262 genes - 80%-standardized 2 -1 value **-1** --2 **-**\_ 00 <u>\_</u>10 D3.1 Lime\_boint

245 genes – 80%-standardized 0 value **-1-**-2 **-**\_ 00 D3.1 <u>\_</u> \_9Q D7\_ Lime\_boint

243 genes - 80%-standardized 2 value 0 -**-1-**

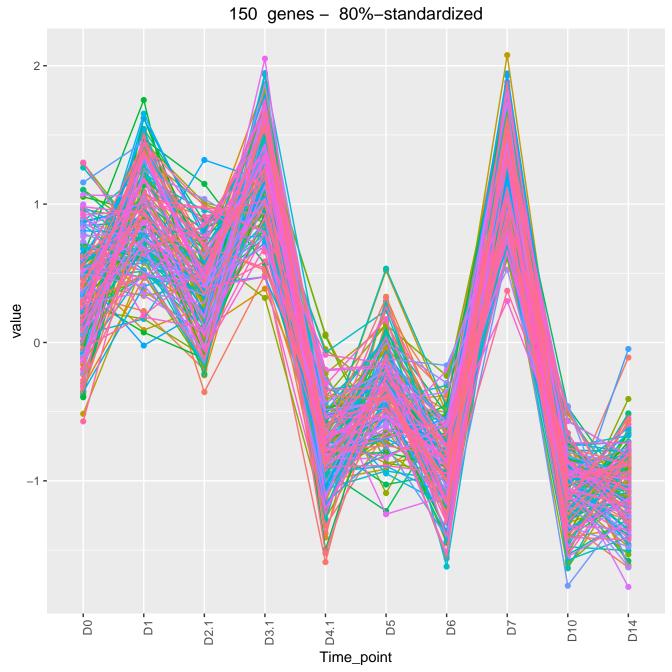
\_9Q

D7

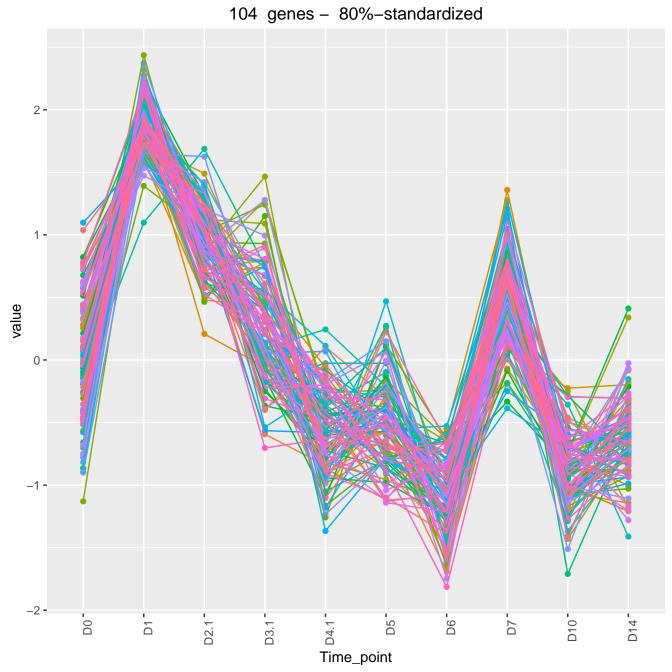
\_ 00 <u>\_</u>10

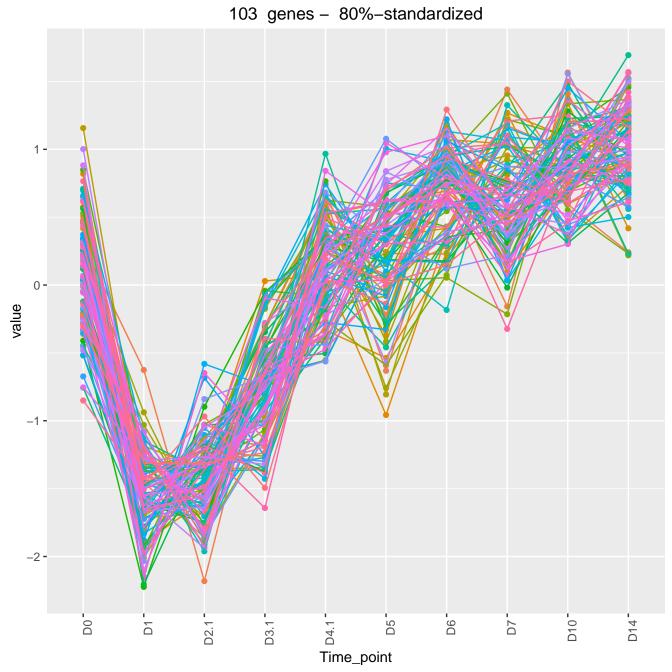
D3.1

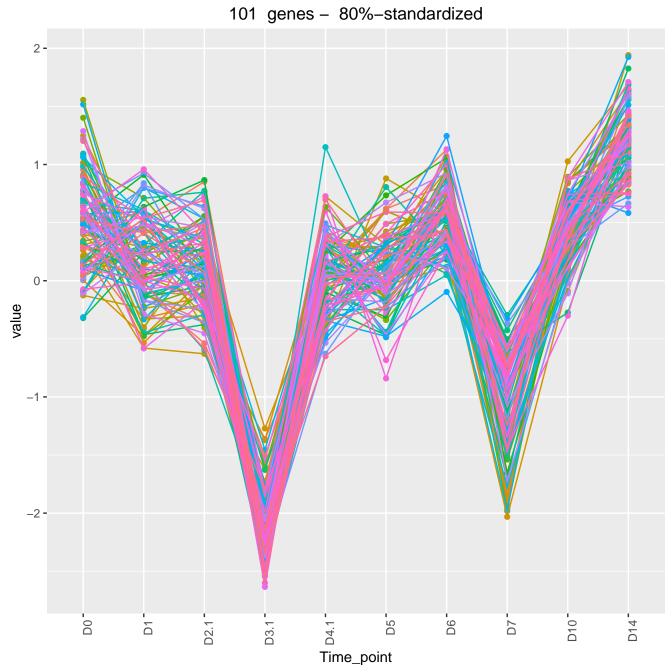
Lime\_boint

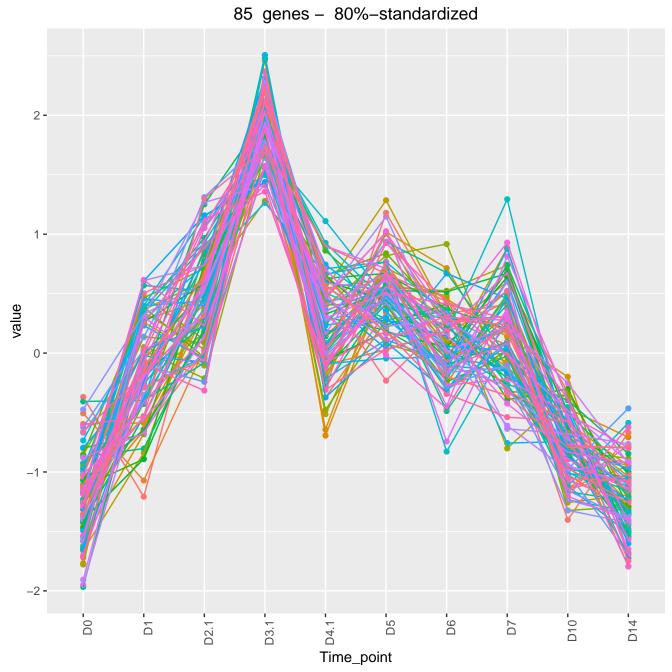


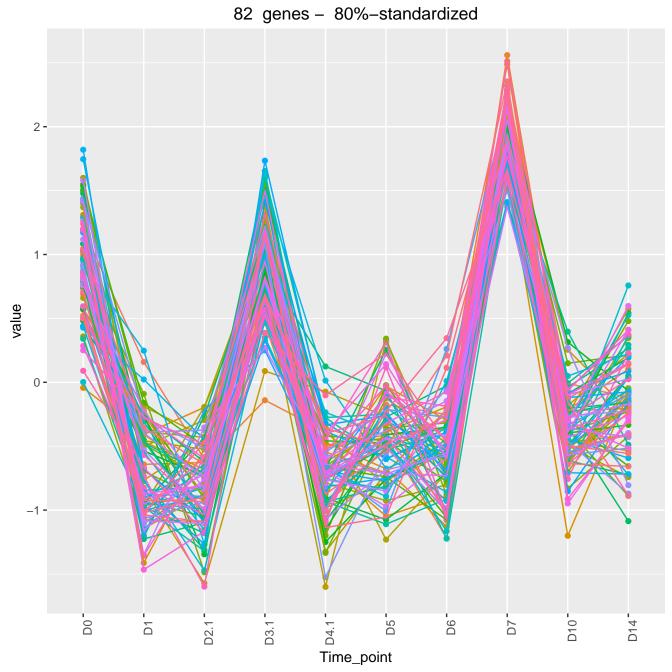
125 genes – 80%-standardized 2 -1 value 0 --1 **-**-2 **-**\_0Q \_ \_ D3.1 D5 -\_9Q D7\_ Lime\_boint

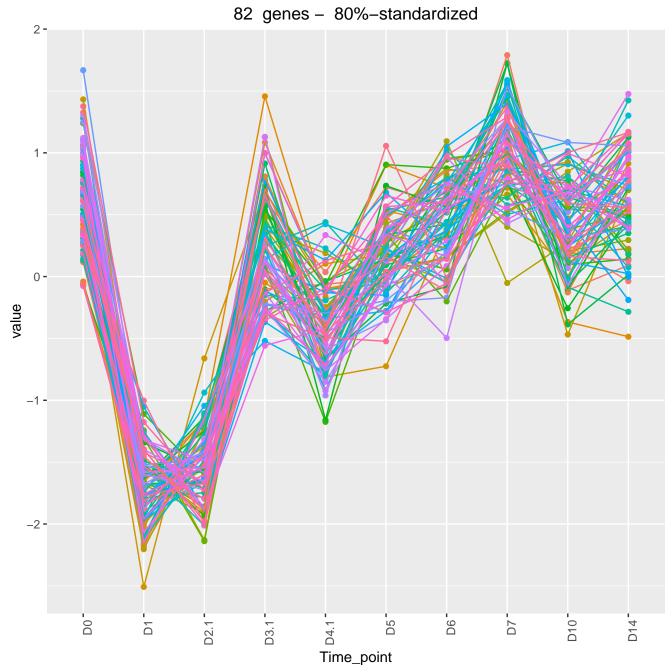


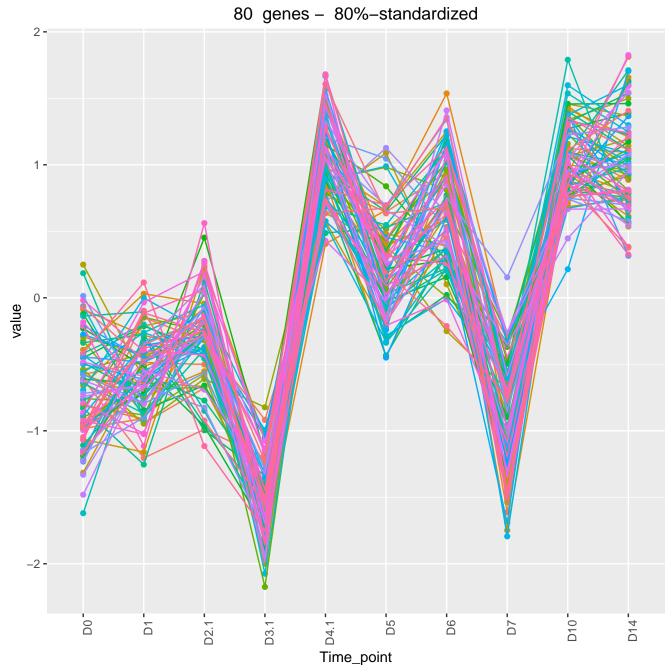


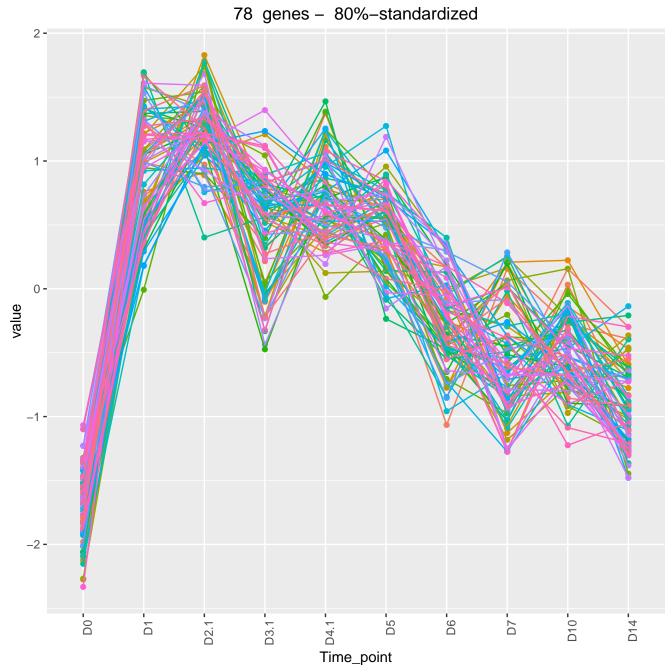






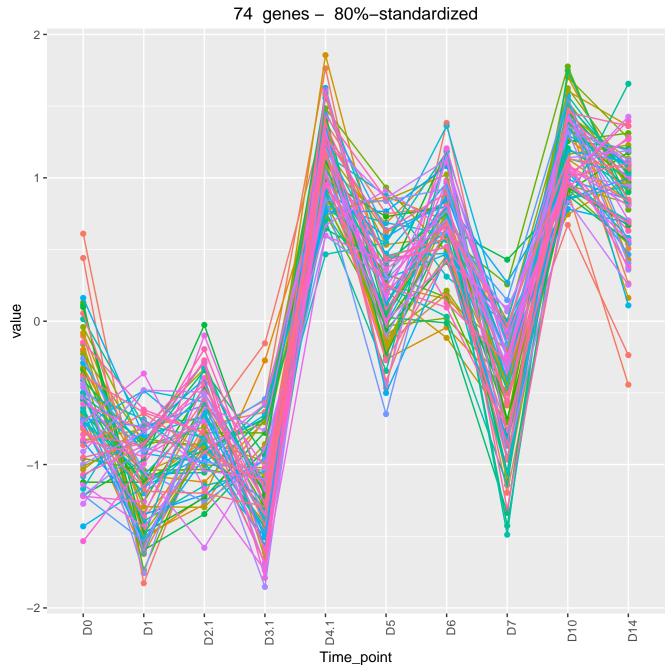




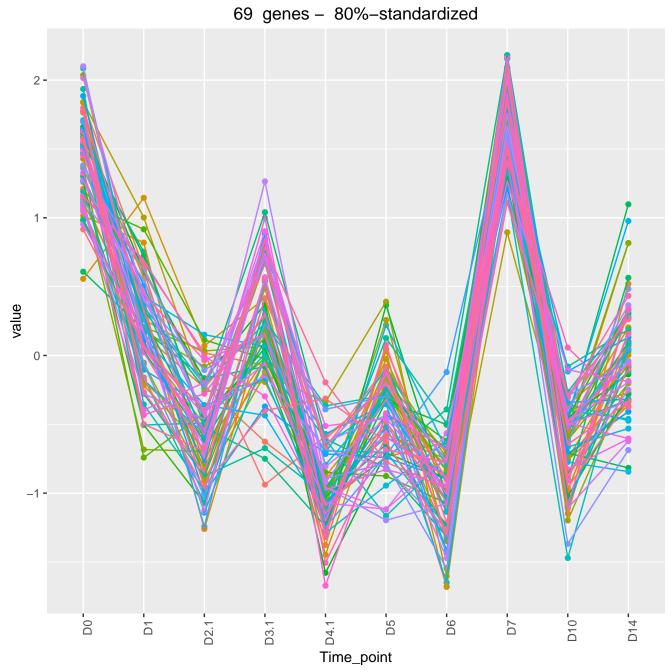


75 genes - 80%-standardized 2-1 value -1 **-**-2 **-**\_ 00 <u>\_</u>10 D3.1 \_9Q D7\_ Lime\_boint

74 genes - 80%-standardized 2 value o --1 **-**-2 **-**\_ \_ D3.1 Lime\_boint

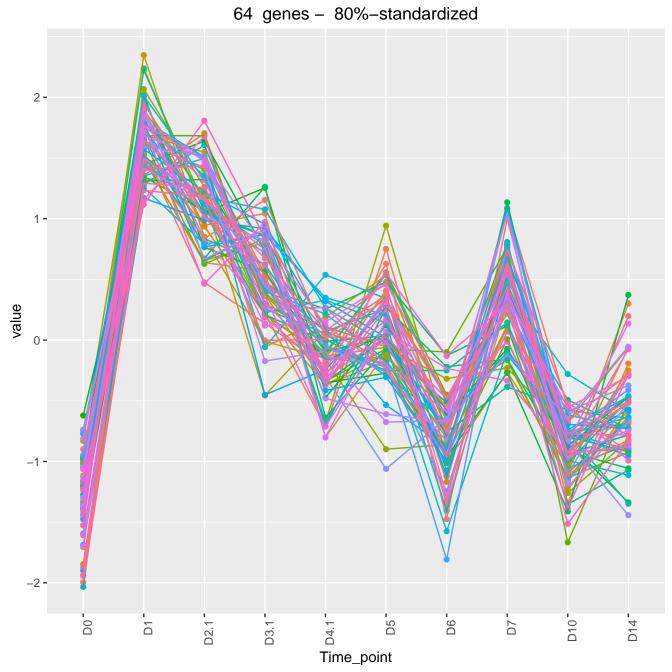


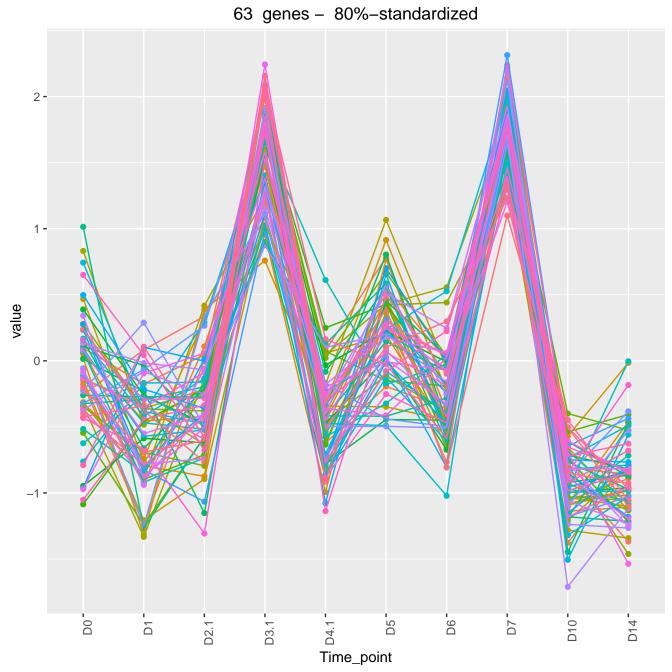
73 genes - 80%-standardized 2 -1 value 0 --1 **-**-2 **-**<u>\_</u>10 D3.1 \_9Q D7\_ Lime\_boint



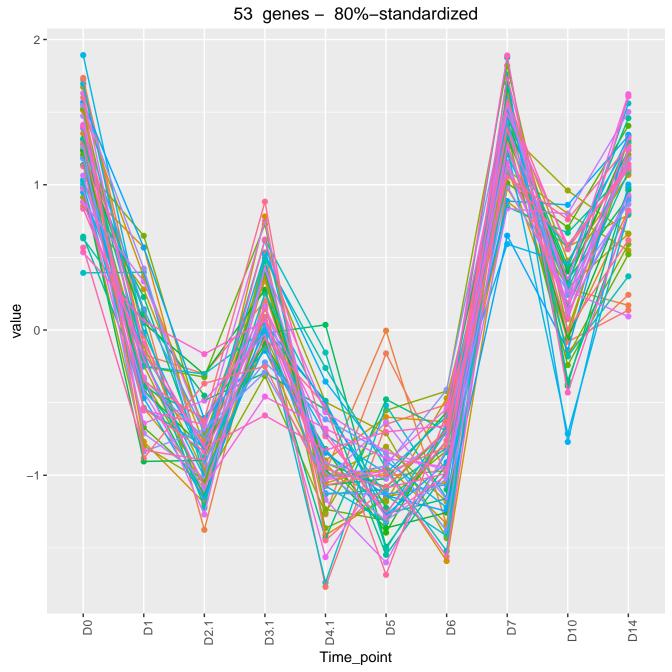
65 genes - 80%-standardized 2 -1 value 0 -**-1-**-2 **-**D3.1 \_9Q D7\_ Lime\_boint

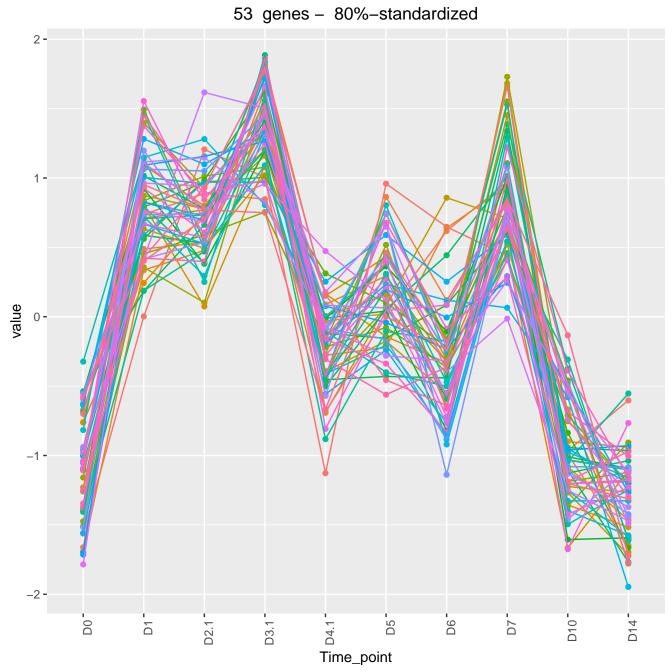
64 genes - 80%-standardized 2 -1 value o -**-1-**-2 **-**\_ 00 D5 -<u>\_</u>10 D3.1 \_9Q D7 Lime\_boint



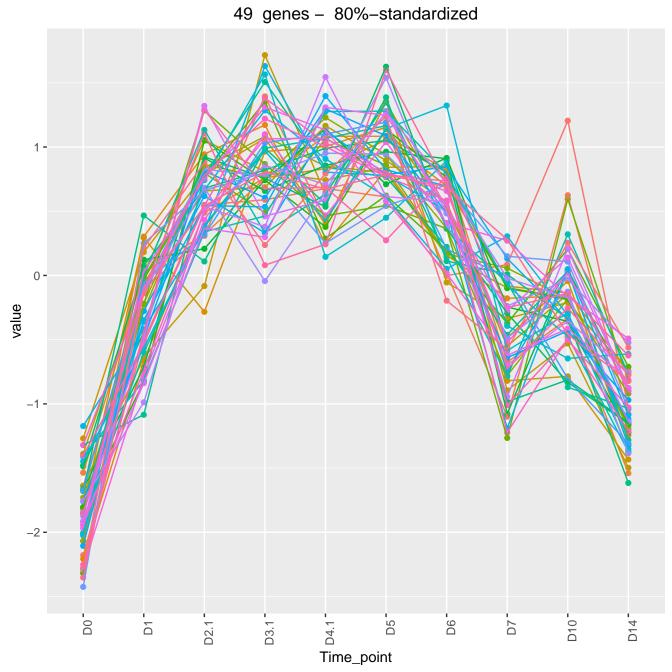


56 genes - 80%-standardized 2 value 0 --1 **-**-2 **-**\_0Q <u>\_</u>10 D5 -D3.1 \_9Q D7\_ Lime\_boint

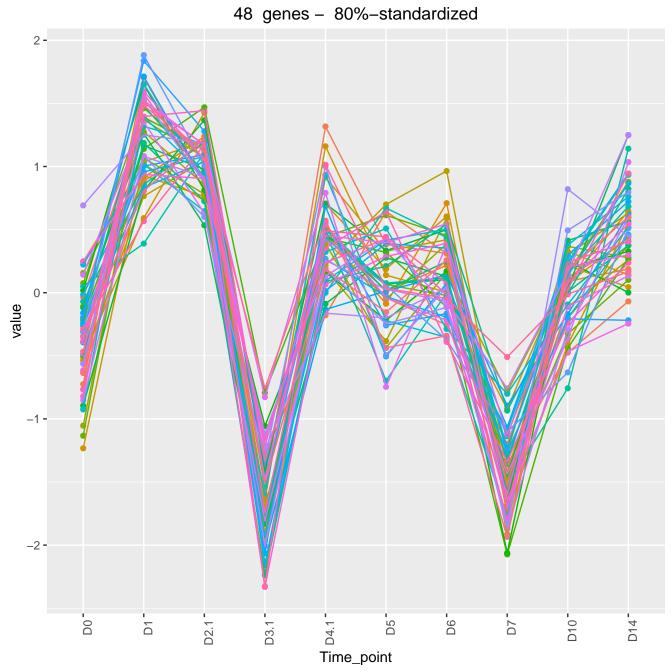


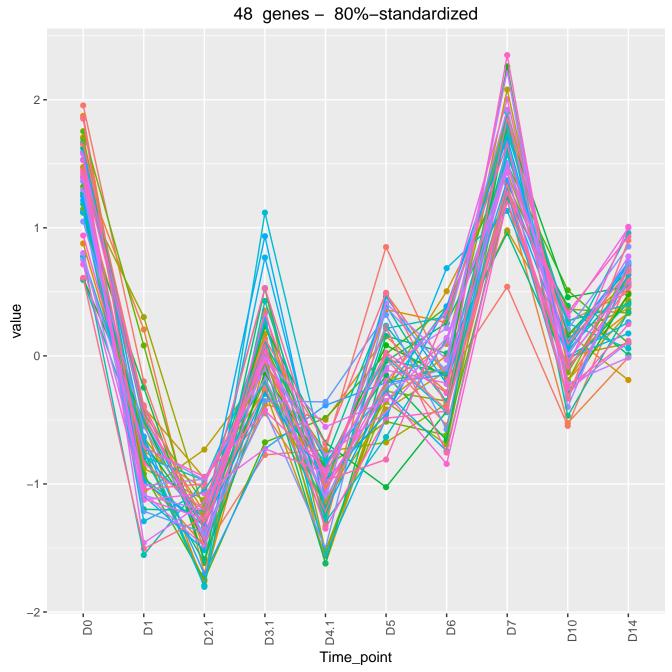


52 genes - 80%-standardized 0 value **-1 -**-2 **-**D5 -<u>\_</u>10 D3.1 \_9Q D7 Lime\_boint

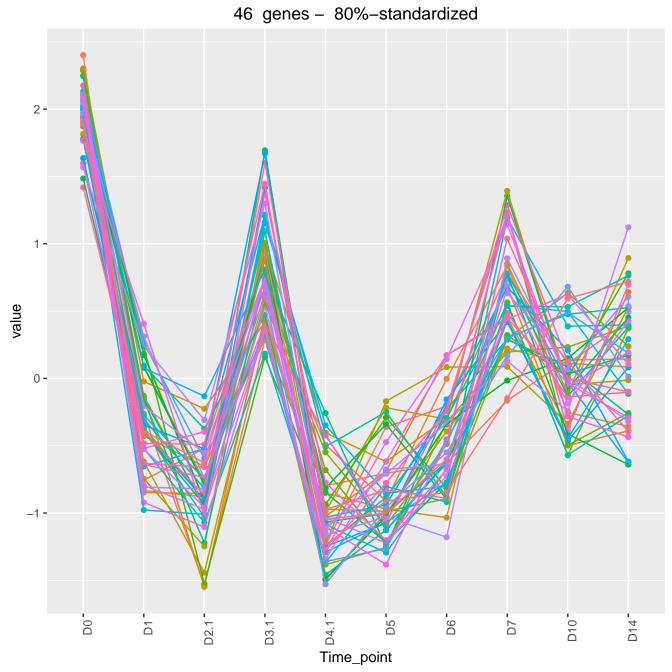


49 genes - 80%-standardized 2 -1 value 0 --1 **-**-2 **-**\_0Q \_ \_ D5 -D3.1 D7\_ Lime\_boint \_9Q

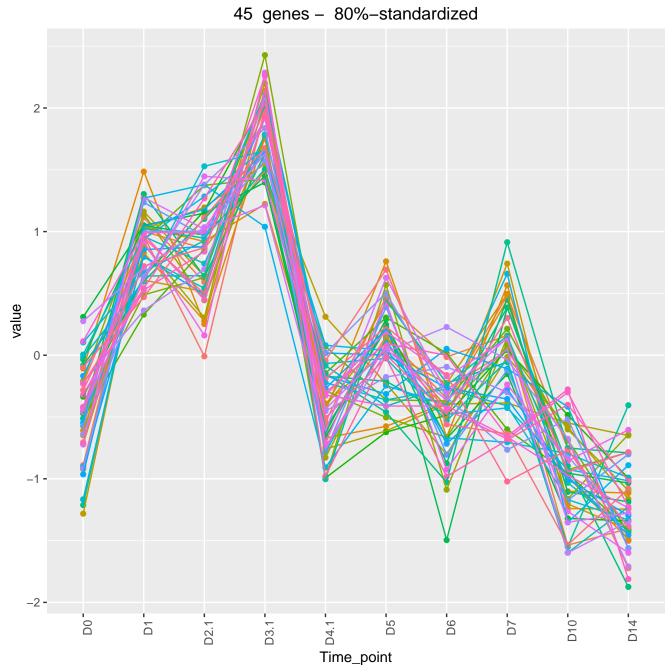


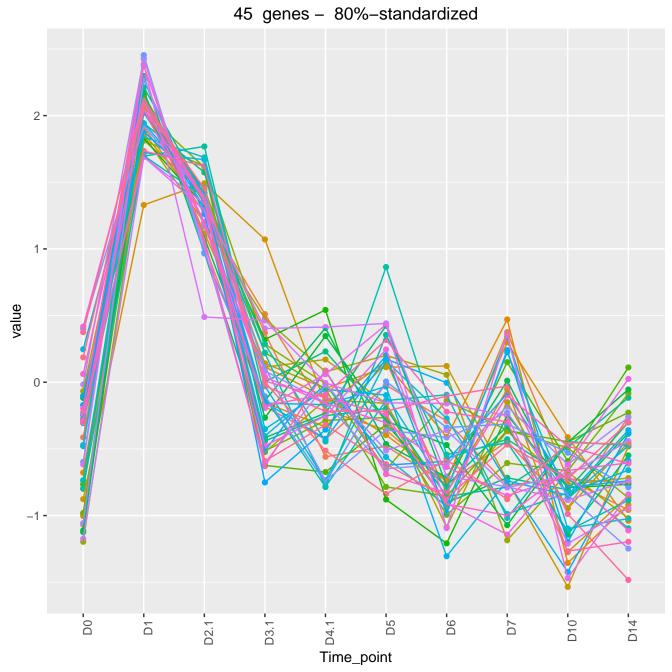


47 genes - 80%-standardized 2 value 0 -**-1-**\_ 00 \_ D2\_ <u>\_</u>10 D3.1 \_9Q D7 Lime\_boint



45 genes - 80%-standardized 2 -1 value - 0 **-1** --2 **-**\_0Q Time\_point D3.1 <u>\_</u>10 \_9Q D7<sup>-</sup>



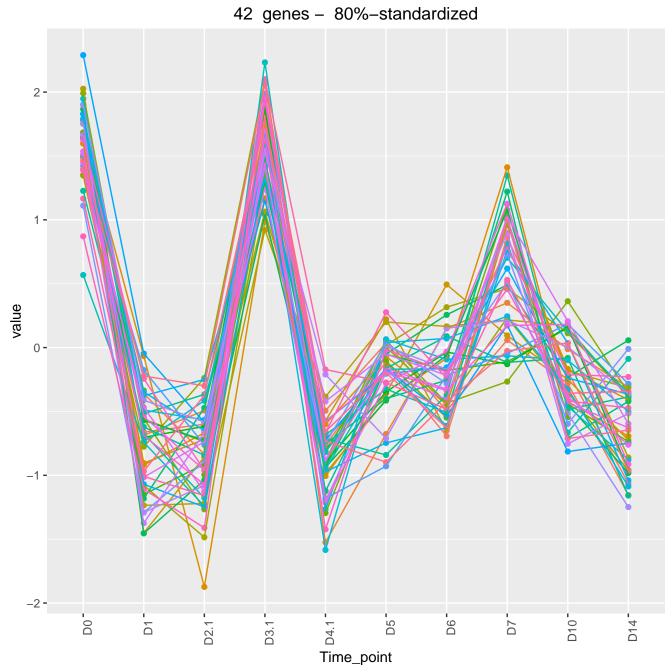


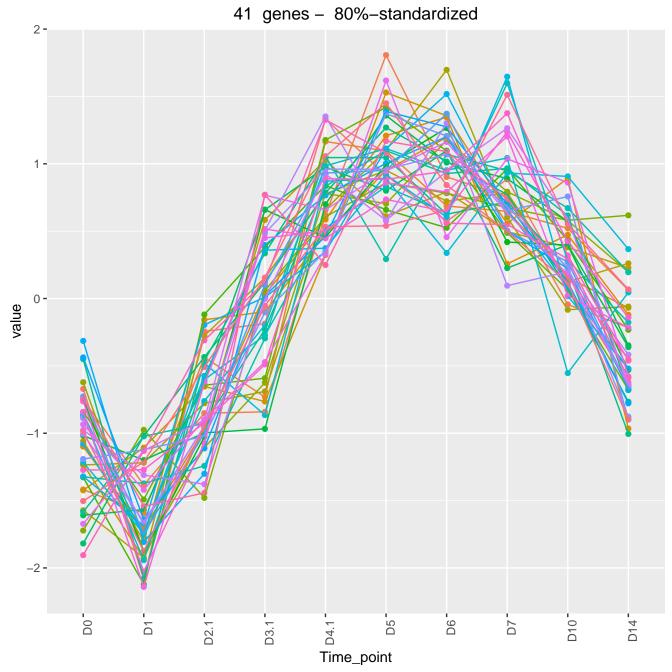
45 genes - 80%-standardized 2 -1 value **−1 -**-2 **-**\_ 00 <u>\_</u>10 D3.1 \_9Q Lime\_boint

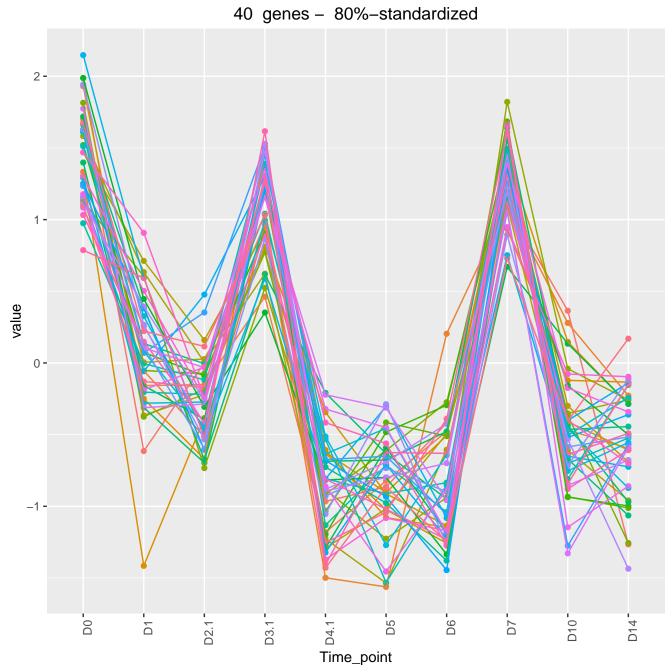
42 genes - 80%-standardized 2 -1 value 0 --2 **-**\_ 00 \_ D2\_ <u>\_</u>10 D3.1 \_9Q D7\_ Lime\_boint

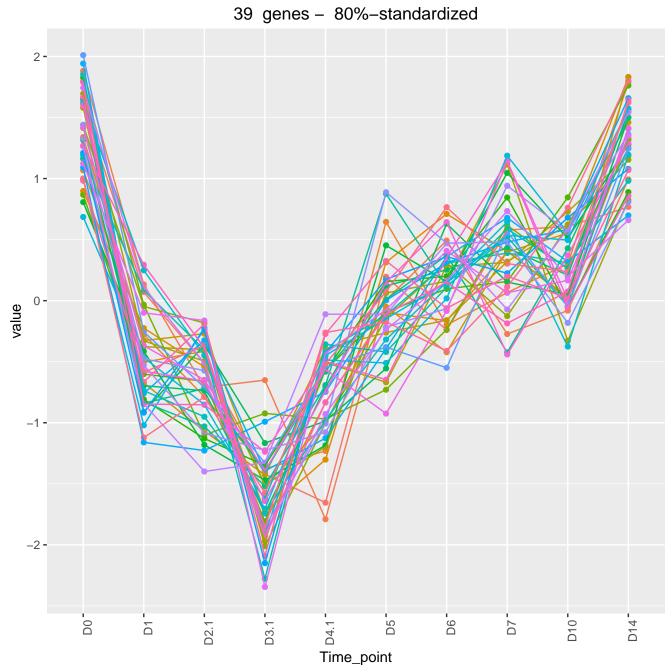
42 genes - 80%-standardized 1 -0 value **-1-**-2 **-**\_0Q \_ \_ D5 -D3.1 \_9Q D7<sup>-</sup> Lime\_boint

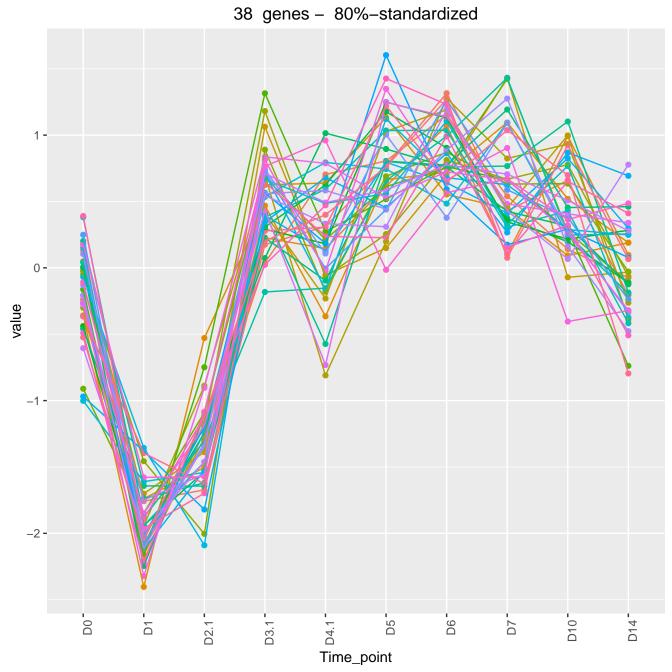
42 genes - 80%-standardized 1 -0 value \_1 **-**-2 **-**\_ 00 D5 -<u>\_</u>10 D3.1 \_9Q D7 Lime\_boint

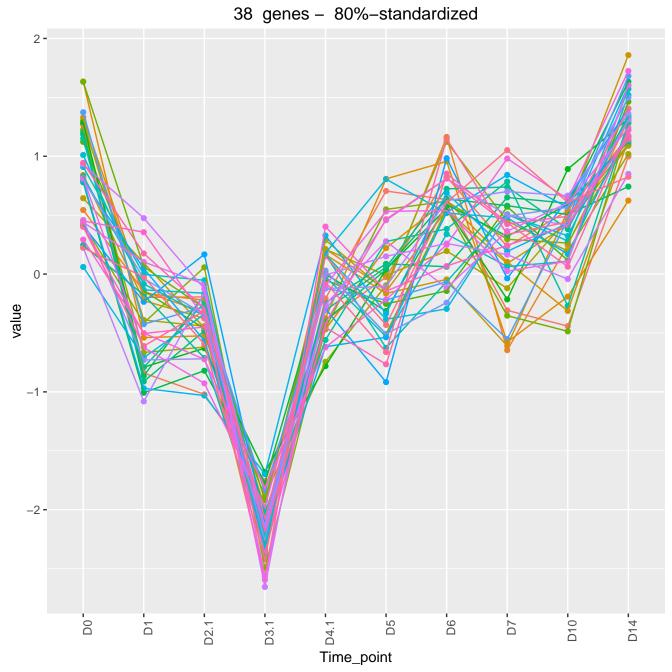


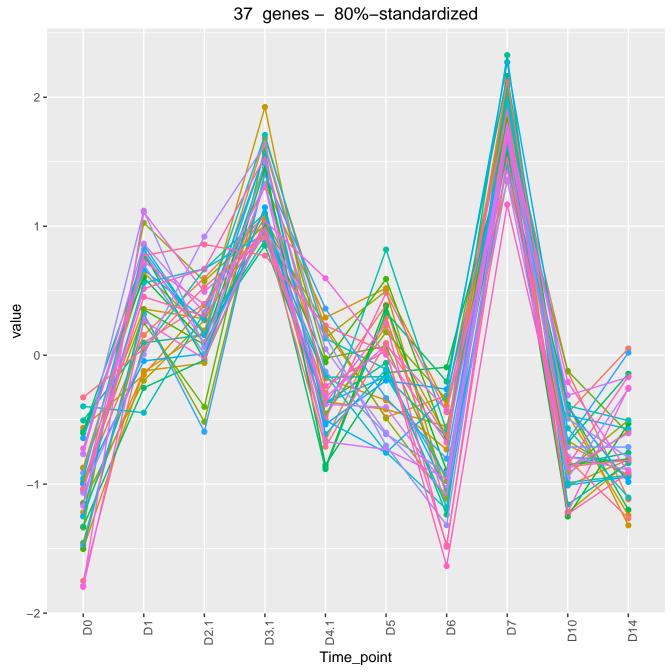


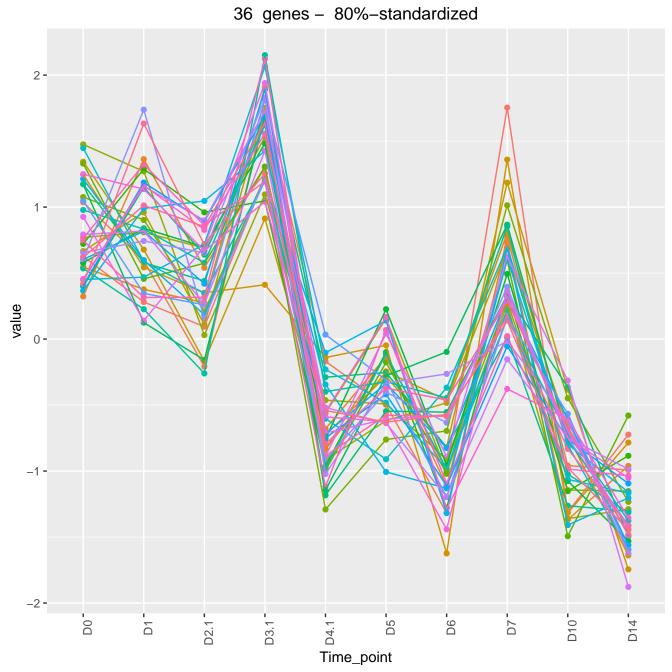


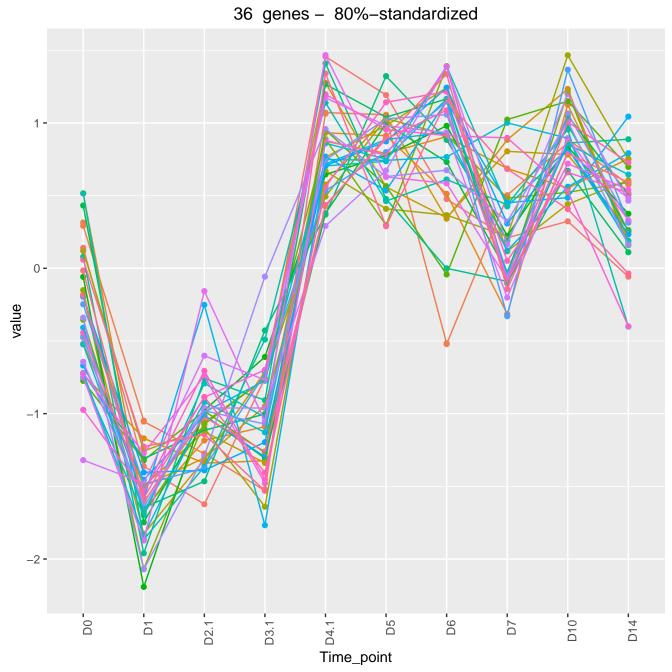




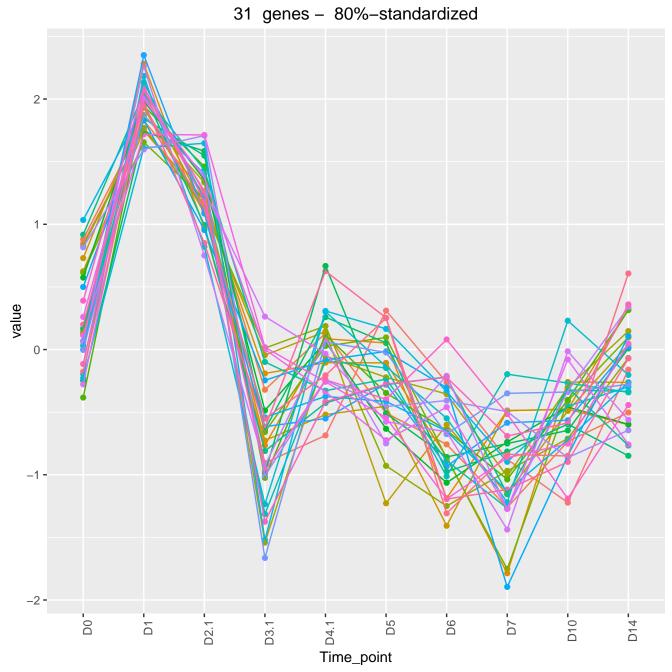


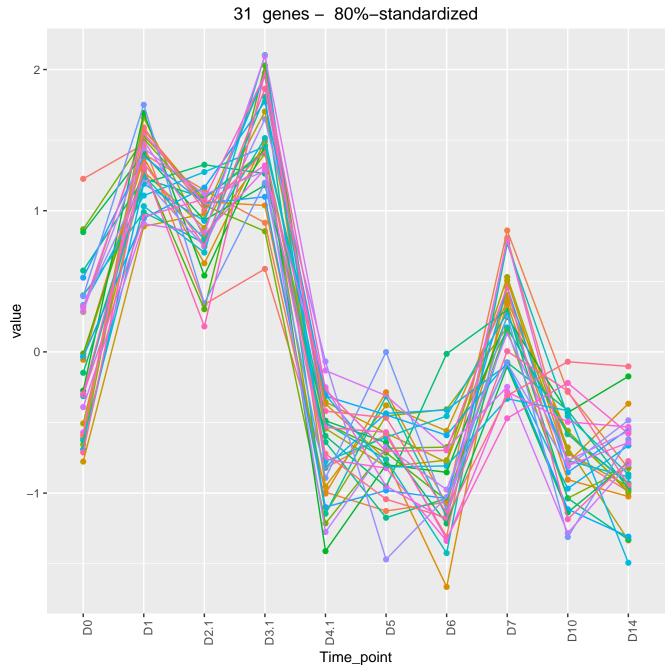


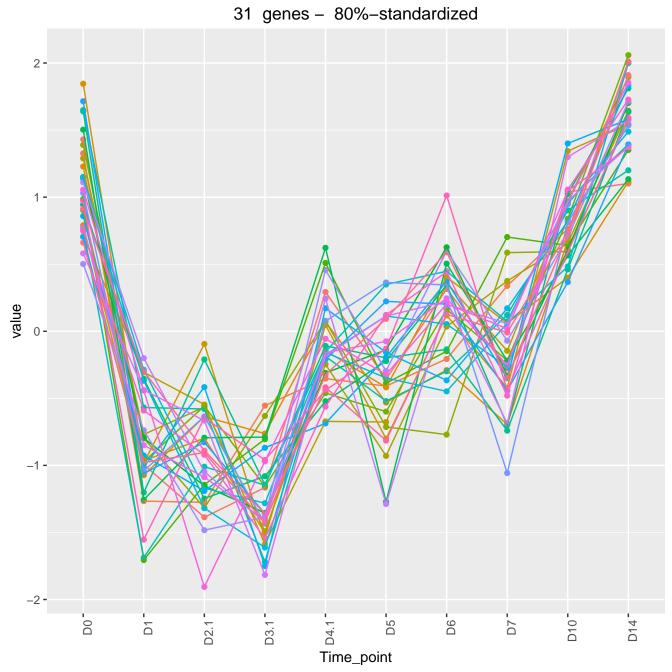


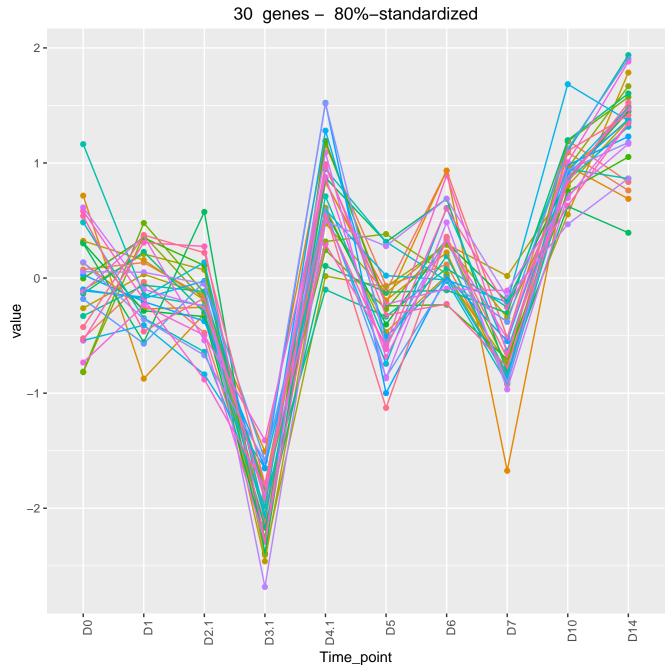


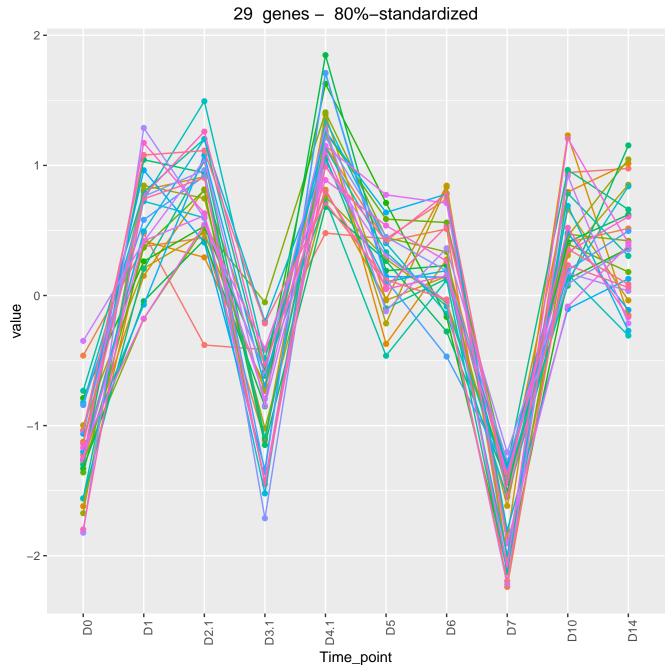
34 genes - 80%-standardized 2 -1 value 0 -\_1 **-**-2 **-**\_0Q D3.1 <u>\_</u>10 \_9Q D7<sup>-</sup> Lime\_boint





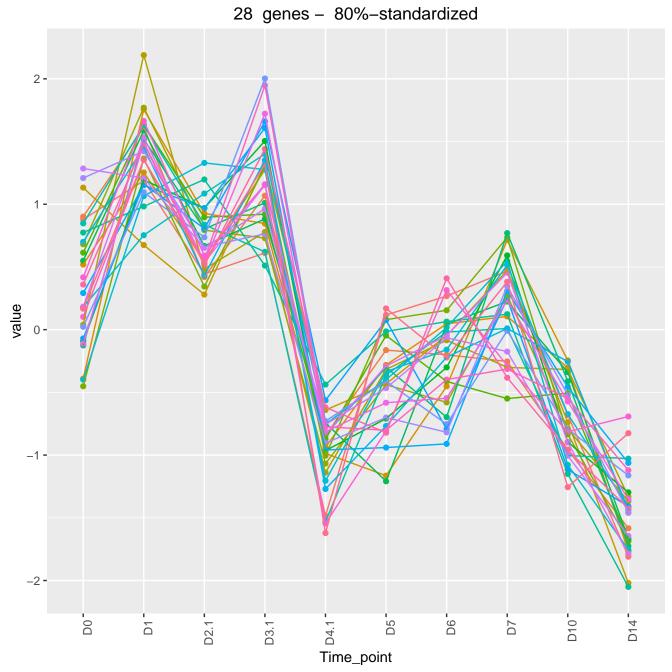


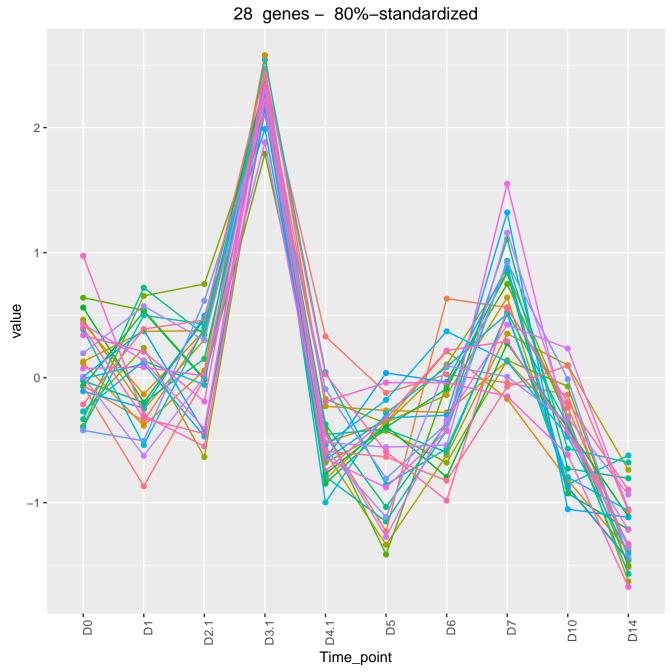




29 genes - 80%-standardized 2 -1 value 0 -\_1 **-**-2 **-**\_0Q \_ \_ D3.1 D5 -\_9Q D7\_ Lime\_boint

28 genes - 80%-standardized 2 -1 value 0 --1 **-**-2 **-**Time\_point D3.1 \_9Q D7\_





28 genes - 80%-standardized 2 -1 value 0 --1 **-**<u>-</u> \_ D2\_ D3.1 \_9Q D7 Lime\_boint

27 genes - 80%-standardized 1 -0 value -1 **-**-2 **-**\_ 00 D3.1 D5 -<u>\_</u>10 \_9Q D7<sup>-</sup> Lime\_boint

27 genes - 80%-standardized 2 -1 value 0 -**-1-**-2 **-**\_0Q \_ \_ D5 -D3.1 \_9Q D7\_ Lime\_boint

27 genes - 80%-standardized 2 -1 value 0 -**-1-**-2 **-**

D5 -

Lime\_boint

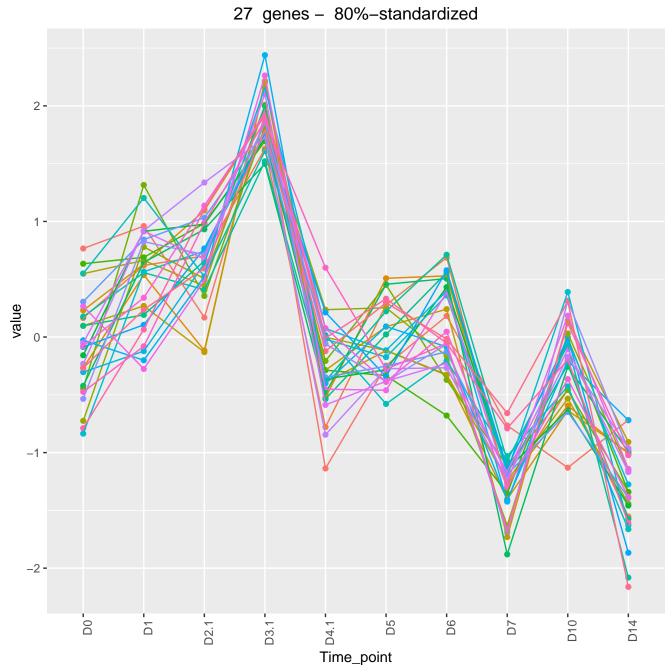
\_9Q

D7\_

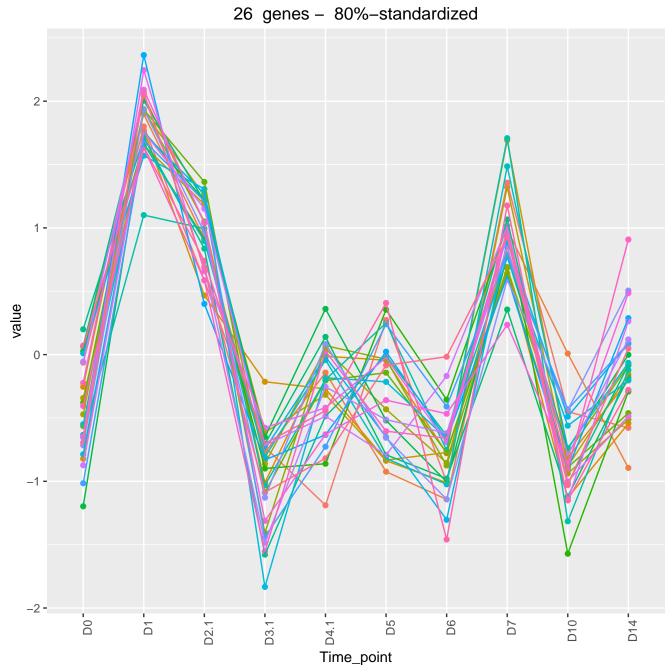
\_ 00

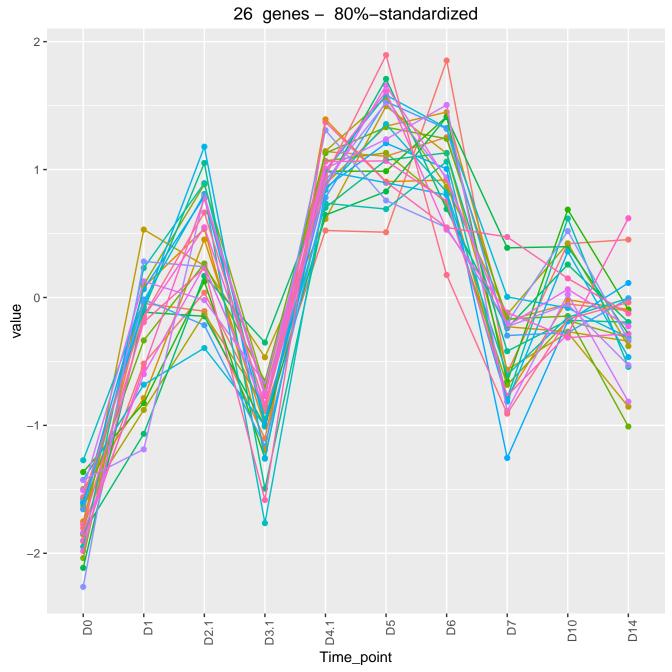
<u>\_</u>10

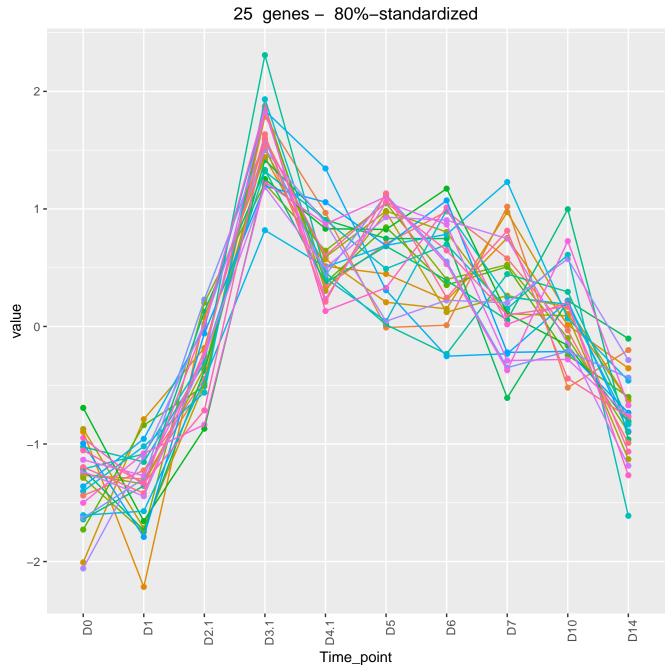
D3.1



27 genes - 80%-standardized 2 -1 value 0 -\_1 **-**\_ 00 \_ \_ D5 -D3.1 \_9Q D10 Lime\_boint D7\_







25 genes - 80%-standardized 1 -0 value **-1** --2 **-**\_ 00 D5 -D3.1 \_9Q D7\_ Lime\_boint

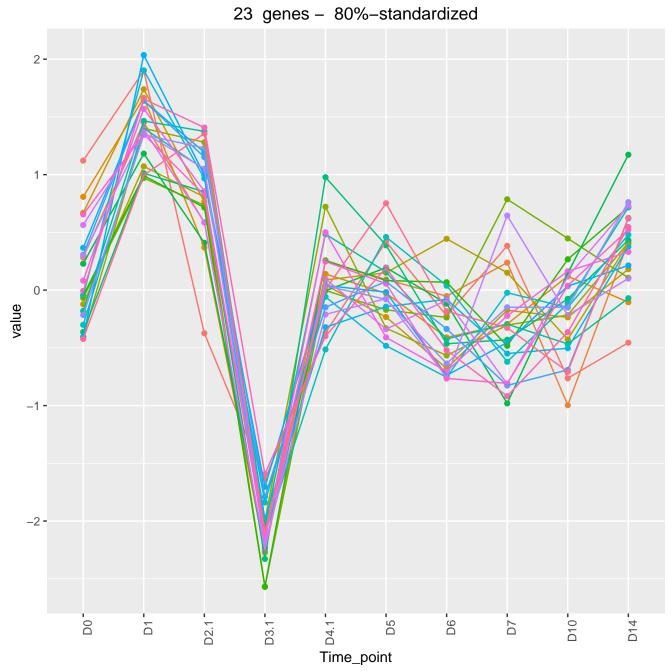
25 genes - 80%-standardized 2 -1 value - 0 -1 --2 **-**\_ 00 D5 -<u>\_</u>10 D3.1 \_9Q D7\_ Lime\_boint

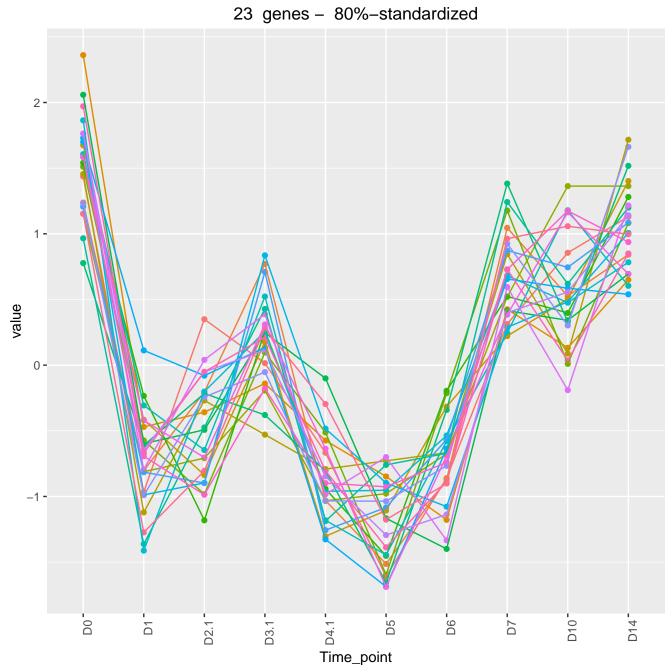
24 genes - 80%-standardized 2 value o-**-1-**-2 **-**\_ 00 \_ \_ D5 -D3.1 \_9Q D7\_ Lime\_boint

24 genes - 80%-standardized 1 -0 value -1 **-**-2 **-**\_ 00 <u>-</u>10 D3.1 D5 -\_9Q Lime\_boint

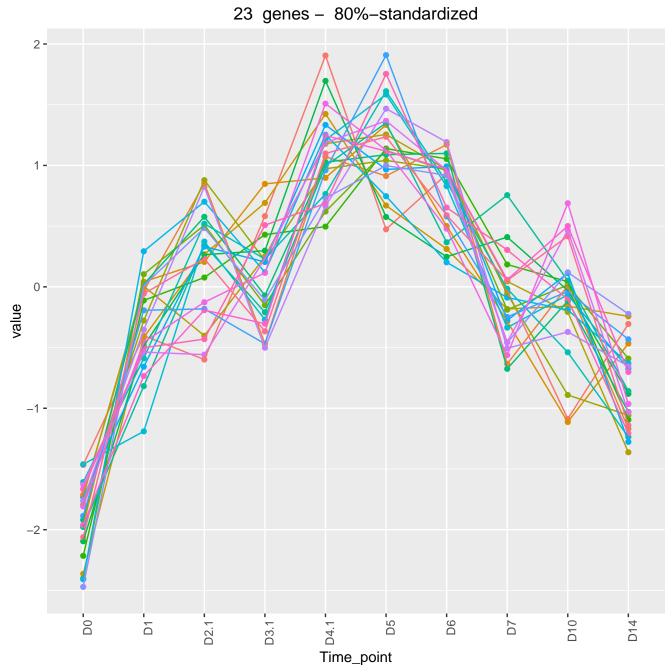
24 genes - 80%-standardized 2 -1 value -1 **-**-2 **-**\_ 00 \_ \_ D5 -D3.1 \_9Q D7\_ Lime\_boint

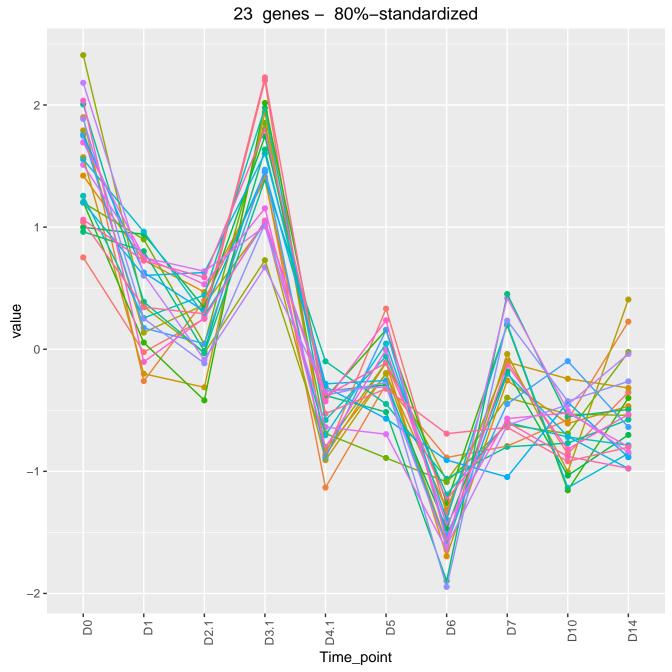
24 genes - 80%-standardized 2 -1 value o--1 **-**-2 **-**\_ 00 <u>\_</u>10 D3.1 \_9Q D7 Lime\_boint

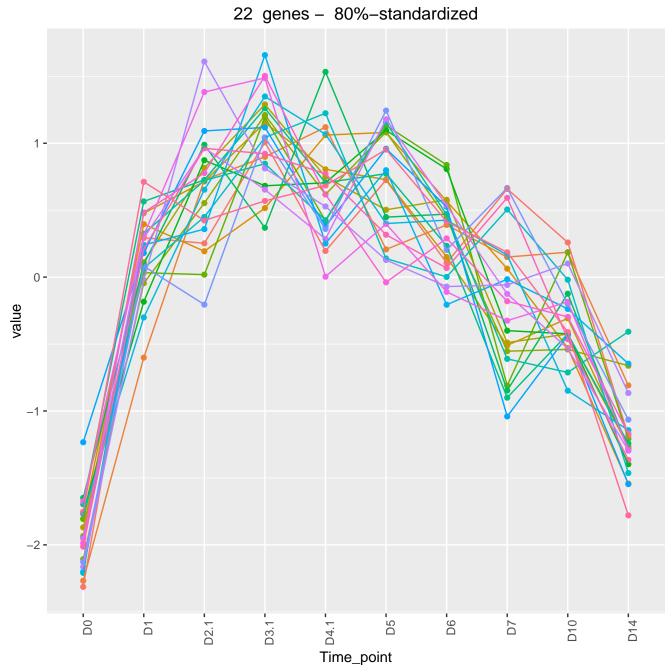


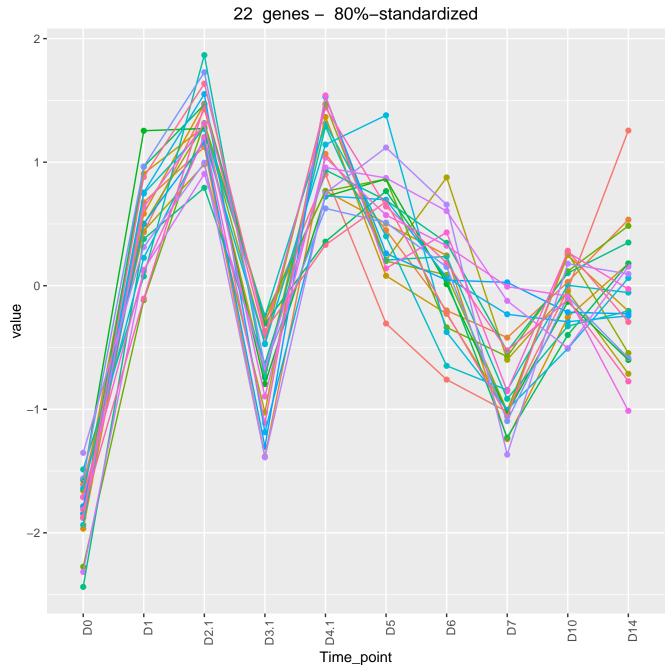


23 genes - 80%-standardized 2 -1 value 0 --2 **-**\_ 0Q D5 -<u>-</u> D3.1 \_9Q Lime\_boint

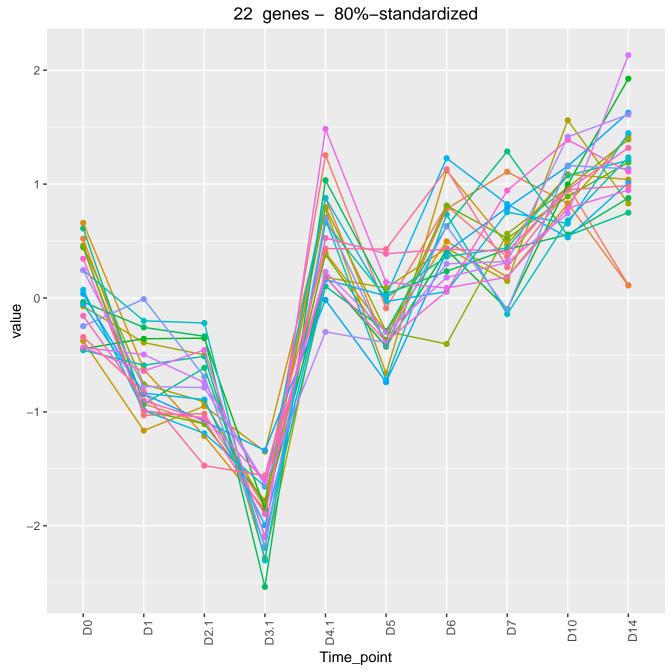




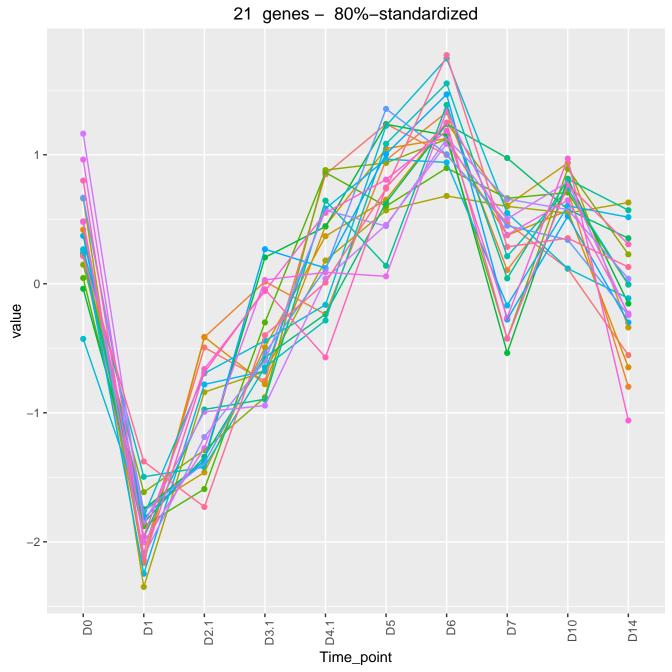


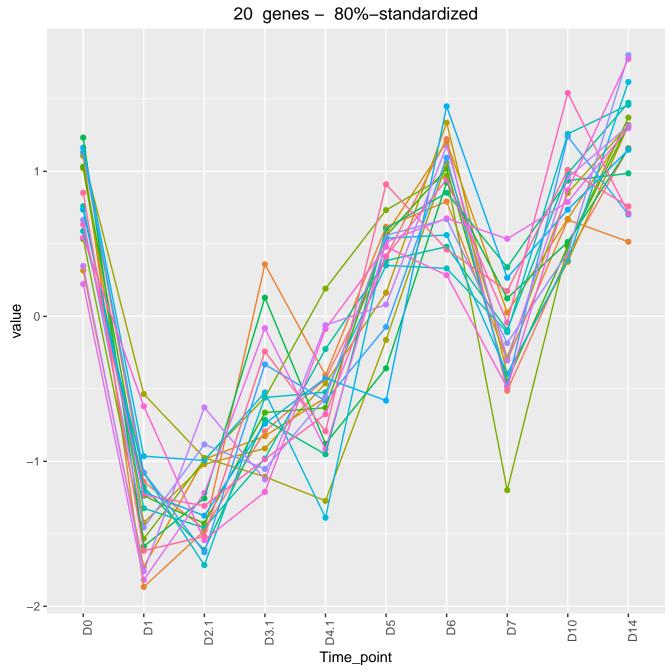


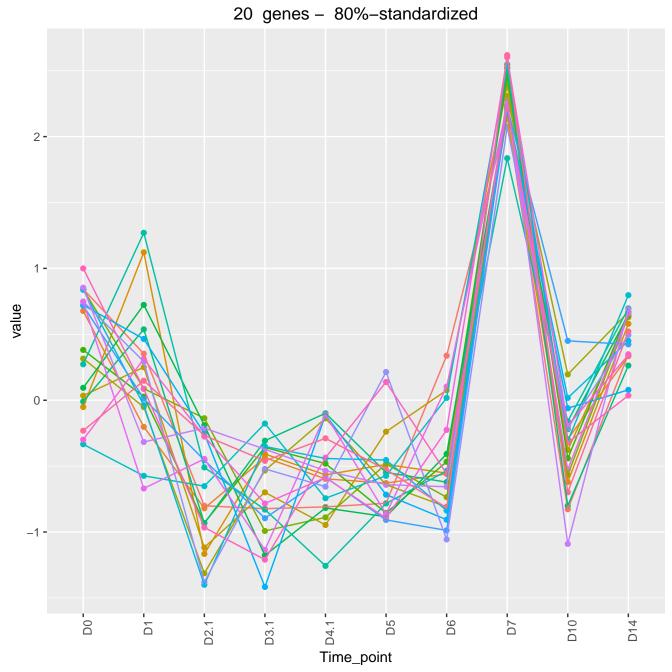
22 genes - 80%-standardized 2 -1 value 0 --1 -\_ 00 D5 -<u>\_</u>10 D3.1 \_9Q D7 Lime\_boint

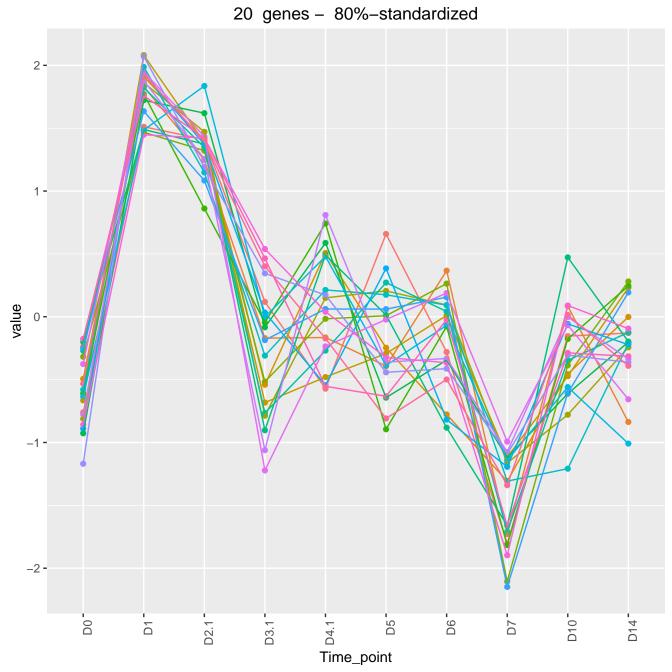


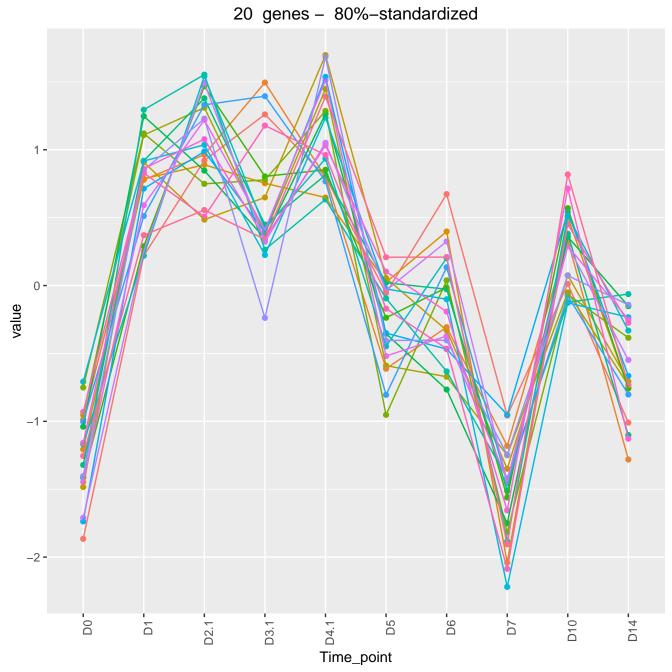
22 genes - 80%-standardized 2 -1 value 0 -**-1** -\_ 00 D5 -D3.1 \_9Q D7\_ Lime\_boint

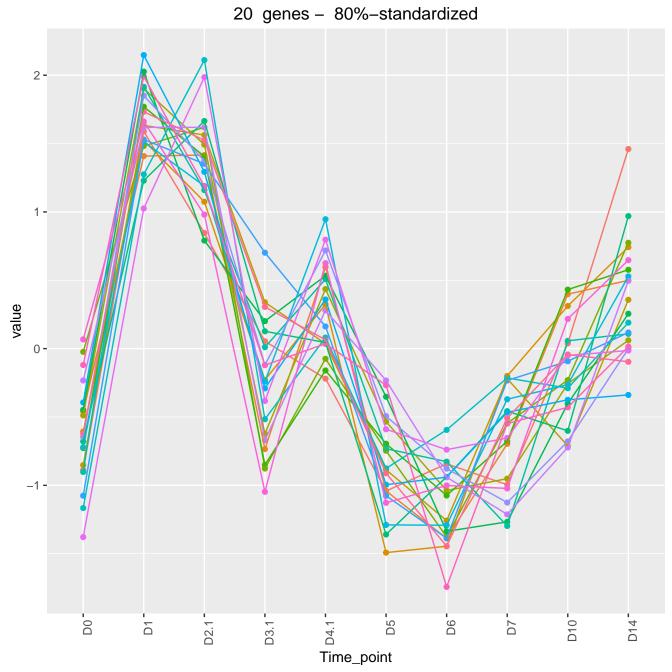


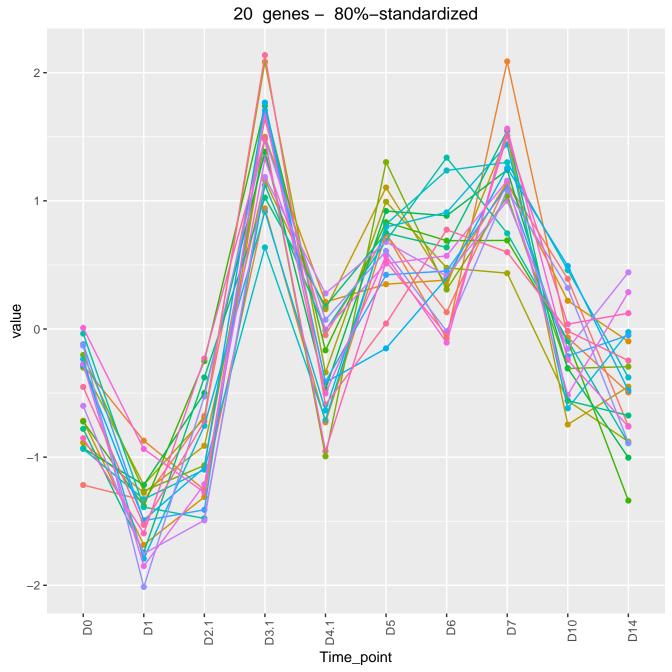




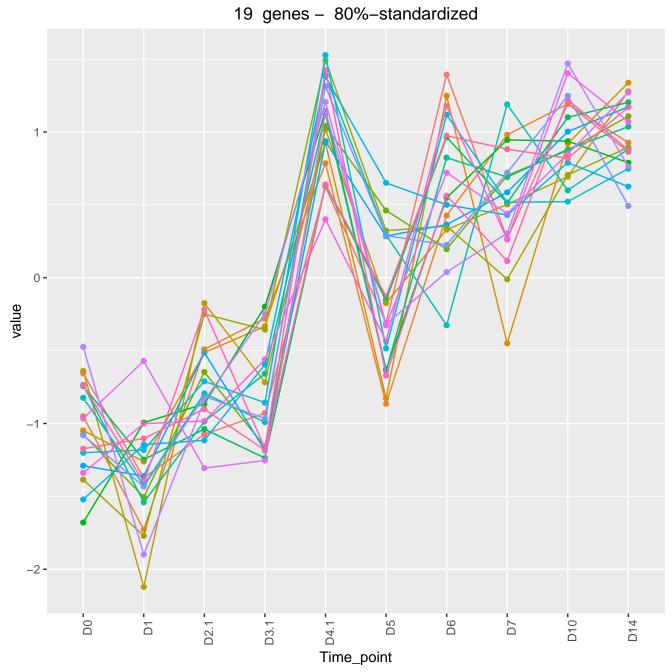


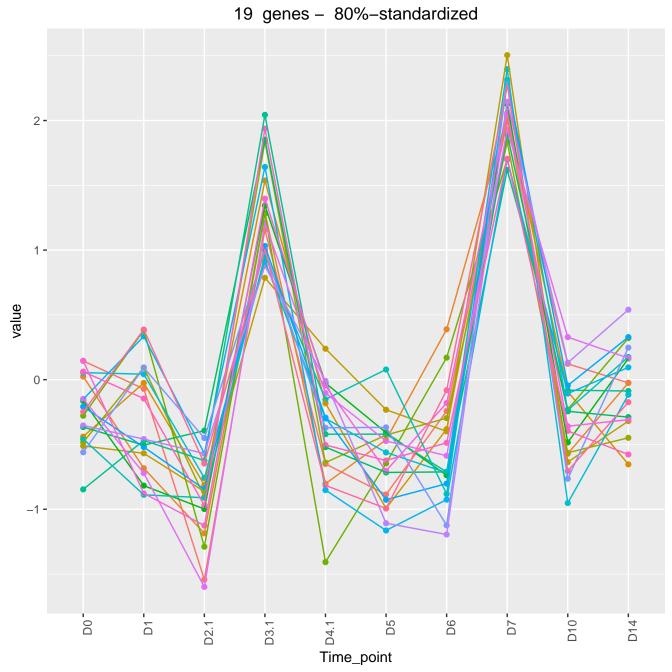


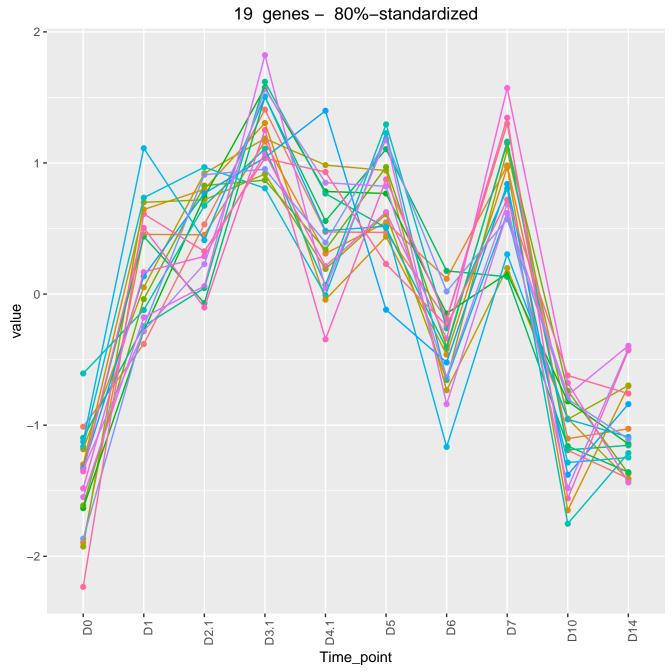


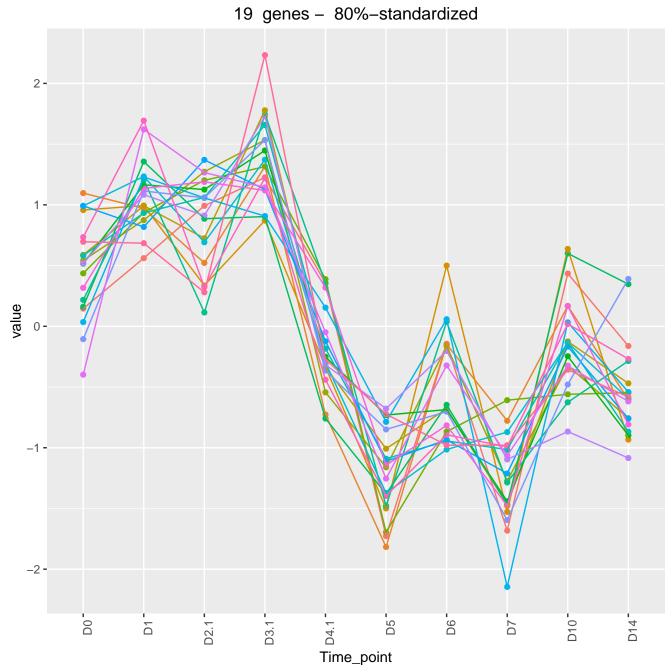


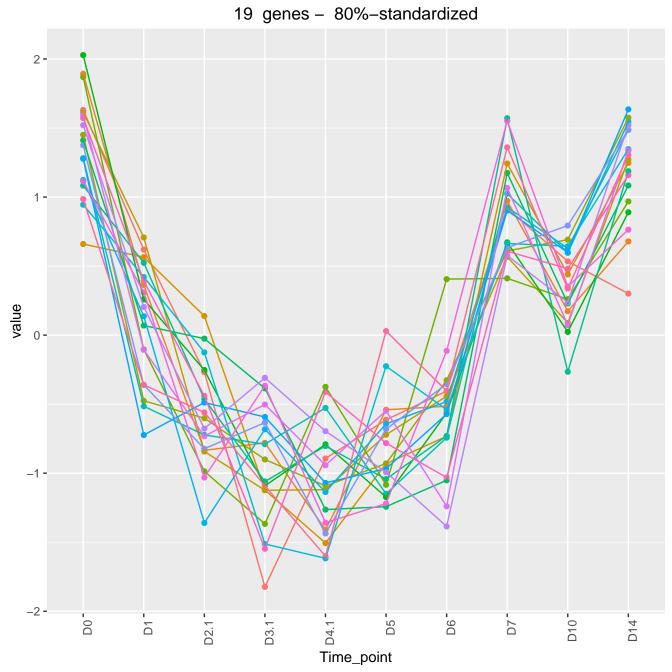
19 genes - 80%-standardized 2 -1 value **-1-**-2 **-**\_ 00 D5 -D3.1 \_9Q D7\_ Lime\_boint











18 genes - 80%-standardized 2 -1 value 0 -**-1-**-2 **-**\_0Q Time\_point D1 D3.1 \_9Q D7<sup>-</sup>

