Supplementary figure S1. Connectivity patterns for TFs and their direct connections with fibrosis markers in A) NM-401, B) NRCWE-26 and C) BLM GRNs. The digital in the tables indicate the number of connections in a network between a TF and its direct target in a network, provided that the target gene is consisted in a list of fibrosis markers and this gene is differentially expressed. The red color scale denotes the magnitude of the values. M represents the module number to which a TF belongs.