Figure S1: exclusion criteria for 2 ERM and 2 PDR samples from study. A) Principal component analysis (PCA) of all samples showed nice clustering for most samples of each patient group. ERM_10 and PDR_10 were identified as outliers and thus excluded from further analysis. Despite bad clustering of dryAMD_1 and dryAMD_2, these samples were kept for further analysis, due to the very low sample number in the dry AMD group; B: Scaling factors applied for signal level normalization. Exclusion of PDR_11 and ERM_11 (red bars) due to biggest scaling factors, reducing the total scaling factor from 33.4 to a factor of 11.8; excluded samples marked with red box.