

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Transcript abundance per biotype upon random priming reverse transcription.

File Name: Supplementary Data 2

Description: Analysis of differentially expressed gene between the two major clusters FUS/PDDA and Dhh1/Lysine

File Name: Supplementary Data 3

Description: Chemically synthesized motif sequences

File Name: Supplementary Data 4

Description: Amino acid sequence of recombinant proteins