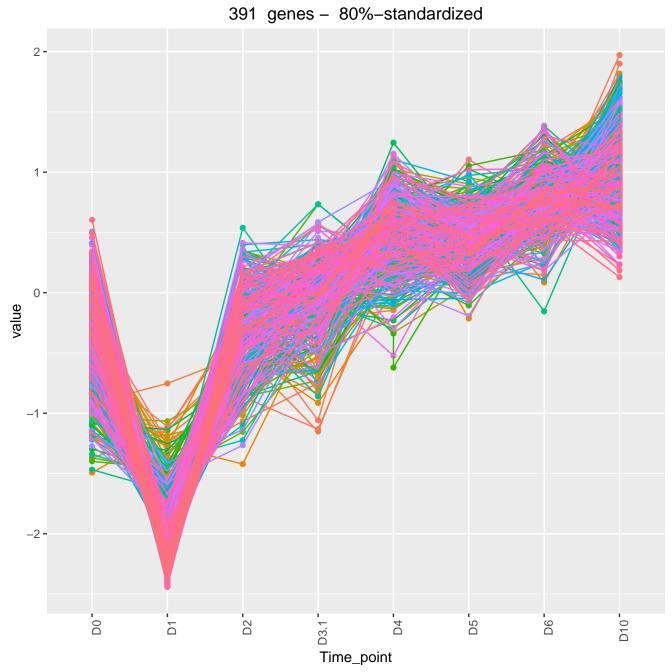
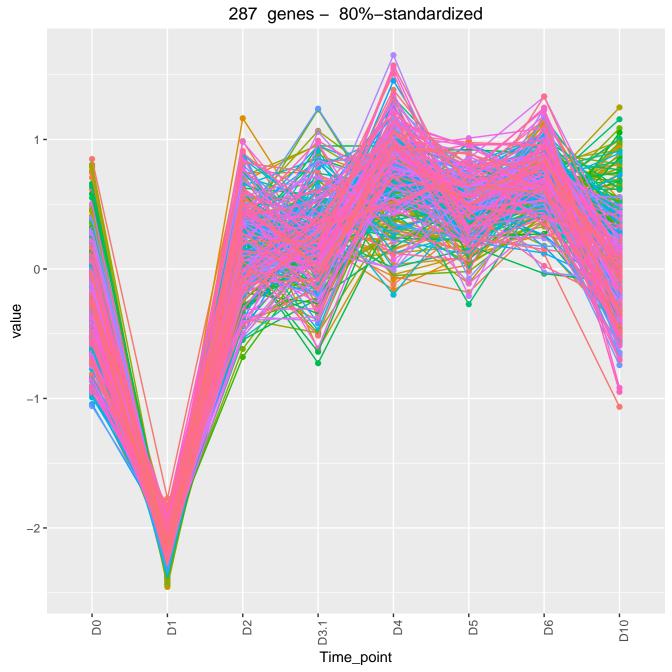
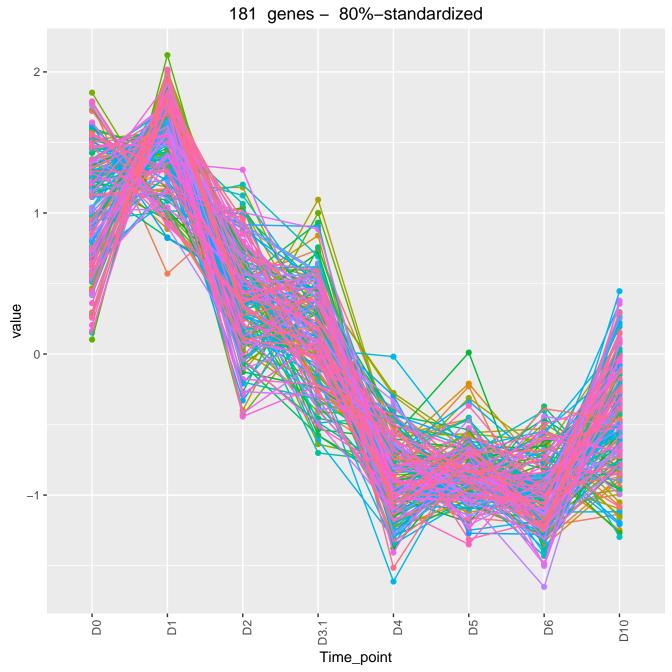
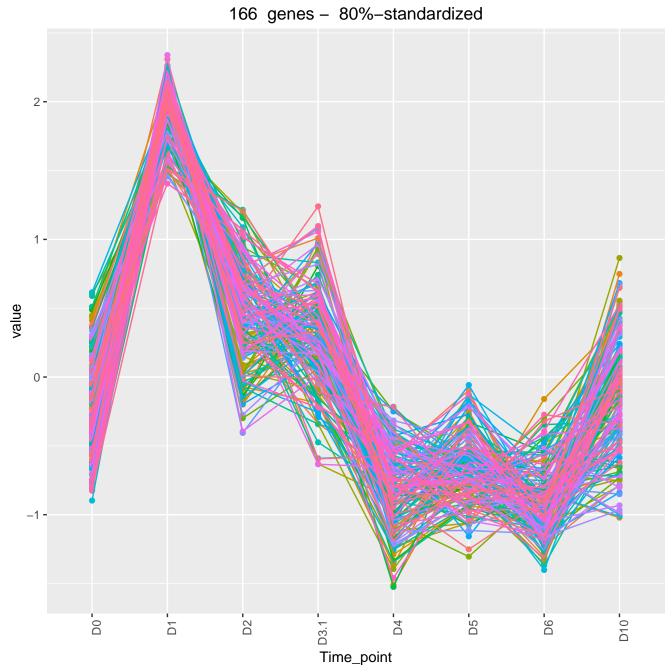


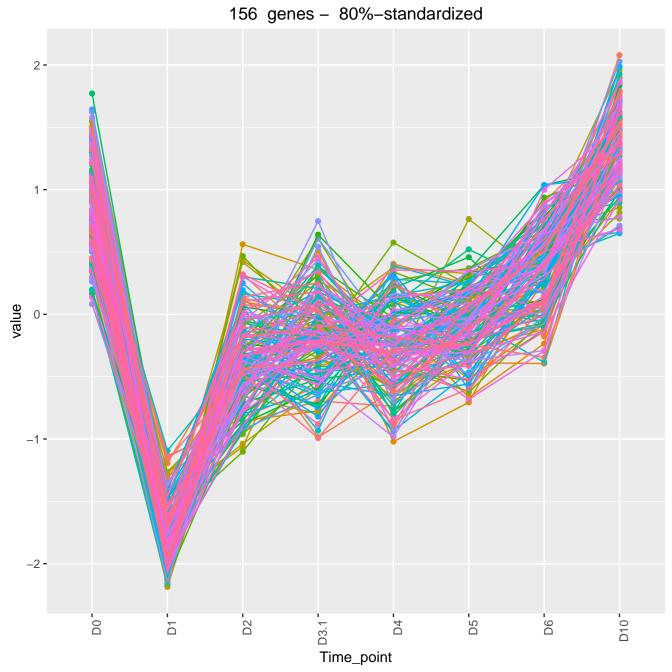
538 genes - 80%-standardized 2 -1 value 0 --1 **-**Time\_point \_ 00  $\overline{\Box}$ 

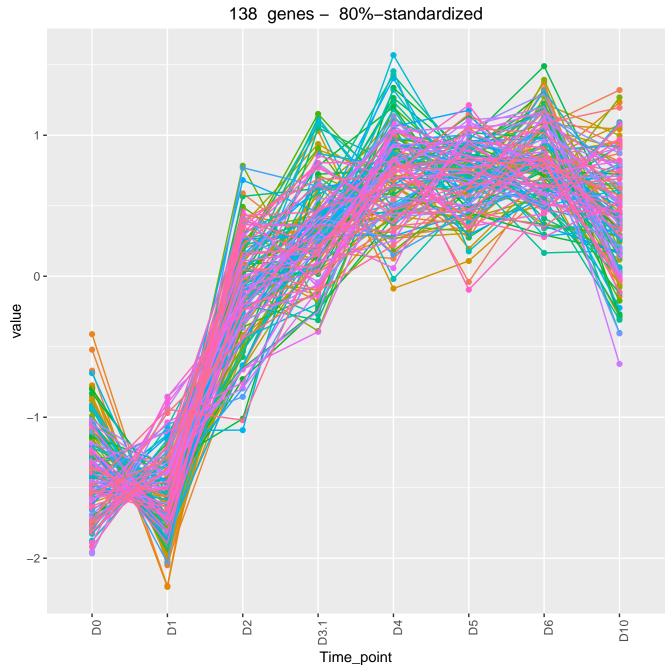


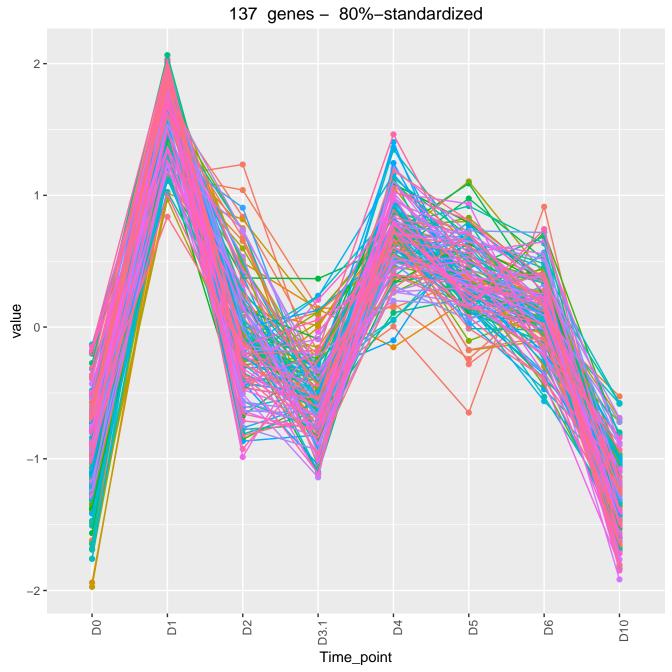


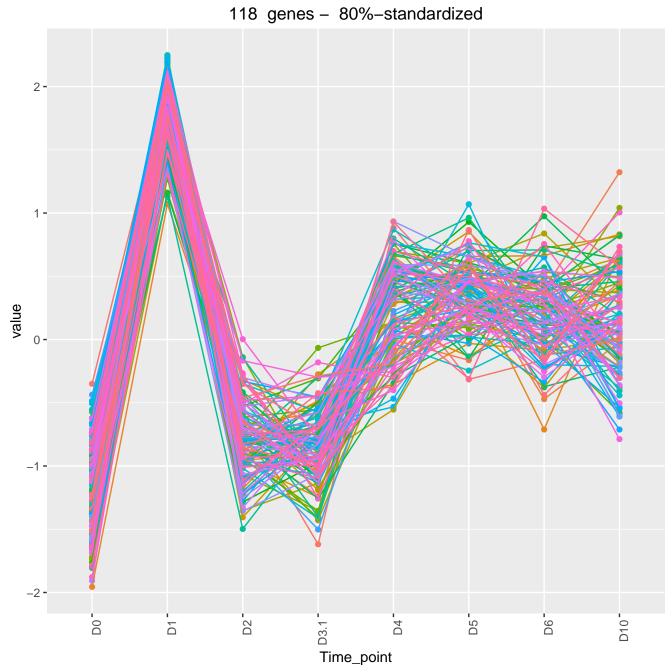


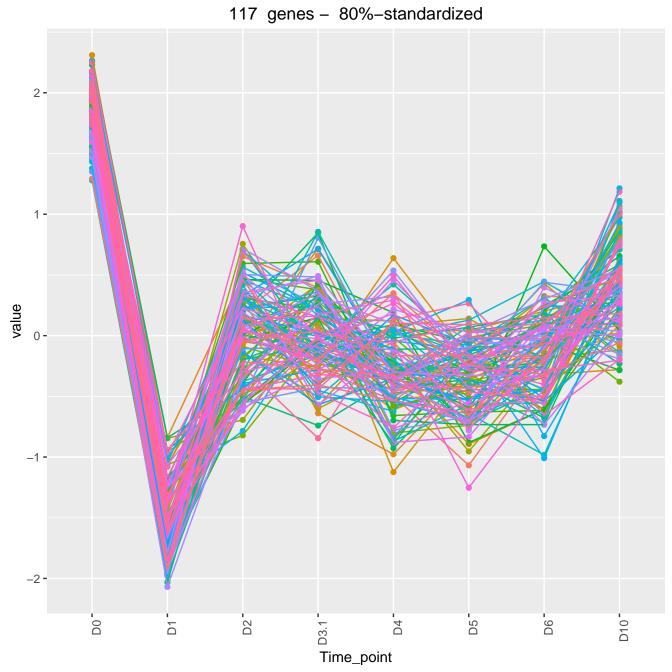


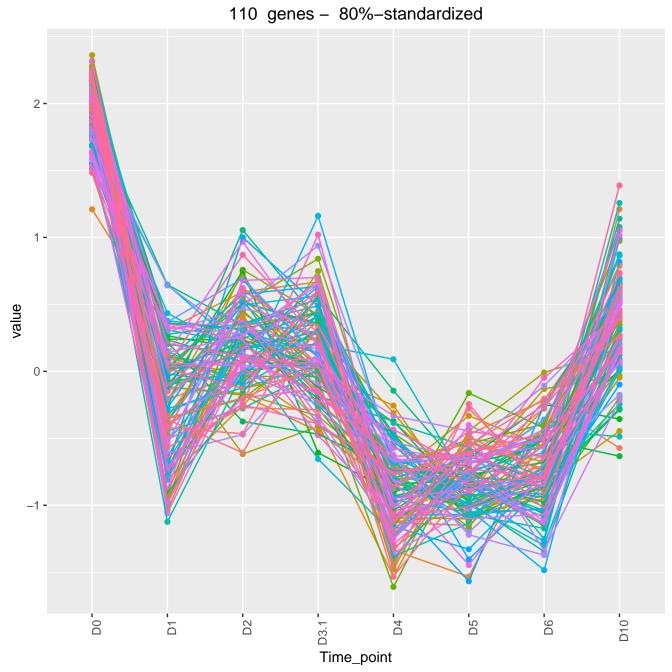


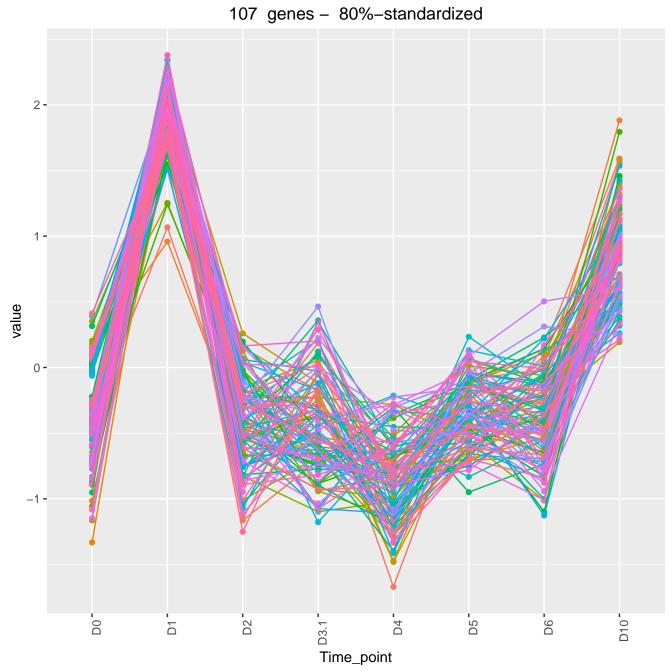


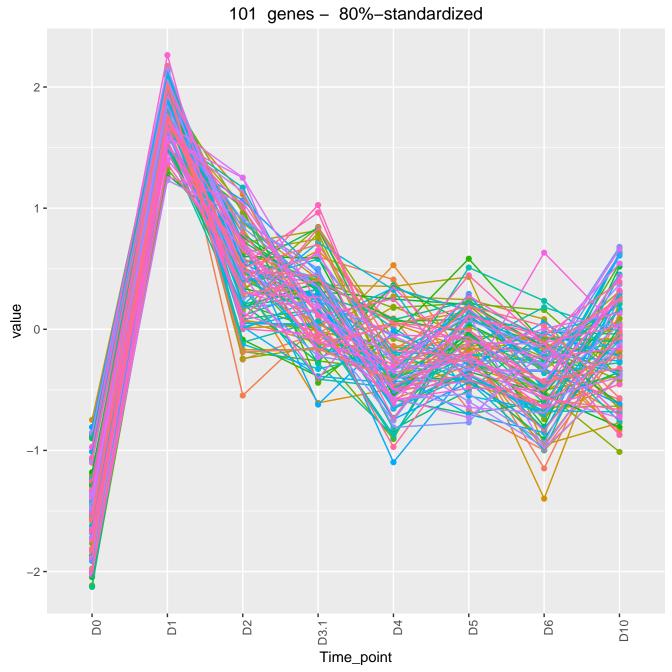


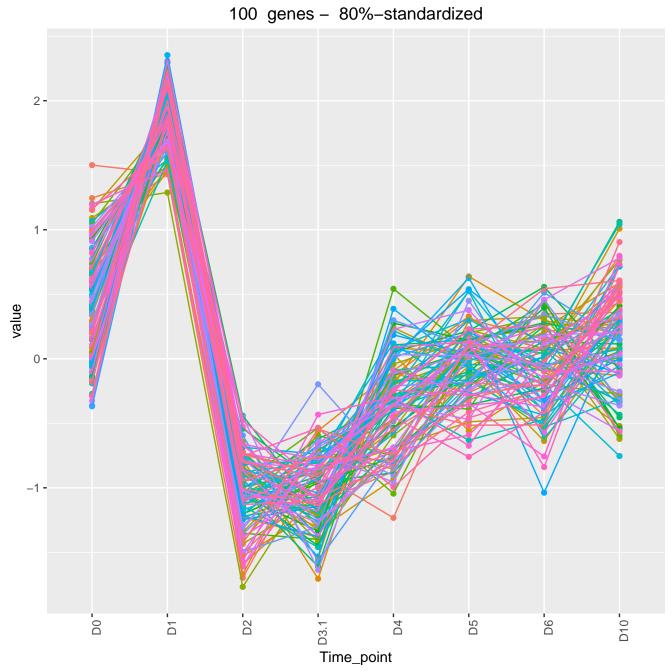


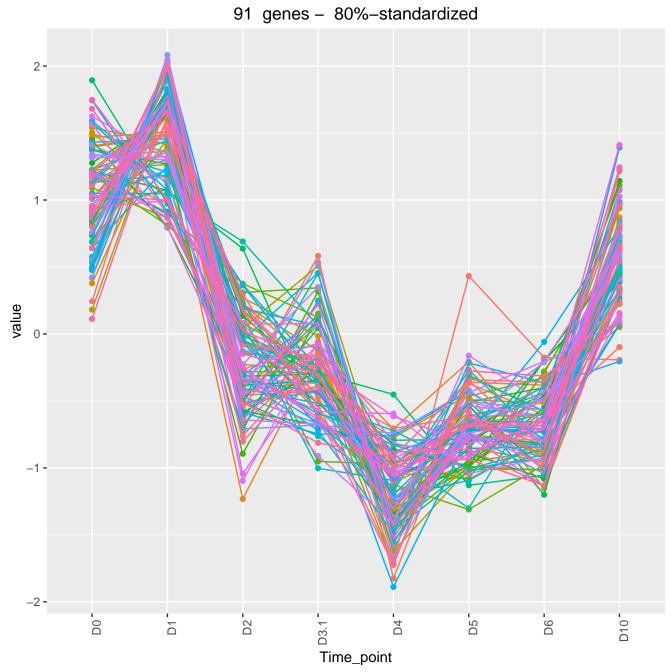


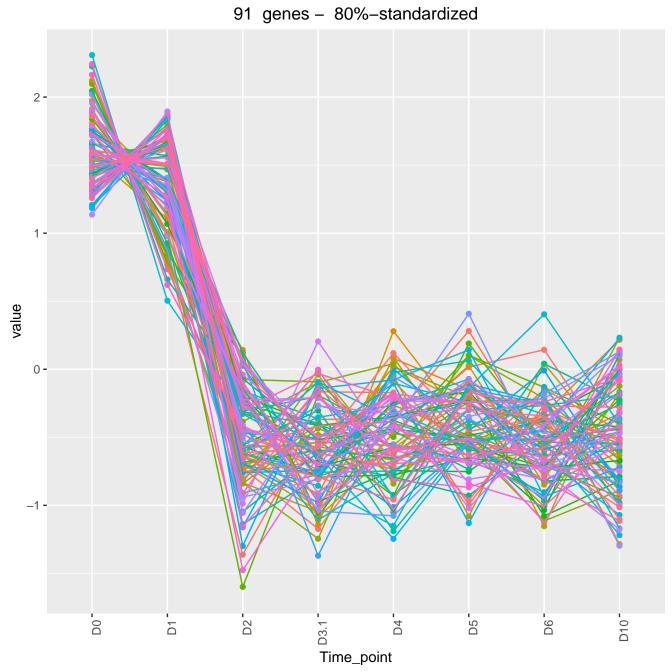


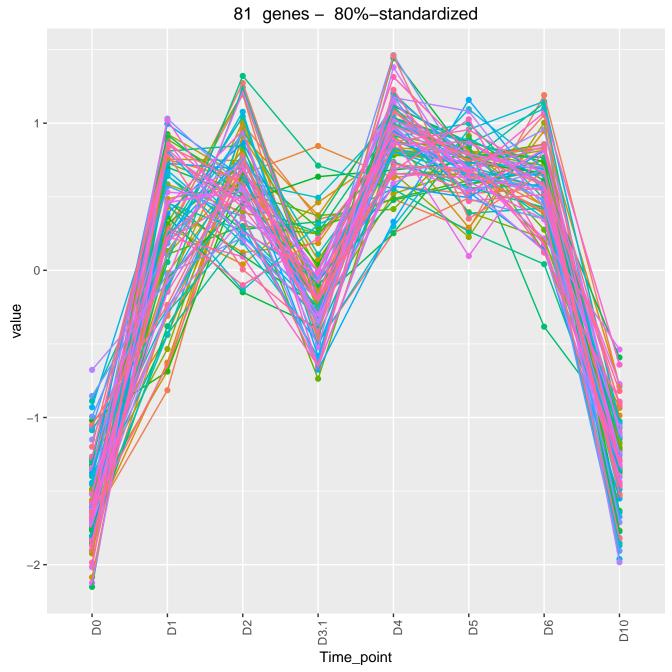


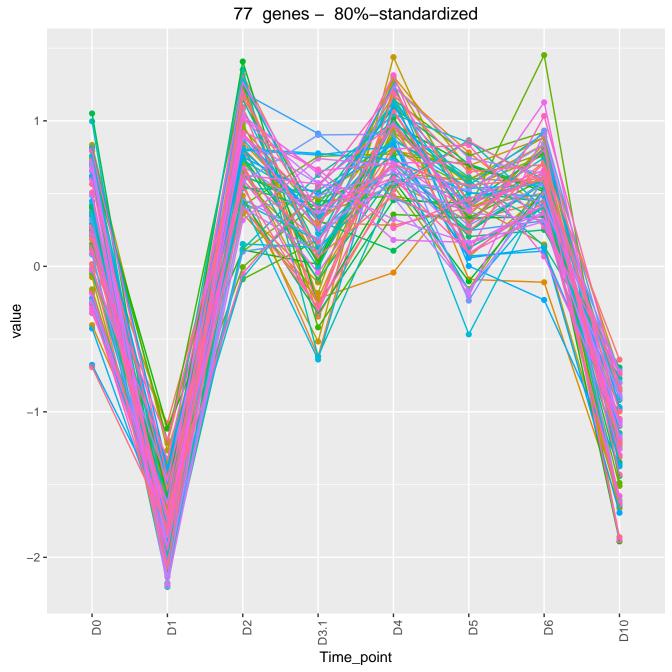


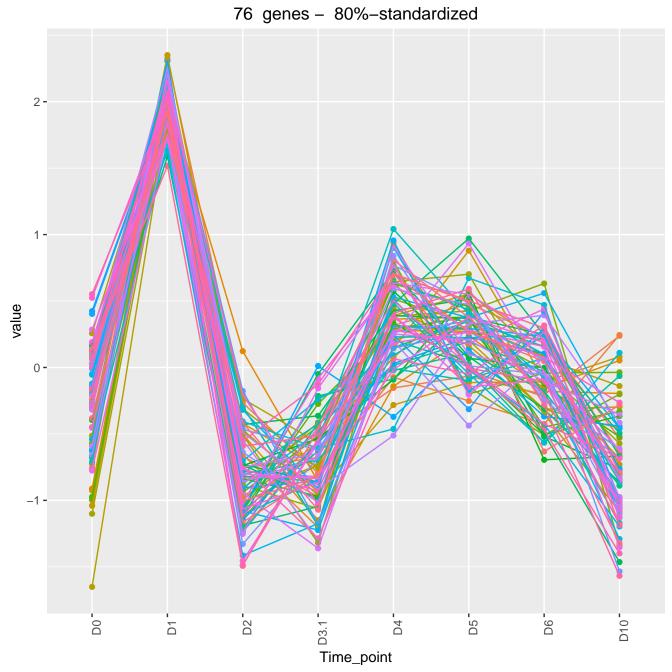




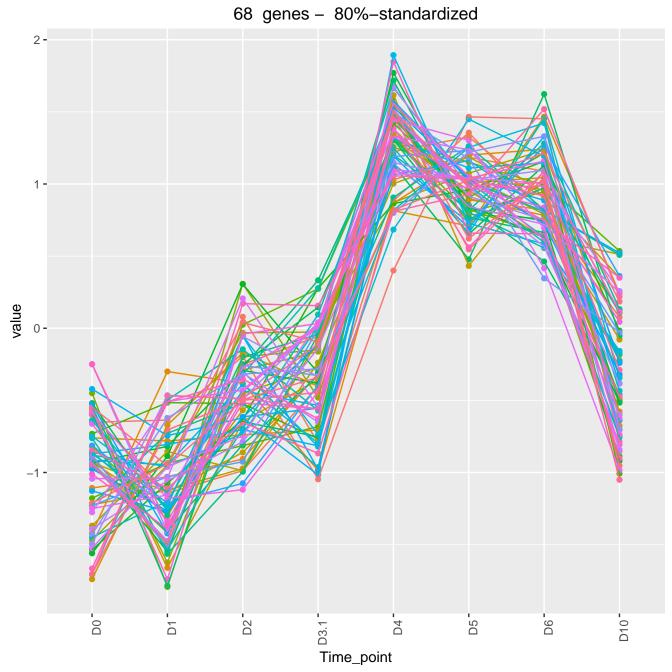




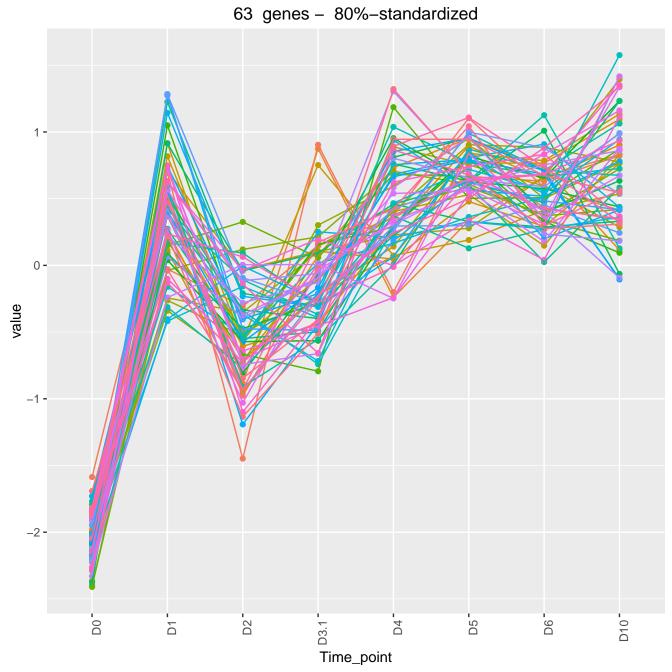


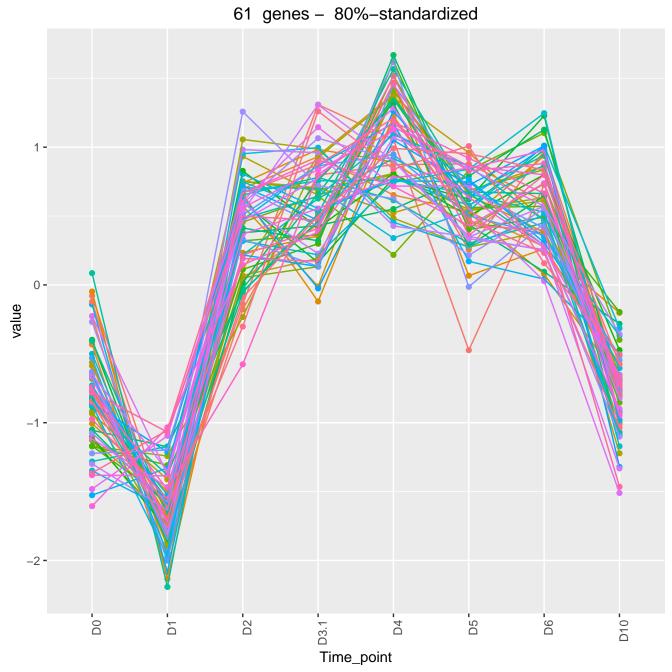


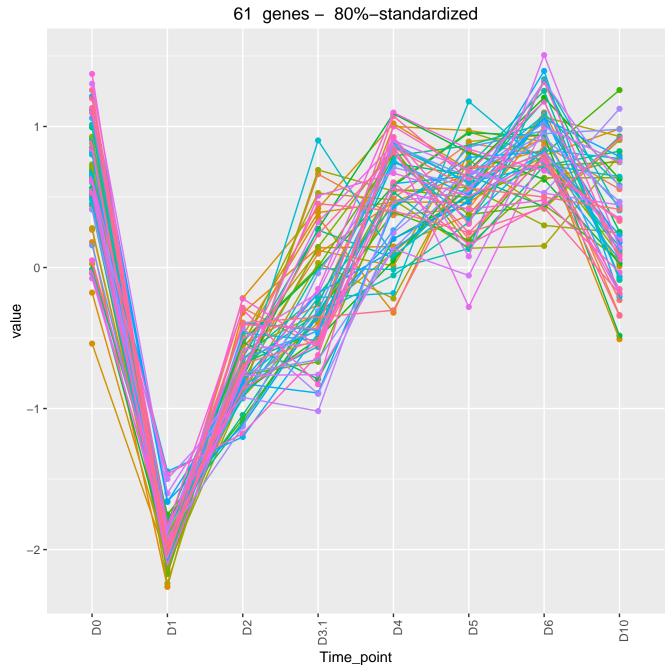
71 genes - 80%-standardized 2value -2 **-**Time\_point \_0Q .

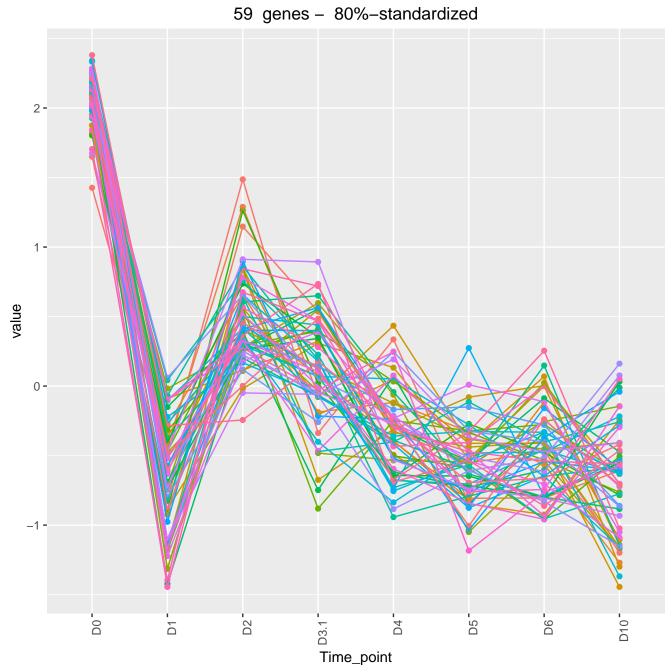


67 genes - 80%-standardized 1 value o -\_1 **-**-2 **-**\_0Q Time\_point D2 7

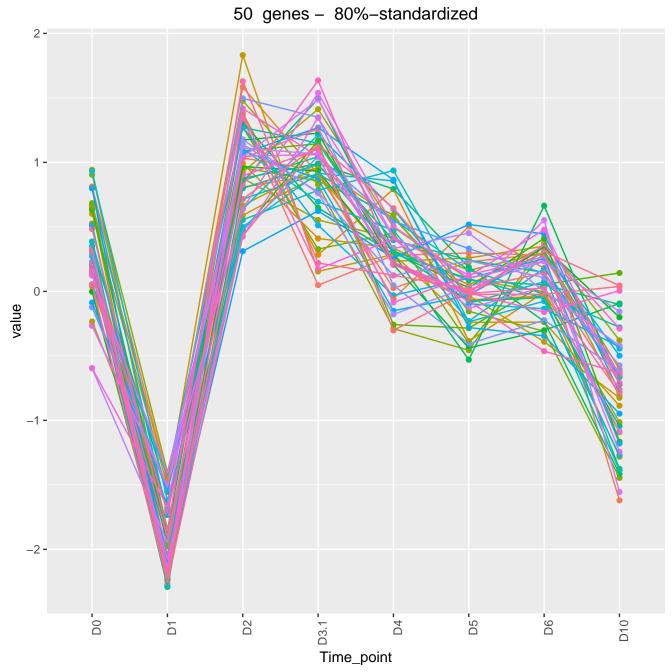


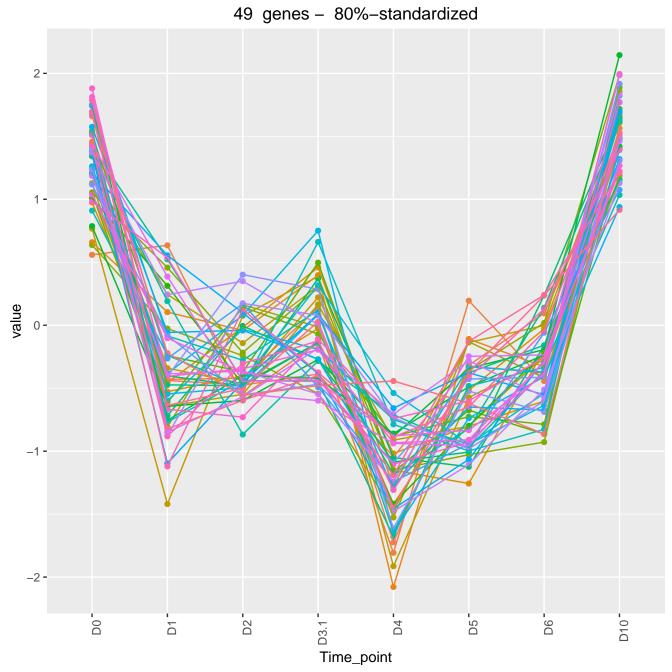


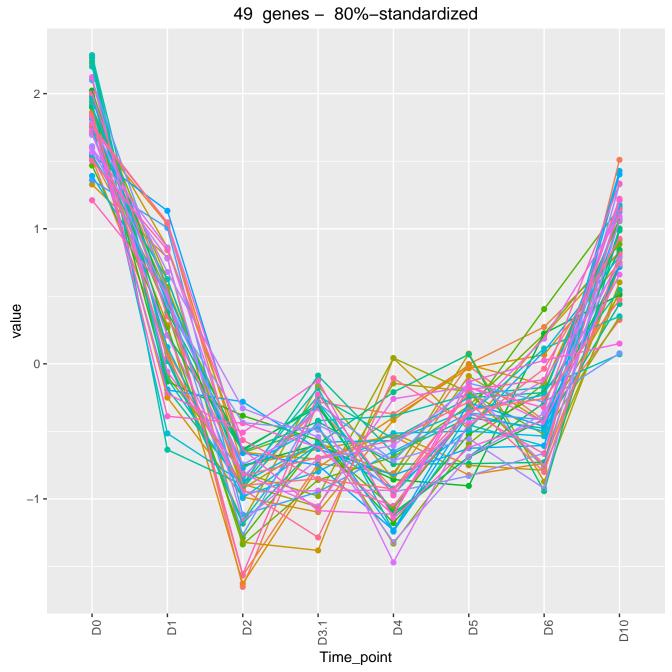


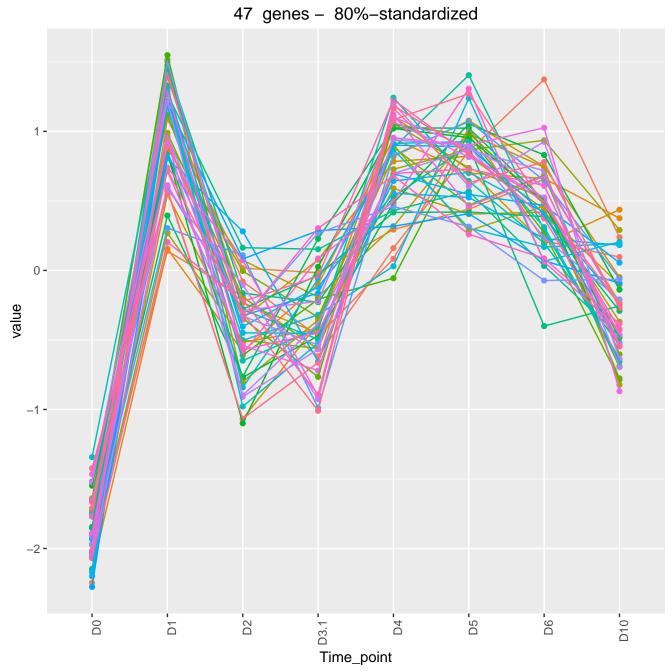


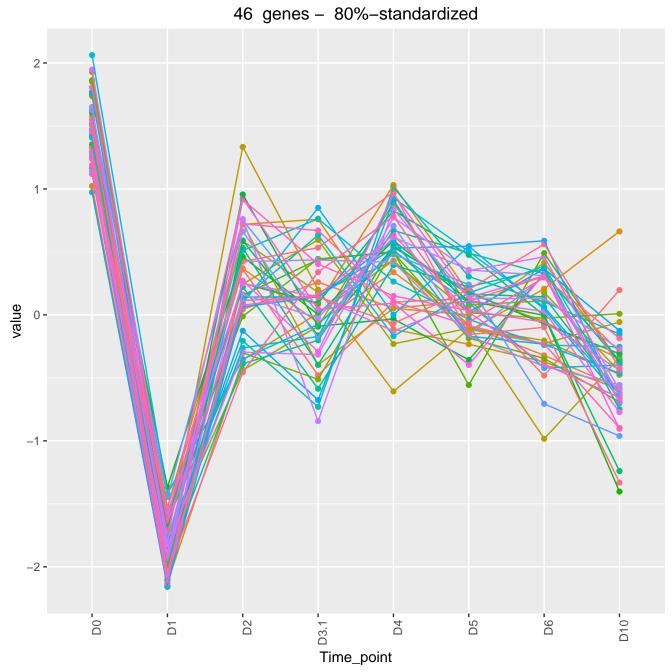
57 genes - 80%-standardized 2 -1 value **-1-**-2 **-**Time\_point D2 D5 7

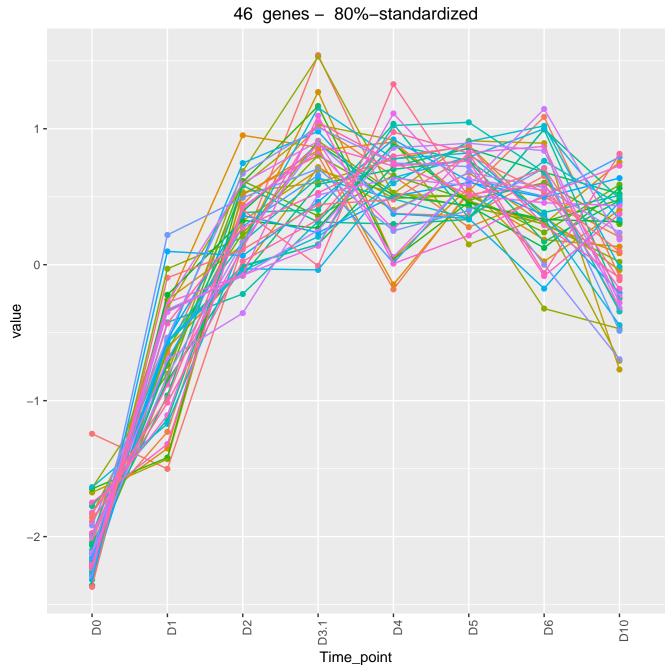


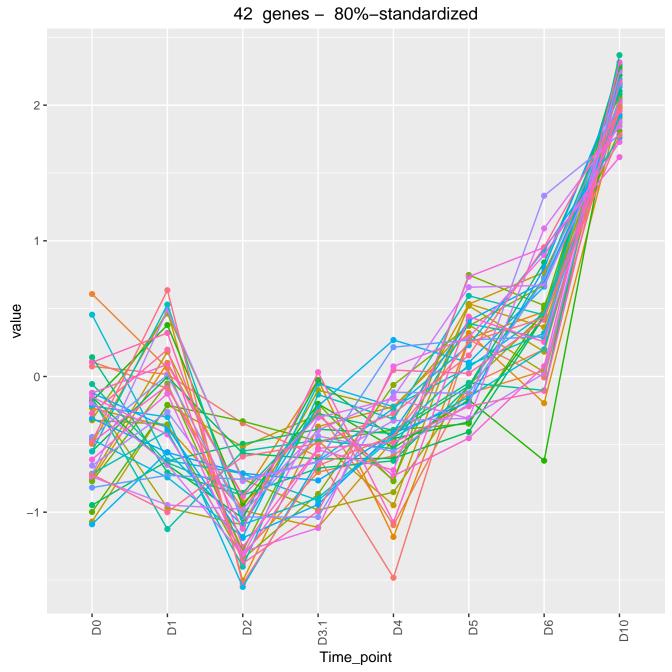


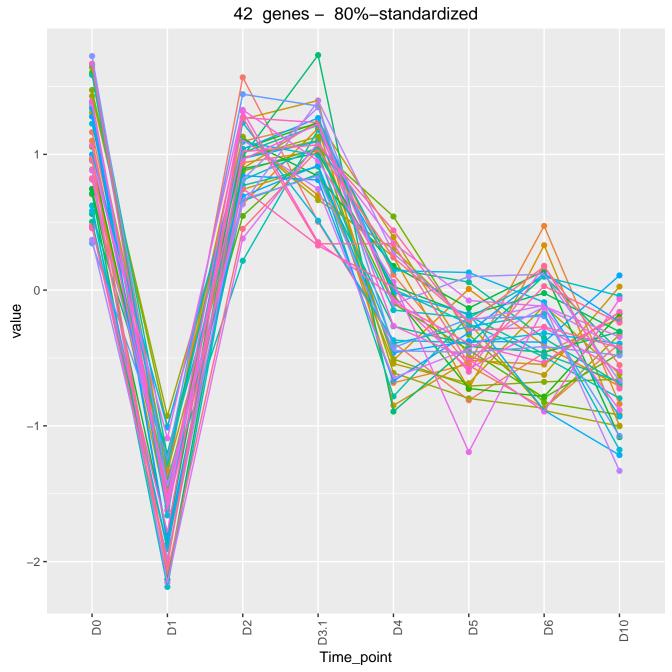


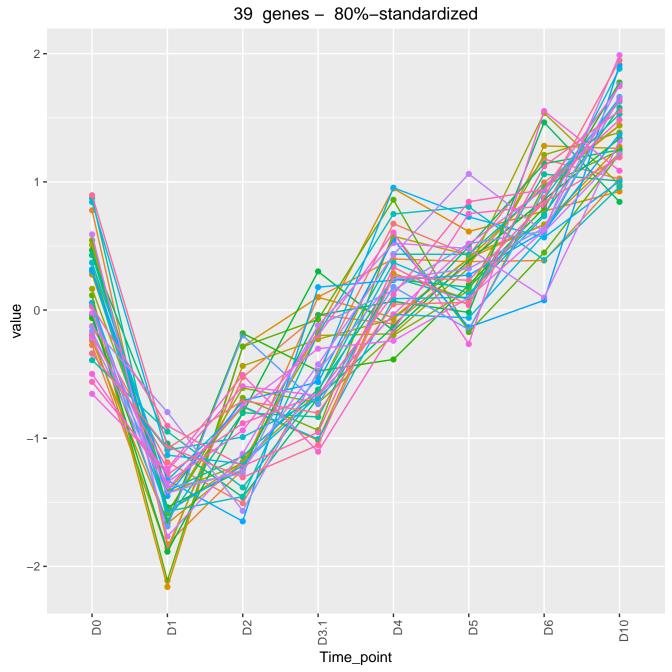


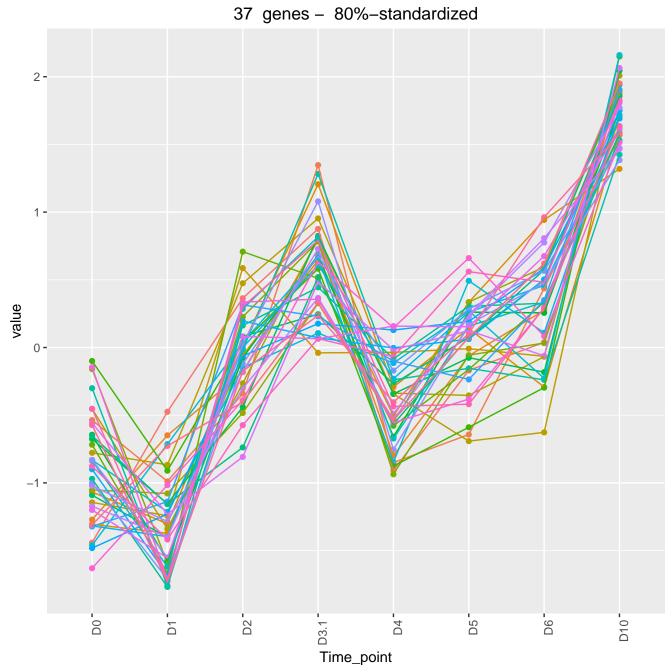


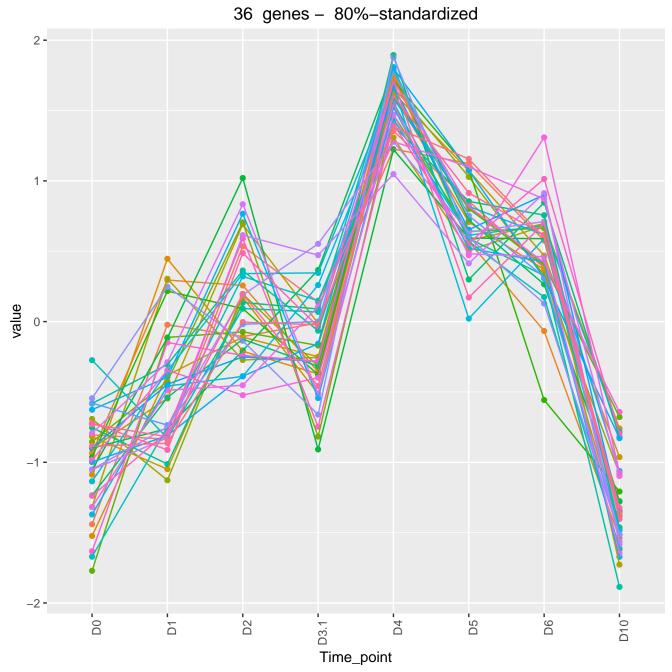


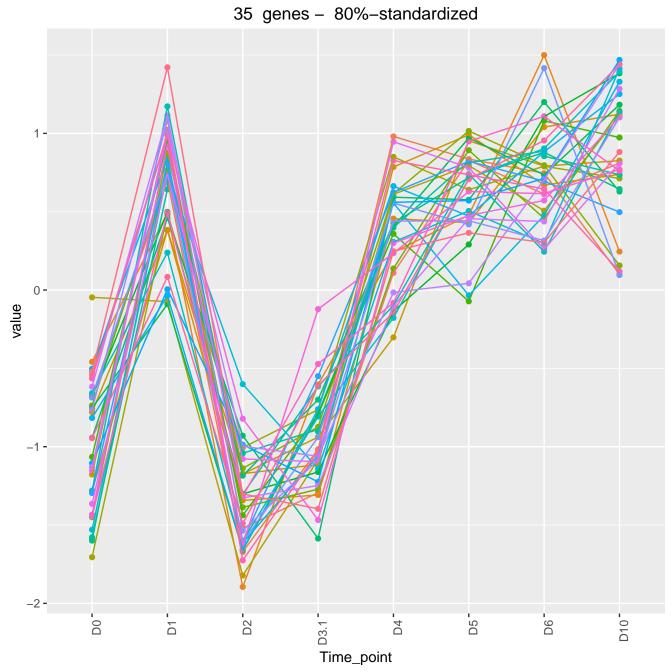




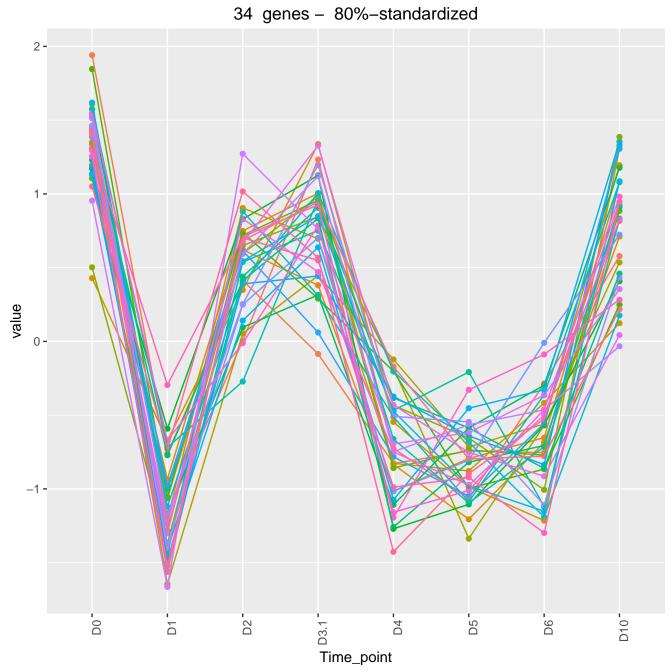




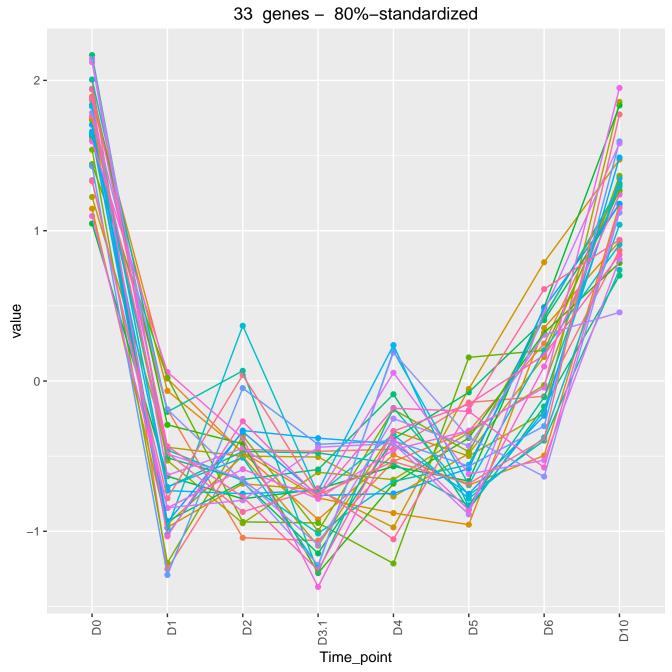


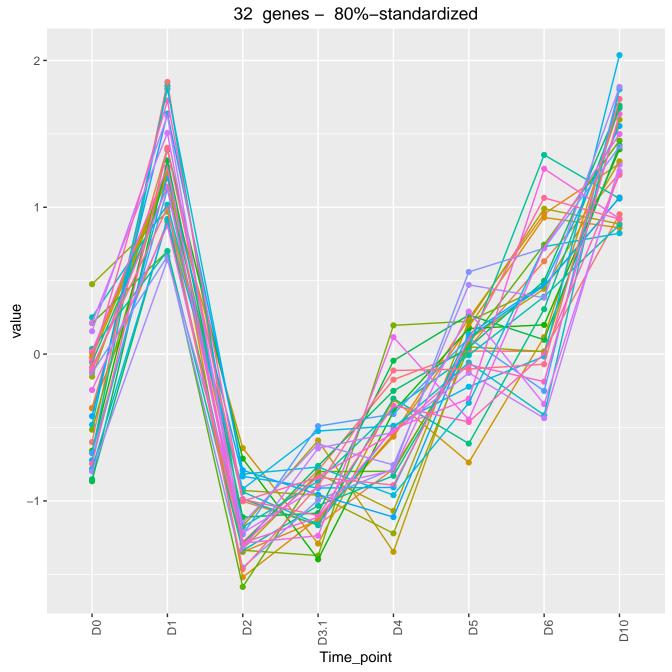


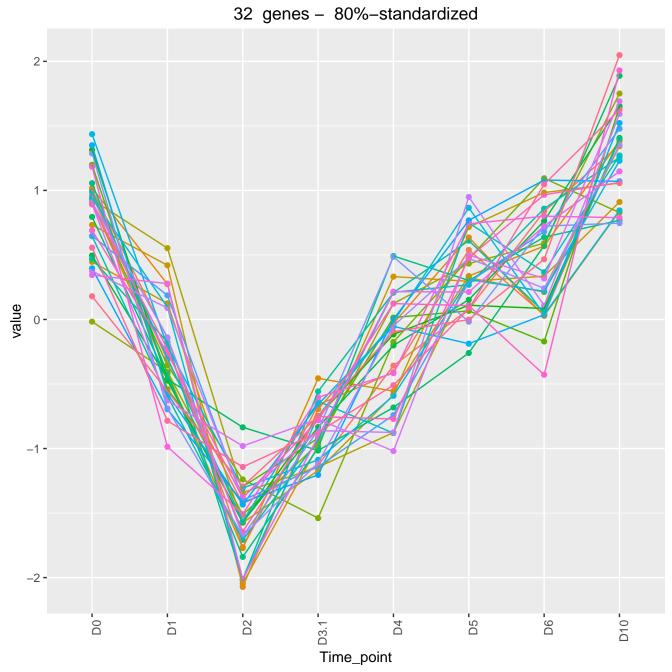
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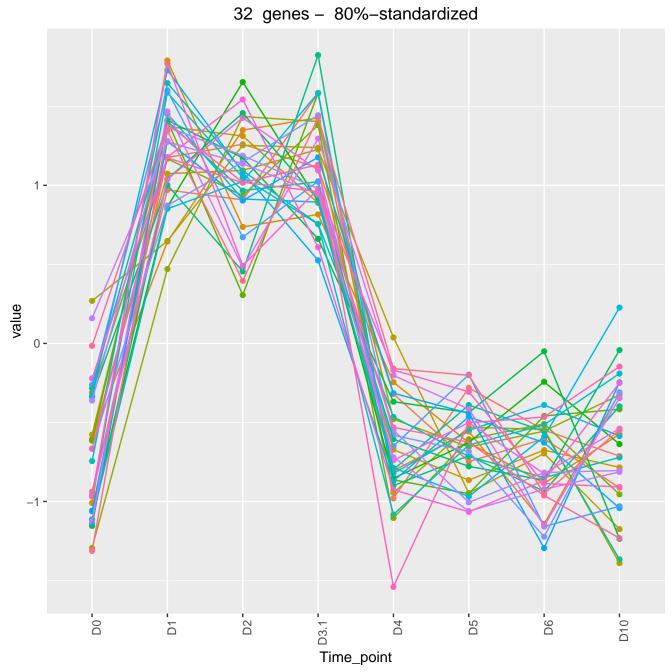


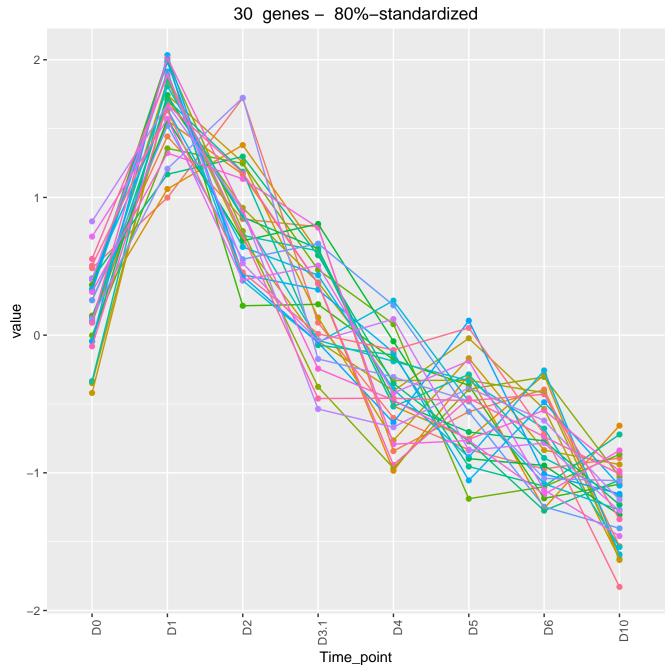
34 genes - 80%-standardized 2 -1 value **-1-**-2 **-**Time\_point \_0Q D2 D5 7

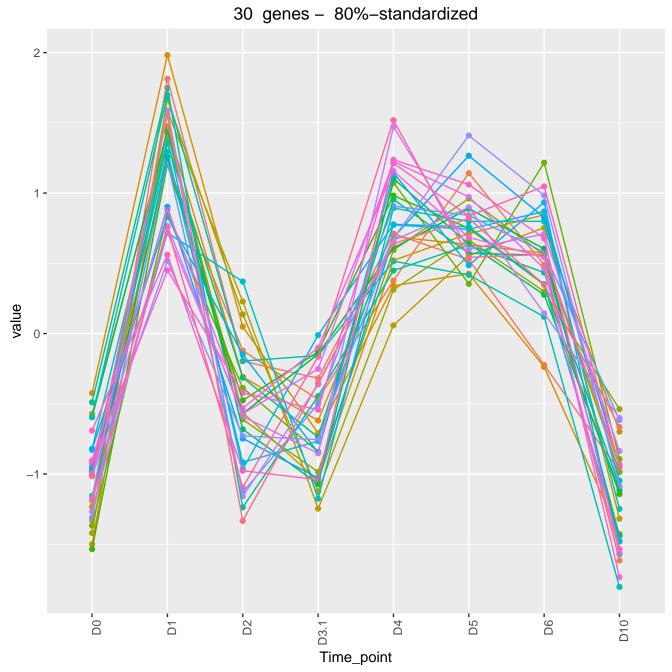


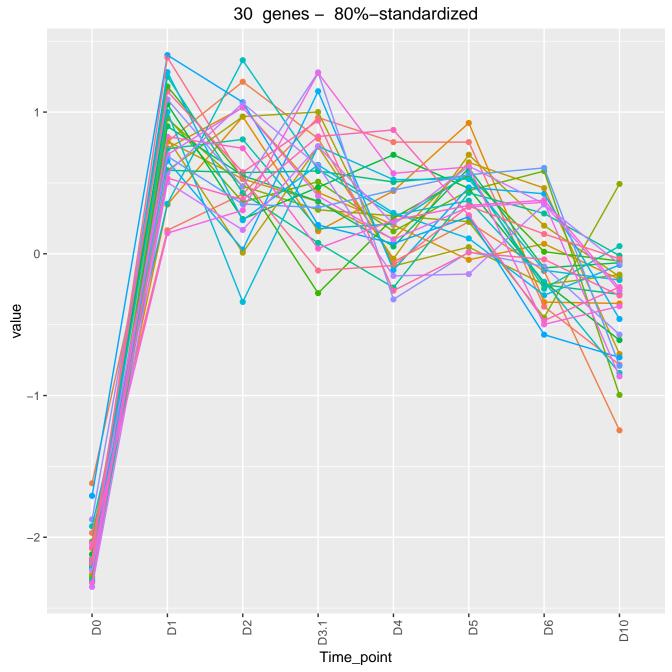


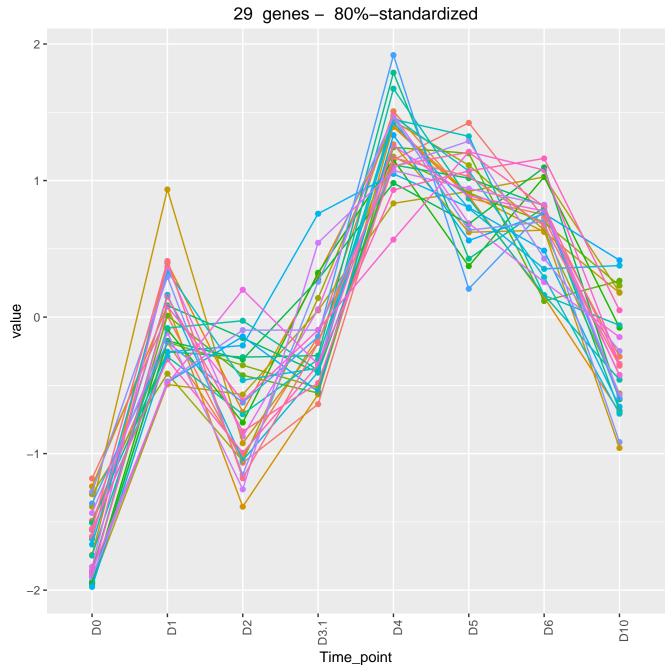


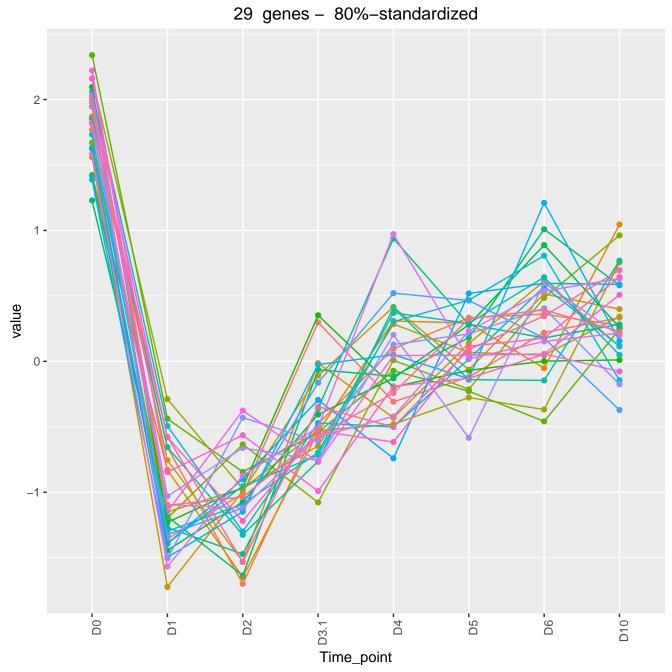


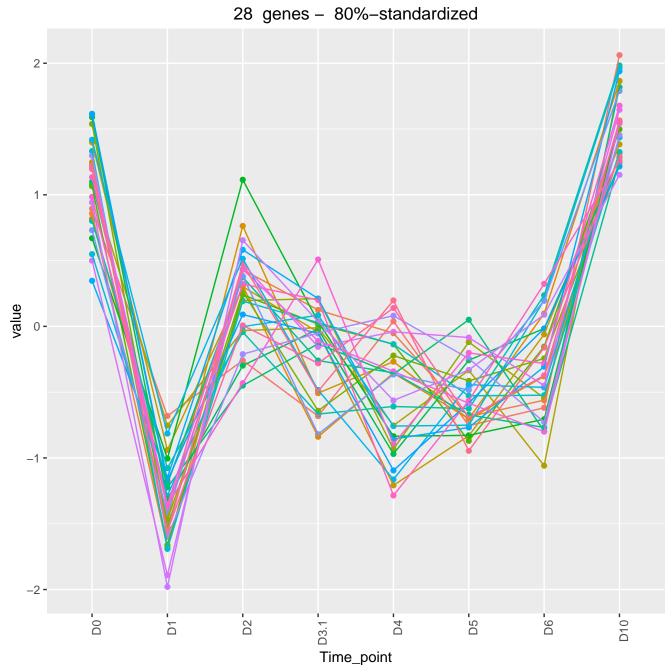


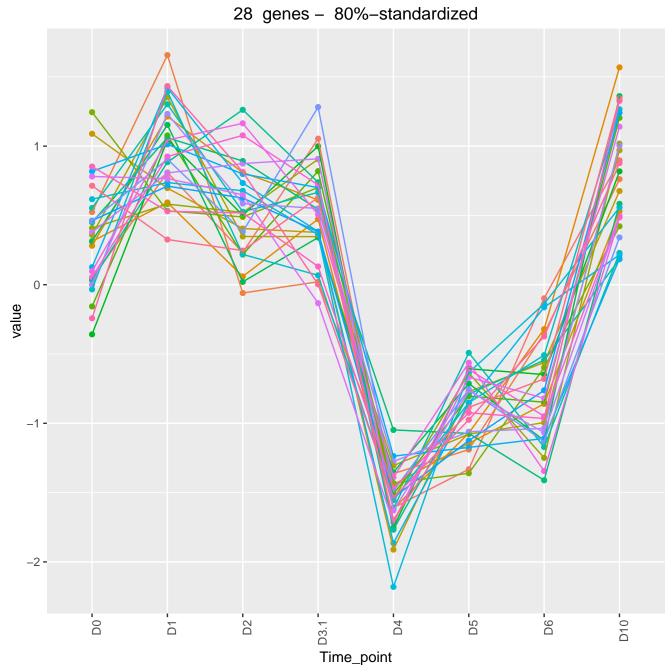


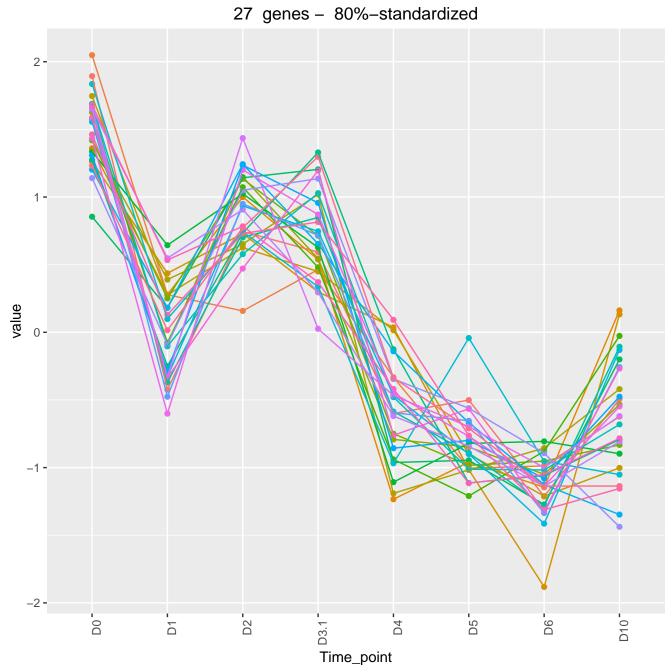


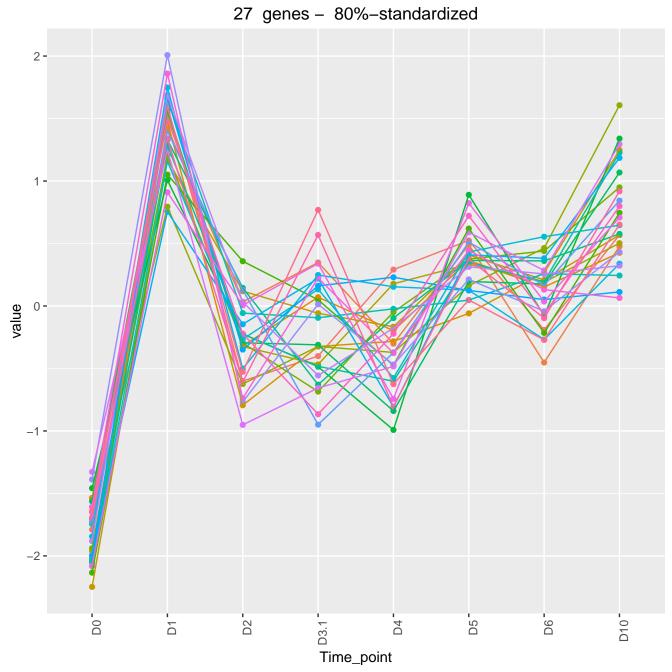


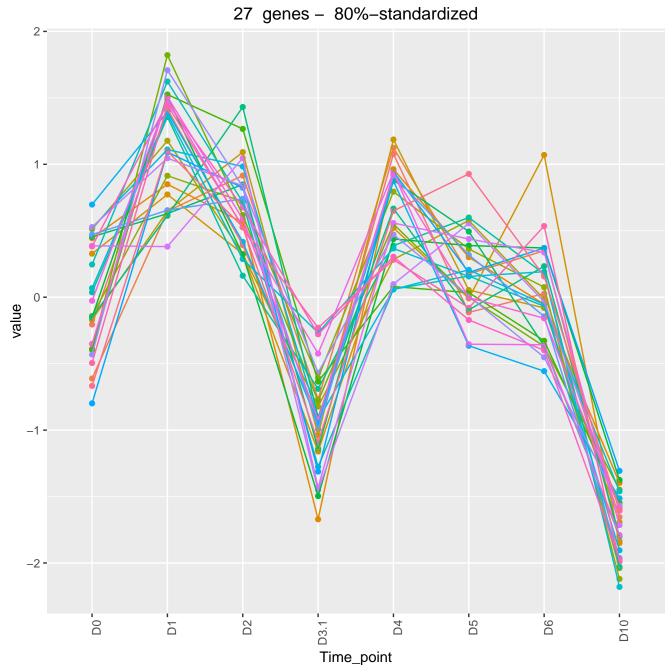


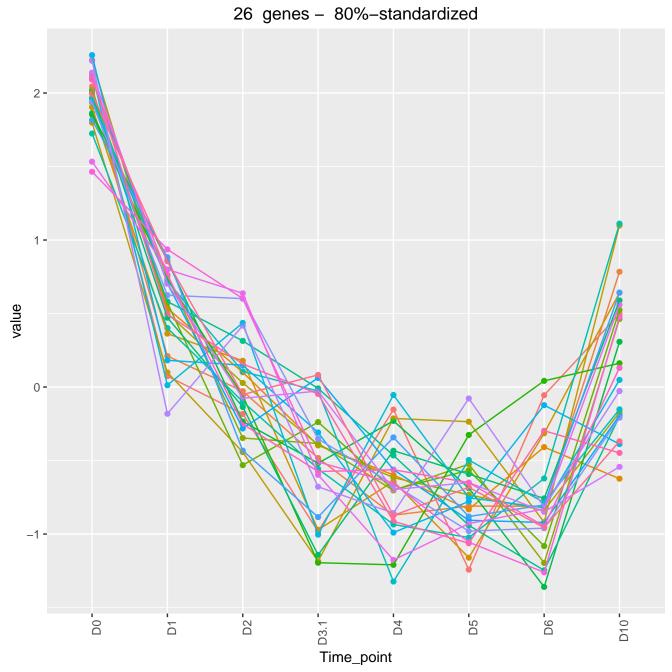


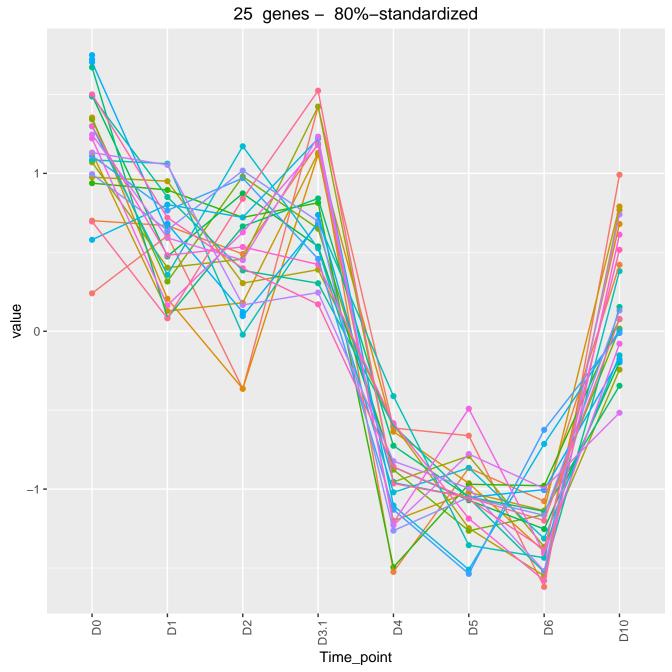


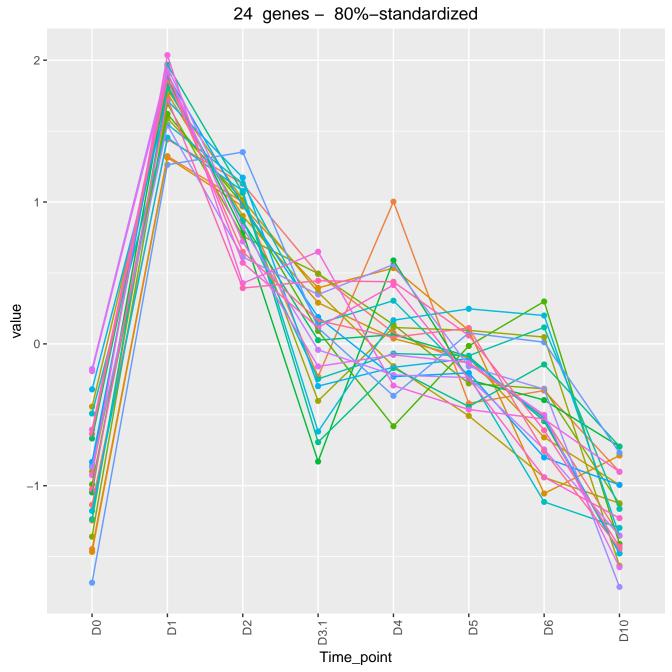


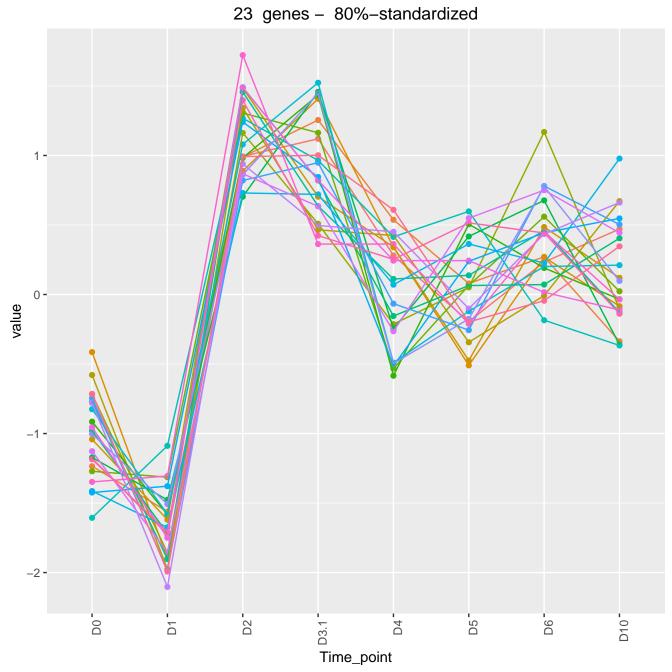


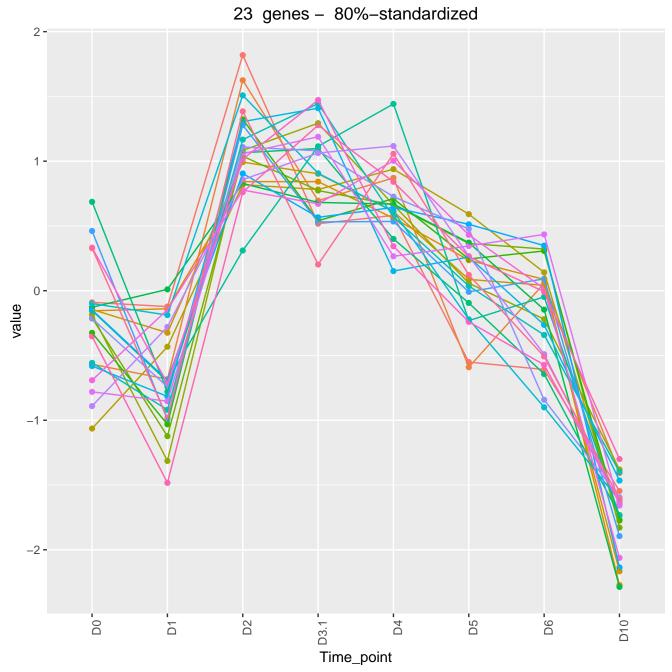


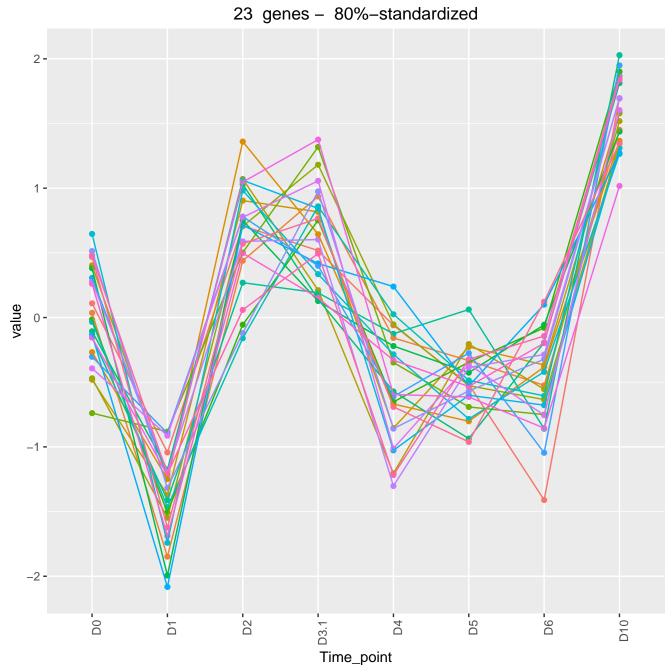


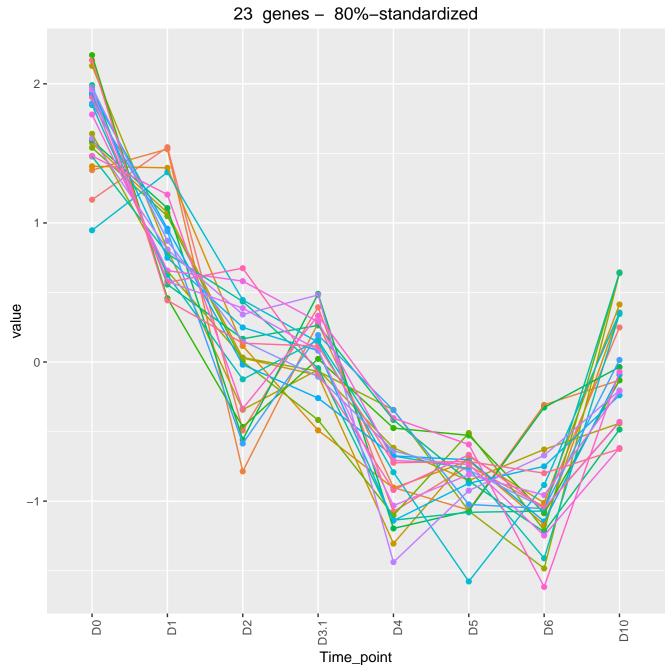


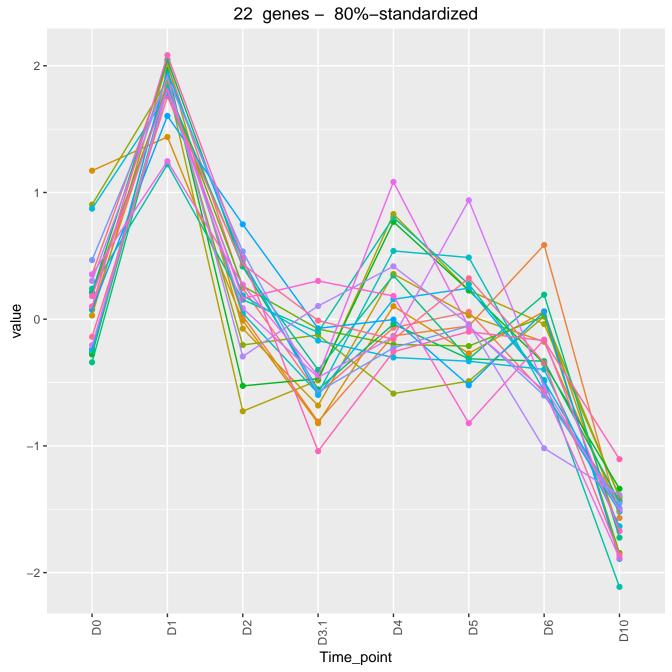


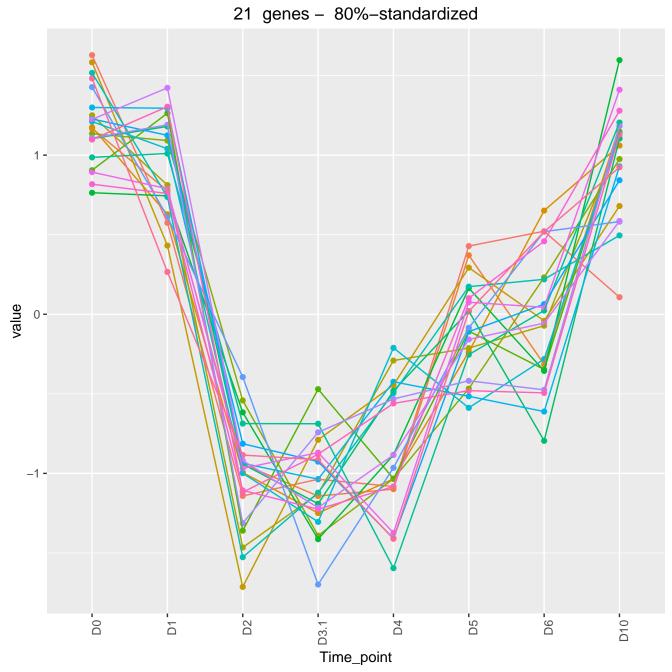


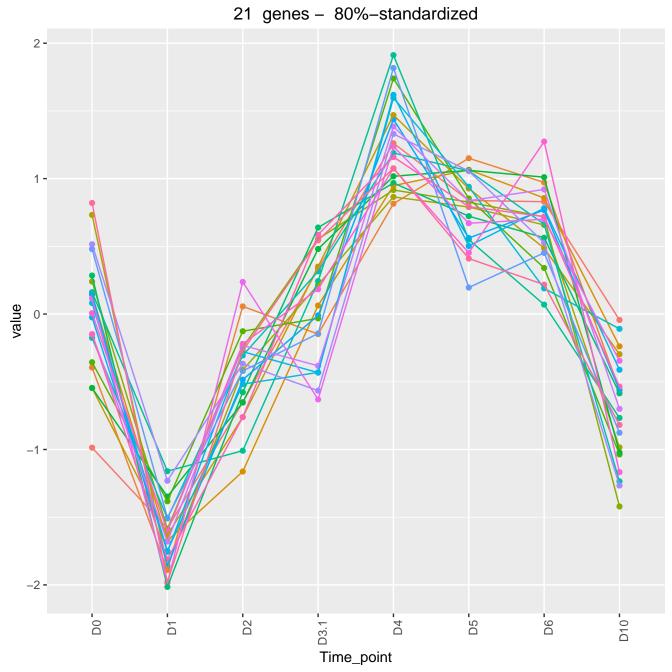


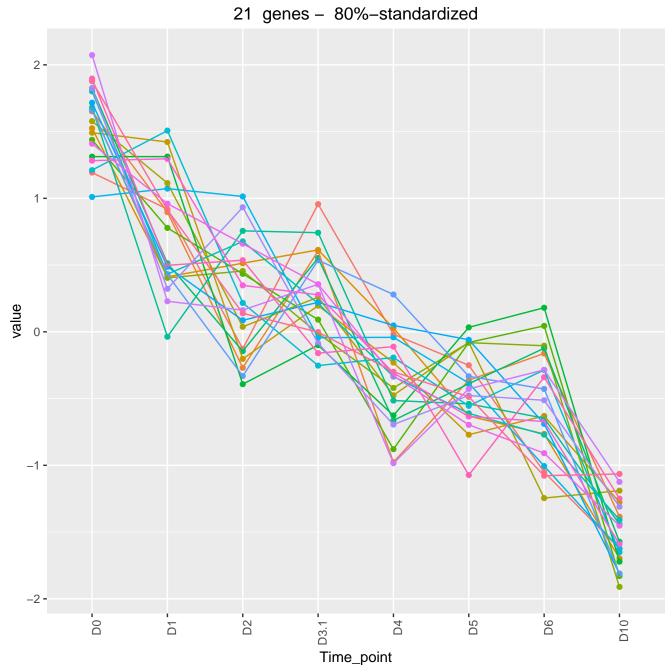


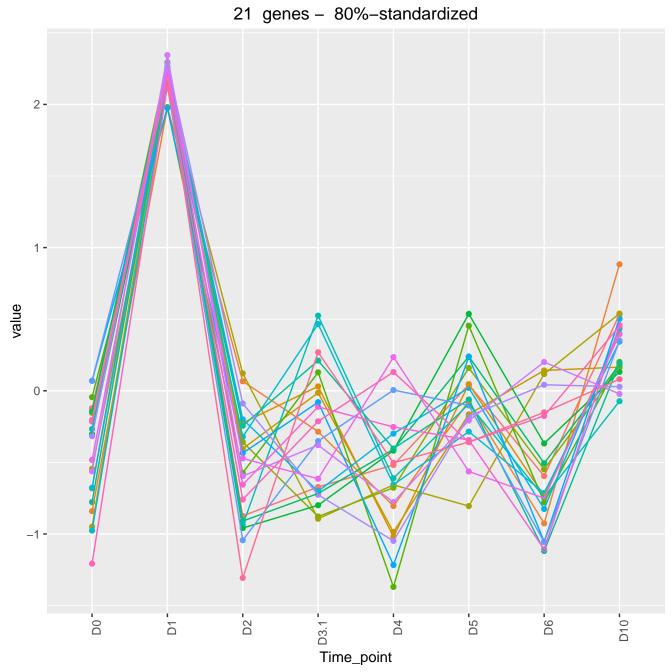


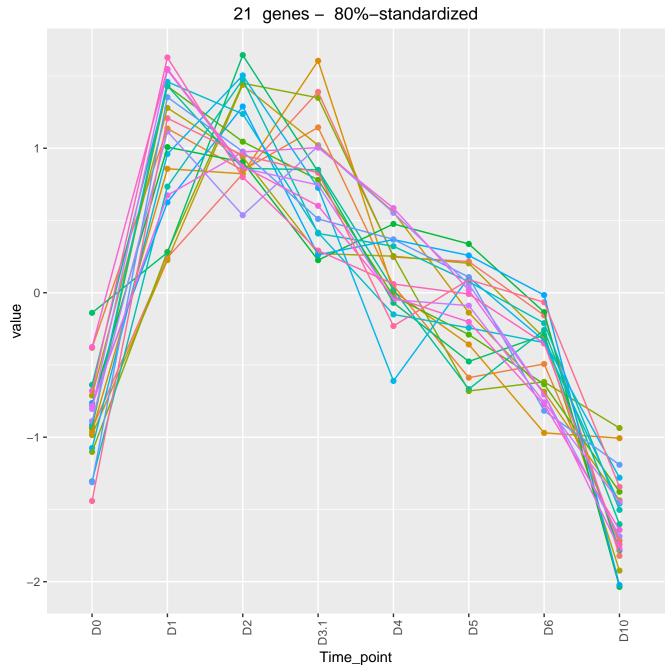


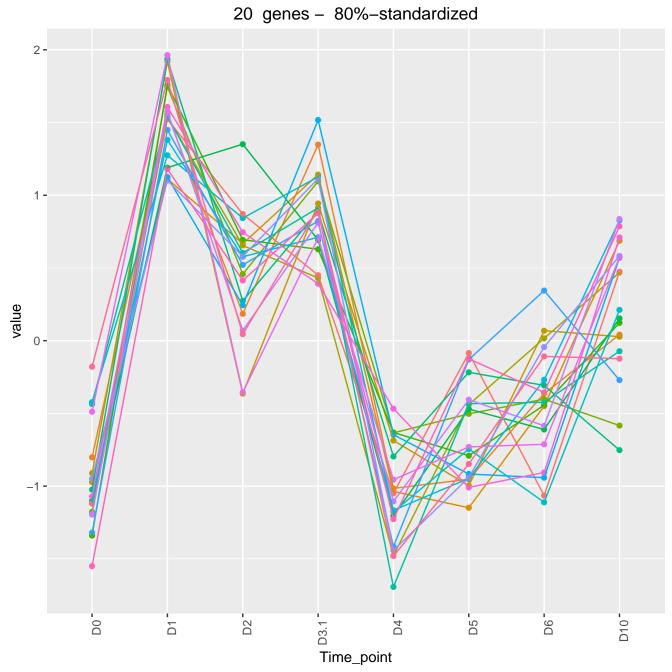


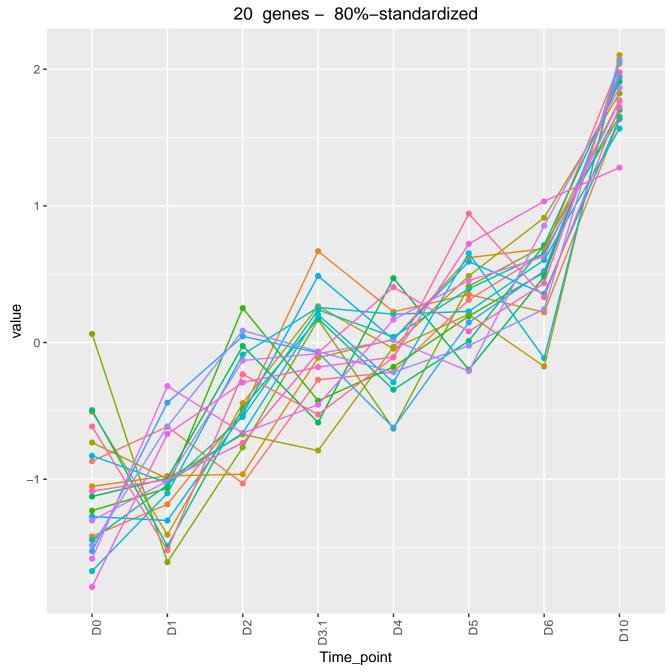


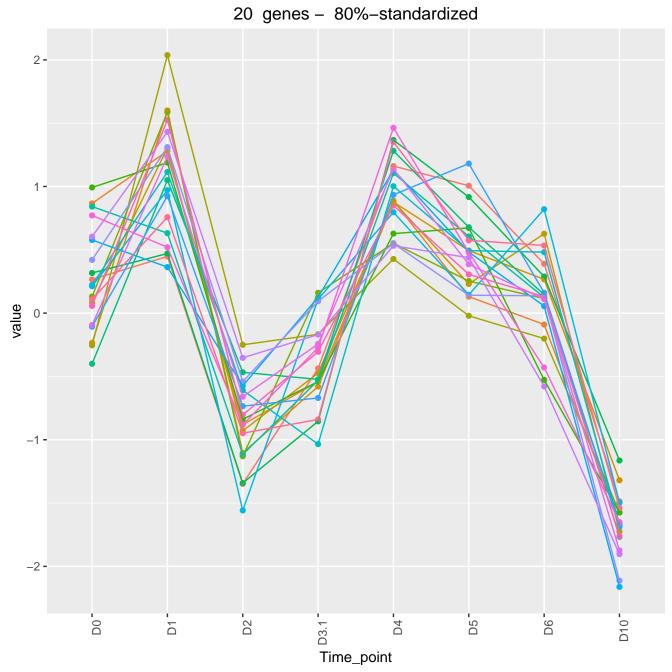


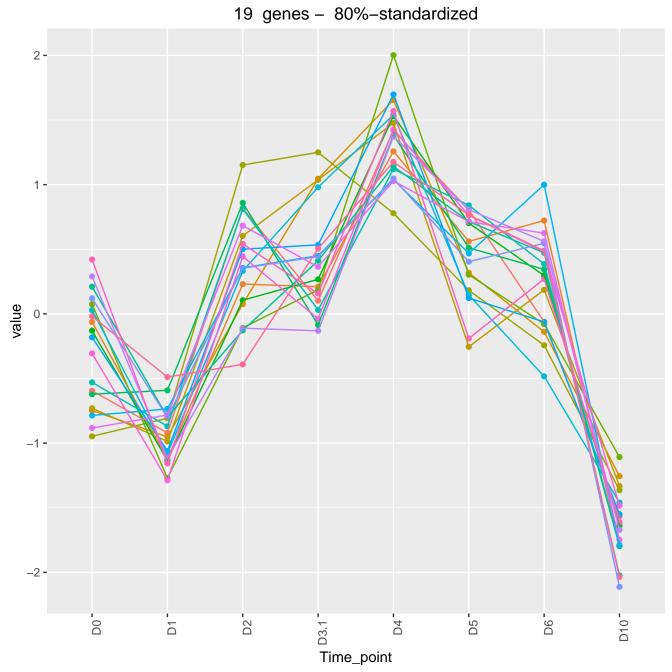


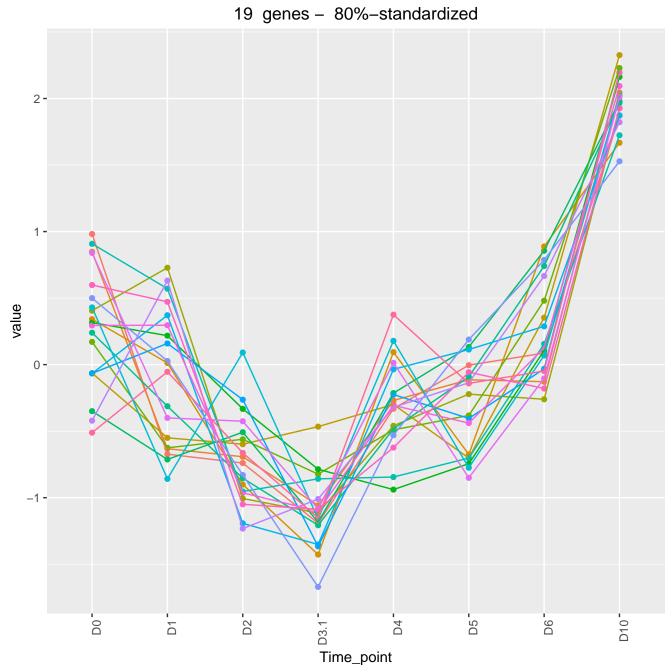


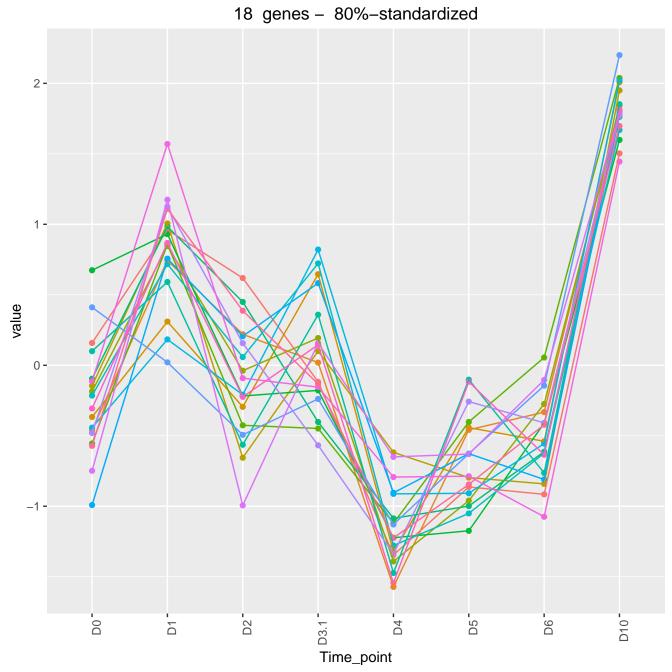


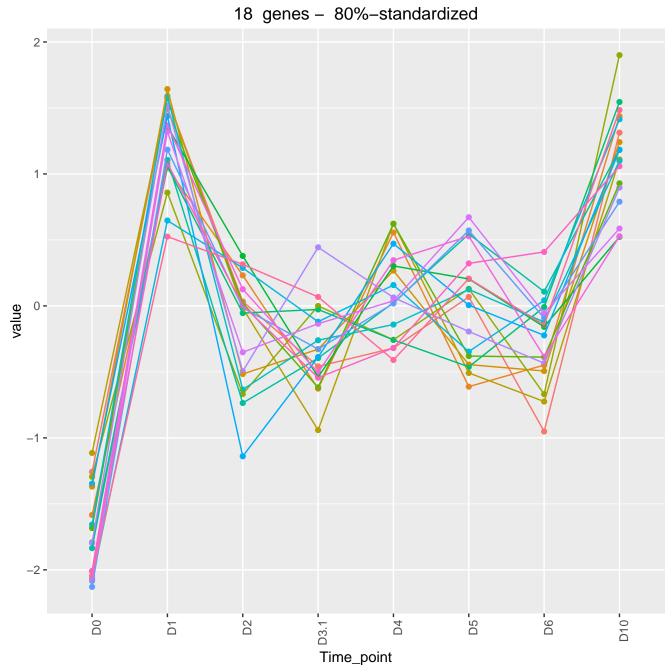


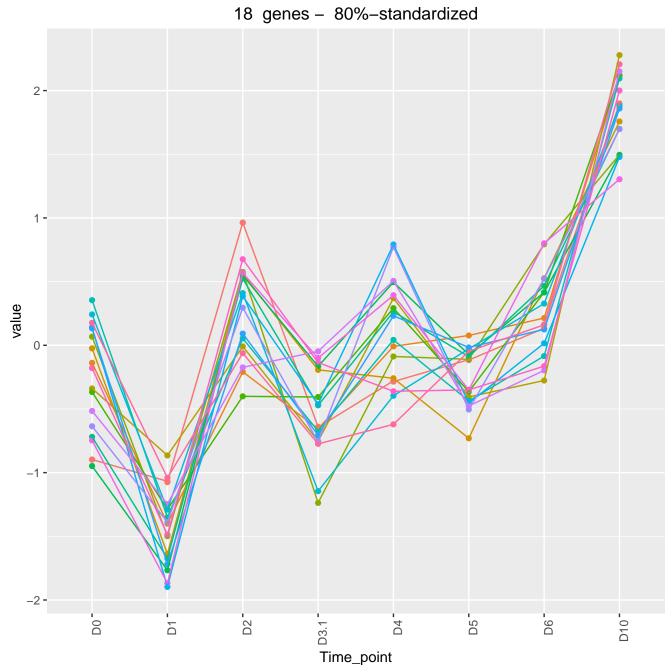


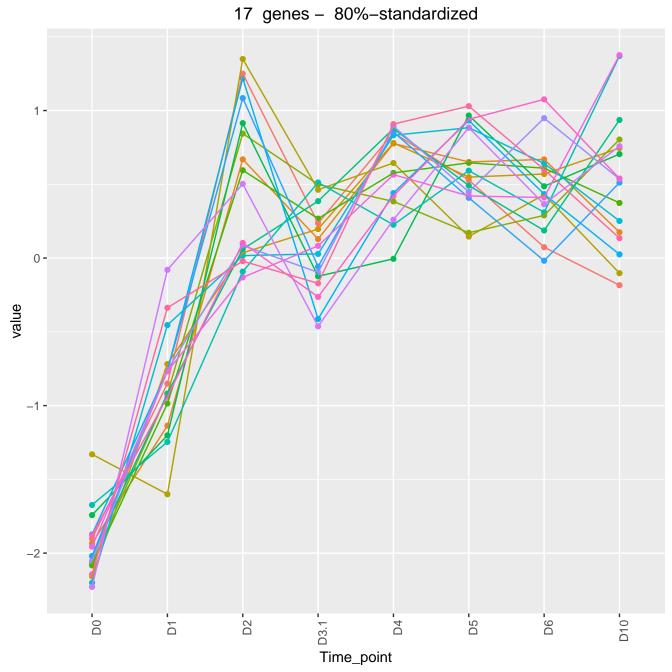


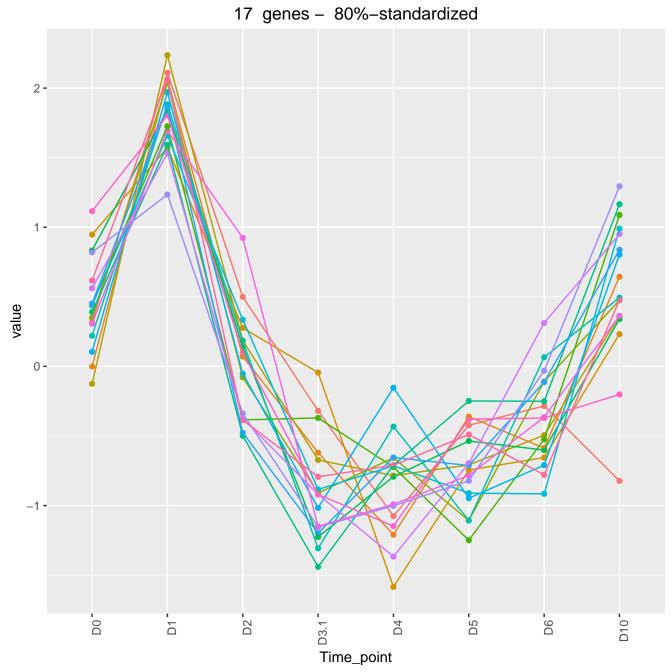


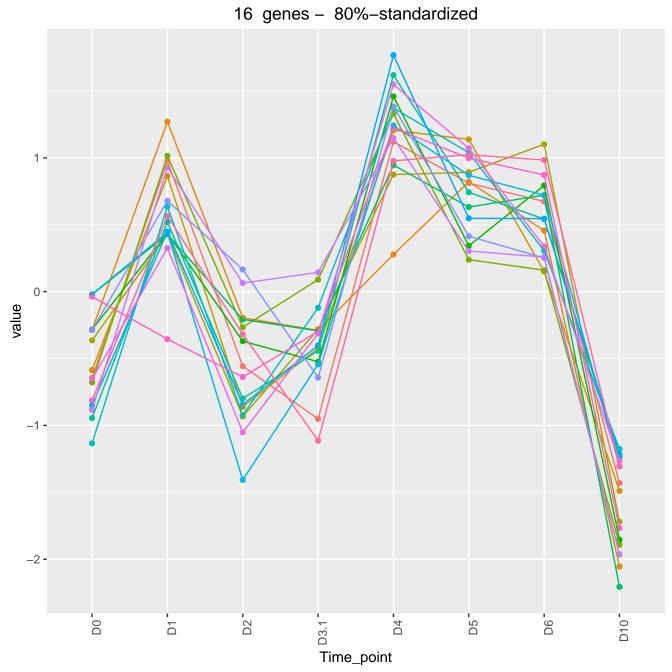


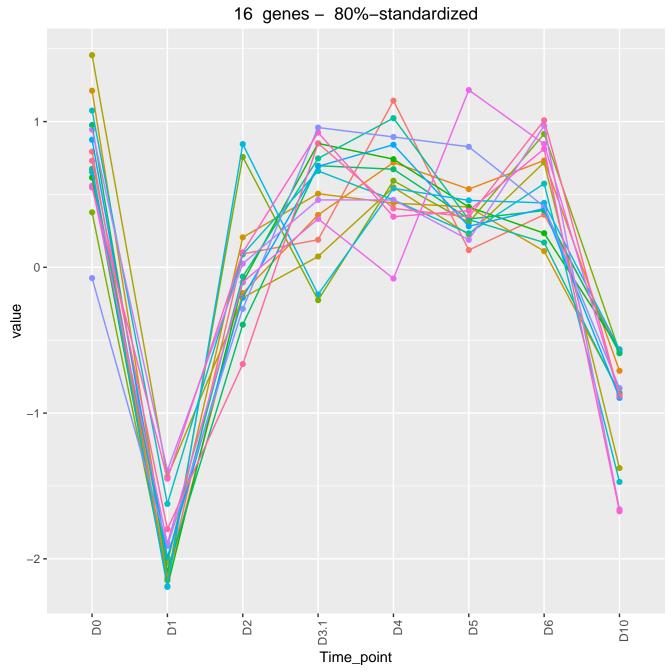


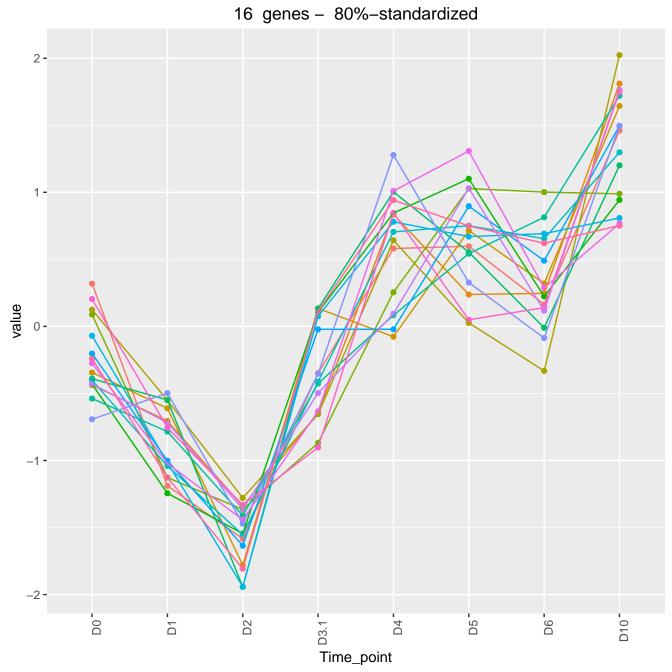


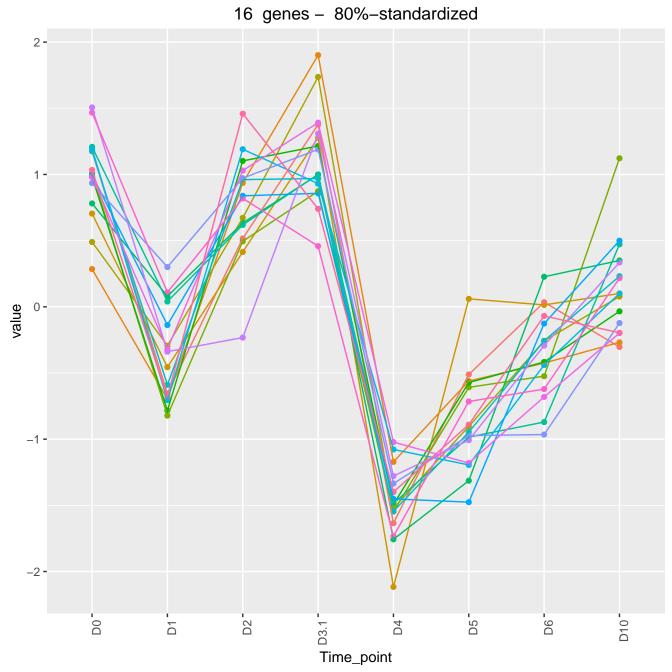


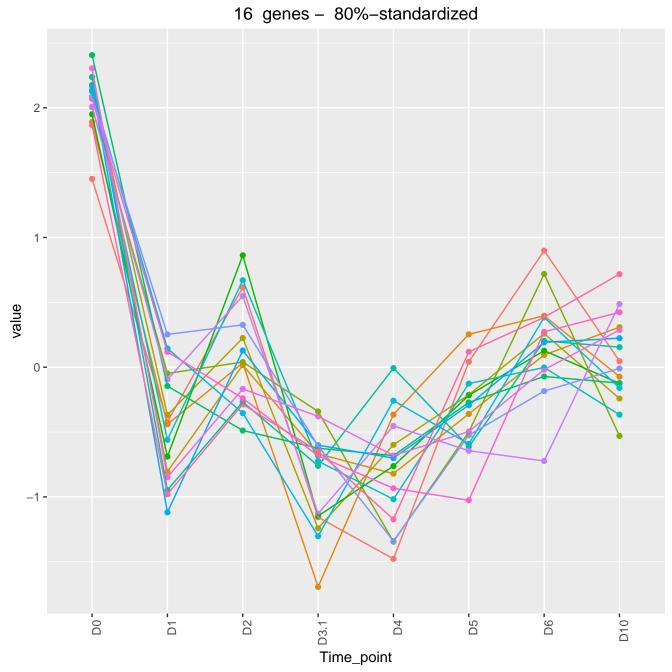


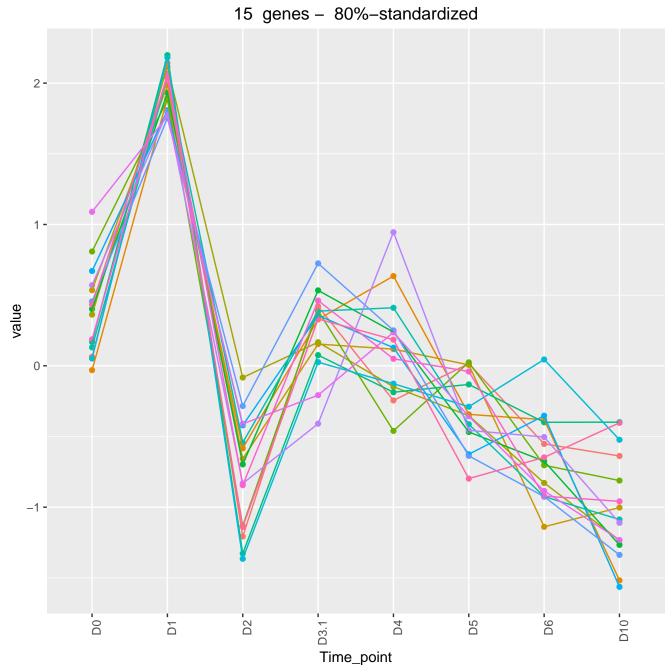


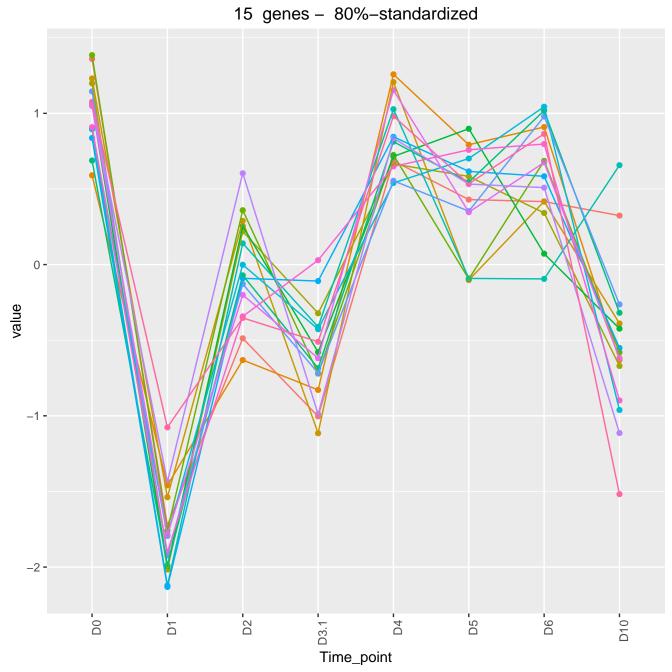


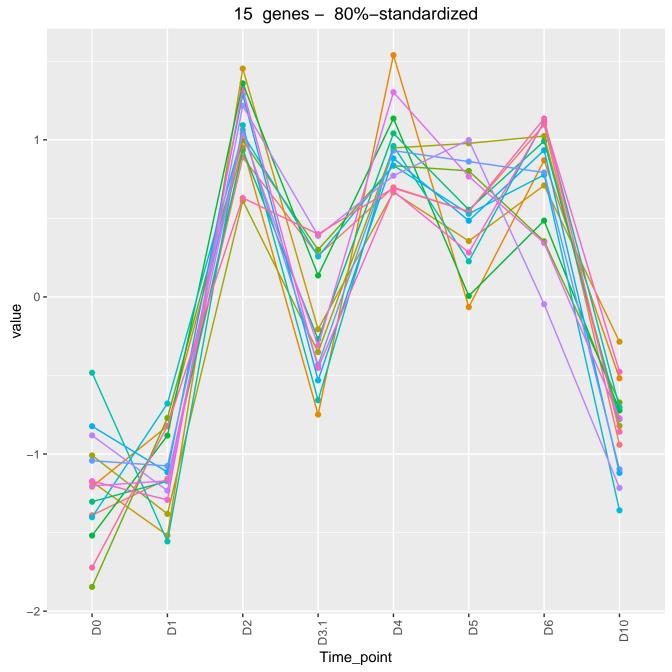












14 genes - 80%-standardized 2 -1 value o -**-1-**-2 **-**Time\_point \_ 00 D2 D5 7

