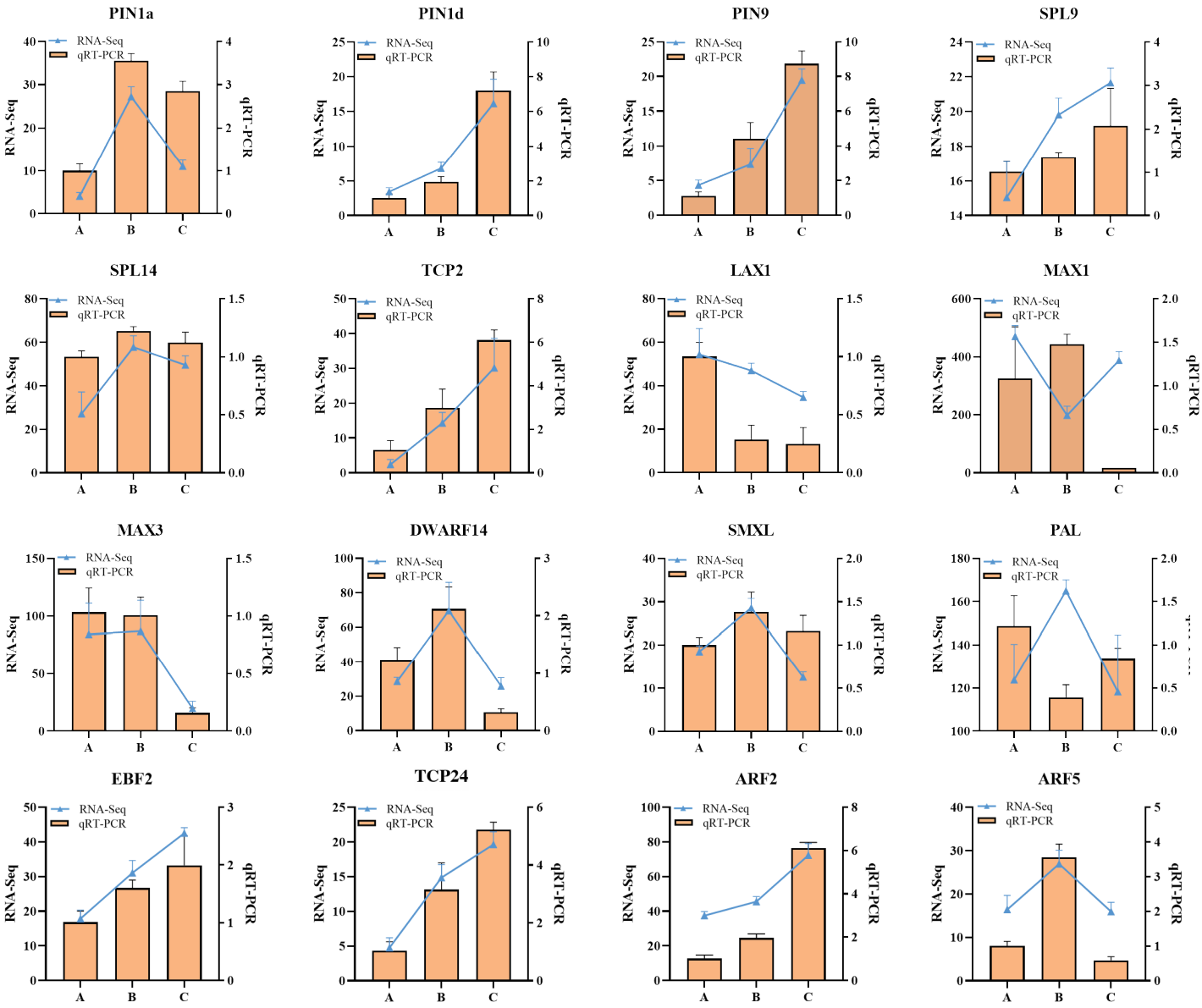

Fig. S7. The analysis of expression patterns of candidate genes. Error bars indicate mean ± SD, n=3