



Figure 4. PCR and restriction digestion analysis to confirm various cloning steps in the construction of pITB441 and pITB450 vectors (see figure 3 for other details).

- A.** *HindIII* digested λ DNA marker
- B.** PCR amplification to check insertion of *rbcS* promoter into pGEMT-Easy (lane 1), NC: Negative control; PC: Positive control. M: *HindIII* λ marker
- C.** Digestion of pGEMT-Easy-*rbcS* with *XbaI* (lane 2), *HindIII* (lane 3) and *BglII-SmaI* (lane 4) showed the presence of expected band corresponding to the size of *rbcS* promoter. Restriction digestion with *NcoI-BamHI* (lane 1) determined that insertion of *rbcS* promoter was clockwise in direction.
- D.** Digestion of pITB250 (lane 1) and pITB450 (lane 2) with *HindIII-BglII* that determined replacement of CaMV 35S promoter with *rbcS* promoter. Note the presence of a ~400 bp band in pITB450 representing the *rbcS*.
- E.** Digestion of pITB441 (lane 1) and pITB440 (lane 2) with *HindIII-NcoI* that determined the cloning of “*uidA*:35S polyA” by replacing “T7 RNAP:35S polyA” Note the presence of a ~2.1 kb band (*uidA*) in pITB441 and a ~2.9 kb band (T7 RNAP) in pITB440.
- F.** Digestion of pBin-HygTX (lane 1) and pBin-HygTX-T7 (lane 2) with *BamHI-SalI* that determined insertion of T7 RNAP into pBin-HygTX.
- G.** Digestion pITB228 (lane 1) and pBin-HygTX-T7 (lane 2) with *HindIII* determine that “T7promoter-*uidA*-T7 terminator” cassette was inserted into pBin-HygTX-T7.