Supplementary Fig. 2. Evolution plots of FQPUT for optimized subsampling of biexponential fitting as more b-values are added in feed forward algorithm.

Annotated numbers on total error curves represent which b-value was added at that iteration of feed forward selection loop. (A) SCC data. (B) Adenocarcinoma data. D = pure diffusion coefficient, D* = pseudo-diffusion coefficient, f = perfusion fraction, FQPUT = fit quality per unit time.