A

% of mutated $BRAF^V600E$ by qPCR

% of mutated $BRAF^V600E$ by pyrosequencing

$R^2 = 0.8834$

B

% of mutated $BRAF^V600E$ in gDNA

% of mutated $BRAF^V600E$ in cDNA

$R^2 = 0.9569$