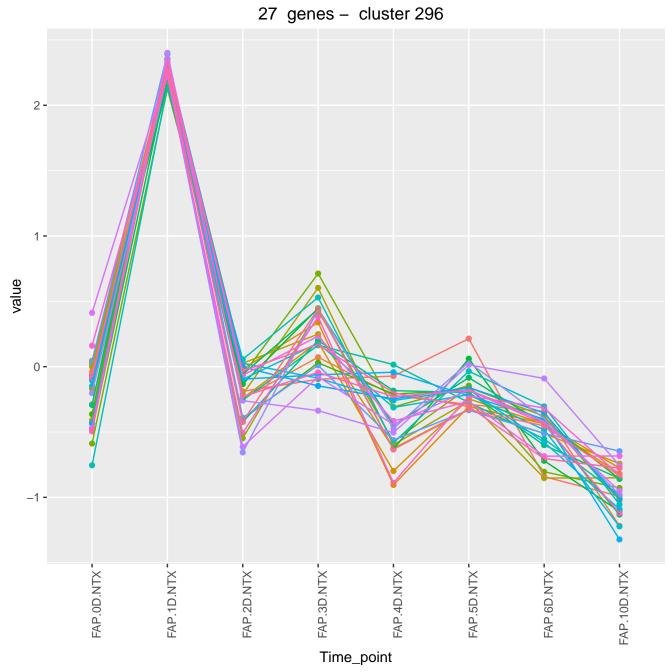


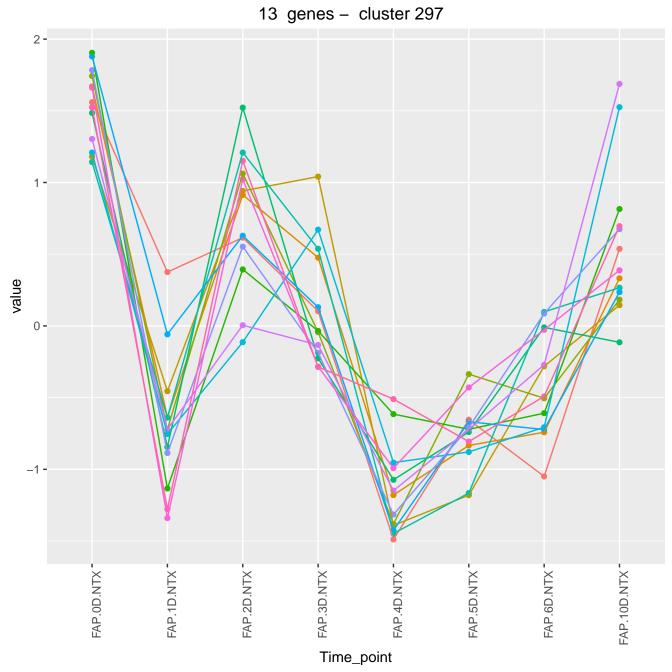


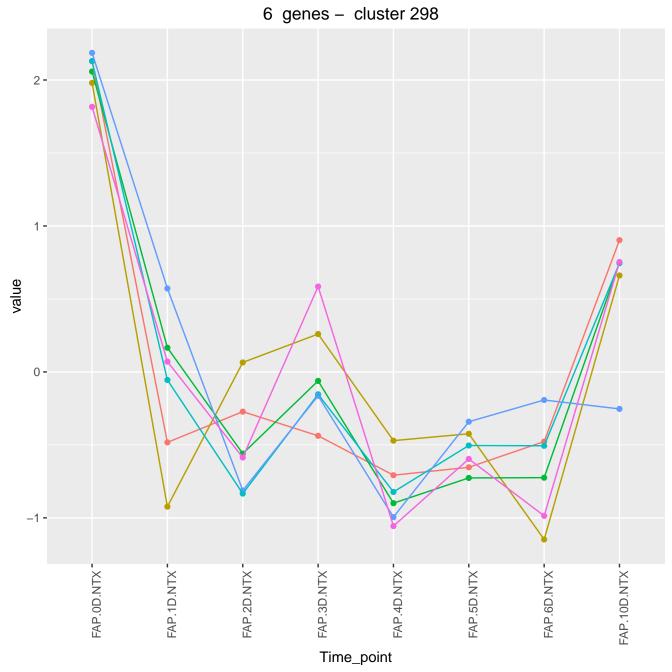


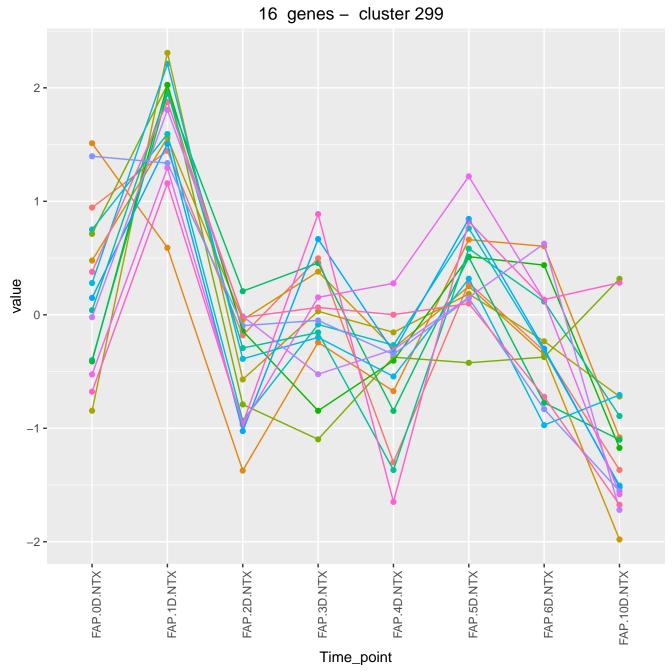


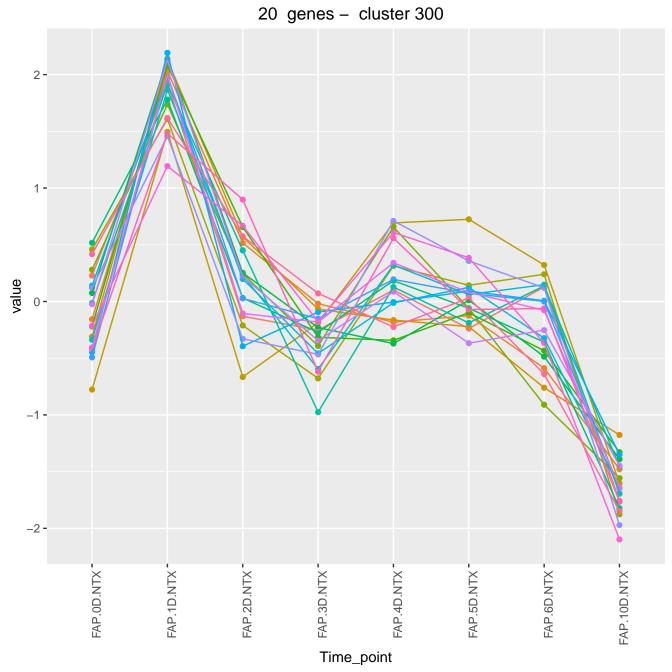


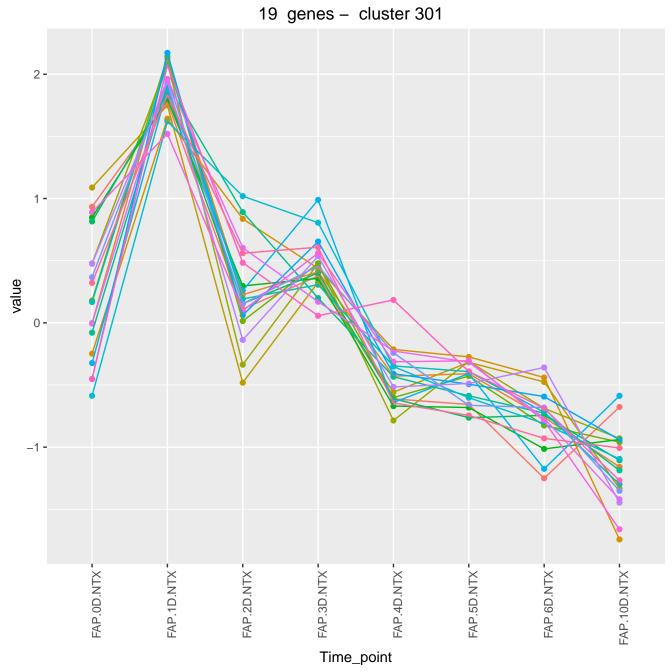


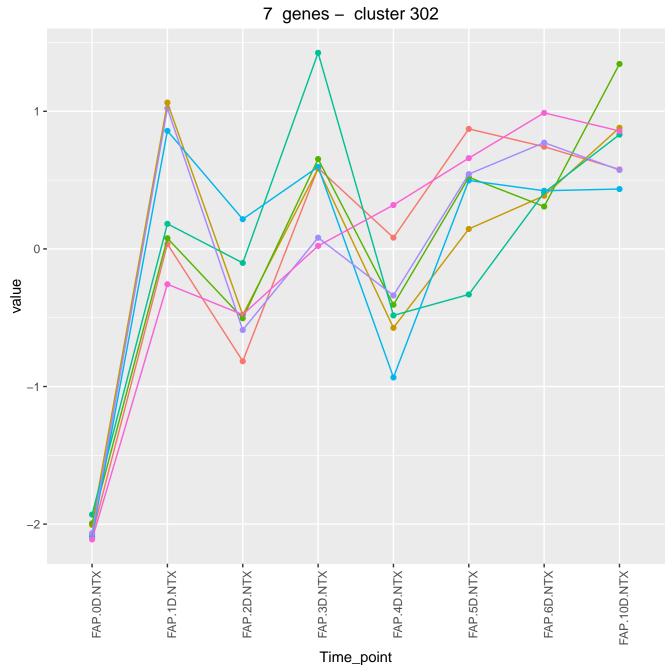




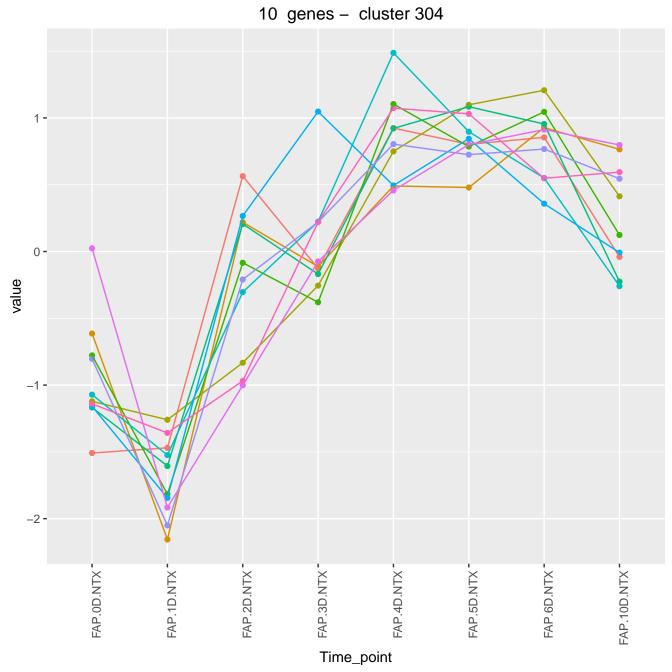


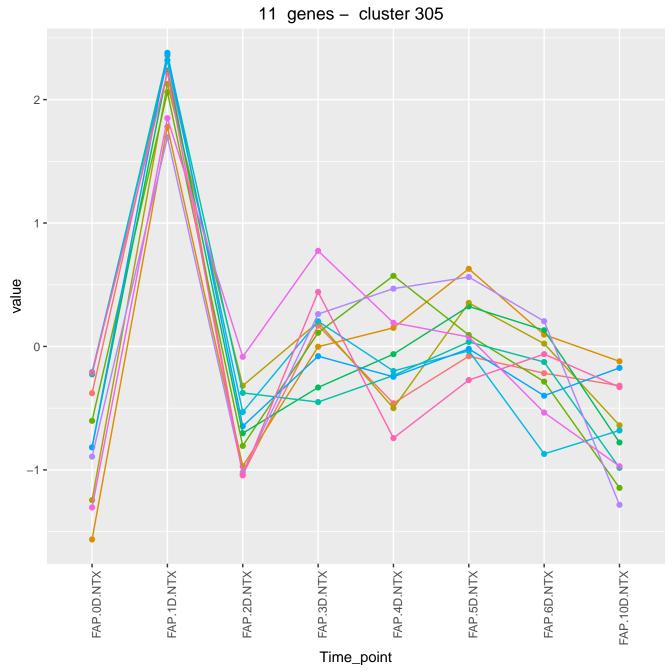


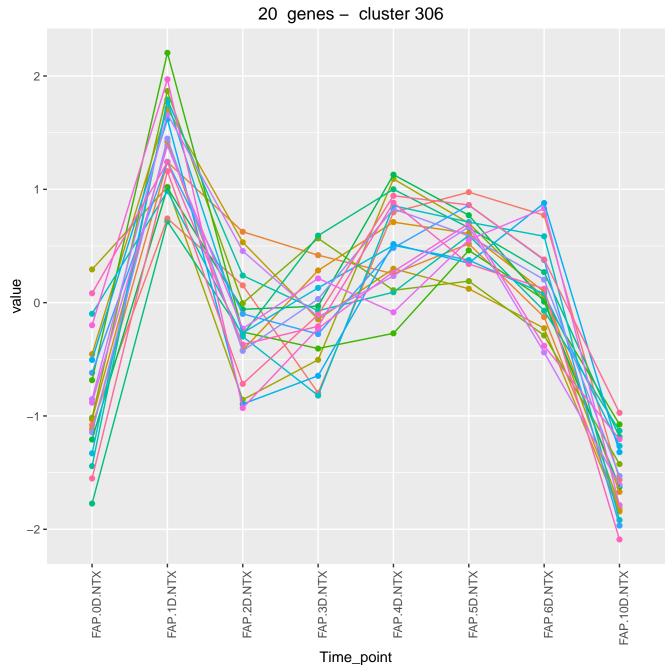


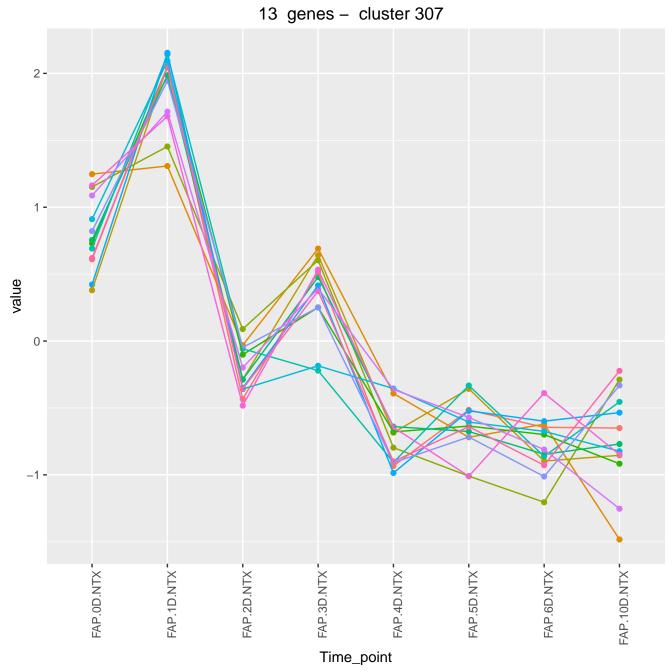


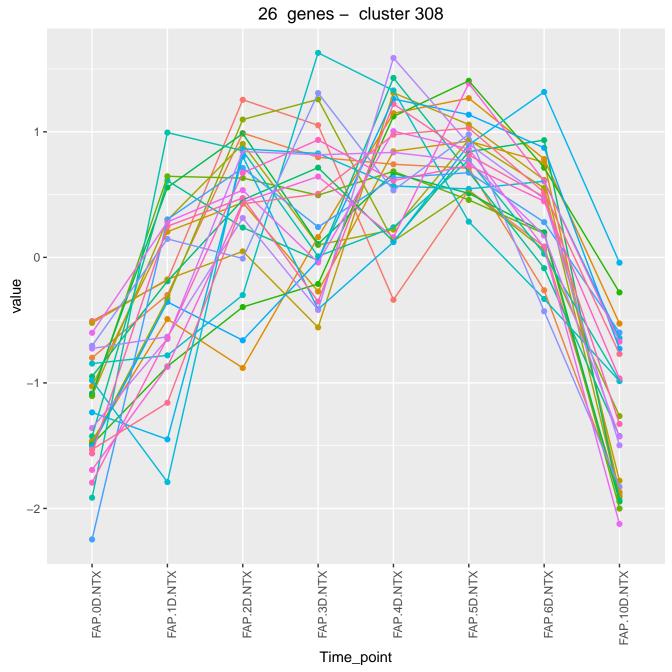
37 genes - cluster 303 2-1 value -1 **-**-2 **-**FAP.0D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.2D.NTX FAP.3D.NTX FAP.1D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

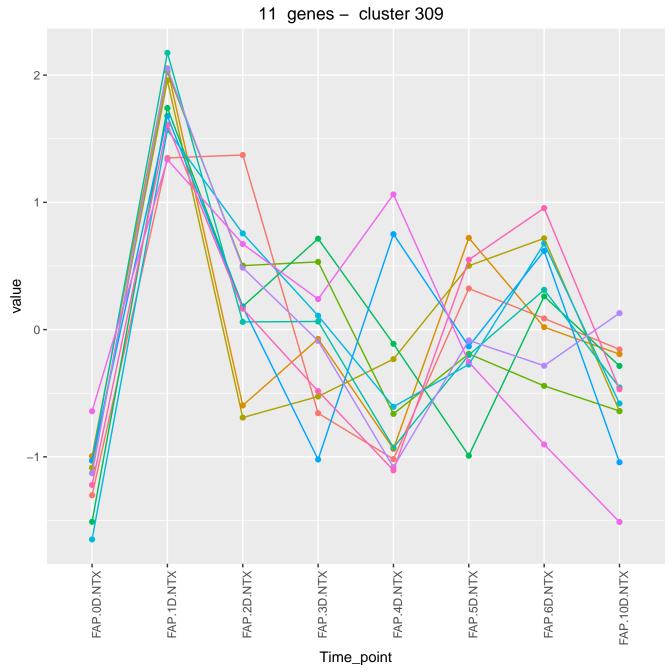


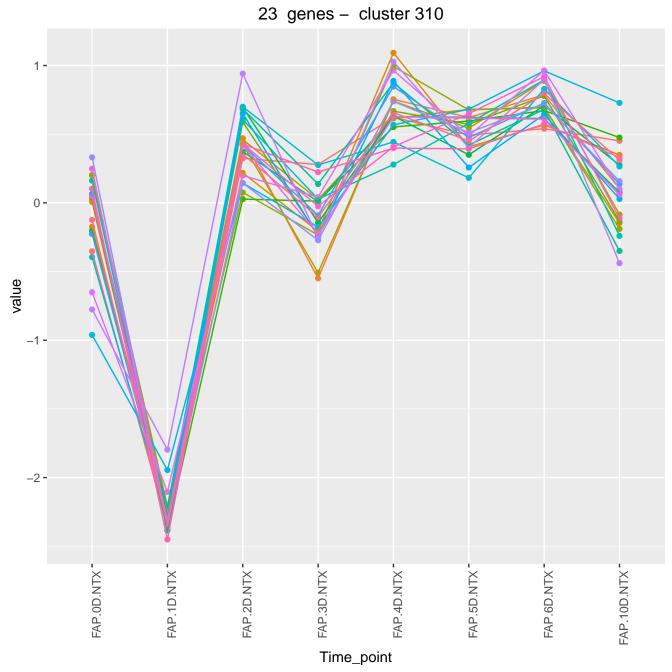


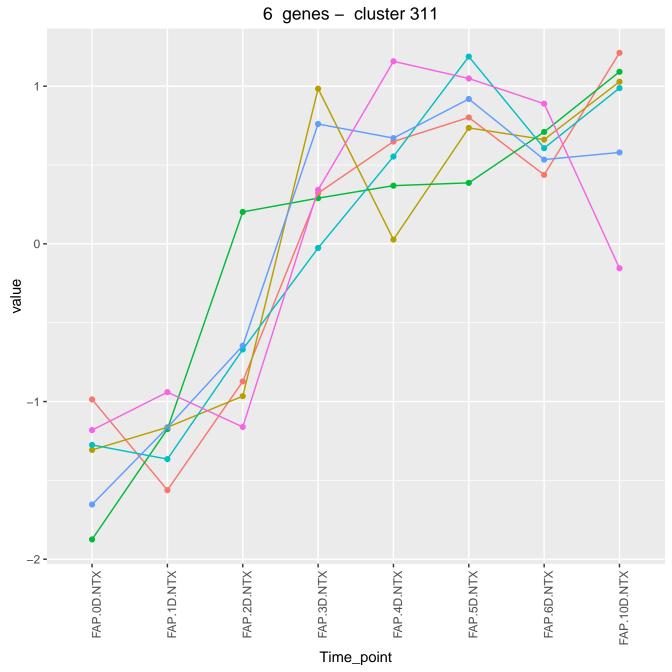


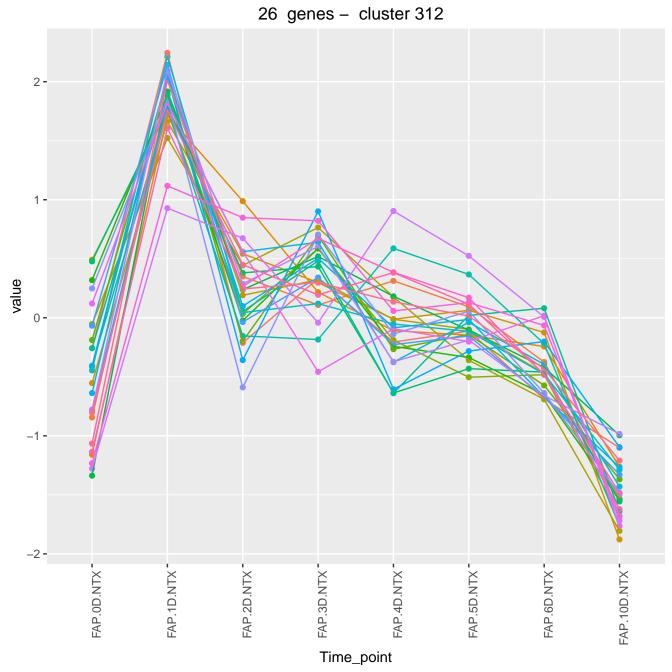


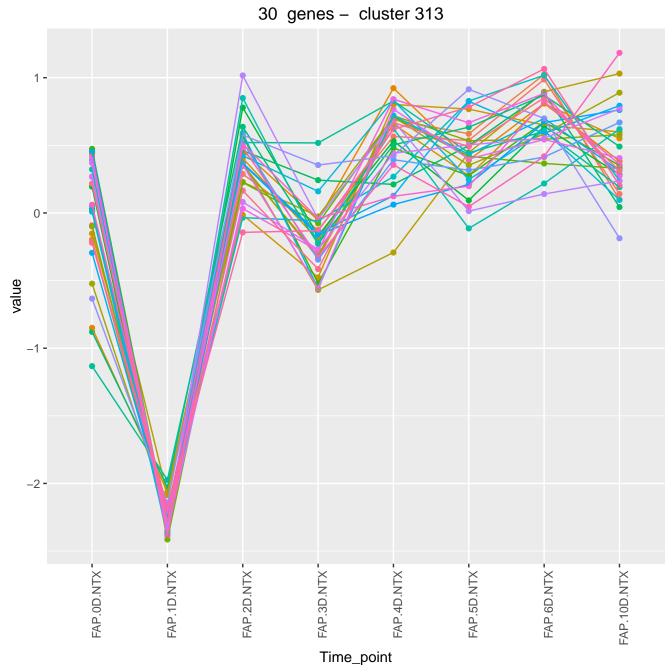


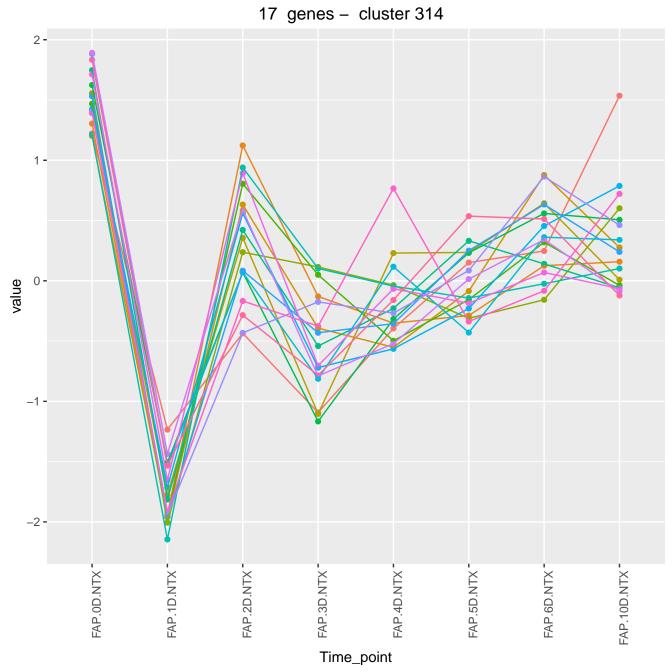


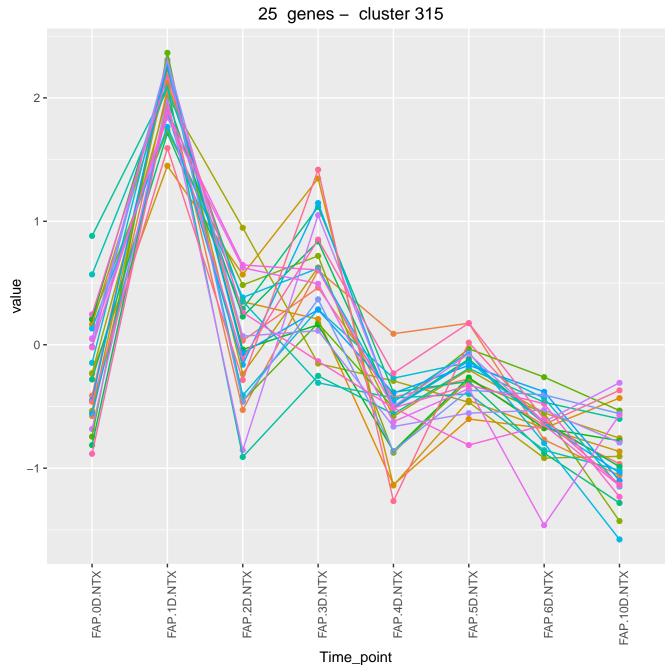


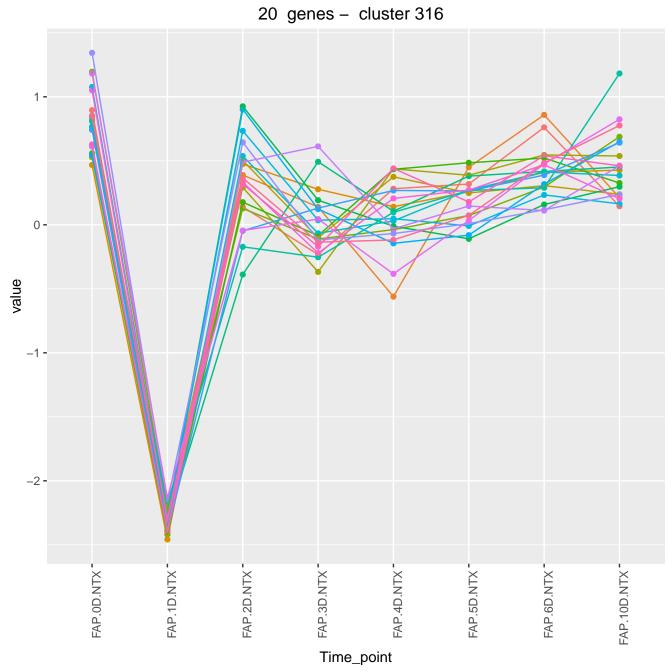


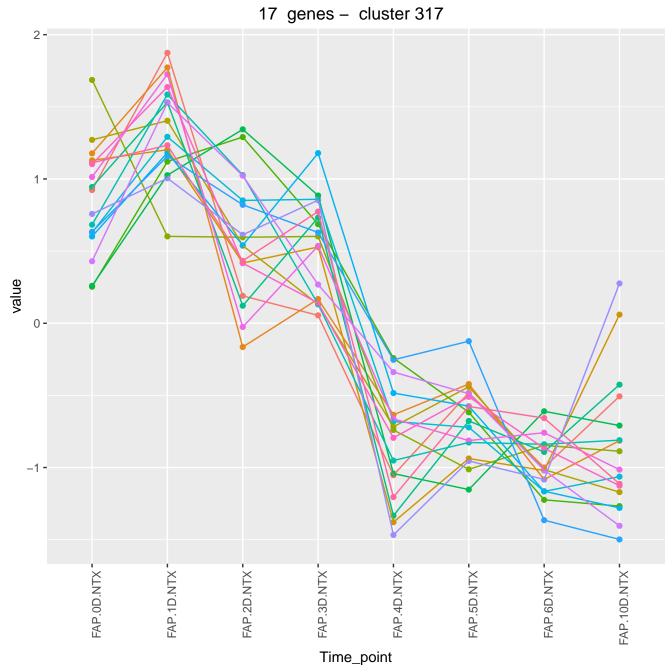


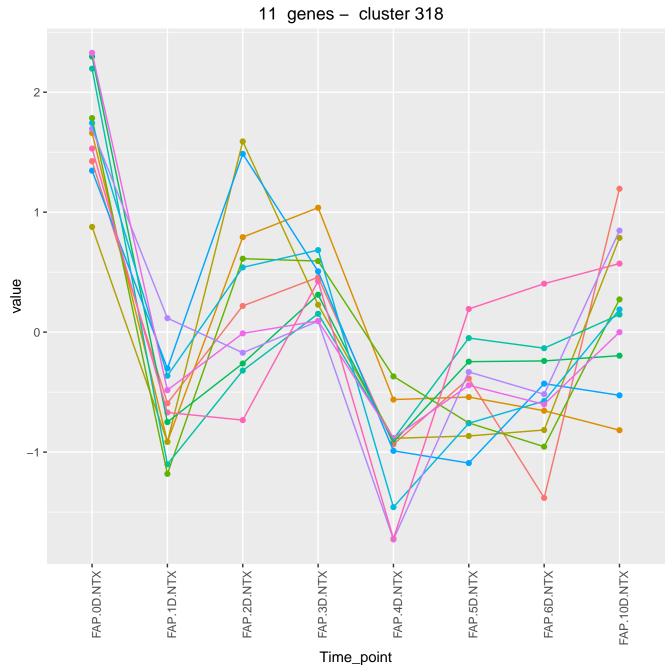


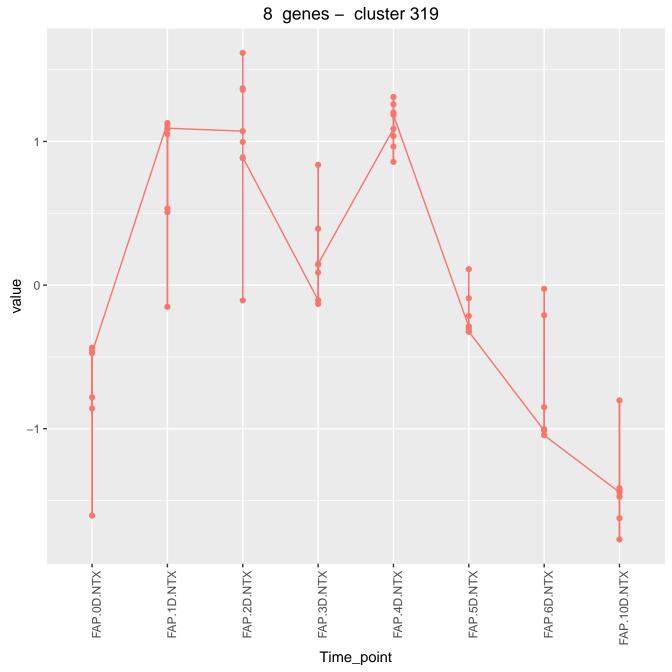


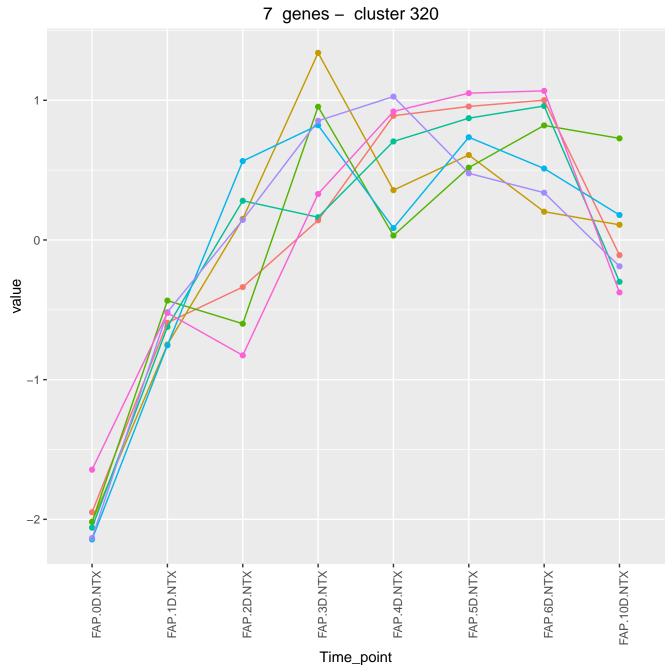


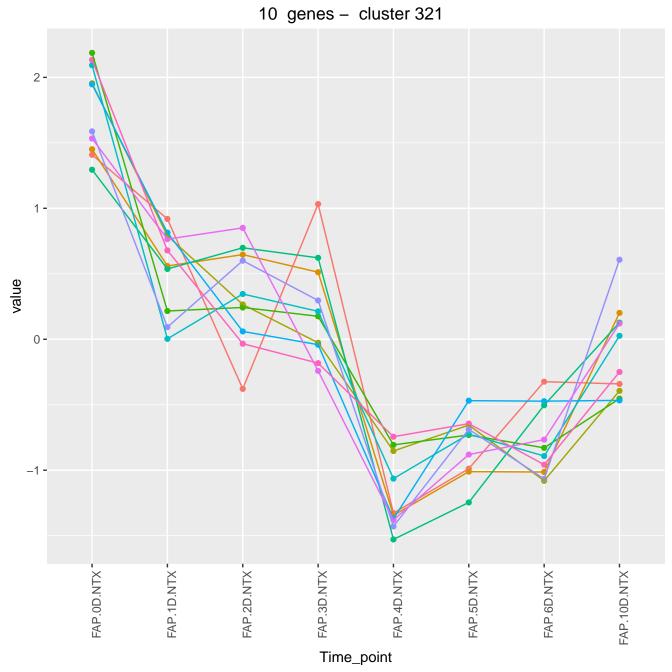


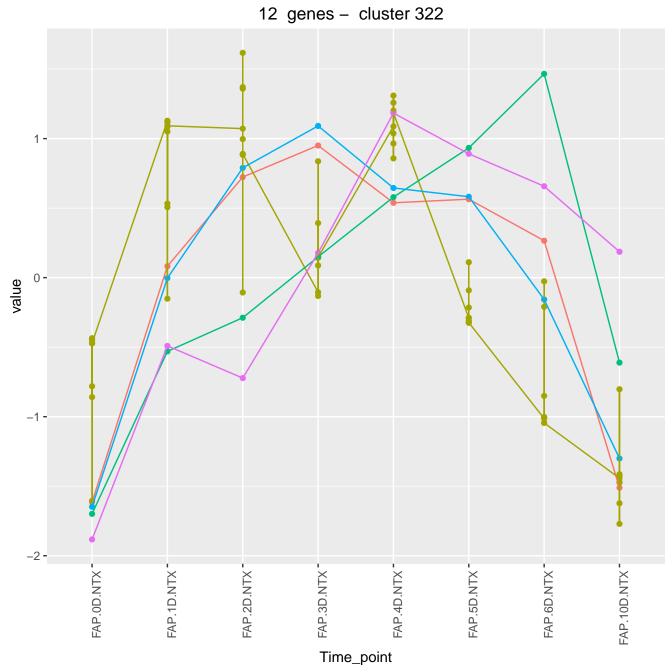


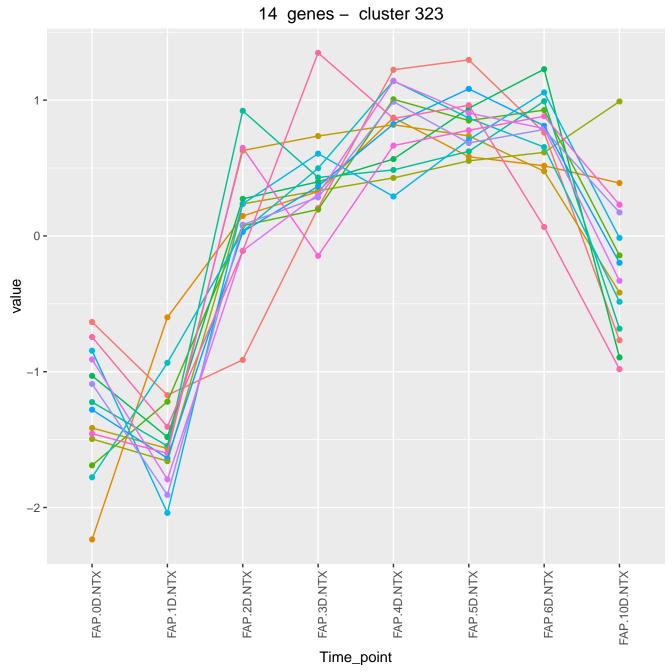


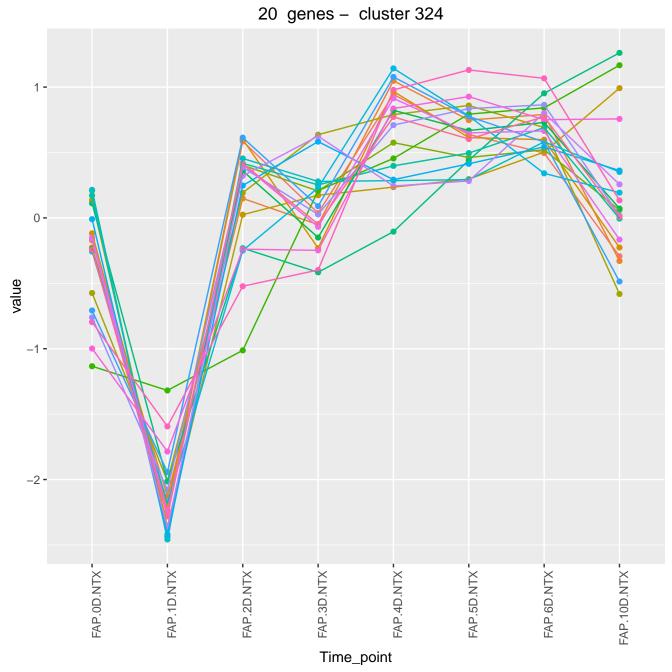




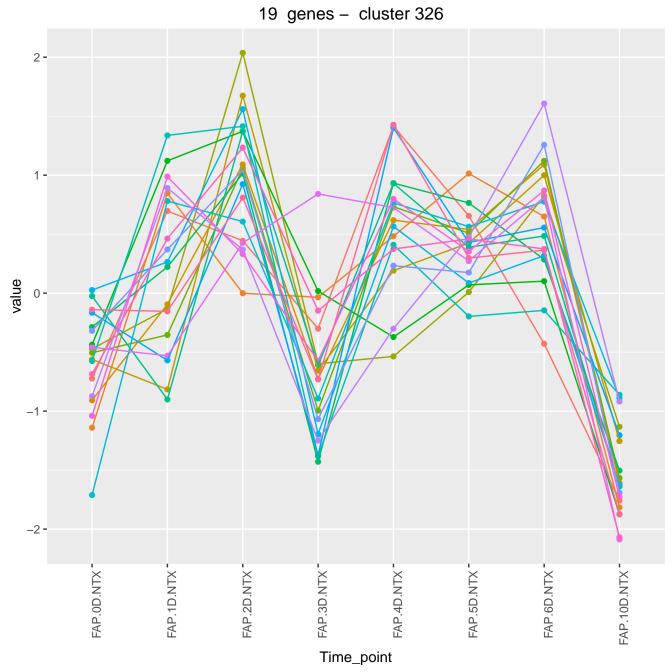


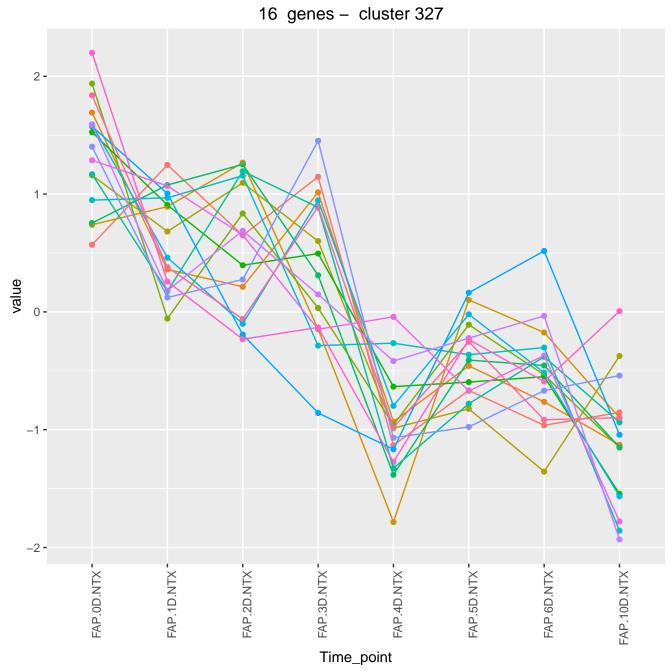


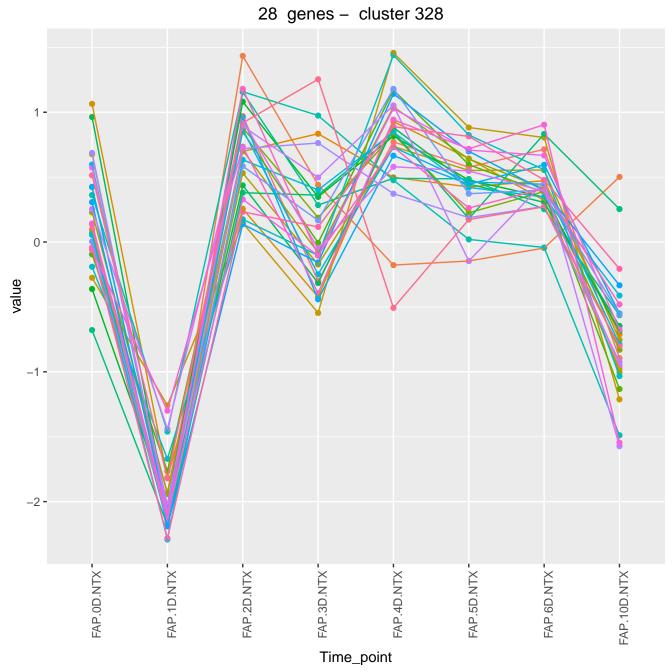


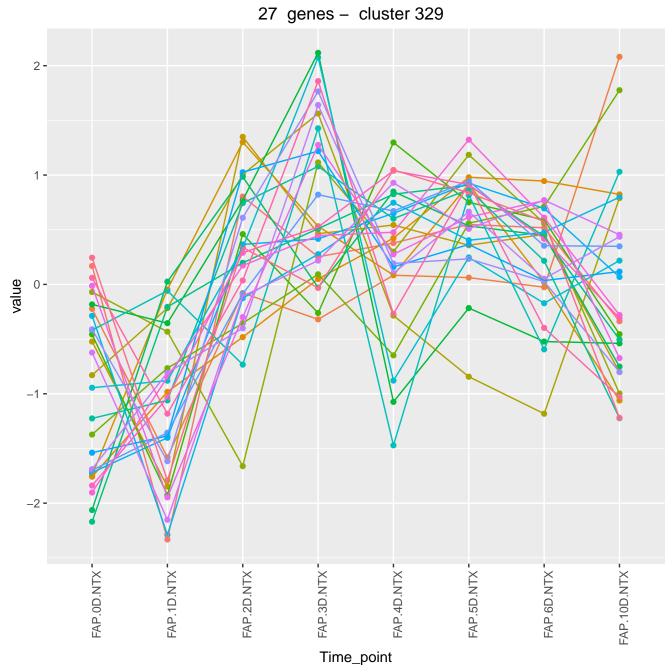


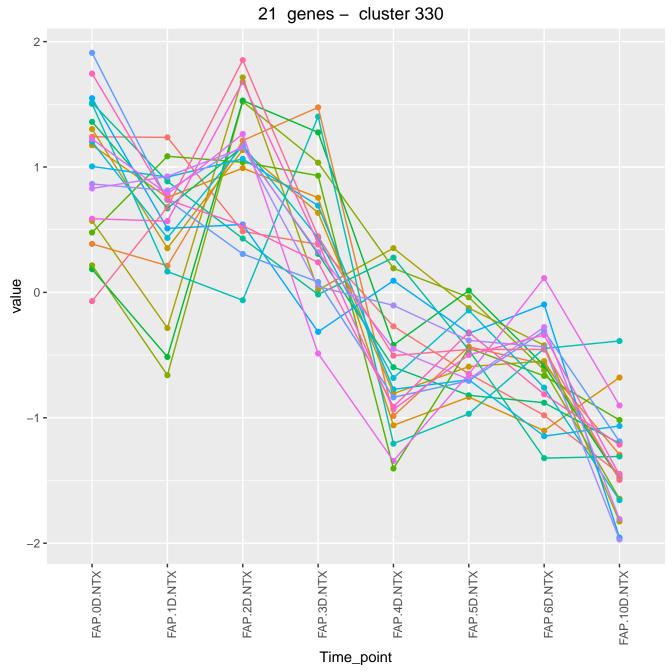
17 genes - cluster 325 1 -0 value **-1** --2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.6D.NTX FAP.4D.NTX Time\_point

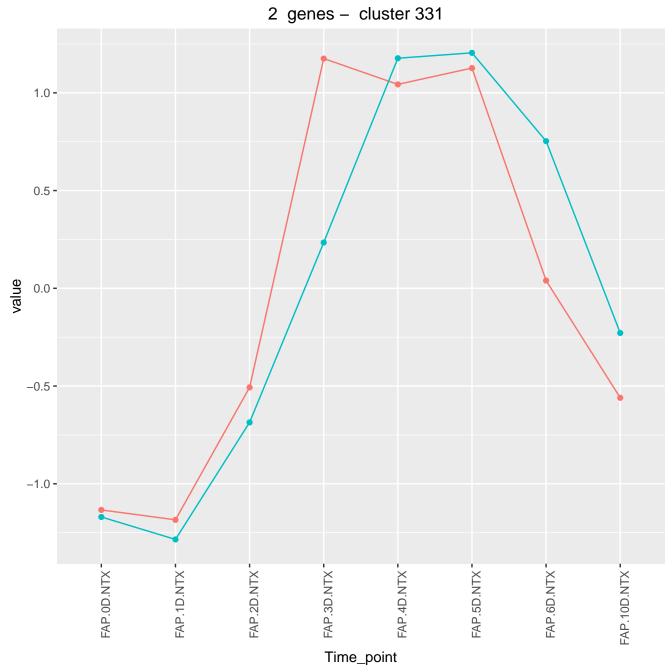


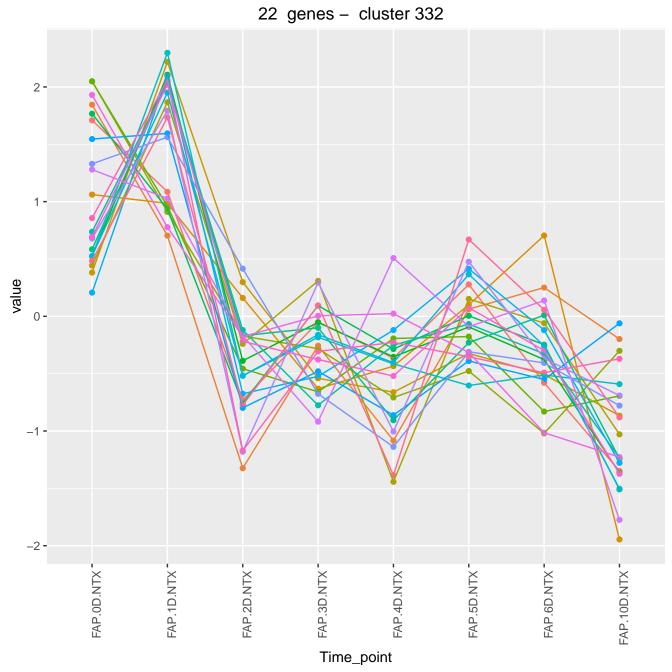


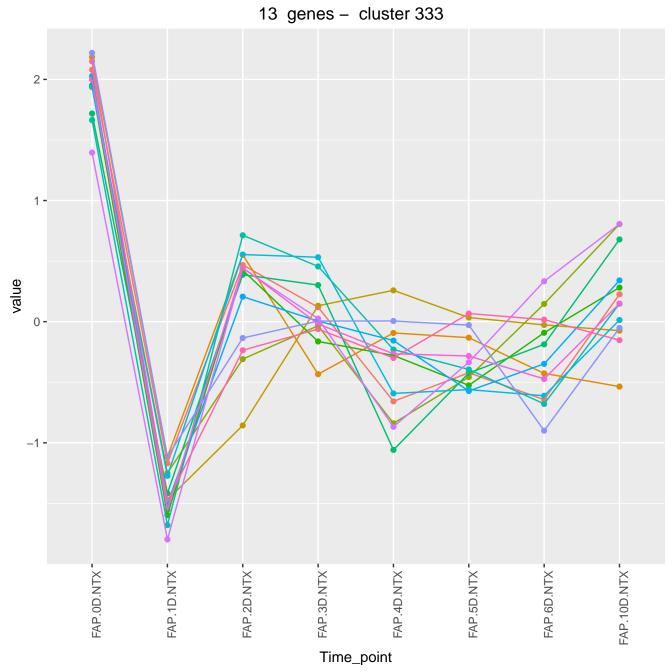


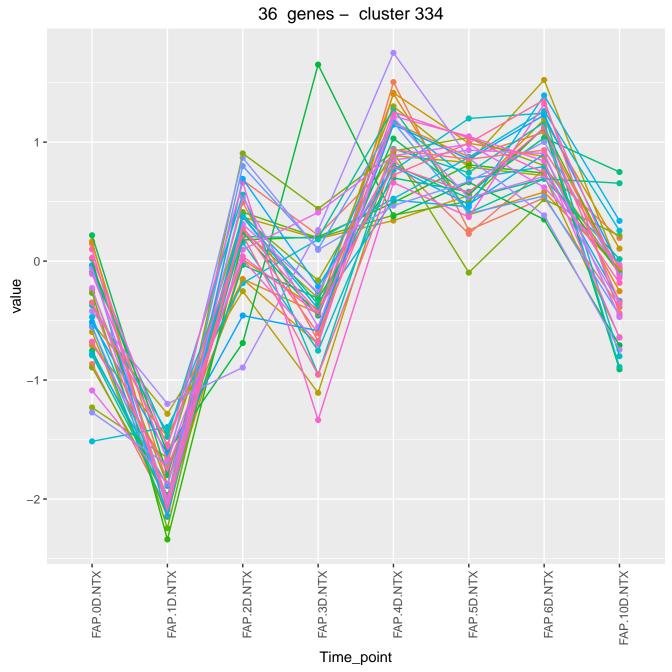


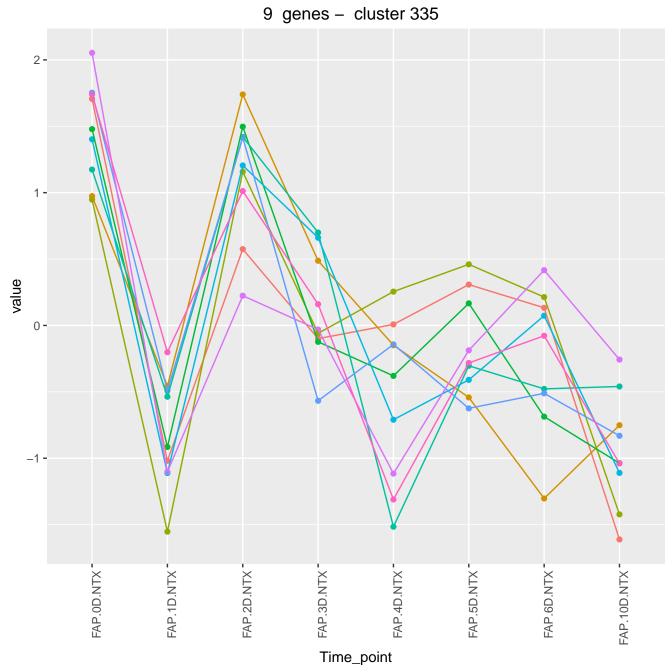


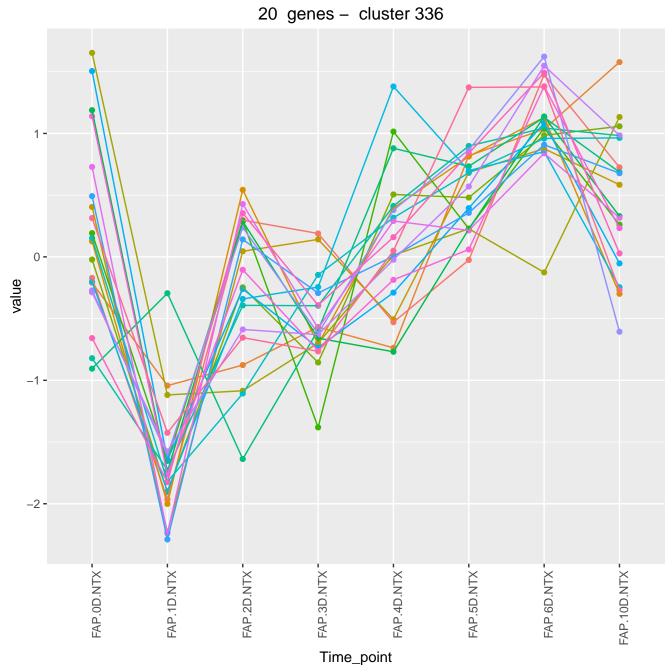


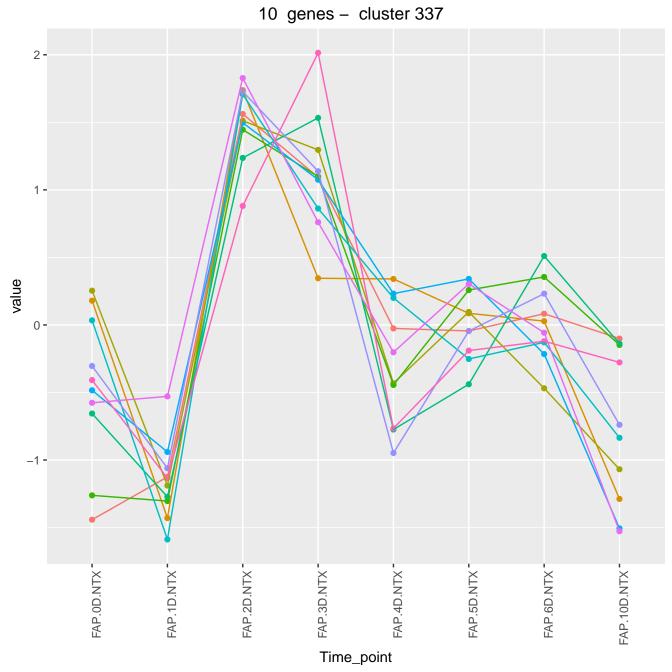


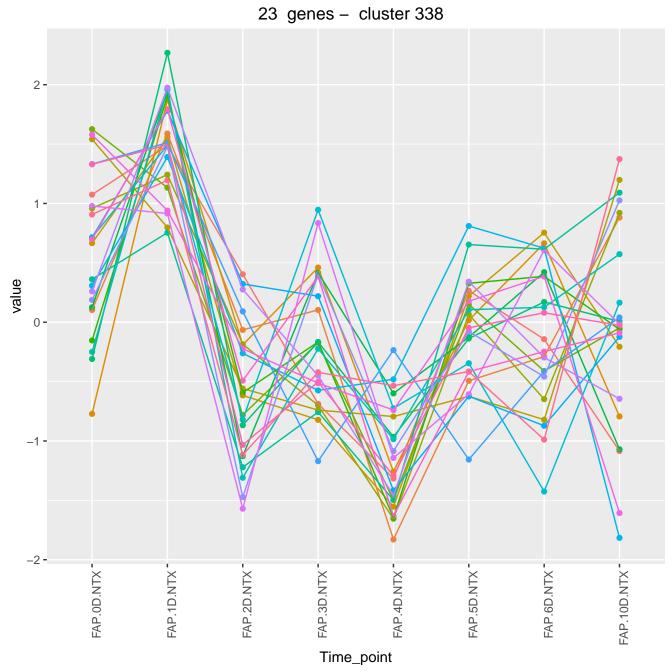




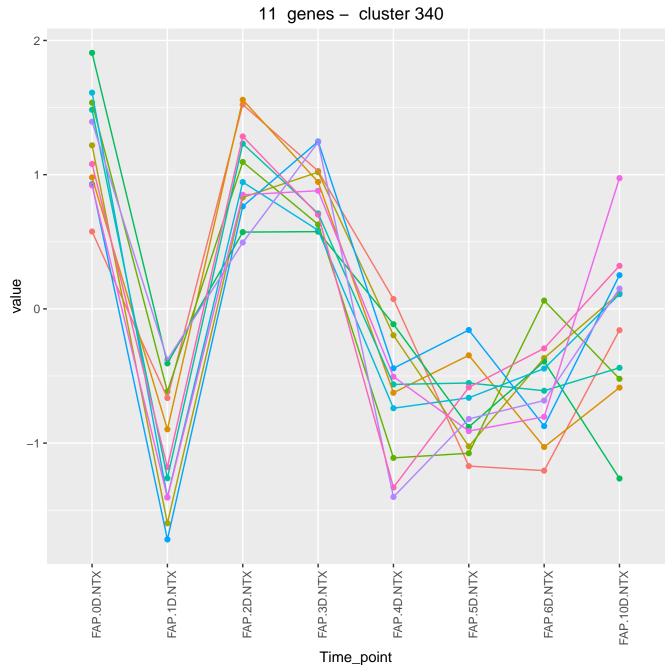


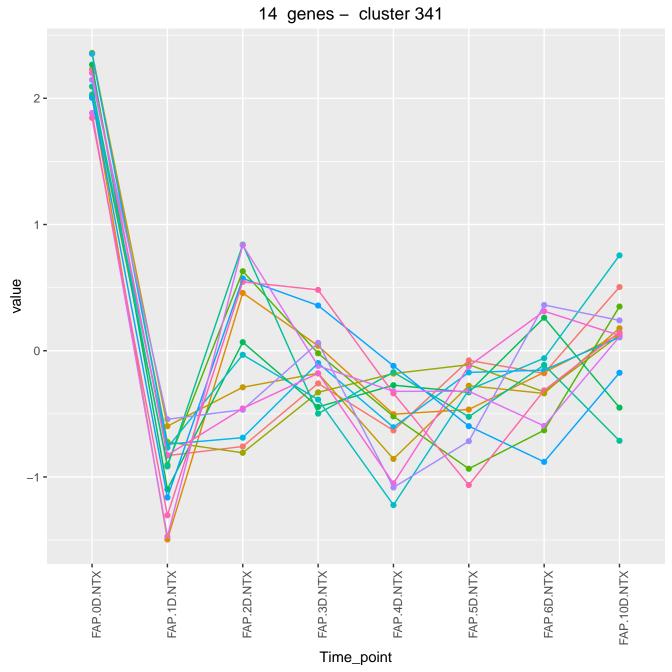


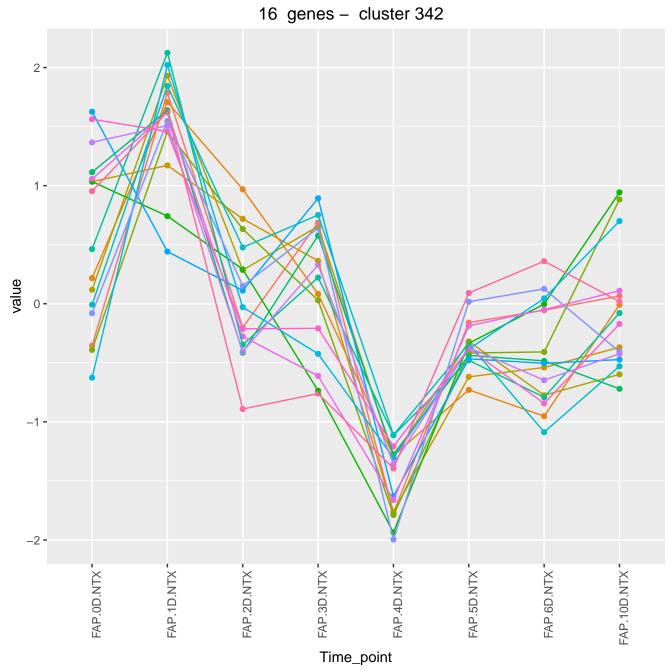


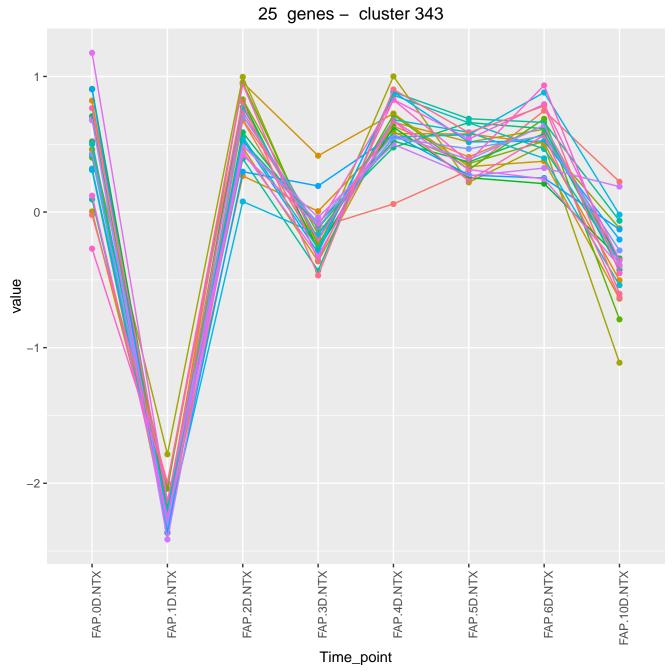


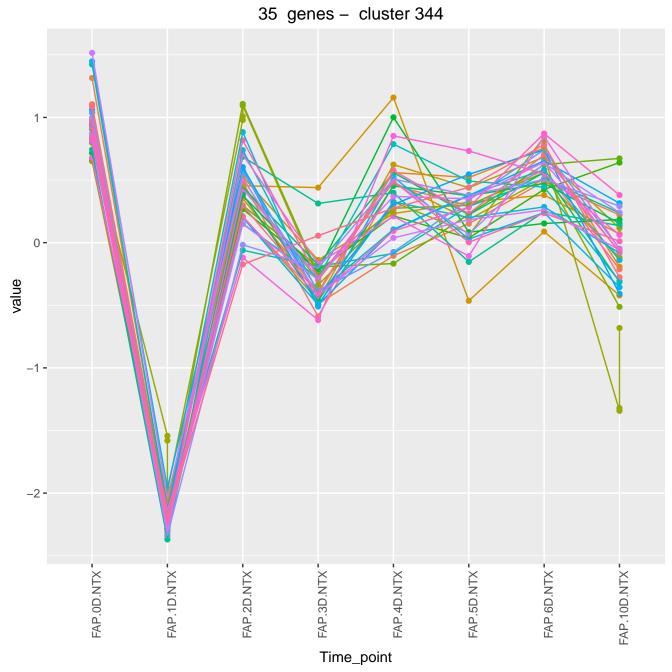
24 genes - cluster 339 2 -1 value 0 -\_1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX FAP.5D.NTX -FAP.10D.NTX FAP.2D.NTX FAP.3D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

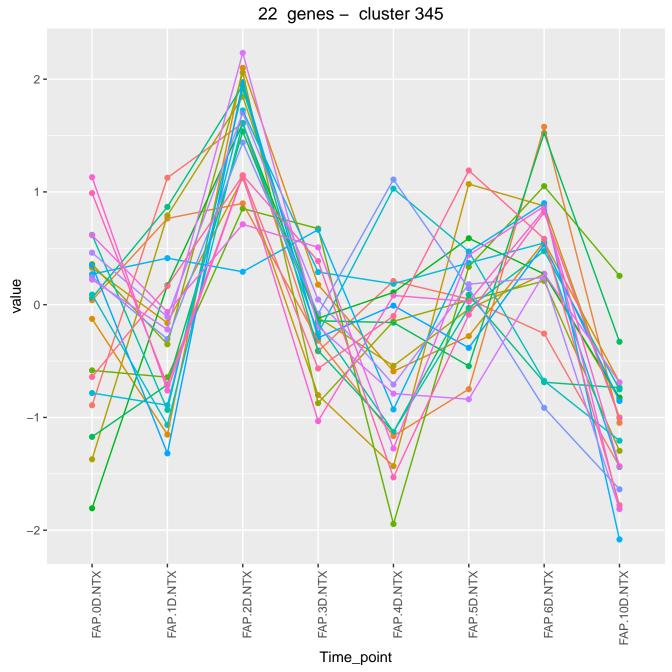


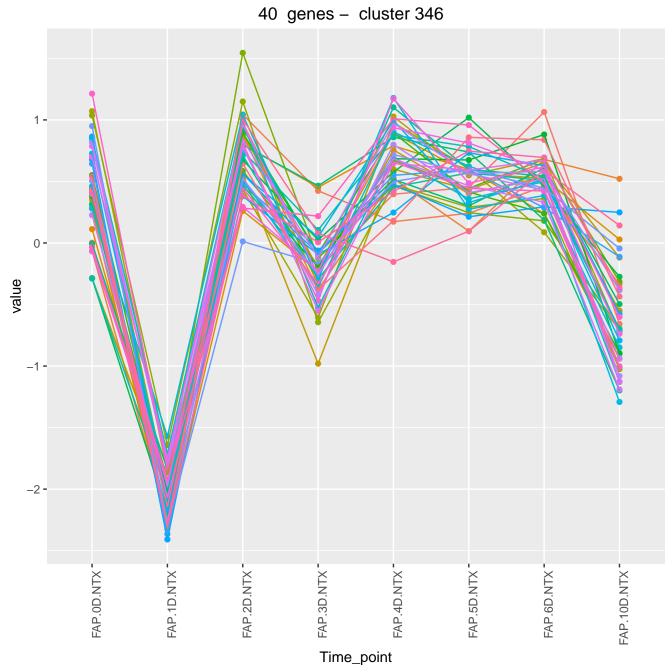


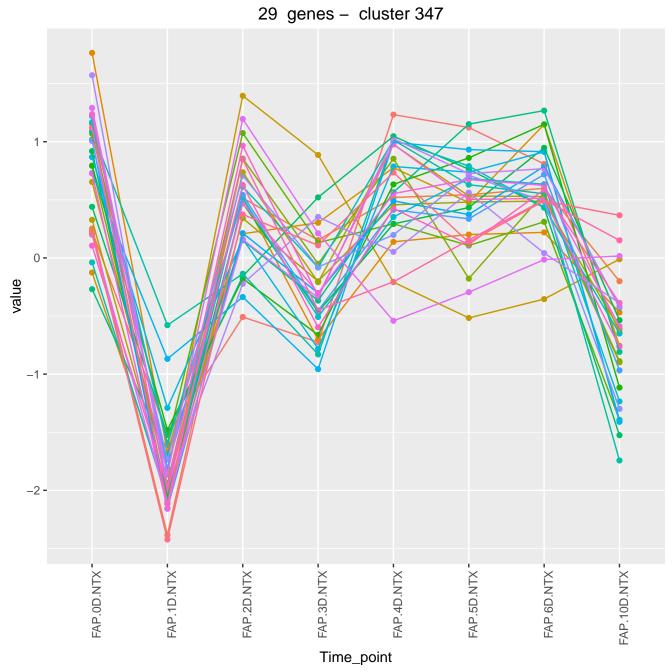


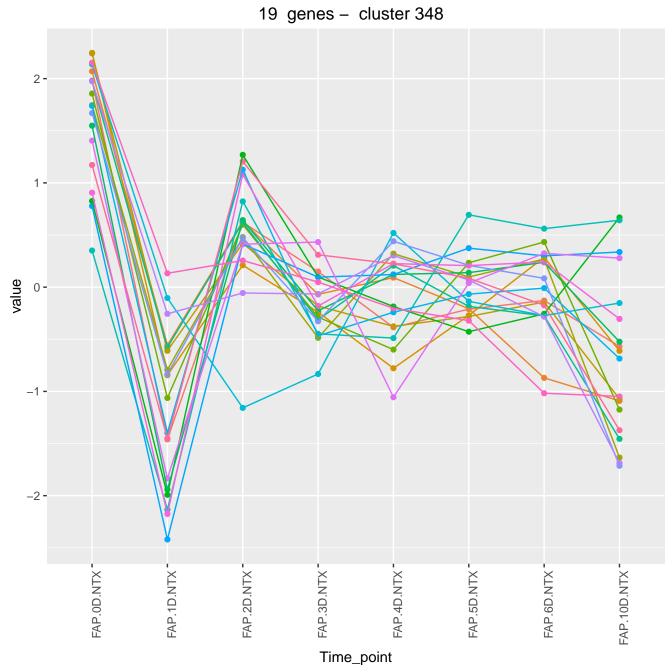


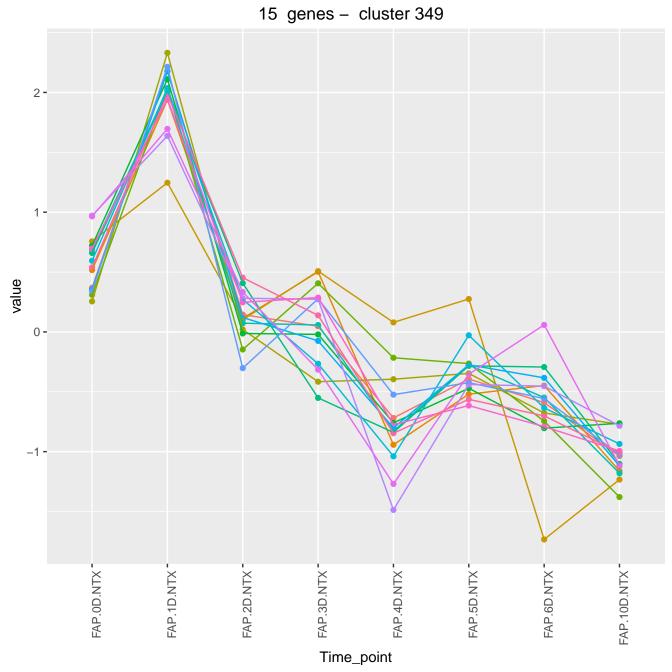


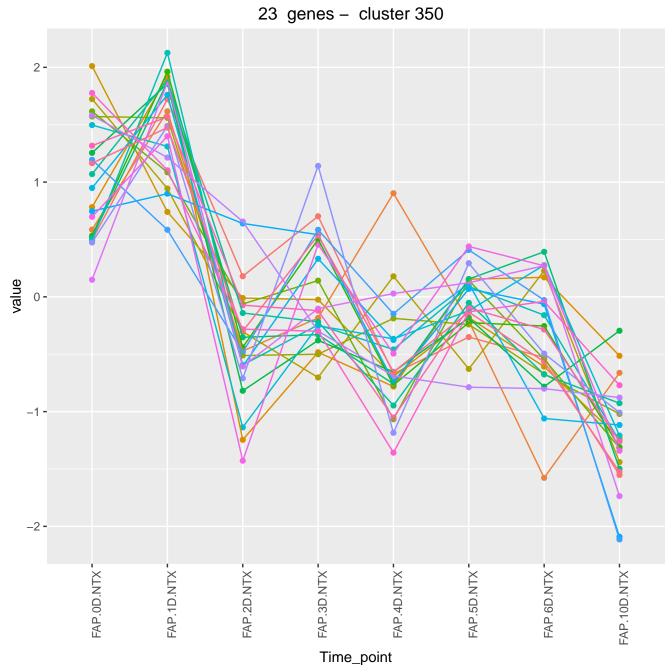


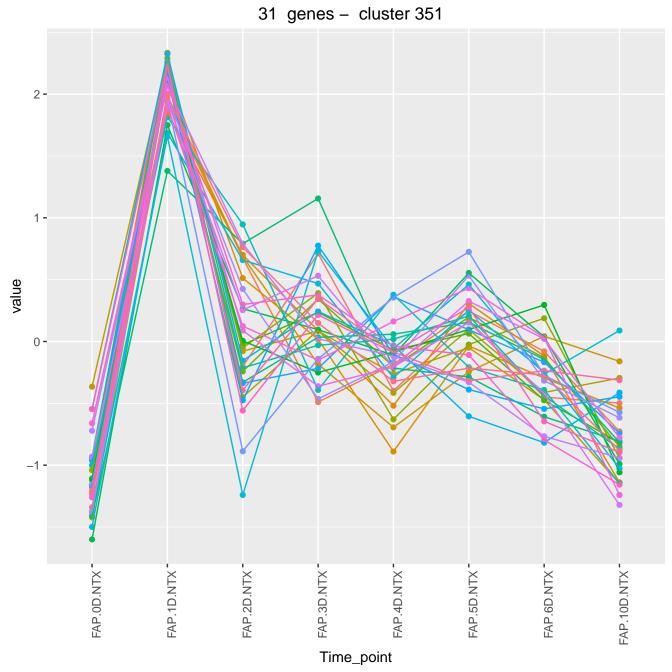


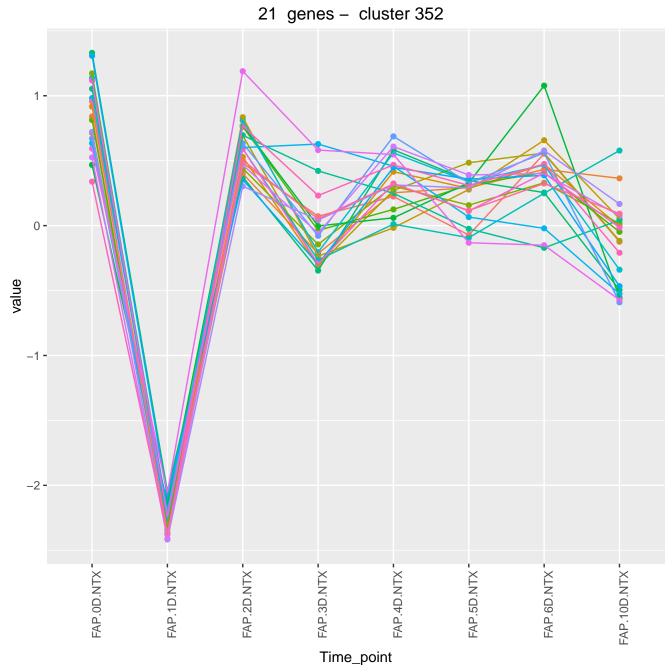


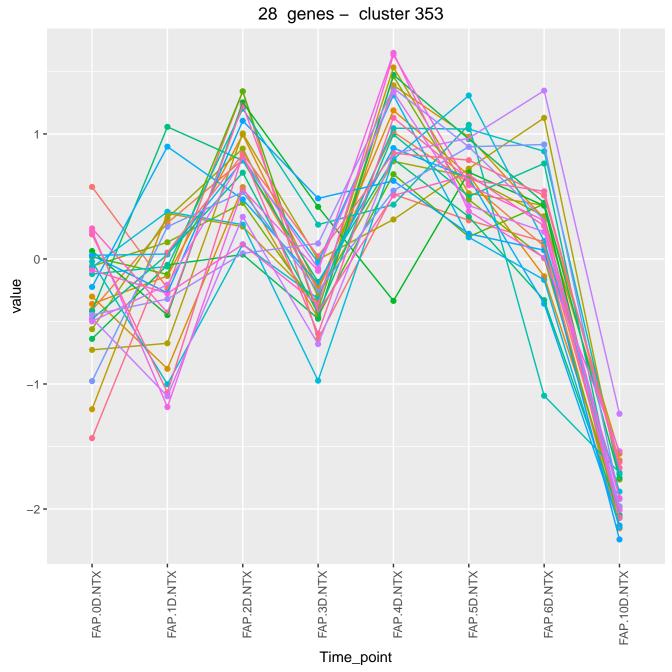


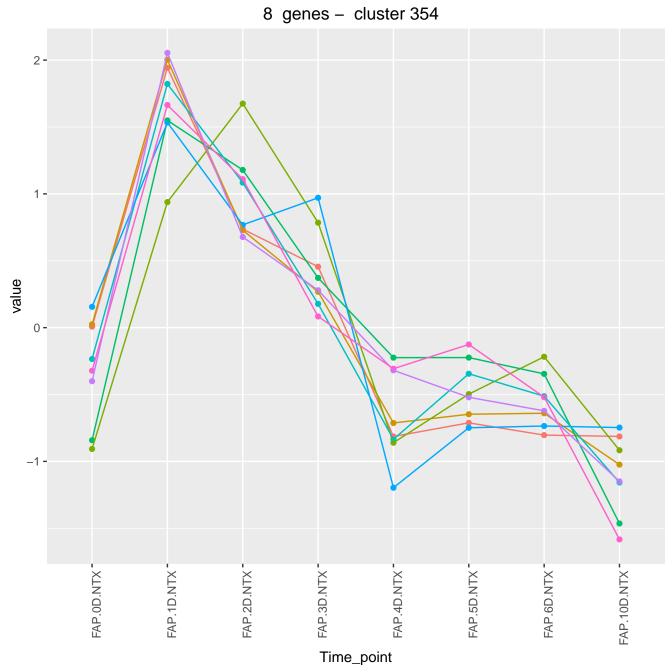


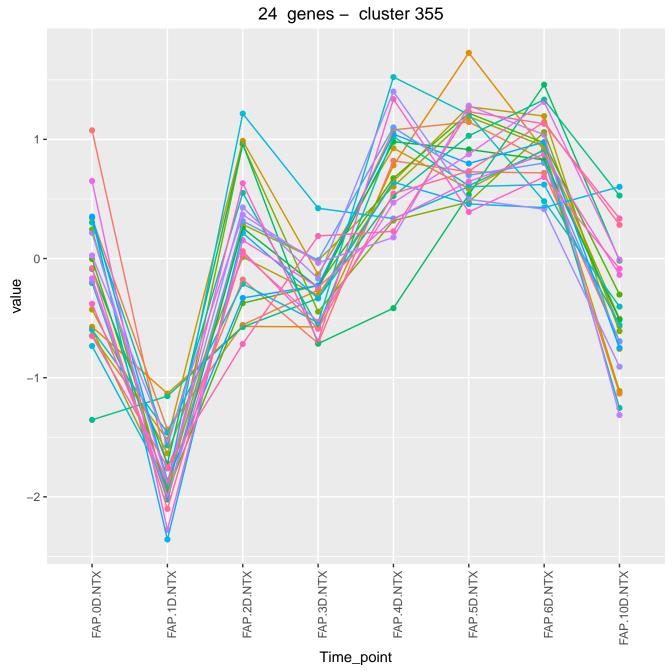


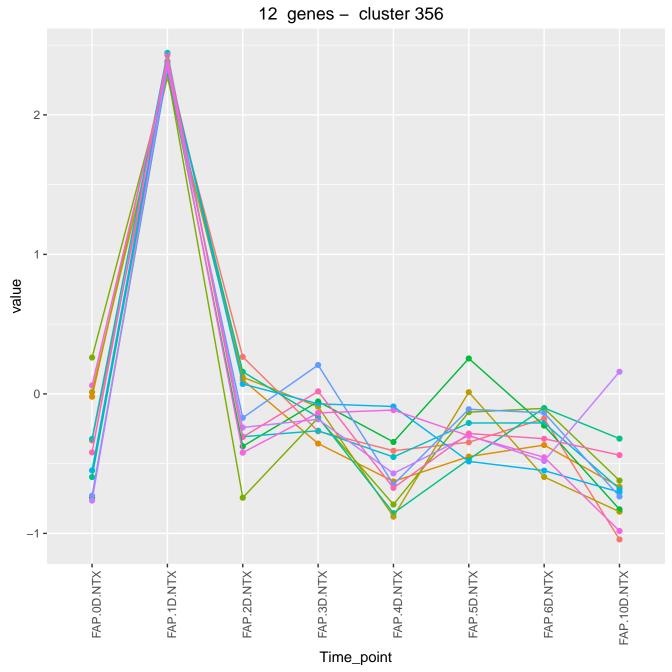


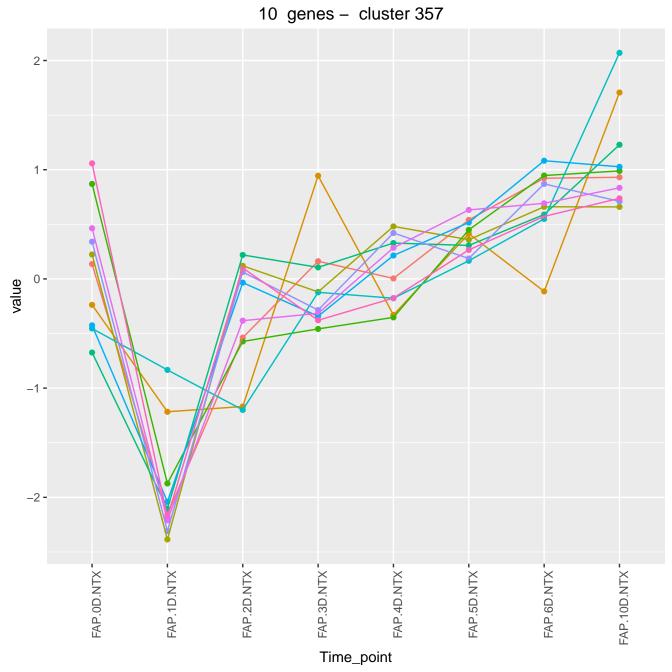


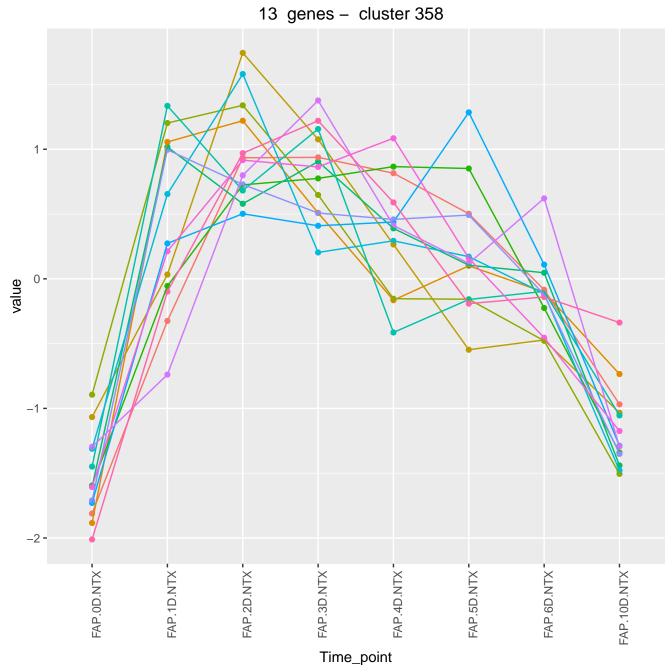


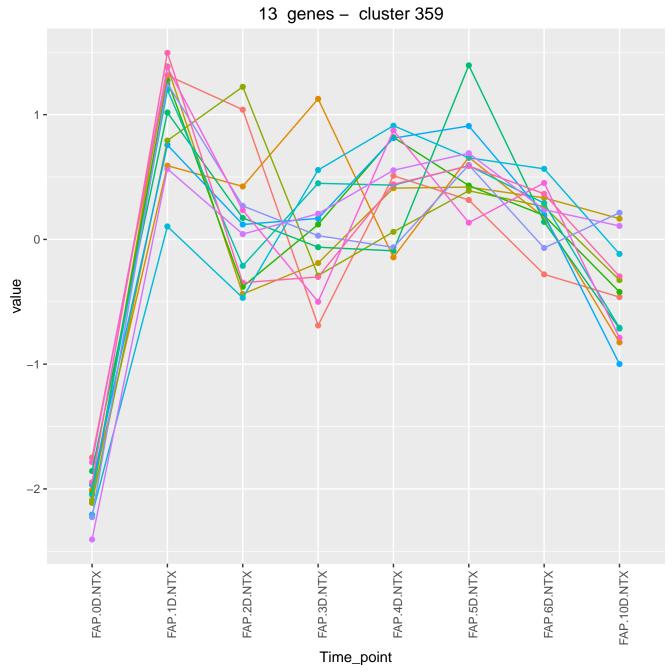




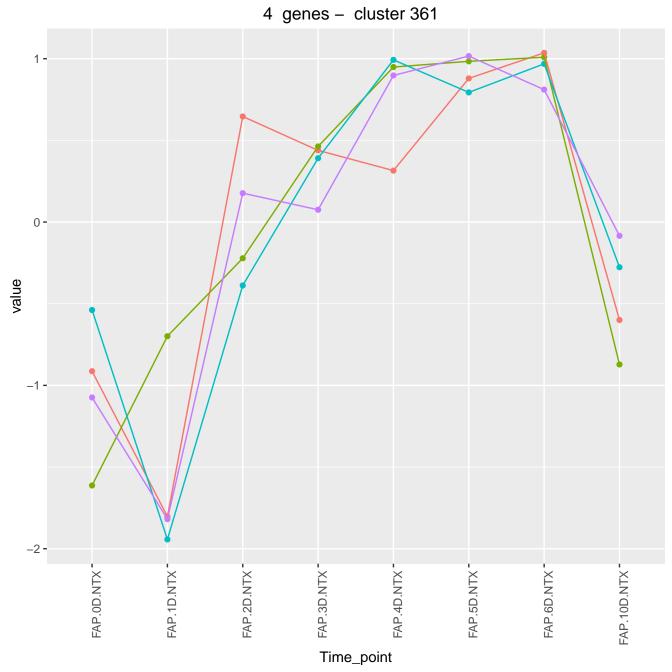


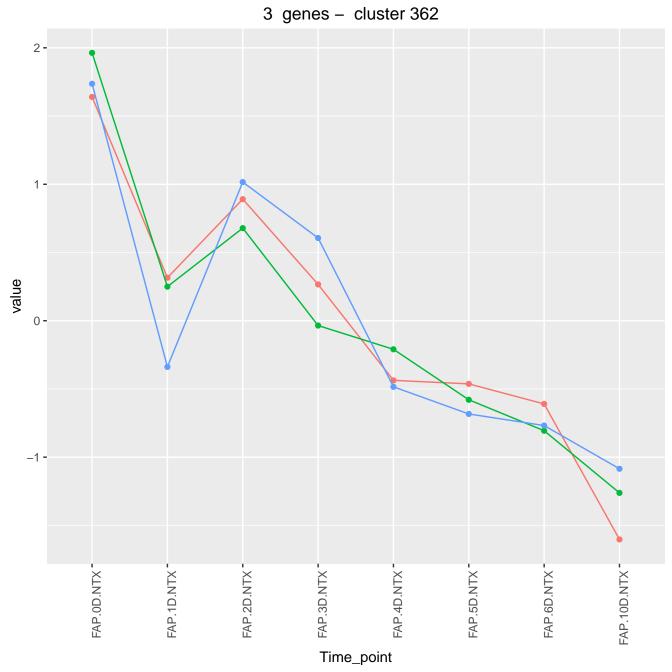




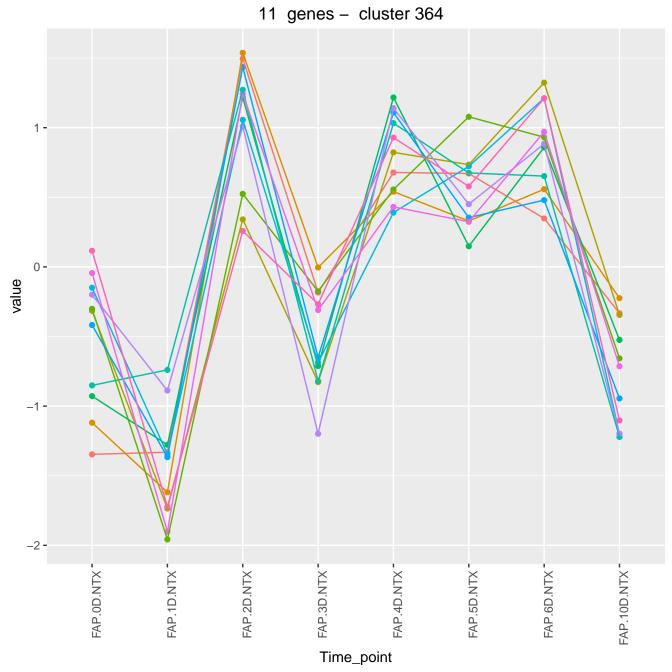


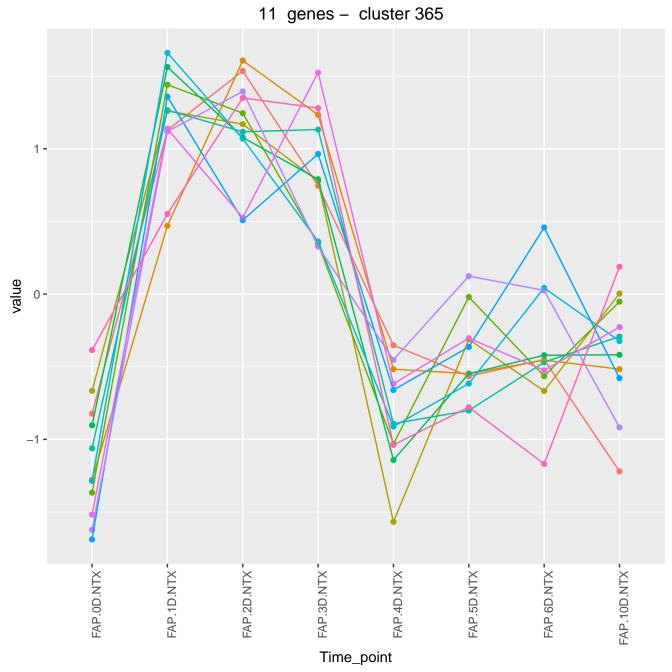
5 genes - cluster 360 2-1 value 0 --1 **-**FAP.0D.NTX 7 FAP.1D.NTX -FAP.2D.NTX <sup>7</sup> FAP.3D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.6D.NTX <sup>7</sup> FAP.4D.NTX Time\_point

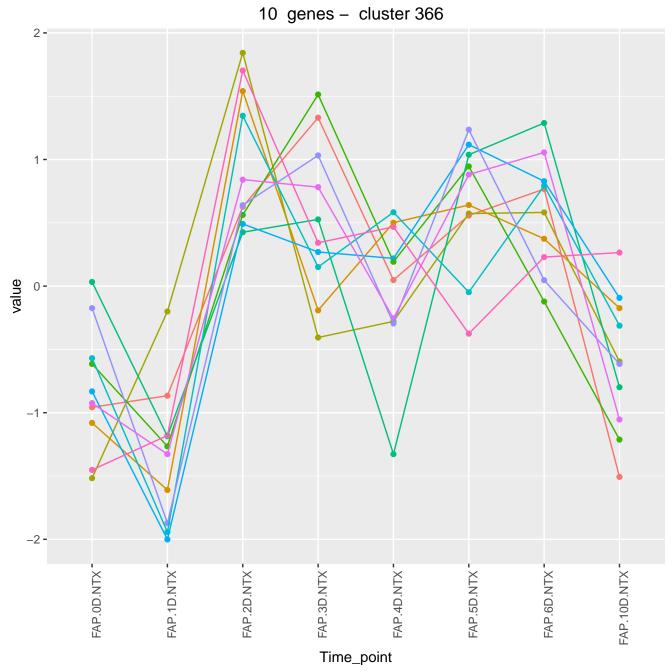


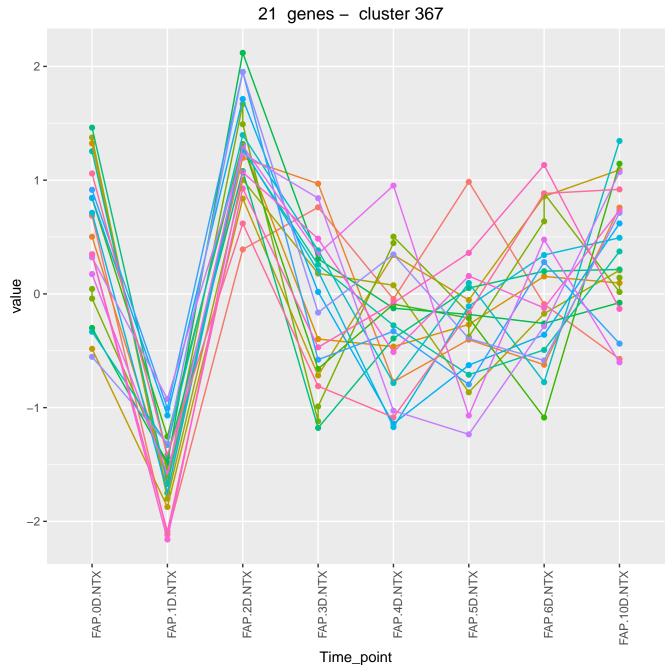


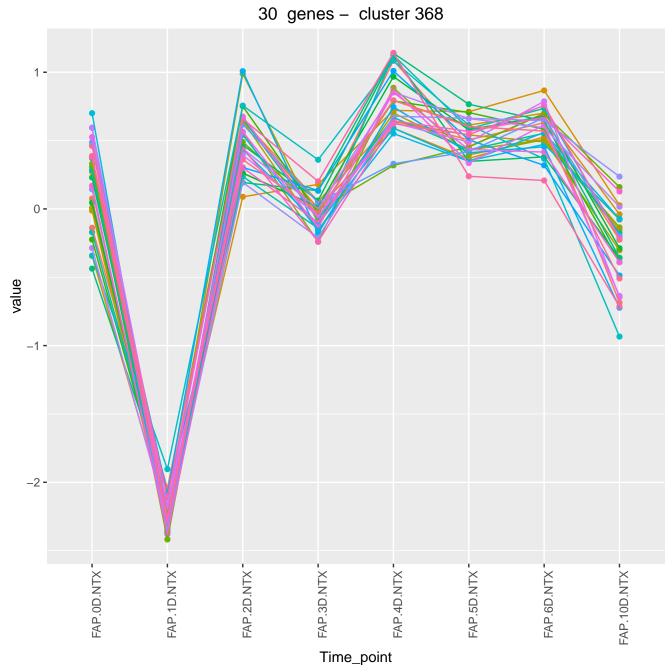
64 genes - cluster 363 2-1 value **-1** --2 **-**FAP.0D.NTX 7 FAP.10D.NTX -FAP.3D.NTX -FAP.5D.NTX -FAP.2D.NTX FAP.1D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

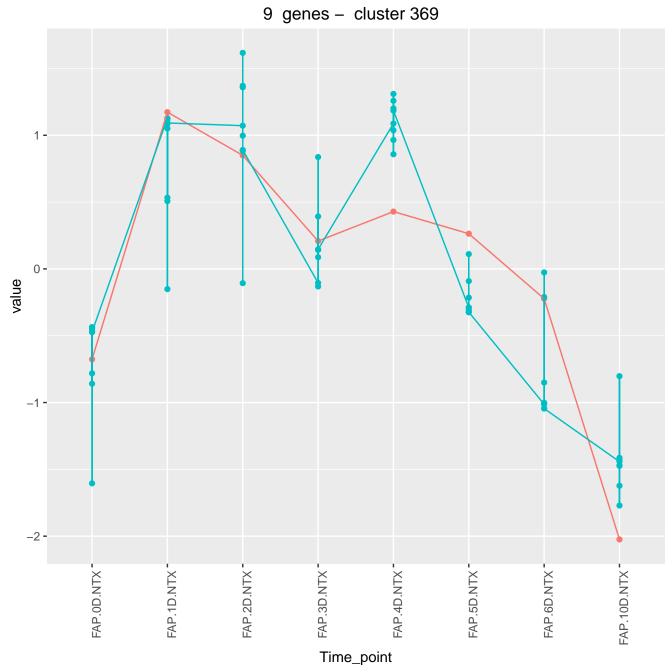


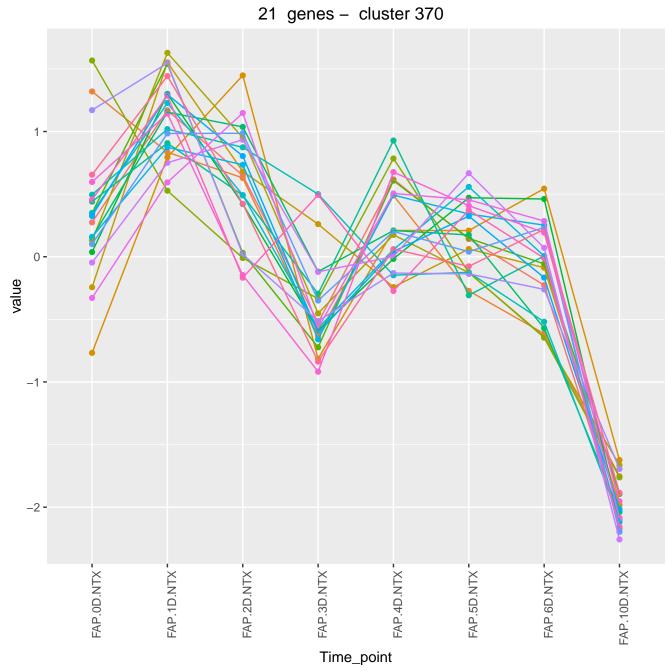


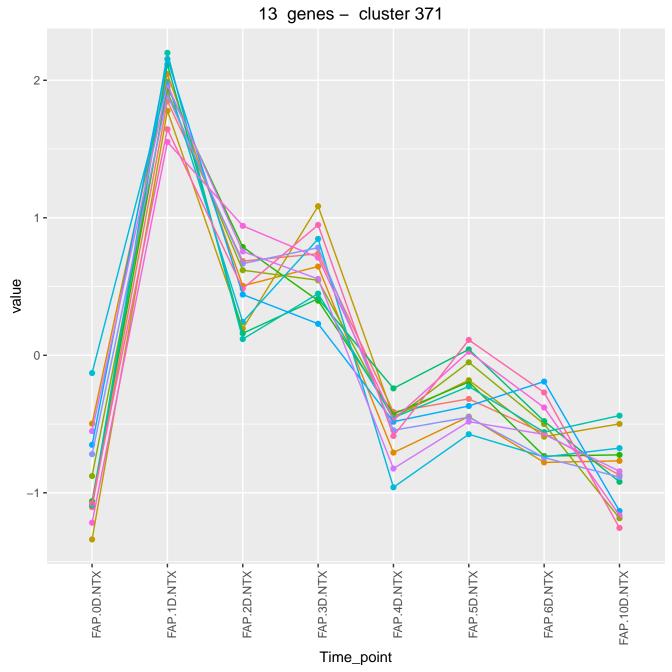


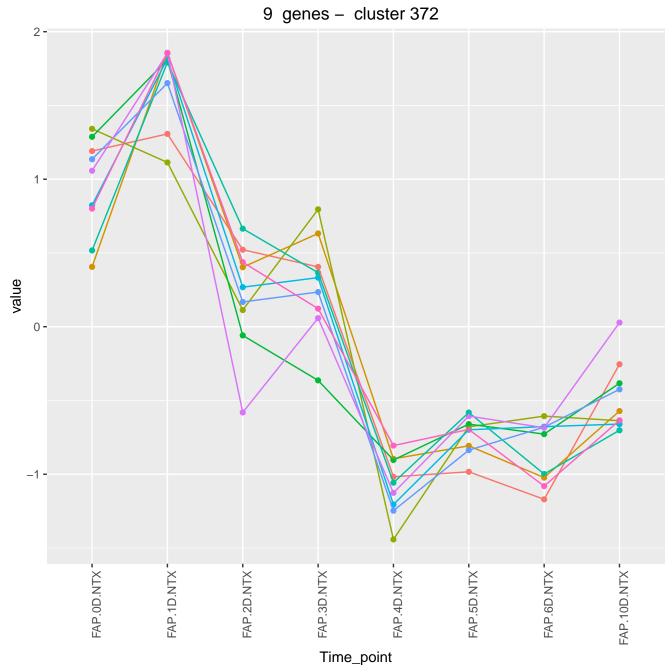


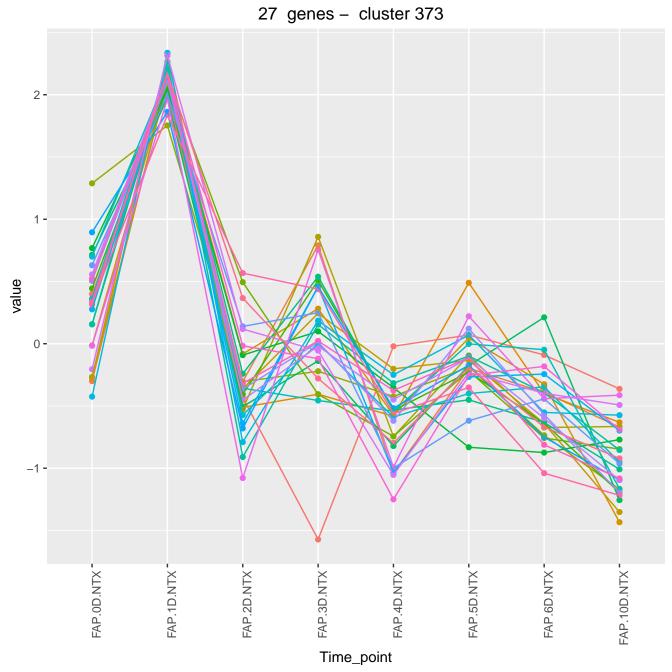


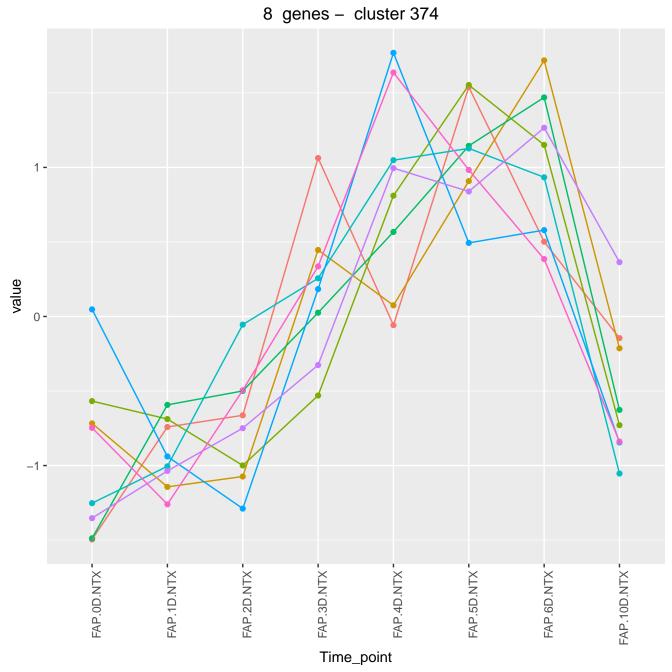


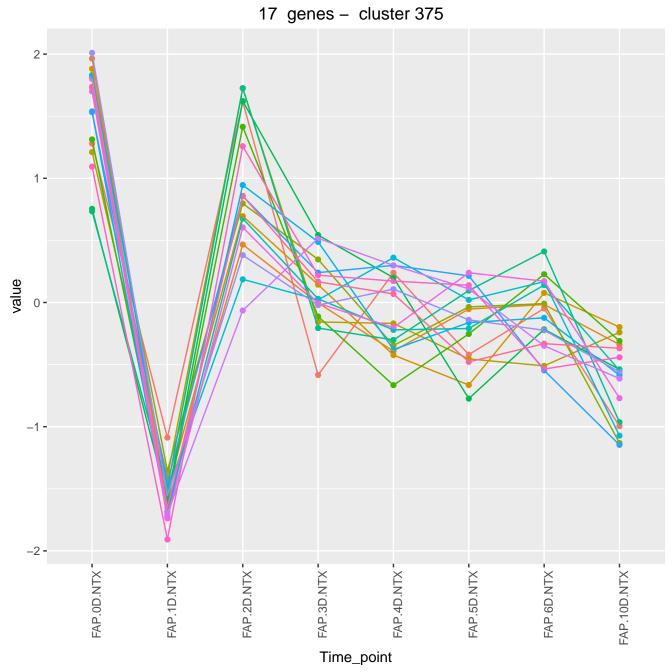


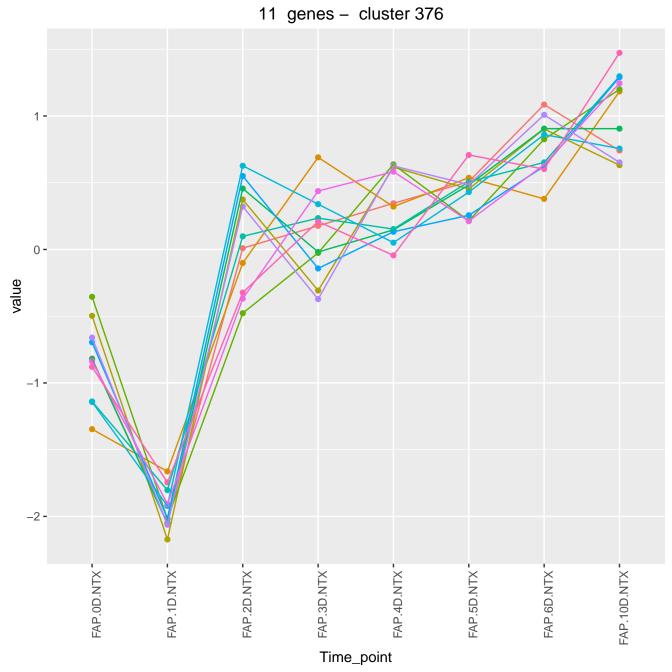


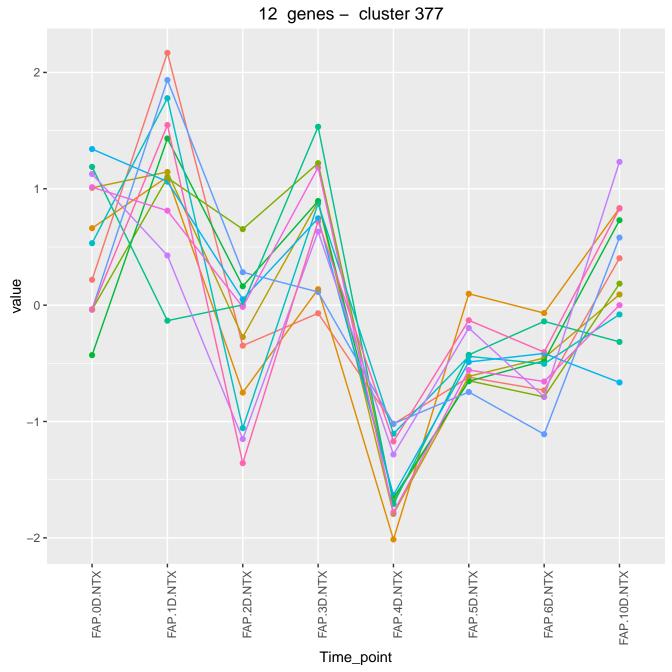


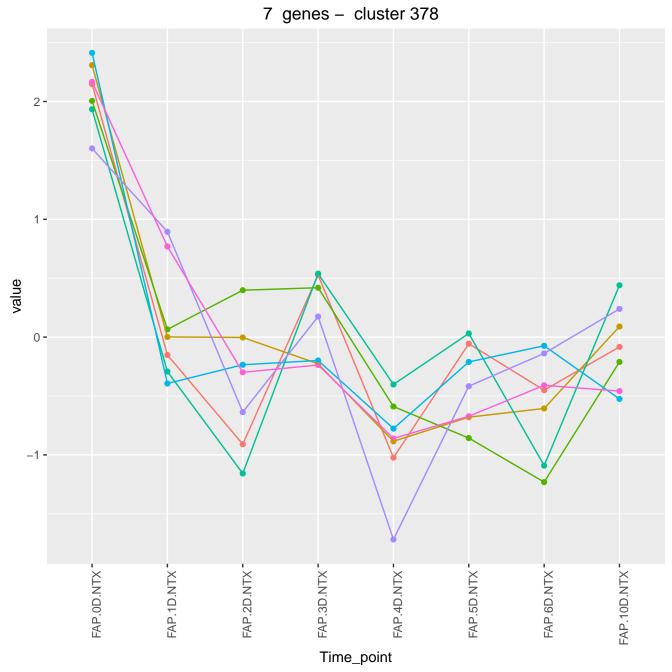


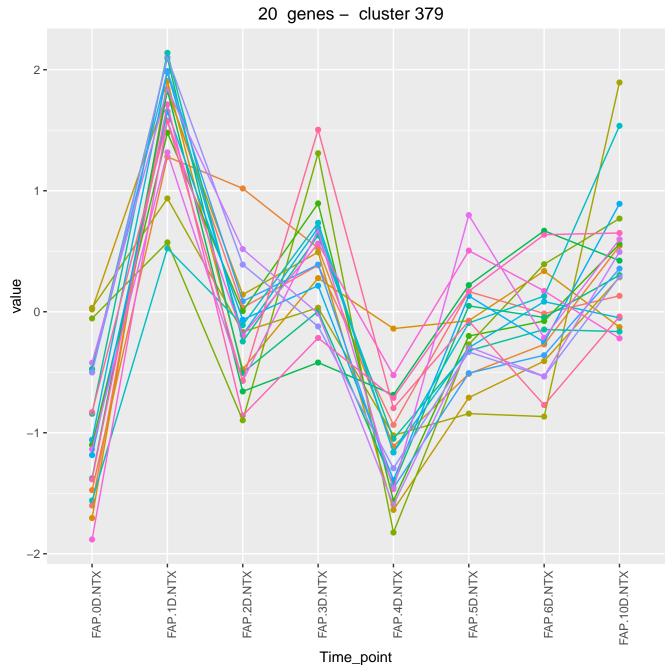


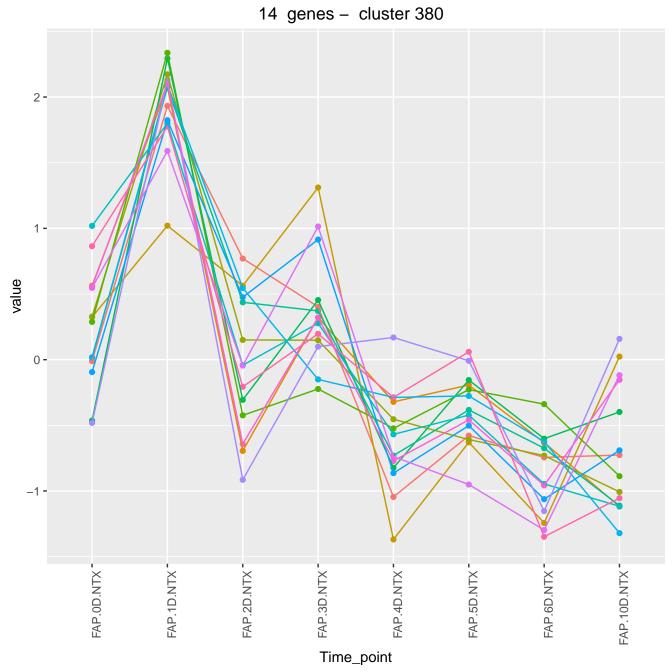


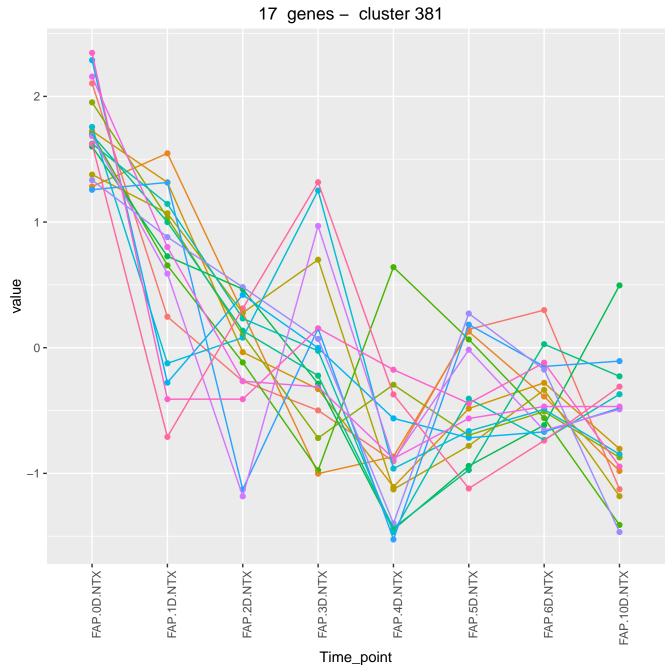




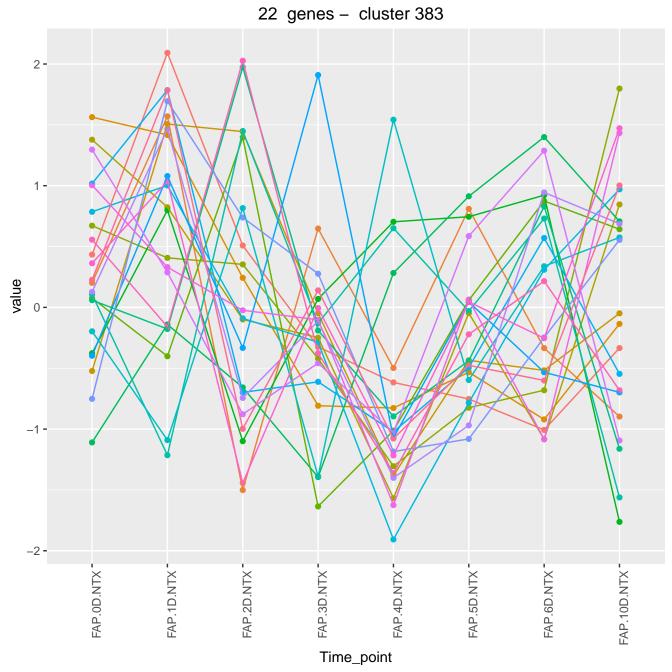




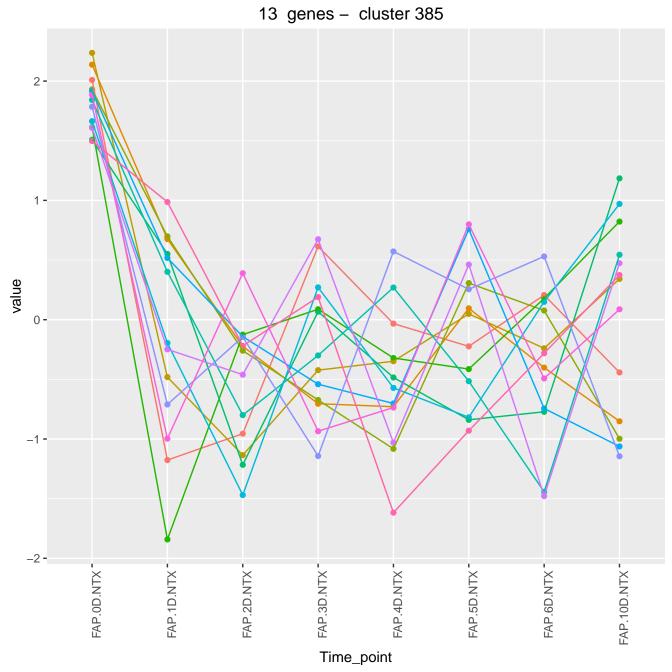


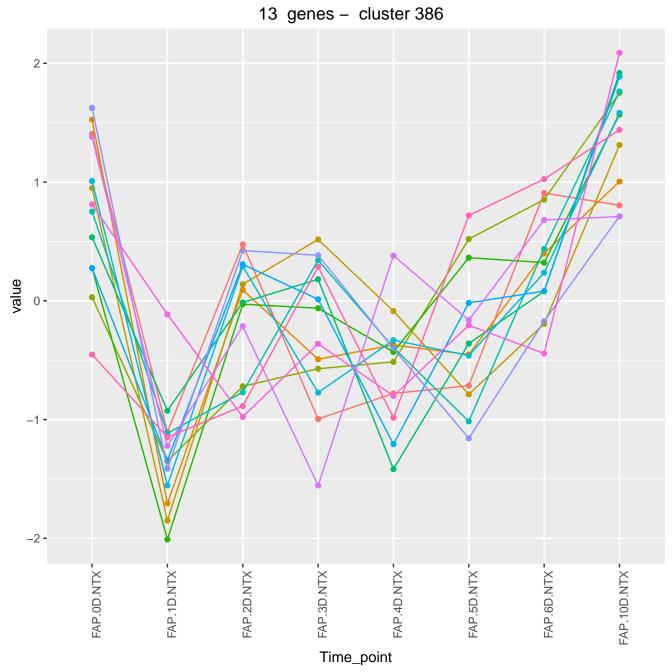


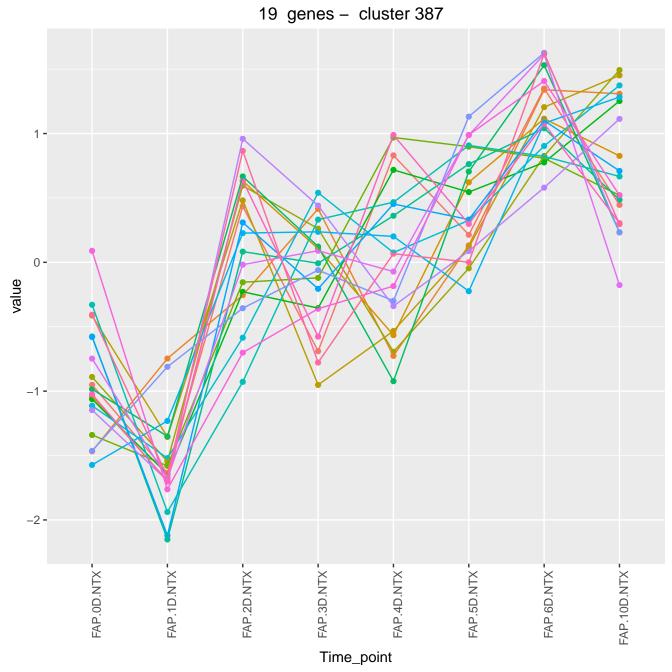
19 genes - cluster 382 2-1 value **-1** --2 **-**FAP.0D.NTX 7 FAP.2D.NTX -FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.3D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

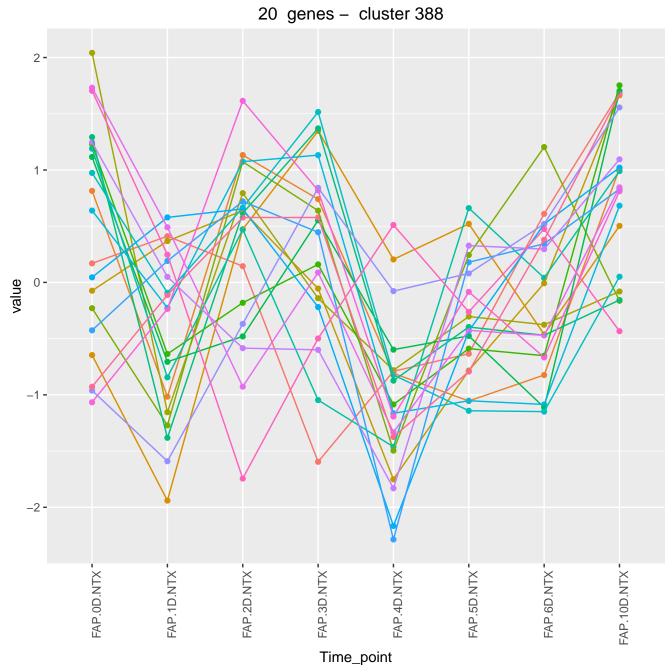


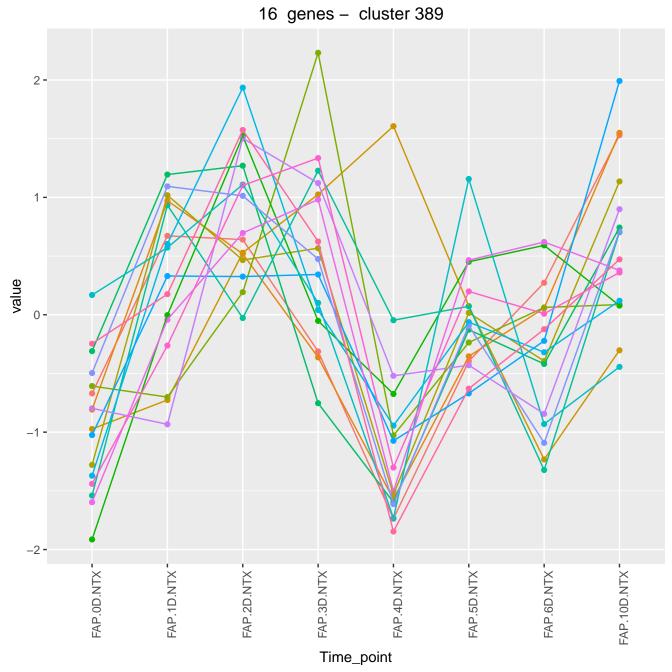
17 genes - cluster 384 2 -1 value 0 -\_1 **-**-2 **-**FAP.0D.NTX 7 FAP.2D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.3D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

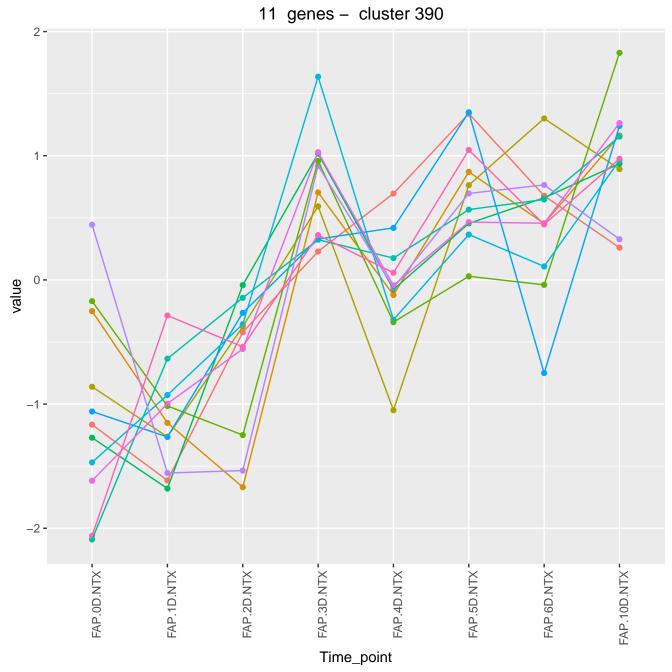


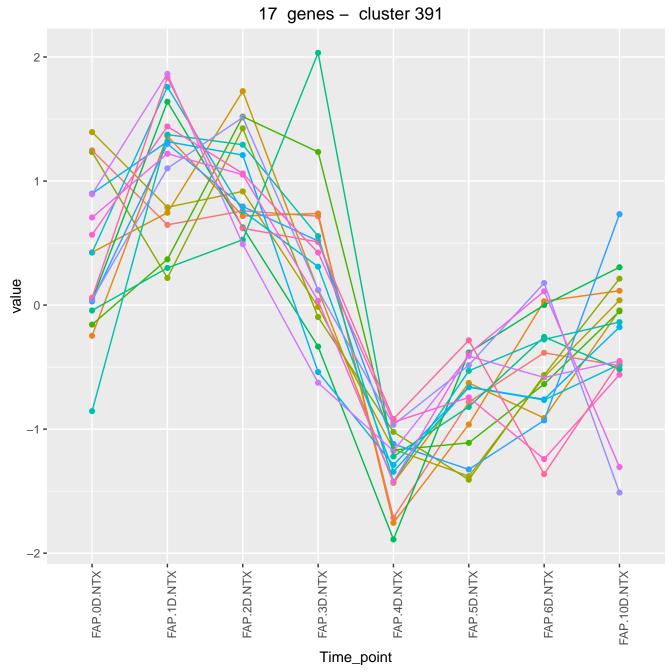




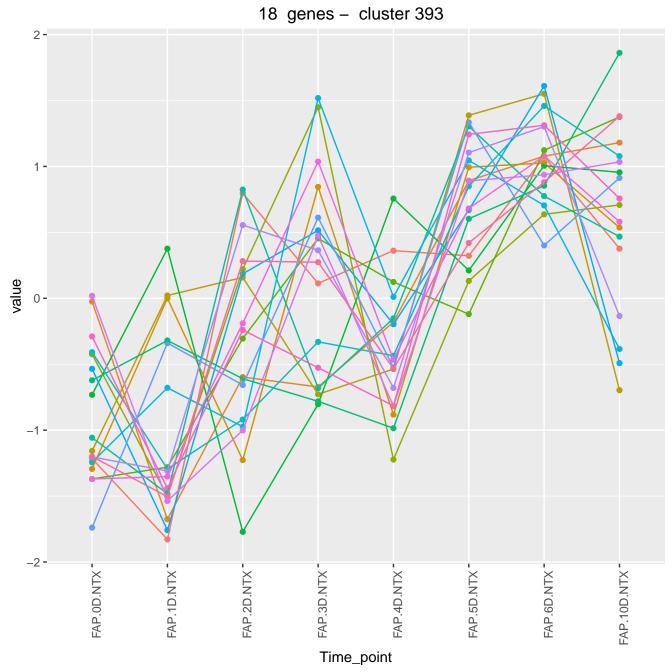


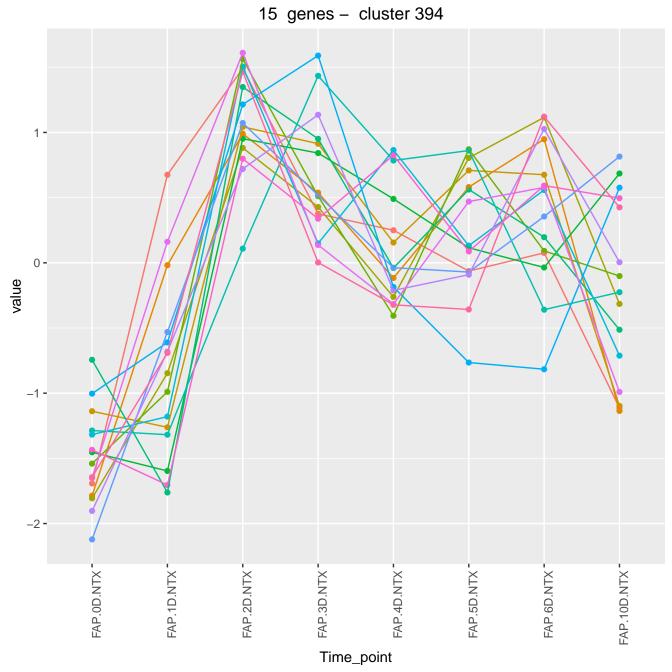


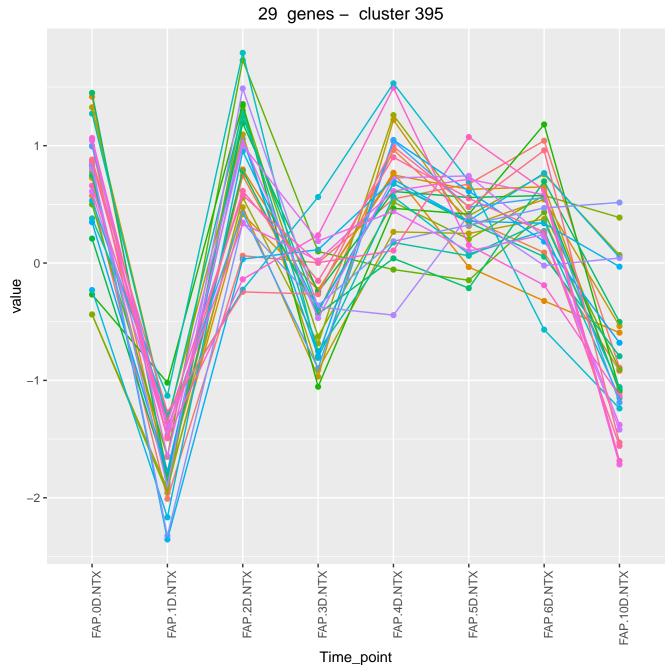


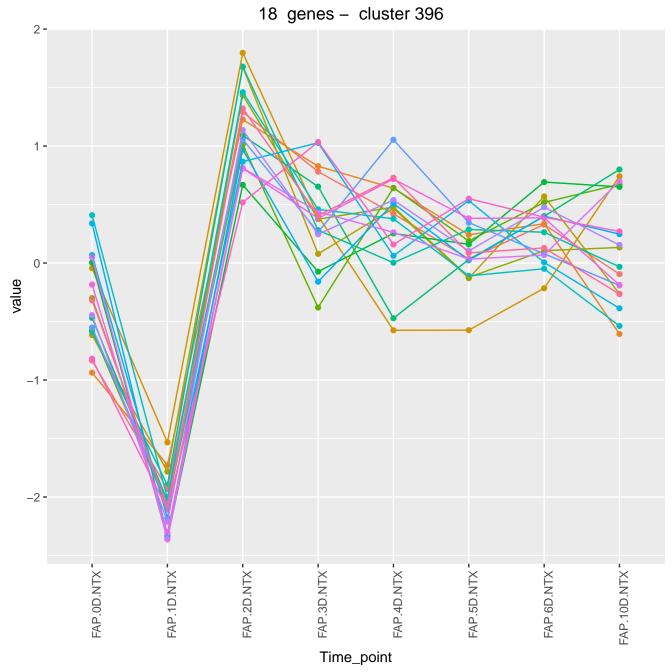


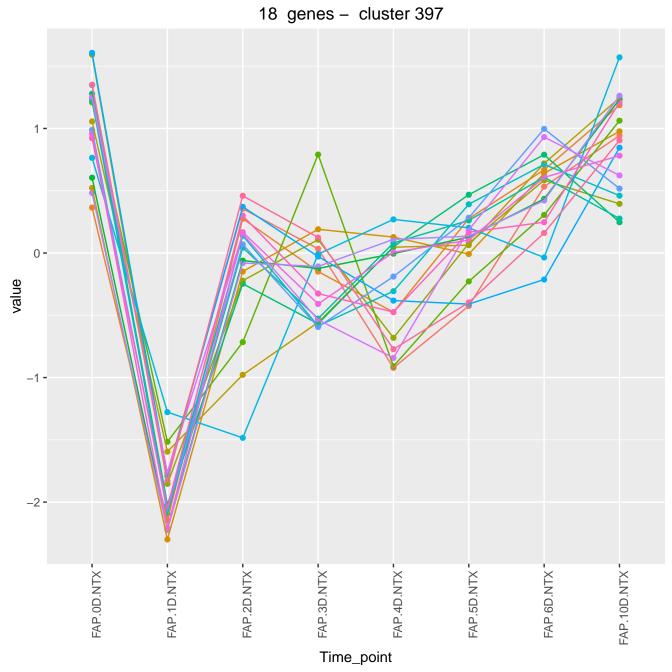
12 genes - cluster 392 2 -1 value 0 -\_1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.4D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.3D.NTX FAP.6D.NTX Time\_point

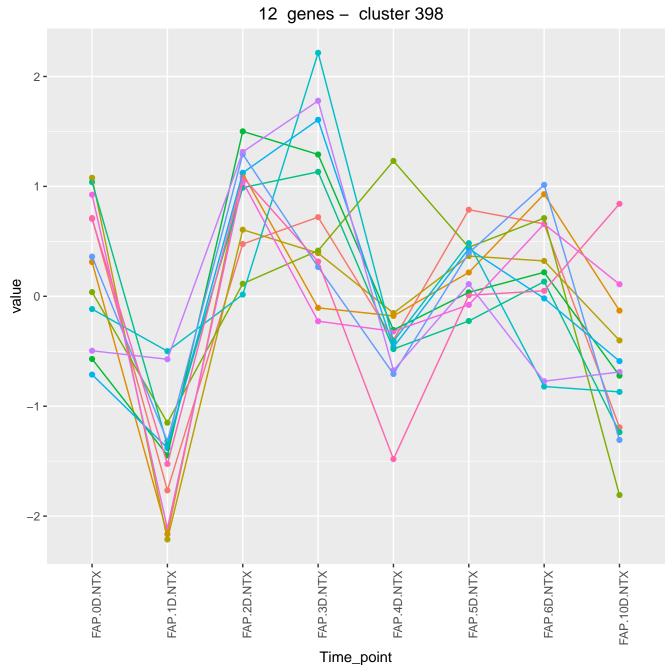


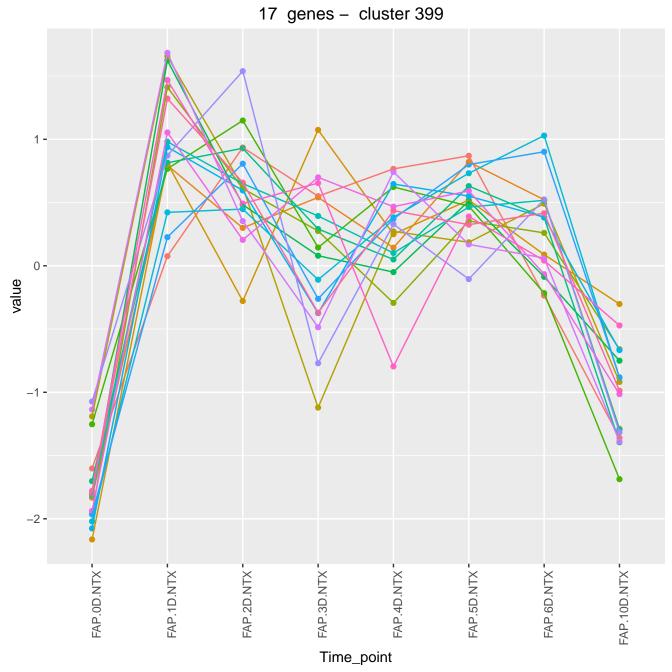


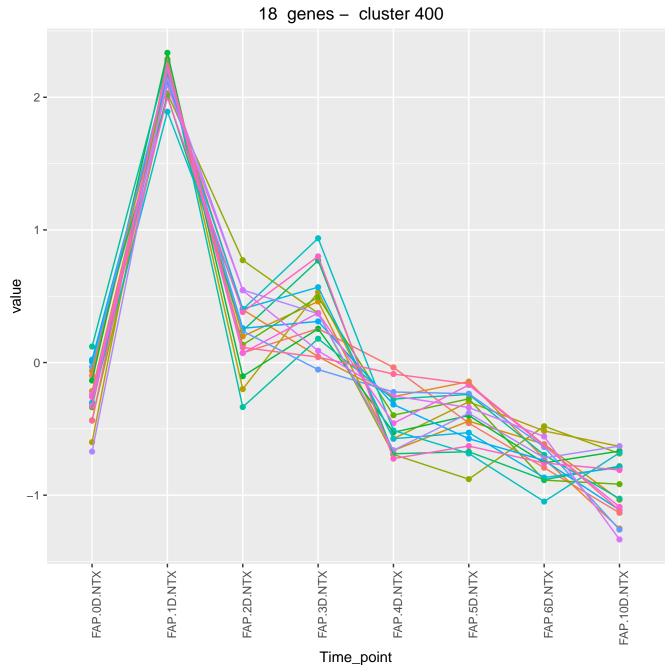


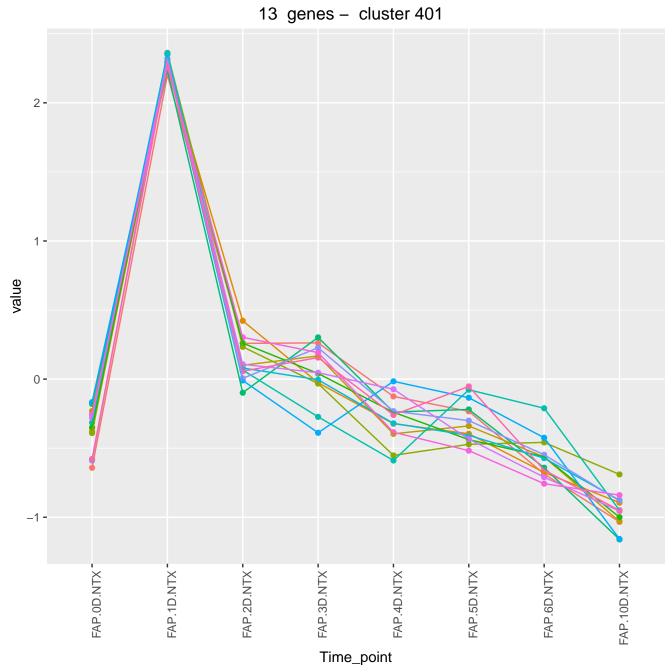


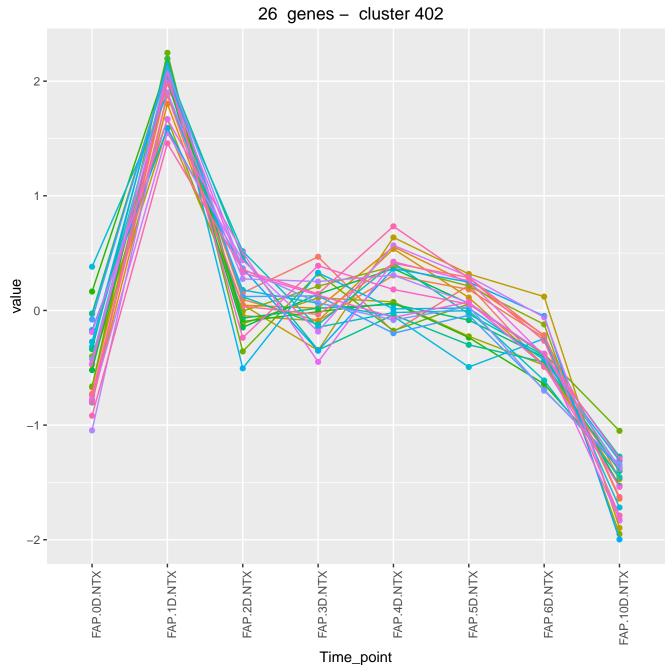


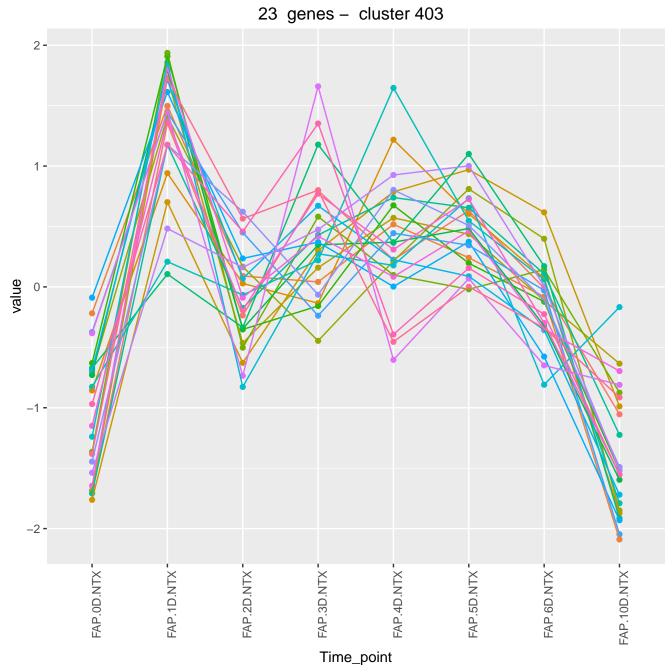


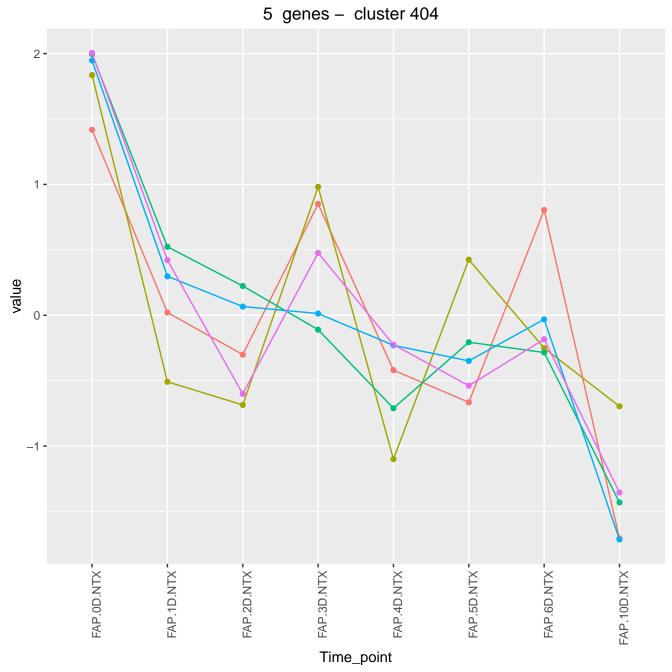


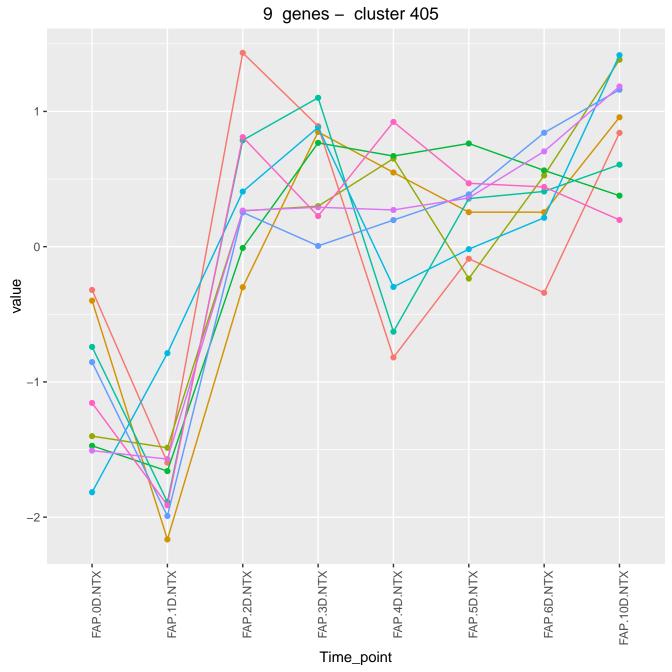


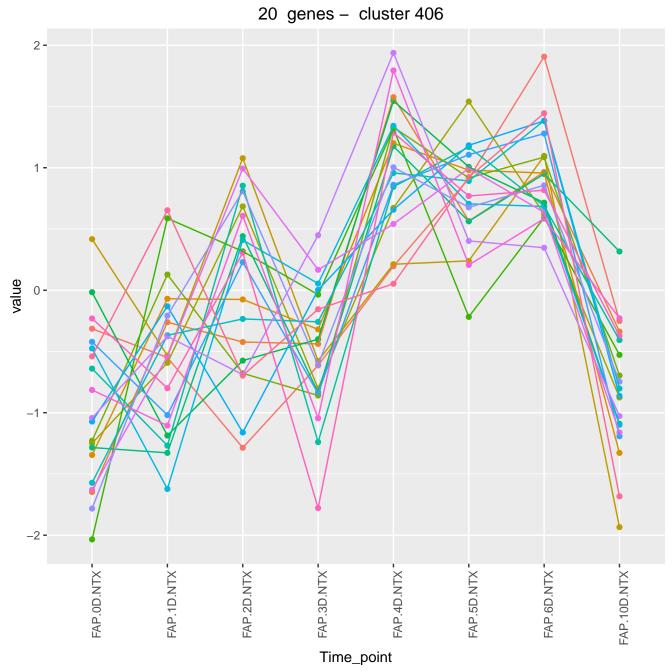


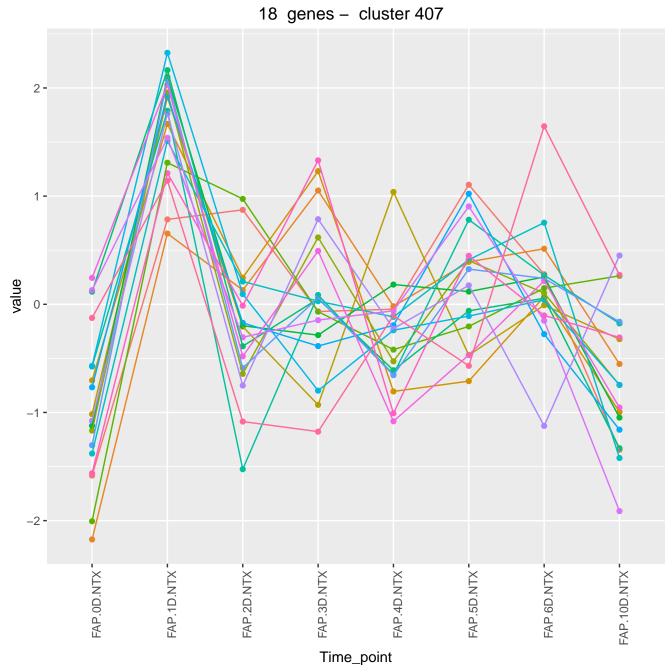


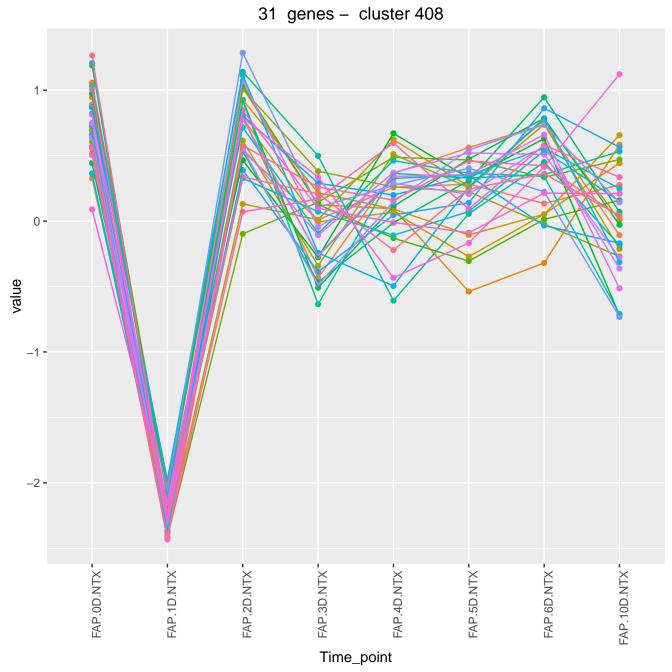


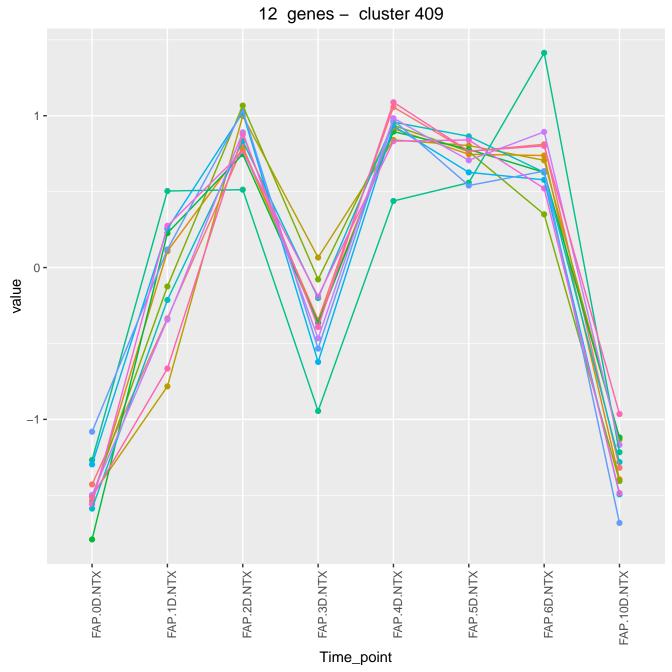


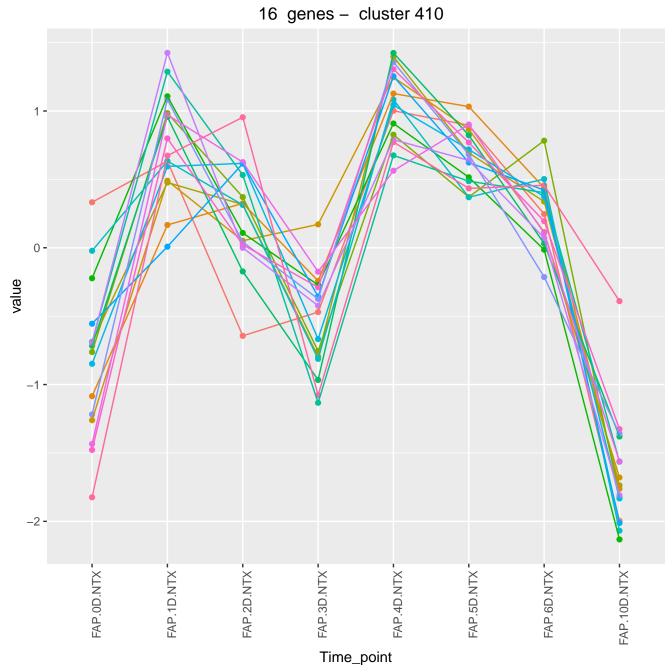


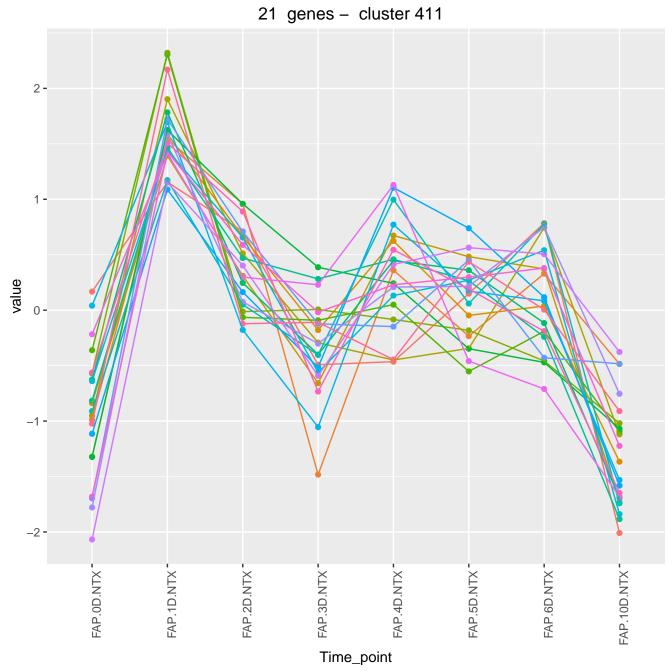


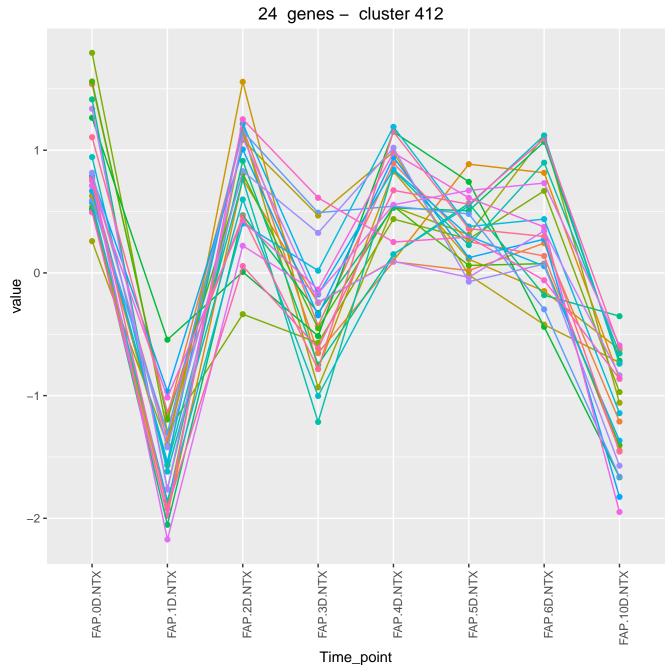


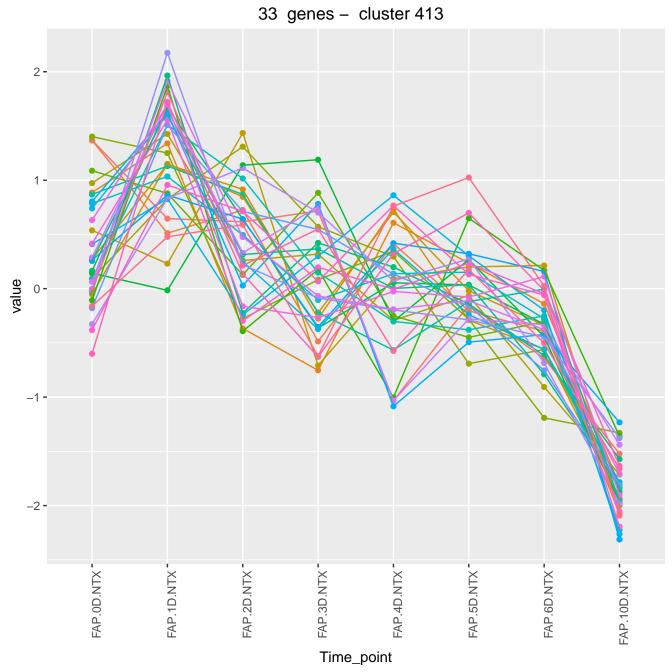


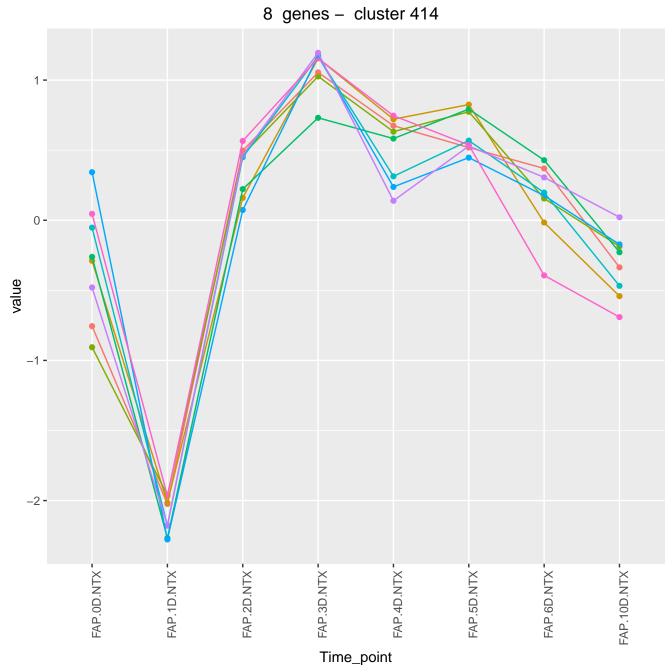


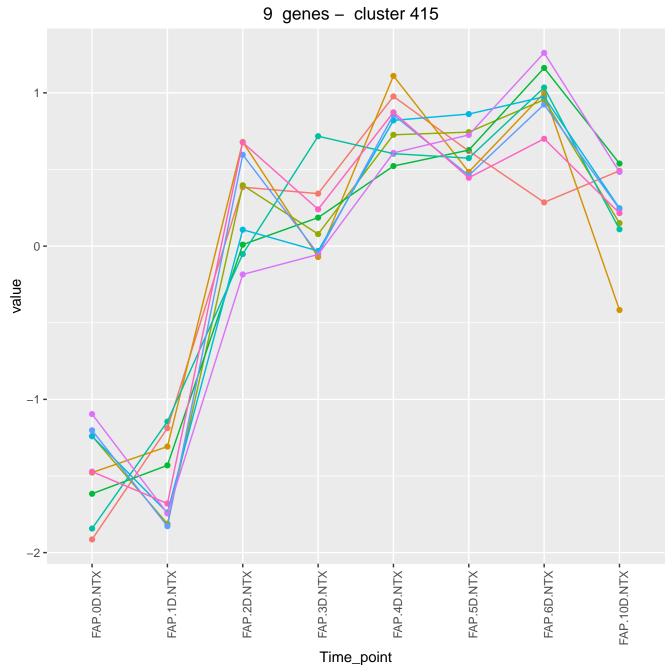


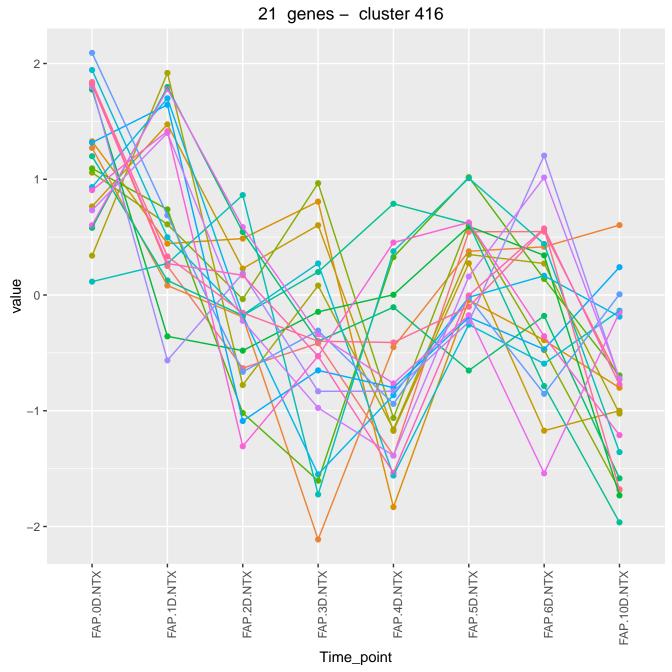


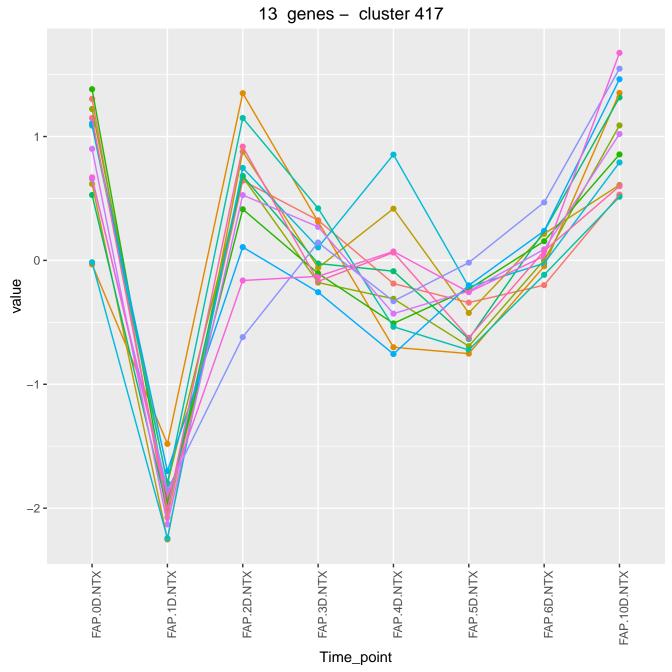




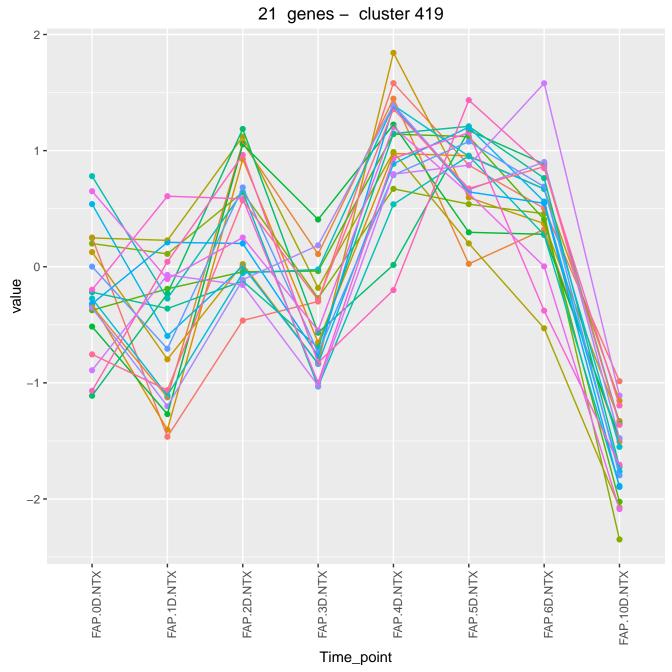


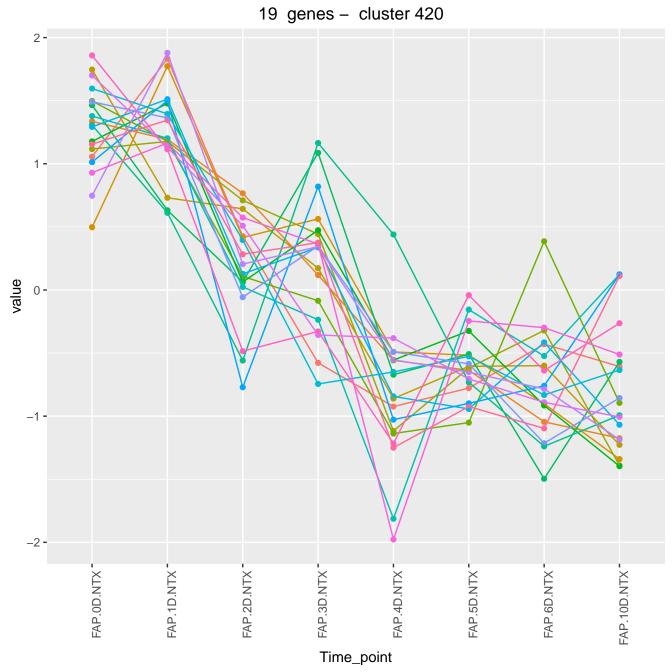


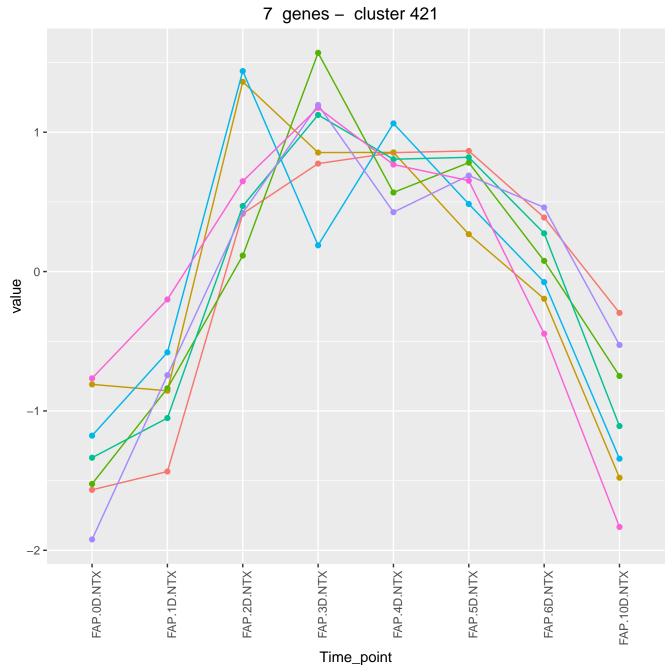


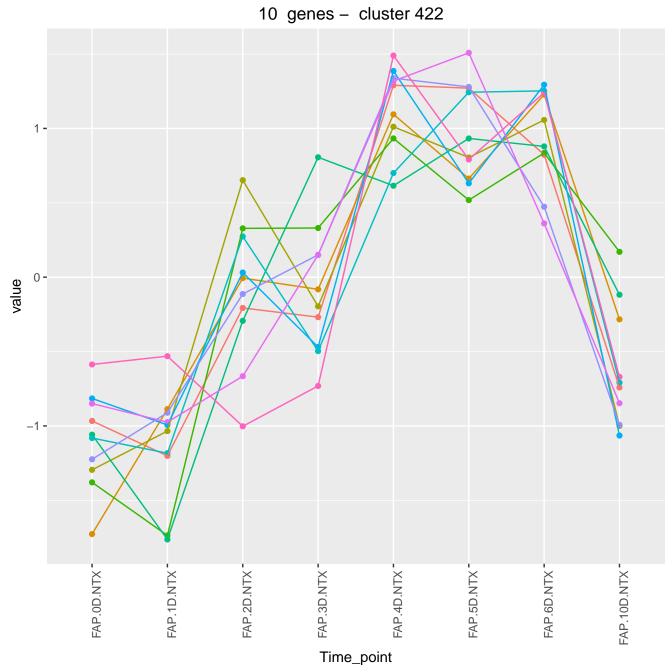


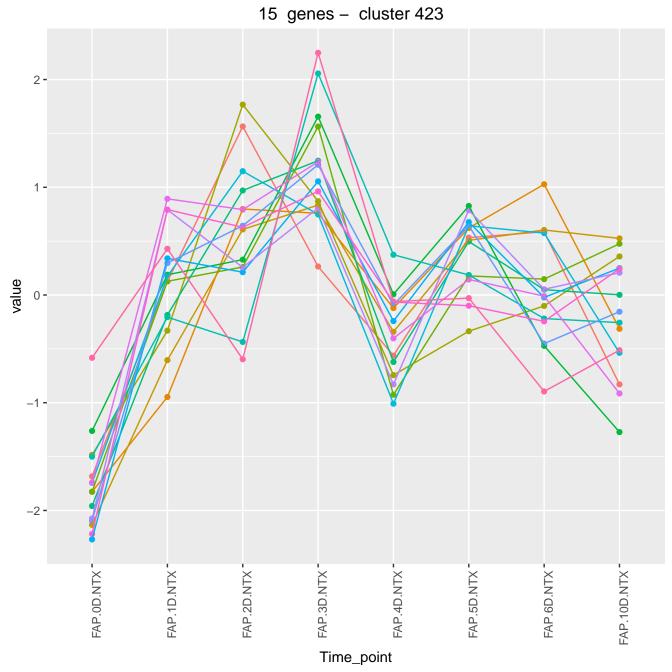
15 genes - cluster 418 2 -1 value -1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.3D.NTX <sup>7</sup> FAP.5D.NTX -FAP.6D.NTX <sup>7</sup> FAP.10D.NTX T FAP.4D.NTX Time\_point

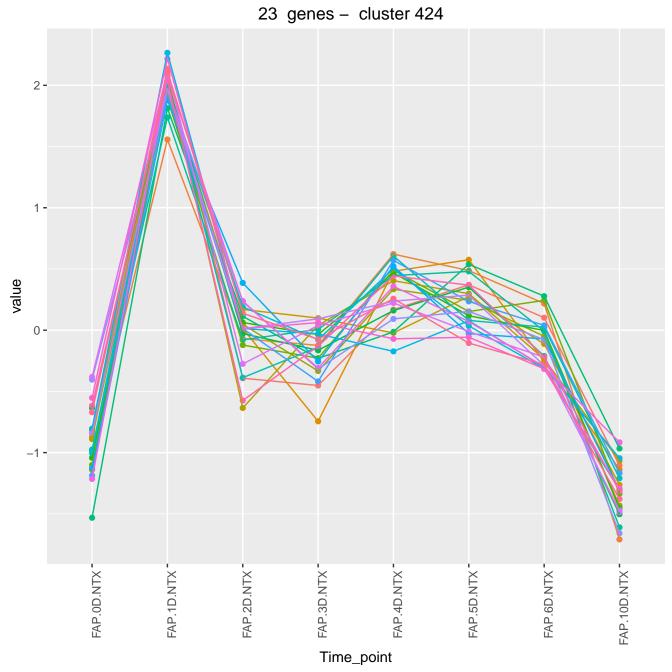


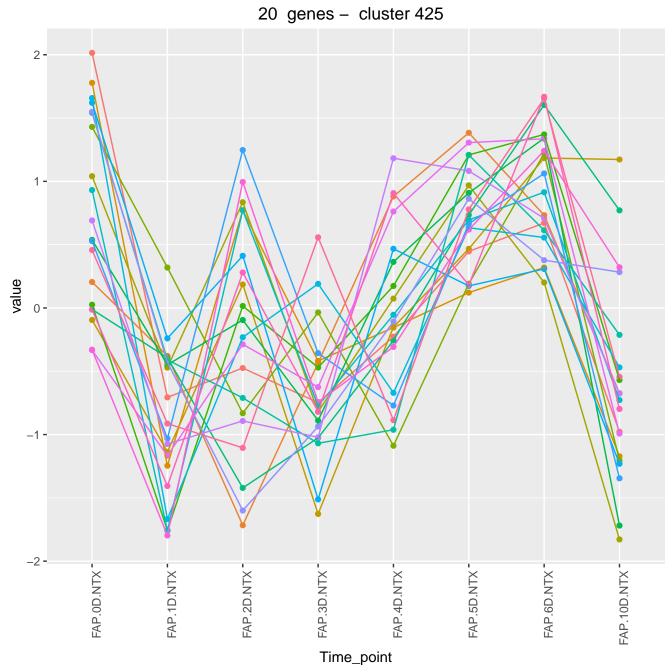


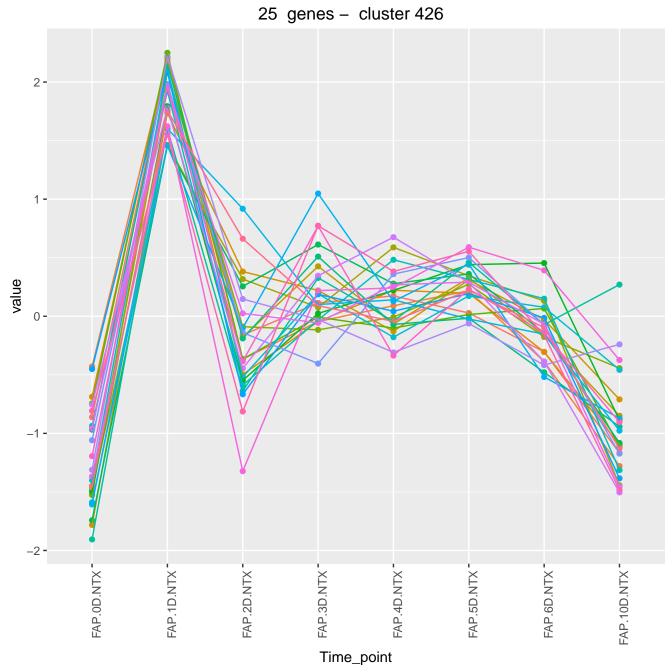


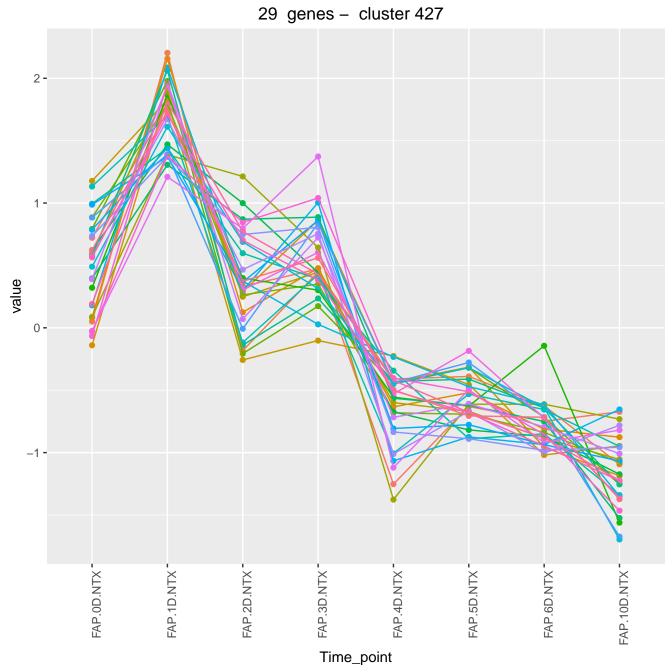




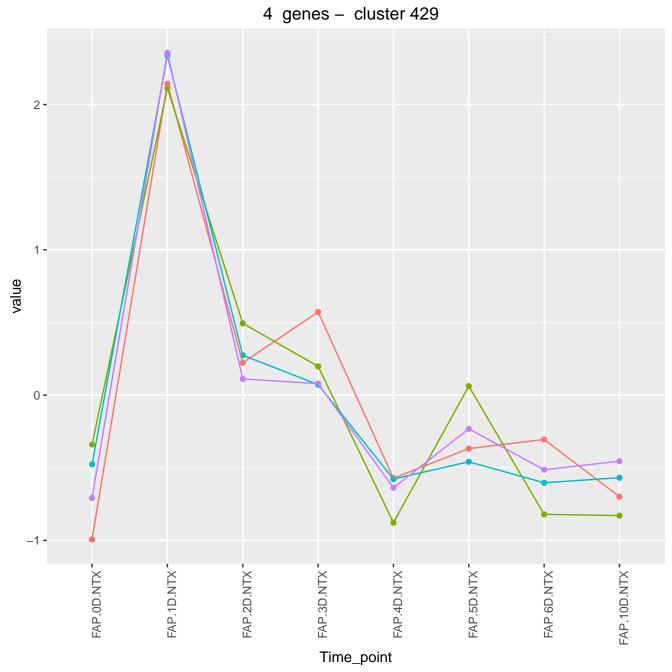


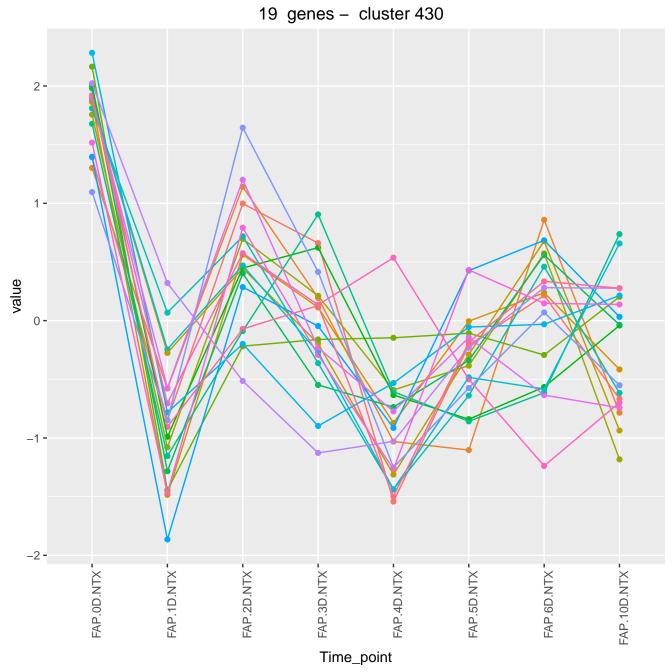


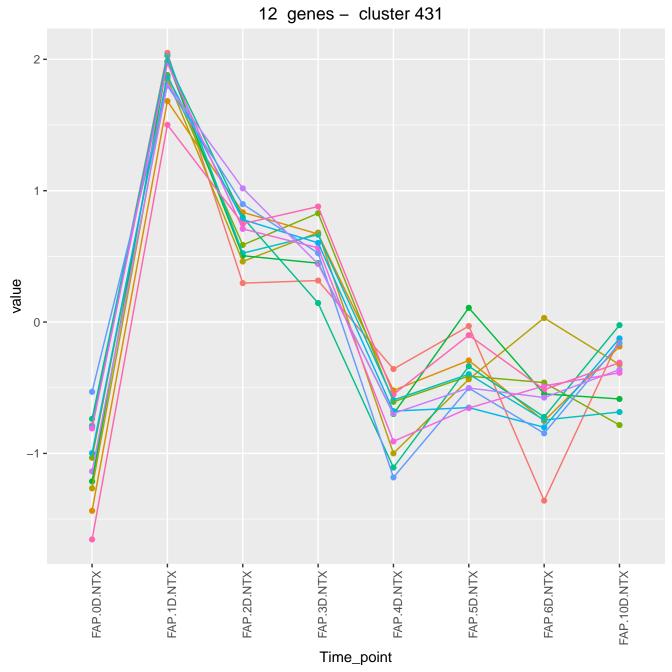


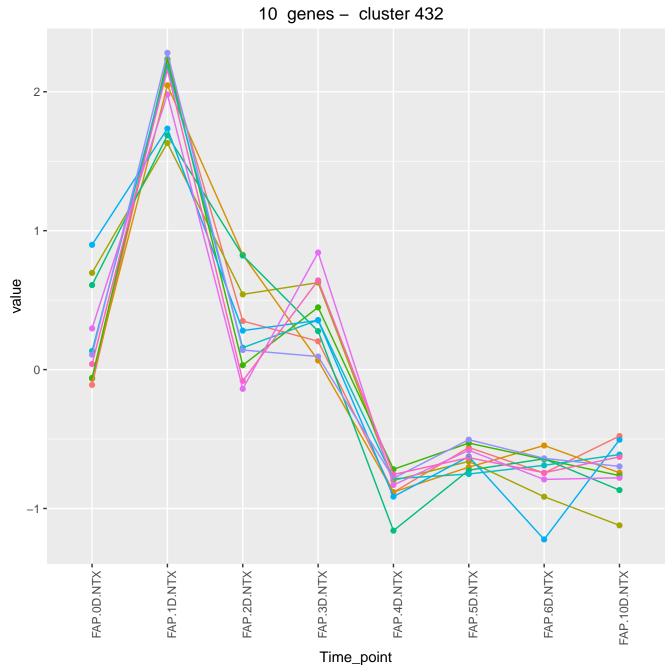


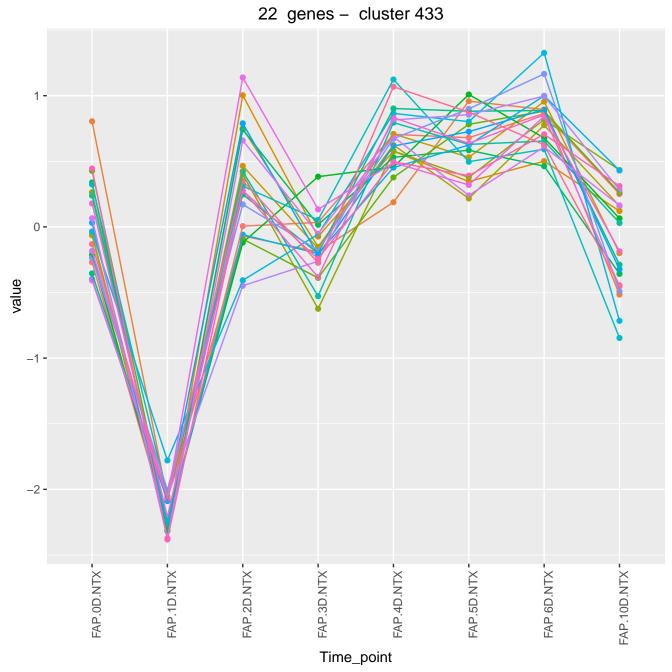
16 genes - cluster 428 1 -0 value \_1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX <sup>7</sup> FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.3D.NTX FAP.6D.NTX FAP.4D.NTX Time\_point

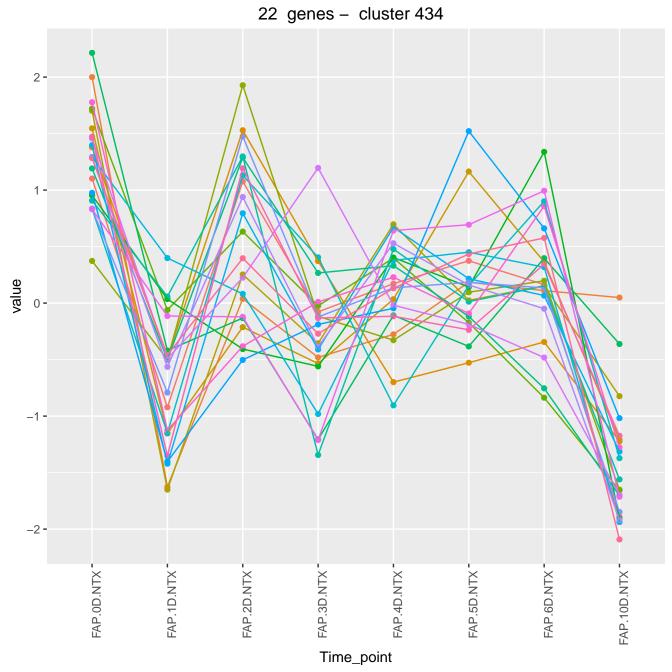


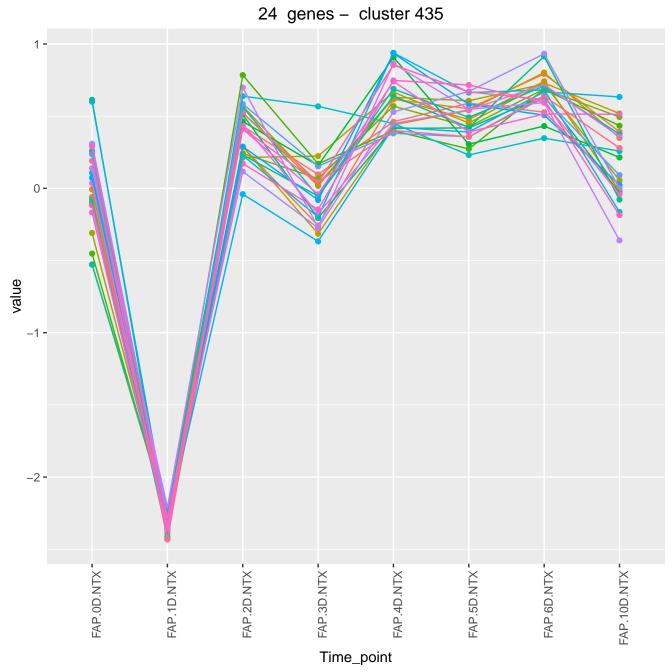


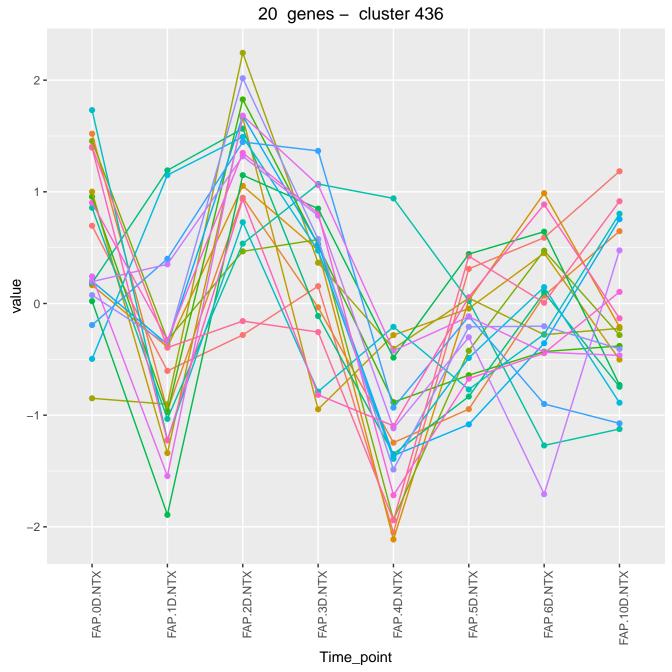




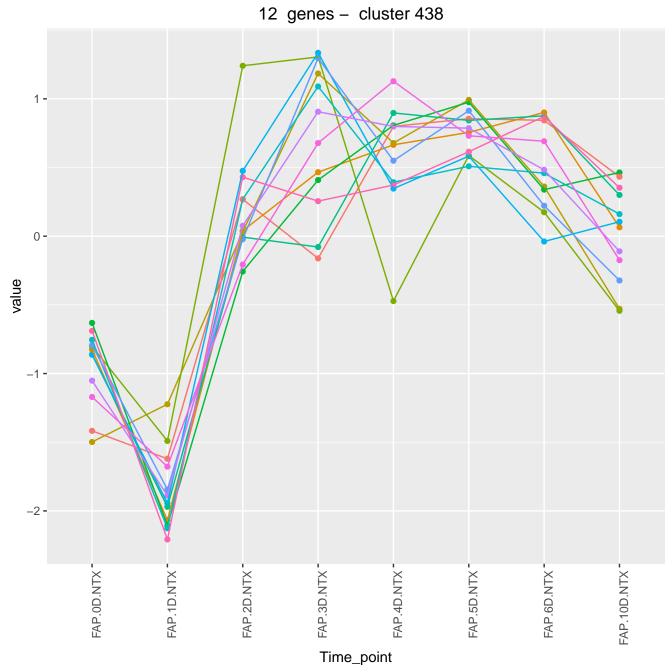


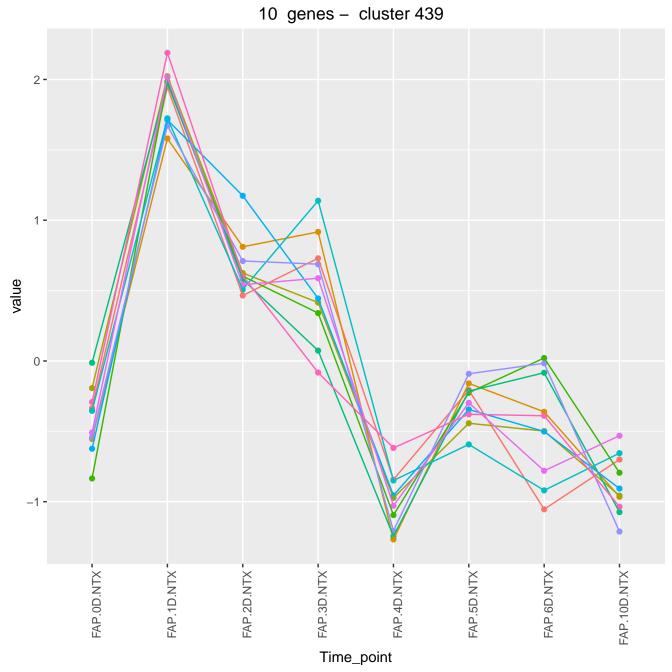


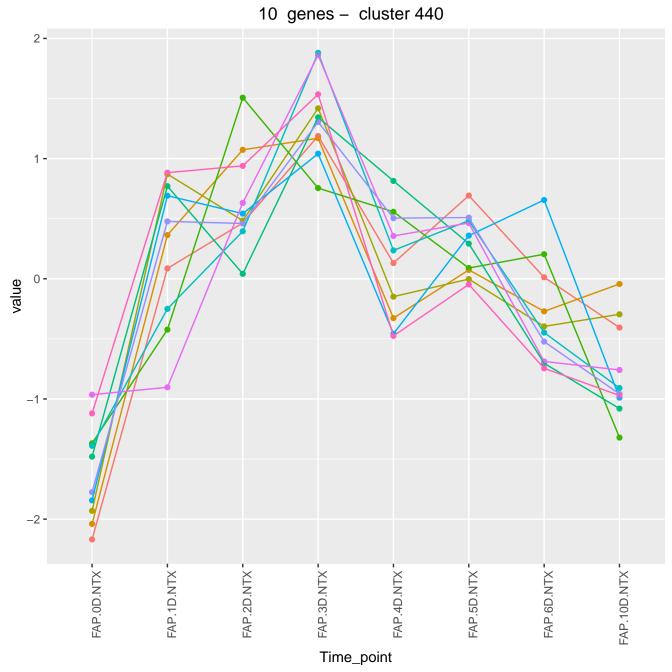


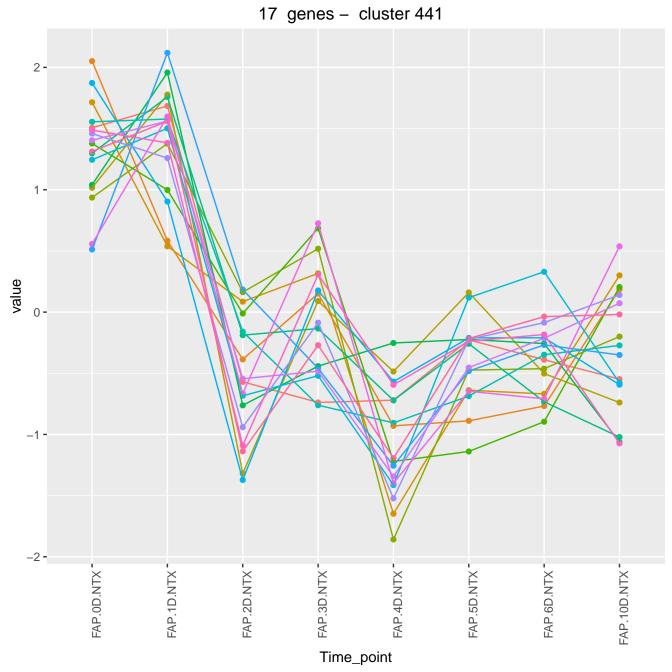


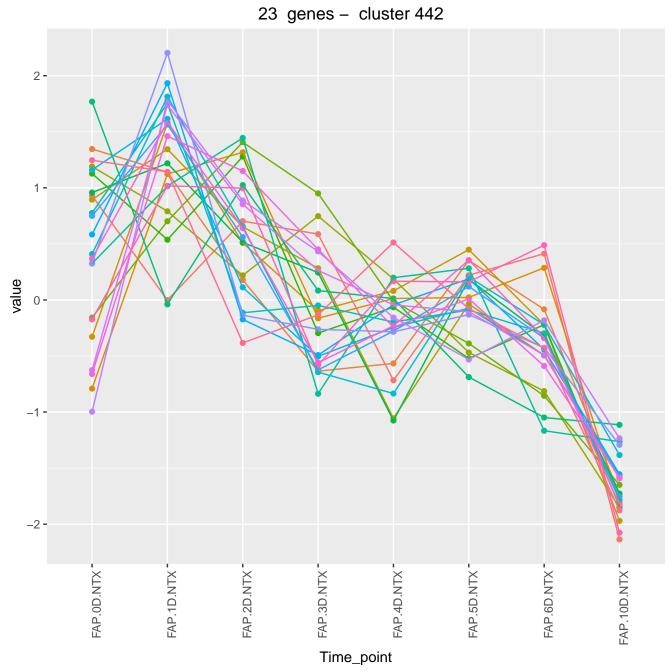
14 genes - cluster 437 1 -0 value \_1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX -FAP.2D.NTX <sup>-</sup> FAP.3D.NTX -FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.6D.NTX FAP.4D.NTX Time\_point

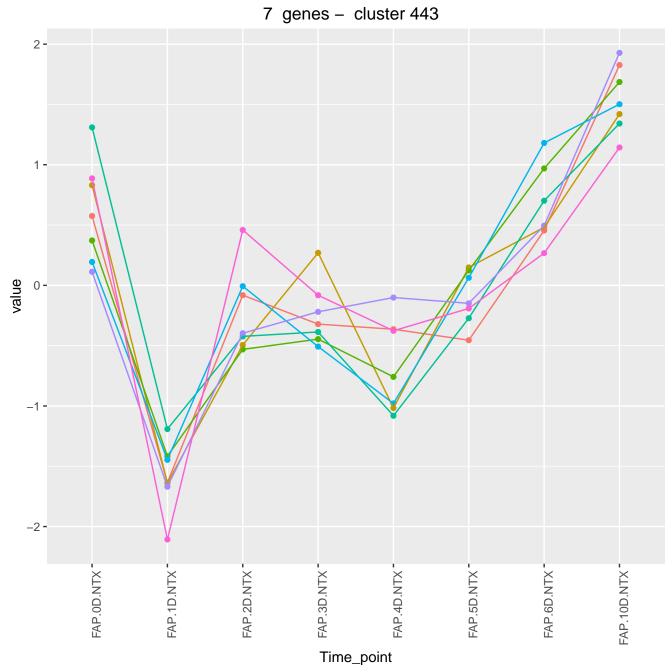




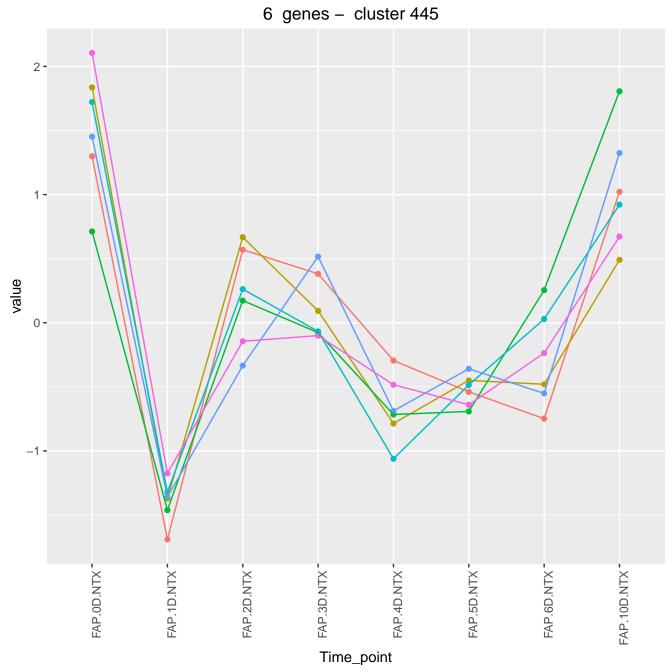


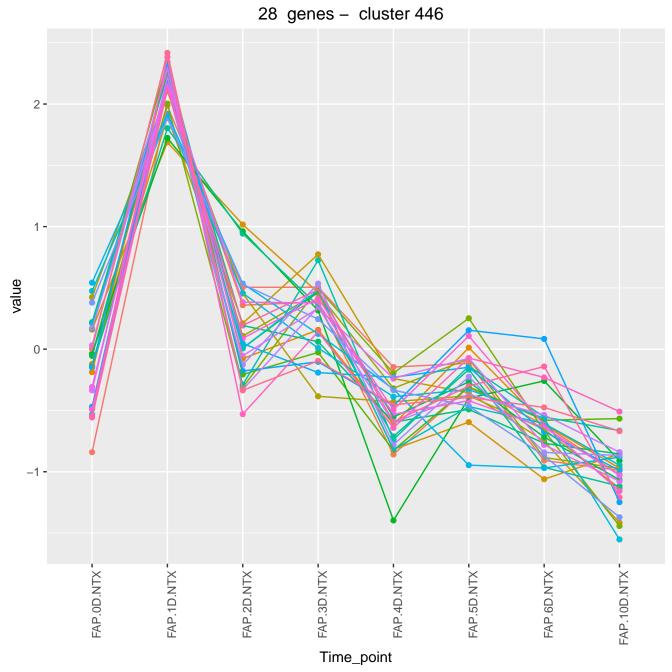


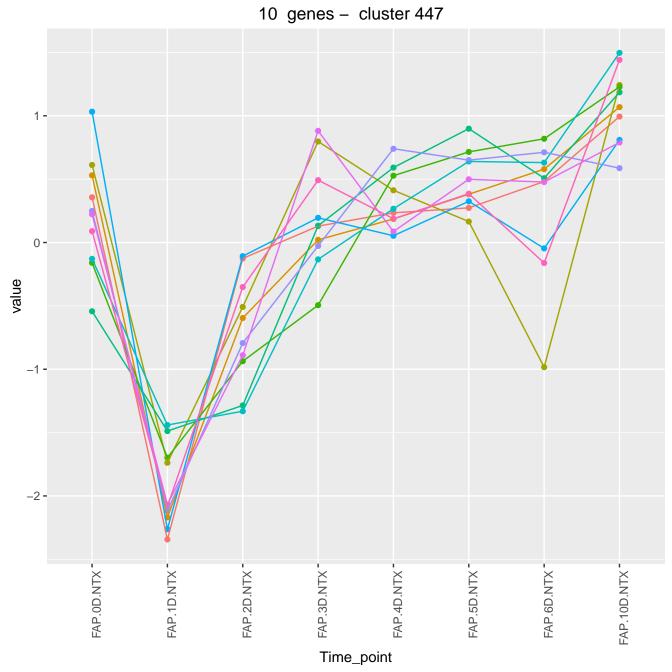


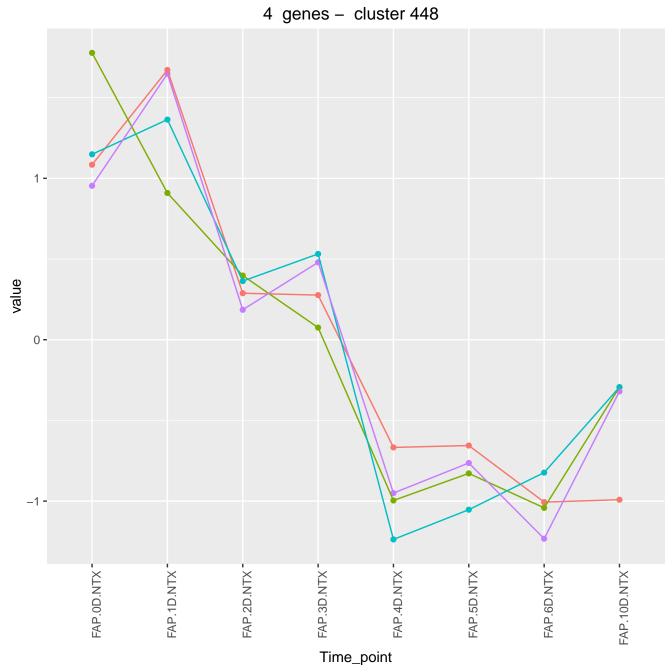


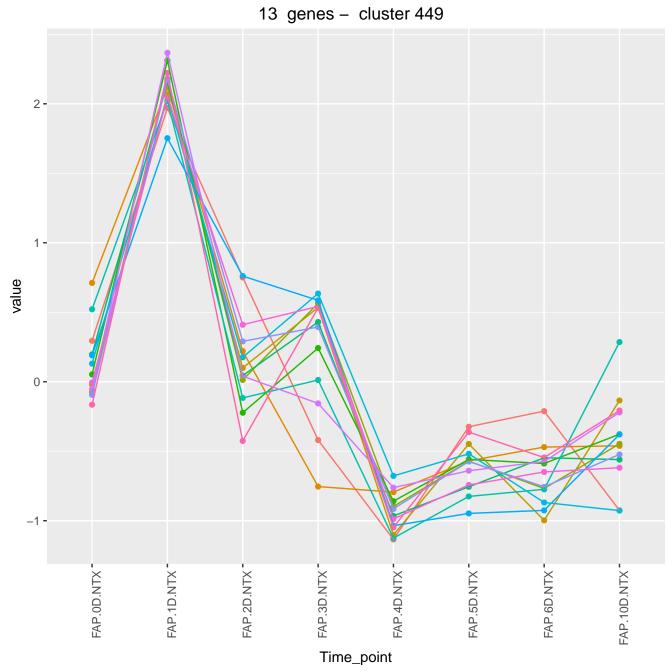
26 genes - cluster 444 2 -1 value 0 -**-1** --2 **-**FAP.0D.NTX 7 FAP.3D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.1D.NTX FAP.2D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

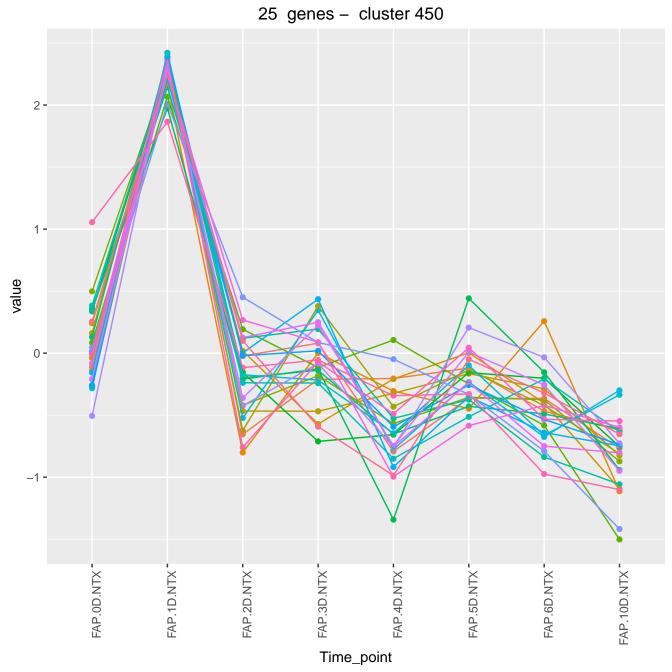


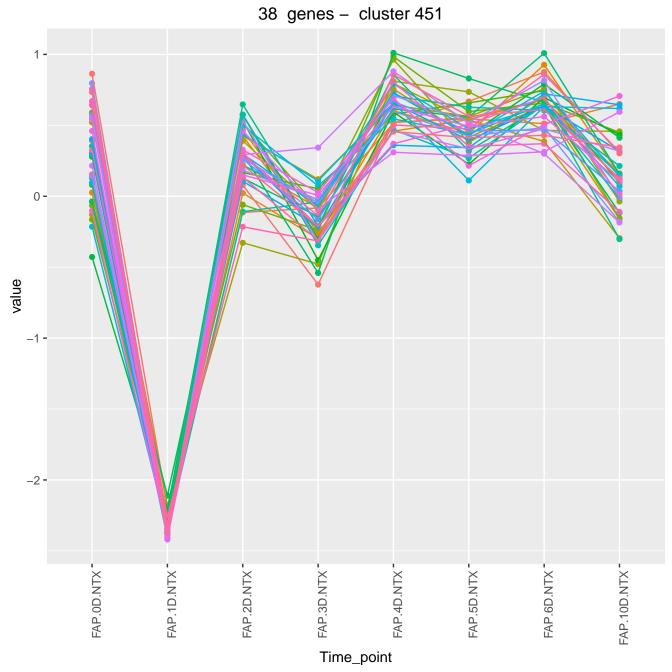


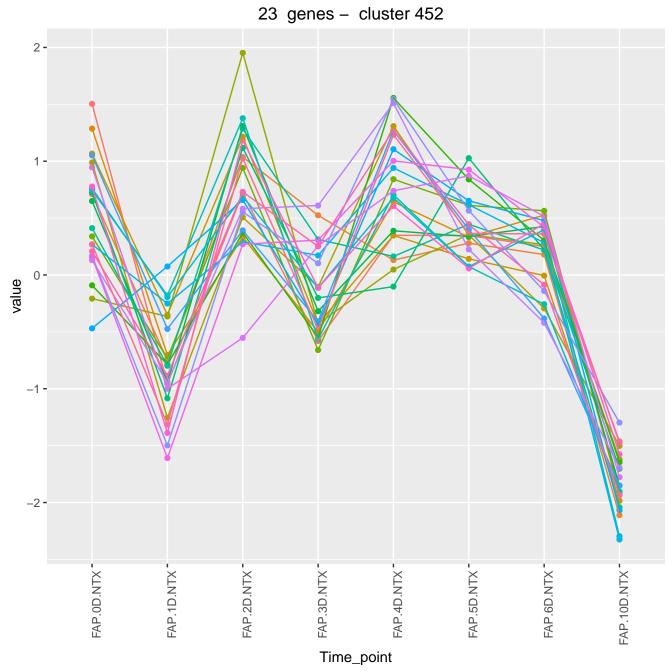




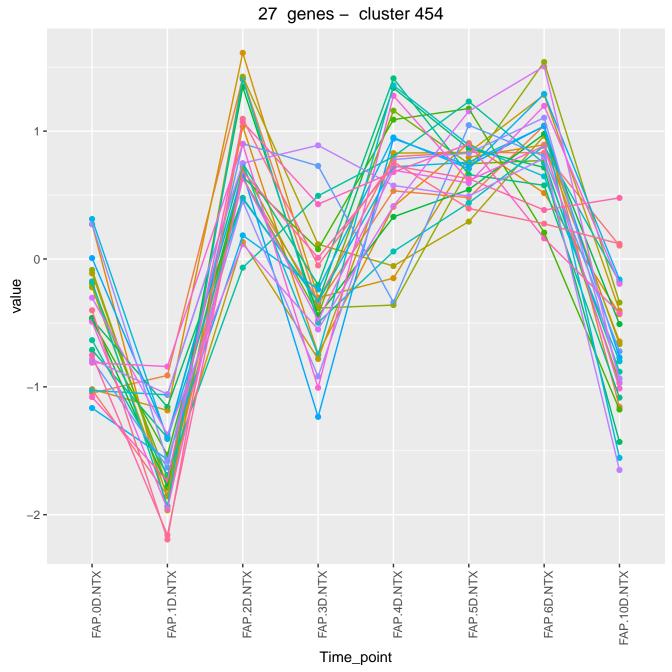


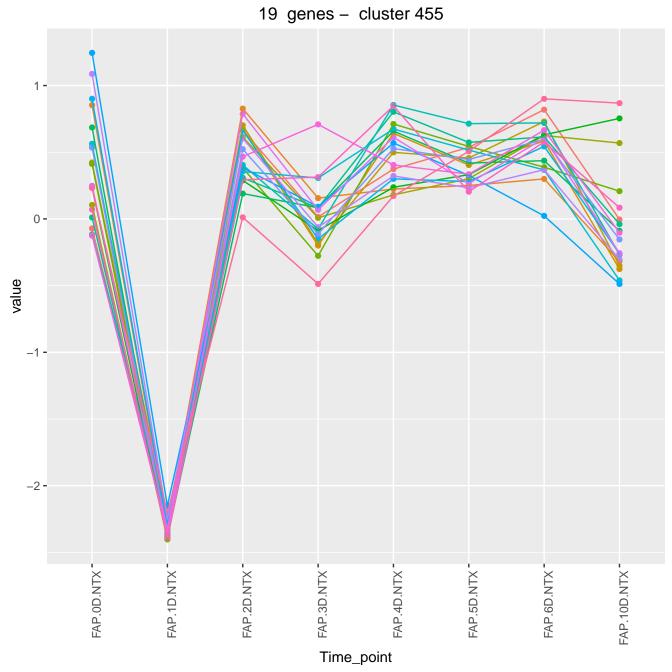


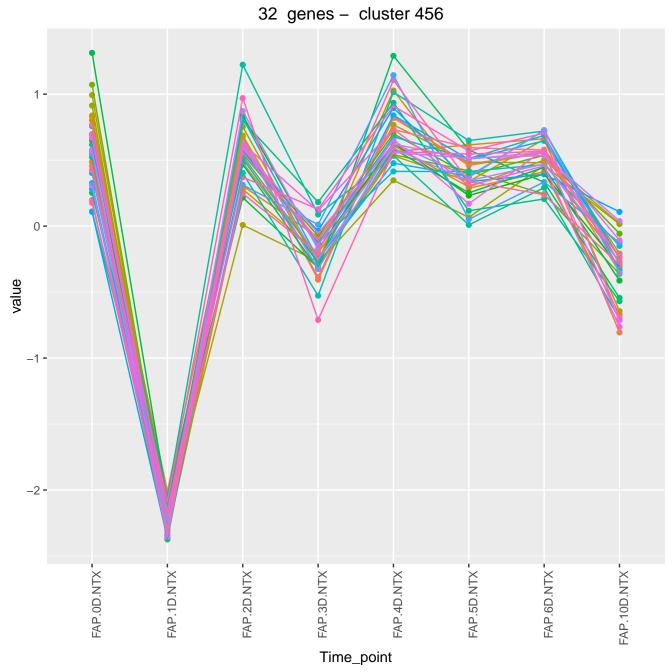


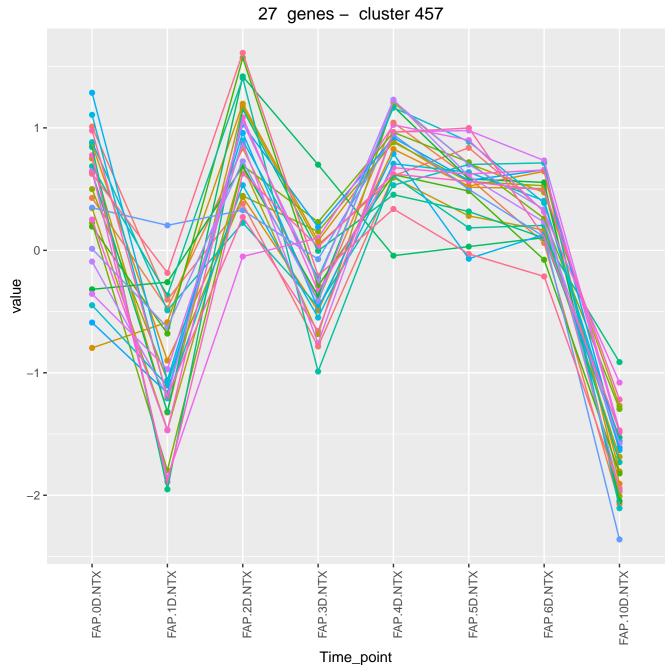


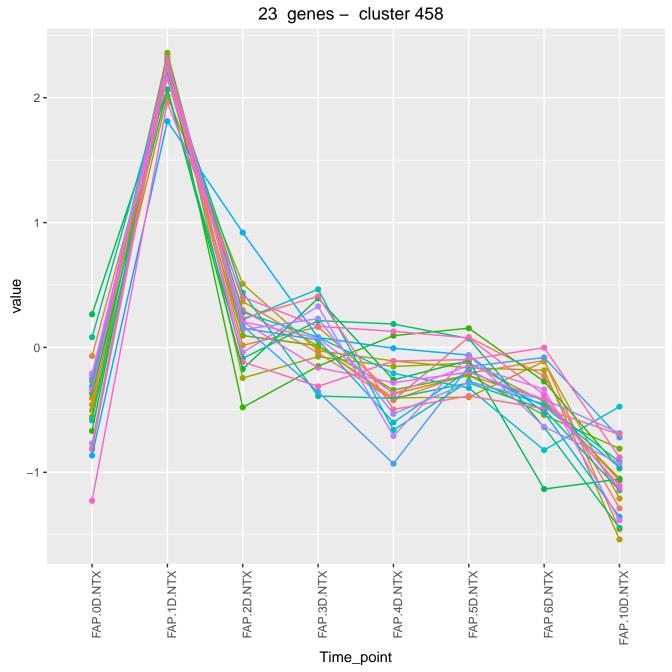
7 genes – cluster 453 1 -0 value -1 **-**-2 **-**FAP.0D.NTX -FAP.1D.NTX -FAP.2D.NTX -FAP.3D.NTX -FAP.5D.NTX <sup>7</sup> FAP.6D.NTX <sup>7</sup> FAP.10D.NTX -FAP.4D.NTX Time\_point

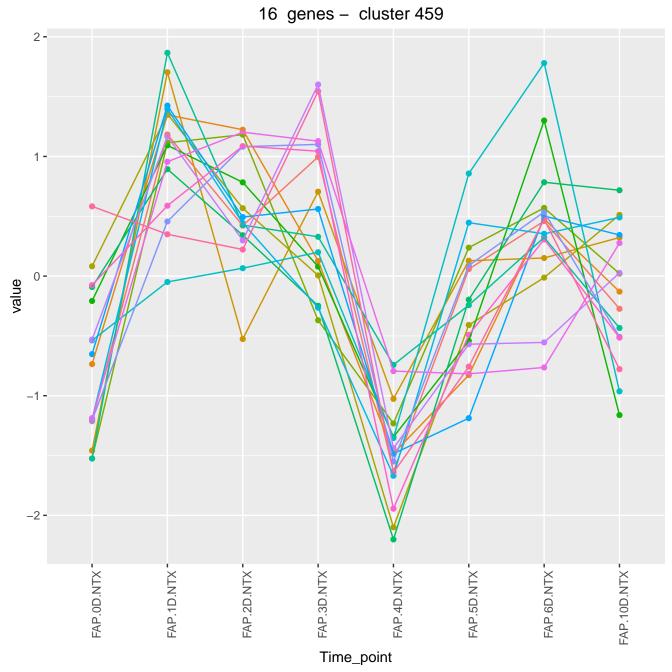


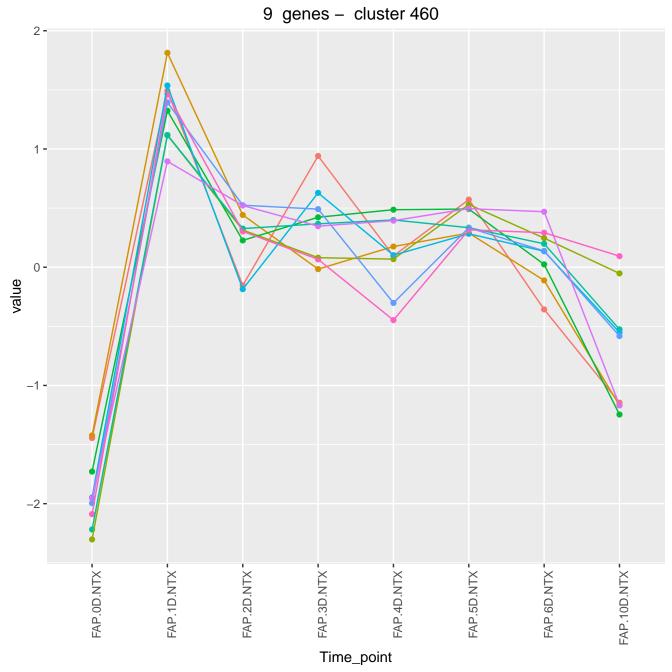


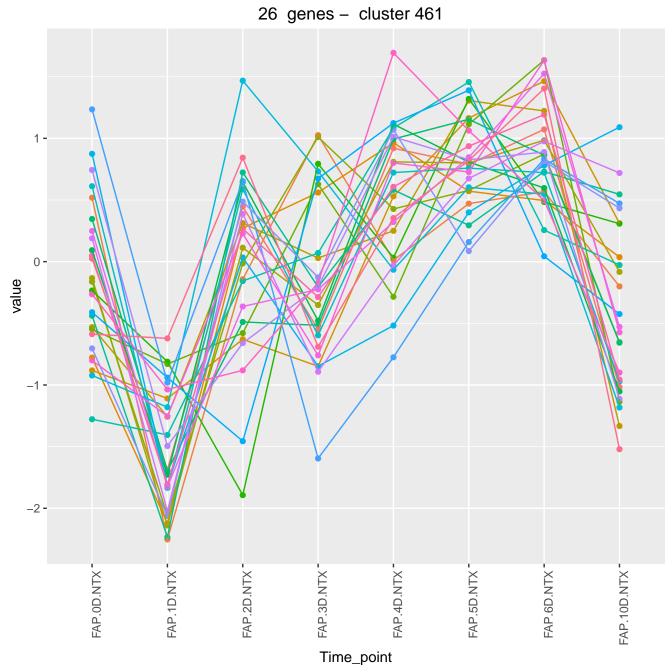




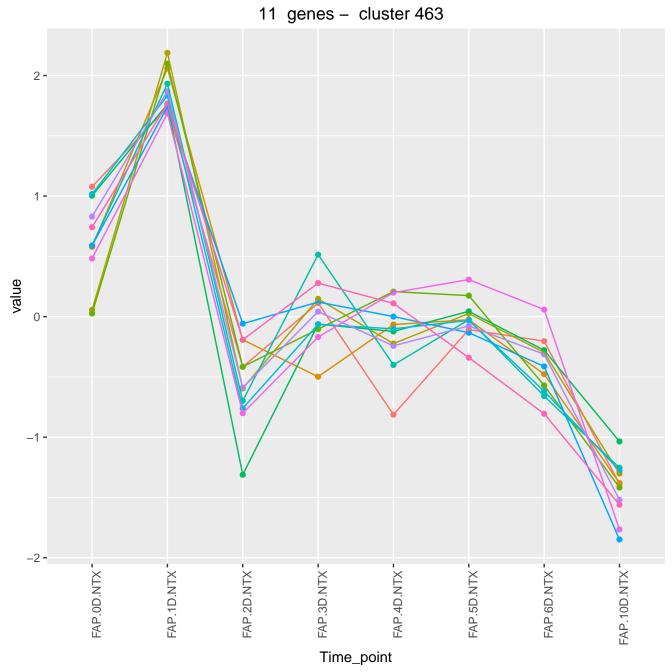


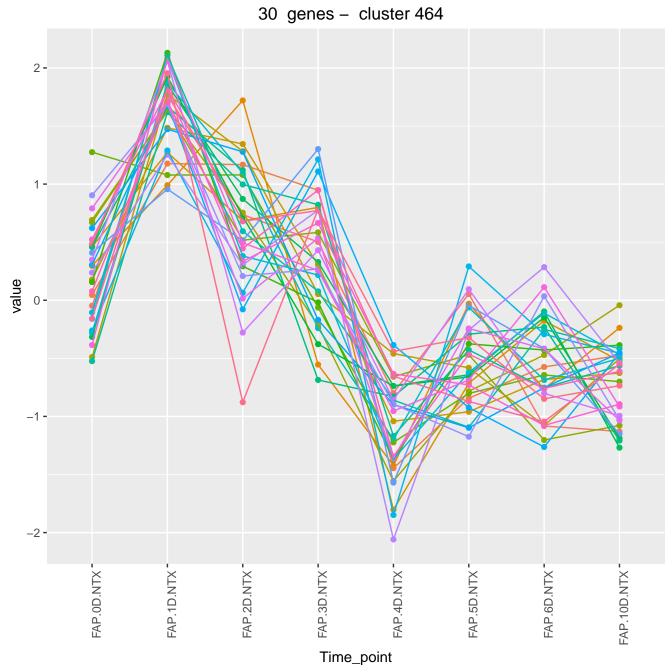


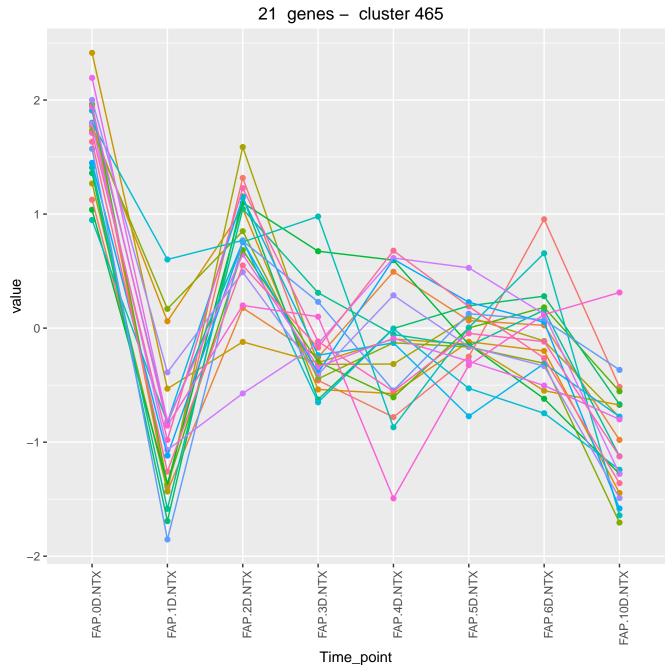


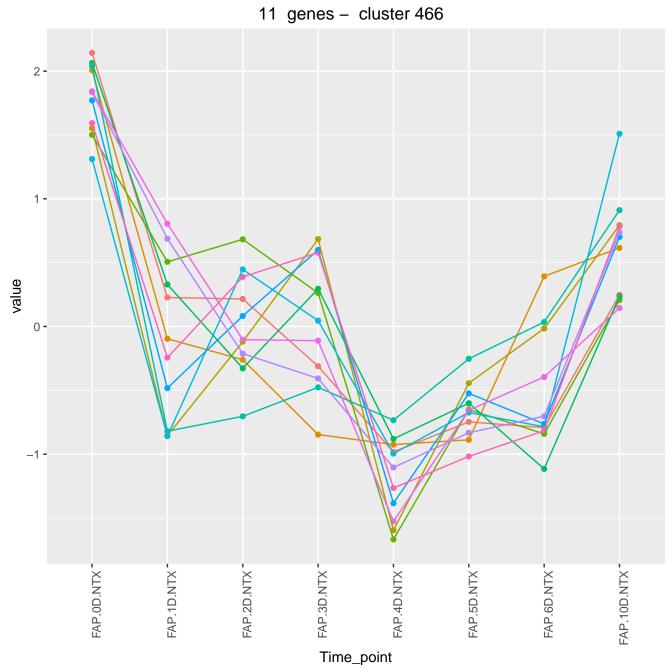


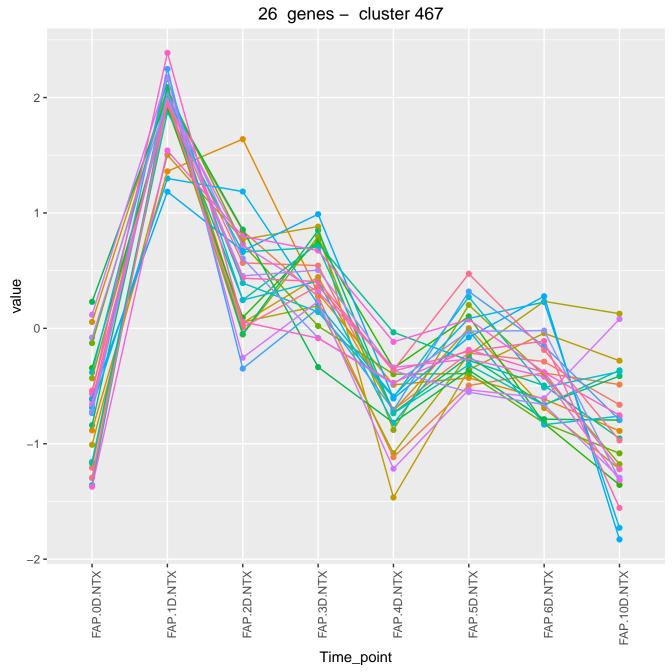
38 genes - cluster 462 1 -0 value \_1 **-**-2 **-**FAP.0D.NTX 7 FAP.10D.NTX -FAP.1D.NTX FAP.2D.NTX FAP.3D.NTX FAP.5D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point

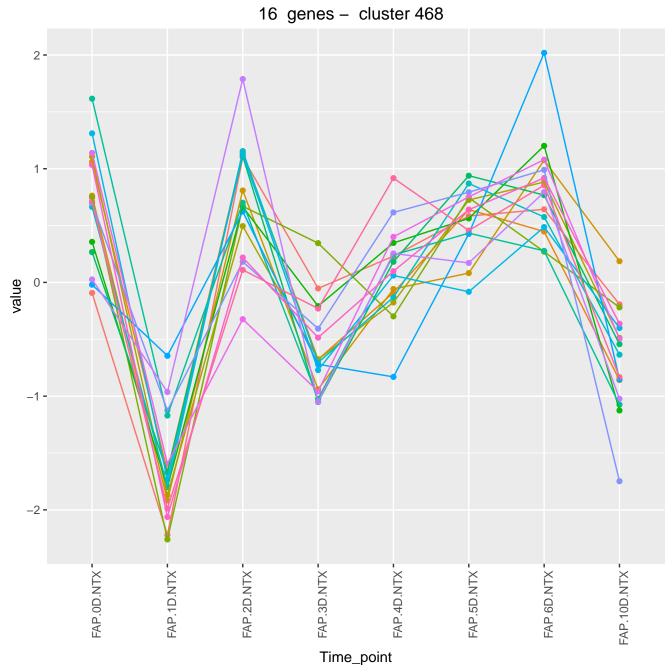




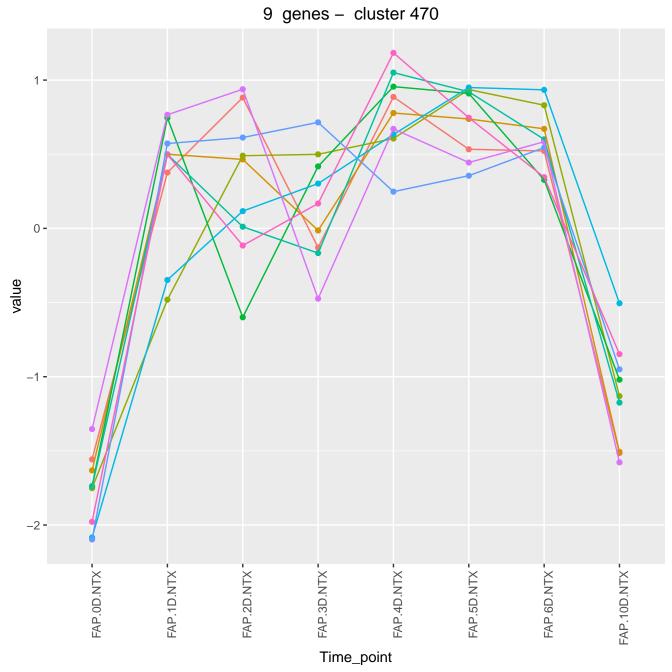


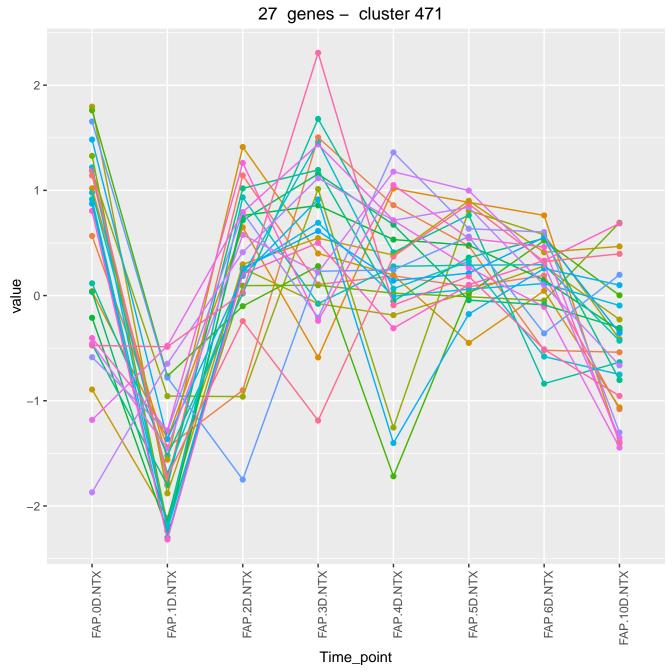


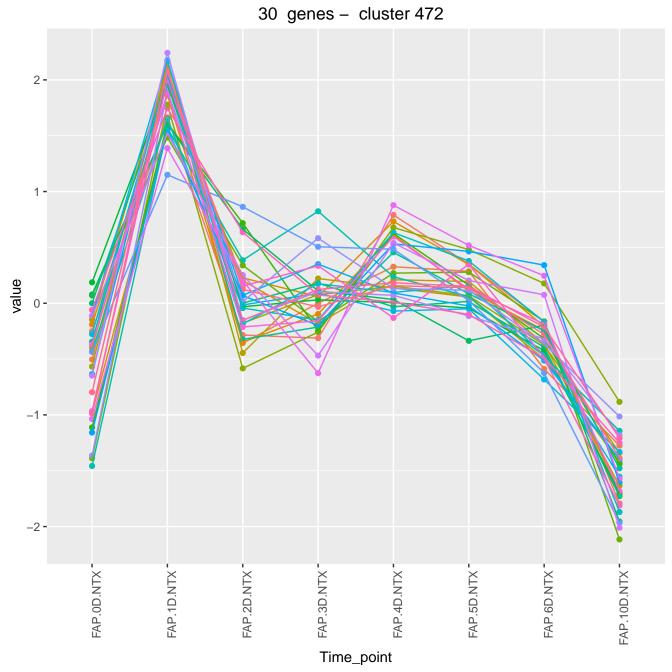




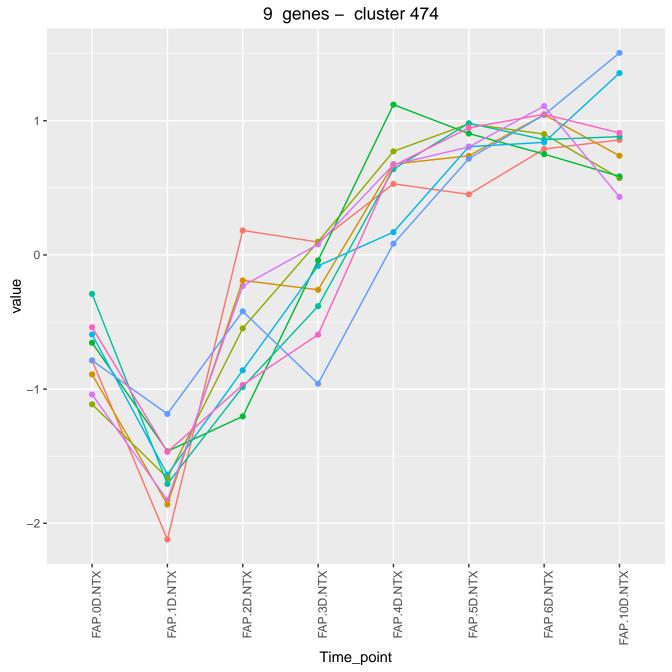
27 genes - cluster 469 2-1 -0 value -1 **-**-2 **-**FAP.0D.NTX 7 FAP.10D.NTX -FAP.1D.NTX -FAP.2D.NTX FAP.3D.NTX FAP.5D.NTX FAP.4D.NTX FAP.6D.NTX Time\_point







32 genes - cluster 473 2 -1 -0 value \_1 **-**-2 **-**FAP.0D.NTX 7 FAP.1D.NTX <sup>7</sup> FAP.2D.NTX -FAP.3D.NTX 7 FAP.5D.NTX <sup>7</sup> FAP.6D.NTX <sup>7</sup> FAP.10D.NTX -FAP.4D.NTX Time\_point



32 genes - cluster 475 1 -0 value -1 --2 **-**FAP.0D.NTX -FAP.5D.NTX <sup>7</sup> FAP.10D.NTX -FAP.2D.NTX FAP.3D.NTX FAP.1D.NTX FAP.6D.NTX 7 FAP.4D.NTX Time\_point

