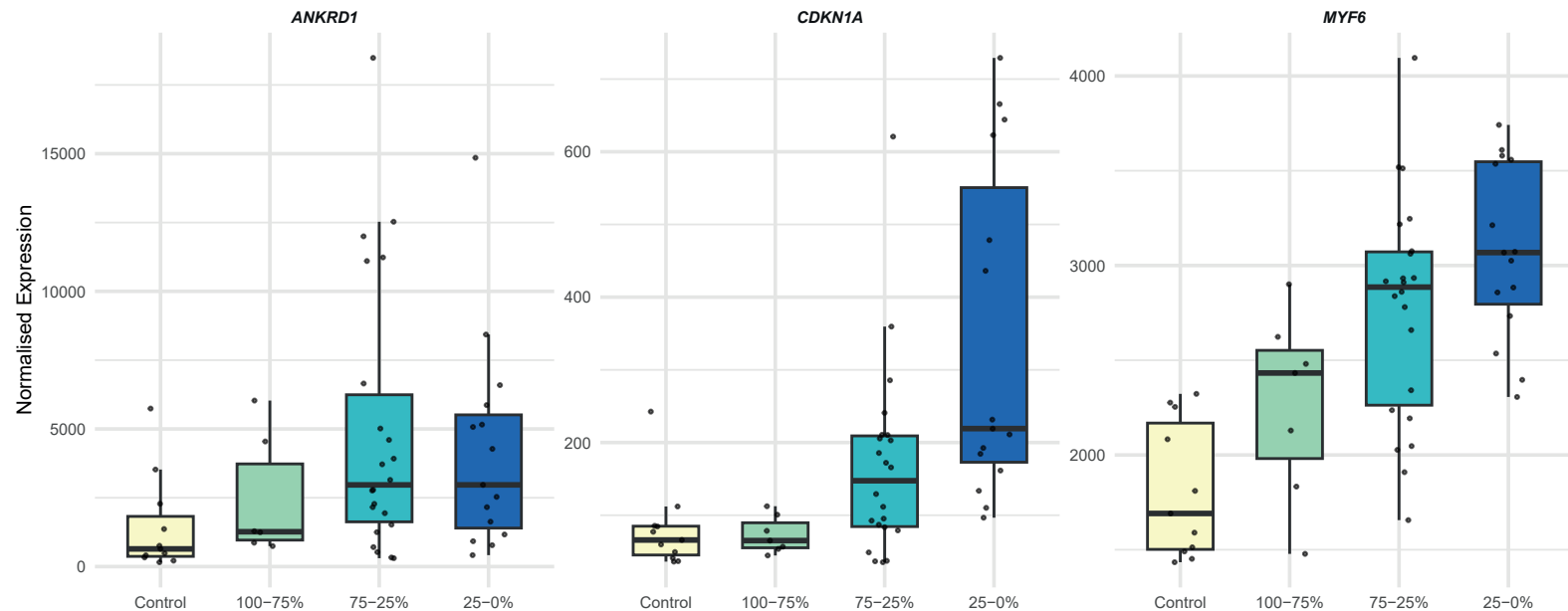# Supplementary Figure S8

Expression of *ANKRD1*, *CDKN1A*, and *MYF6* genes analysed in a bulk RNA-seq dataset from unrelated DM1 patients (9 controls, 43 patients). Patients were stratified by dorsiflexion strength into three clinical groups: 100–75% (mild/proto-DM1), 75–25% (classic DM1), and 25–0% (severe DM1).