JOURNAL-BASED LEARNING EXERCISES



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in biomedical science.

DEADLINE WEDNESDAY 8 JANUARY 2020 The rivaroxaban-adjusted normalized ratio: use of the prothrombin time to Genetic polymorphisms in DNA repair genes and their association monitor the therapeutic effect of rivaroxaban with cervical cancer Kim B, Jang S, Lee YJ, Park N, Cho YU, Park CJ. Br J Biomed Sci 2019; 76 (3): 122-8. Abbas M, Srivastava K, Imran M, Banerjee M. Br J Biomed Sci 2019; 76 (3): 117-21. Assessment No: BJBS1019A Assessment No: BJBS1019B Anticoagulants such as warfarin and heparin are widely used to treat Carcinoma of cervix is the third most common cancer among women worldwide, 01 venous thromboembolism, and do not require frequent monitoring to 01 with approximately 530,000 new cases and 275,000 deaths each year. ensure correct plasma levels. Prothrombin time is not a good candidate for a method to measure the plasma Epidemiologic studies have shown that most cases of cervical cancer are 02 02 caused by the human papillomavirus (HPV), mainly HPV-16 and HPV-18. concentration of rivaroxaban. Among various DNA repair pathways, base excision repair (BER) restores Rivaroxaban is a widely used direct inhibitor of activated factor X (Xa) 03 03 DNA double-strand breaks. Genetic polymorphisms in DNA repair genes may be associated with repair The study cohort comprised 337 healthy individuals (157 males, 180 females). 04 04 efficiency of damaged DNA and influence cancer risk. The Arg188His polymorphism of XRCC1 plays an important role in pancreatic Rivaroxaban (50 mg) was dissolved in 20 mL dimethylsulphoxide (DMSO) to 05 produce a stock solution diluted in phosphate-buffered saline without Ca2+ or Mg2+ 05 and colorectal carcinogenesis. at a final rivaroxaban concentration of 500 mg/mL. A total of 60 rivaroxaban-spiked plasma samples were analysed. Cervical cancer patients (n=230) and healthy age-matched controls (n=270) 06 06 aged between 30 and 70 years were recruited for the study To evaluate sensitivity variations among different thromboplastins, plots were drawn Frozen EDTA blood samples were thawed at room temperature and high molecular 07 07 for the spiked concentration of rivaroxaban and PT of each sample. weight DNA was extracted using a slightly modified salting out method. Three chromogenic anti-factor Xa assay reagents were used in this study. The sample size for each SNP was calculated by QUANTO software using 08 08 major allele frequency and prevalence. Specific rivaroxaban anti-factor Xa activity was measured using two Of the cervical cancer cases, 93.5% were in stages II/III with 6.5% in stages I/IV. 09 09 different instruments. All three thromboplastins tested showed the same sensitivity to rivaroxaban. Compared to the GG genotype, adjusted frequencies of GA, AA and GA+AA 10 10 genotypes were lower in cases compared to controls. The mean (SD) for warfarin-adjusted INR using Neoplastin Cl-plus was 1.80 (0.97). Digested products were visualised on 2% polyacrylamide gel after staining 11 11 with ethidium bromide. Rivaroxaban-adjusted SI values were different from the current warfarin-adjusted Cervical cancer is a complex disease where environmental and genetic factors 12 12 ISI in all tested thromboplastins. play important roles in pathogenesis. Rivaroxaban-specific NR does not reduce the differences between the PT results Women with GA and AA genotypes of XRCC1+399A/G showed 2.8-3.4-fold 13 13 from different thromboplastins and coagulometers. higher risk of cervical cancer. HemoSIL Recombiplastin 2G originates from a human source, so this The raw carriage rates of G (+), G (–) and A (+), A (–) showed significant 14 14 thromboplastin represents the WHO calibration method. association with cervical cancer when compared to controls. Studies have shown a poor correlation between the anti-factor Xa assay and mass Amplification was followed by initial denaturation at 95°C, followed by 35 cycles 15 15 spectrometric measurement of rivaroxaban in normal plasma samples spiked with at 95°C, annealing at 56°C, extension at 72°C, and final extension at 72°C. known amounts of rivaroxaban. The authors found that use of a rivaroxaban-adjusted NR did not minimise The XRCC3+18067C/T genetic variant is associated with cervical cancer. 16 16 inter-thromboplastin variability. Traditional coagulation tests are speedy, widely available and inexpensive, The important molecules of the BER pathway are RAD51, XRCC2 and XRCC3. 17 17 and are suitable for determining plasma DOAC levels. Spiked samples reflect the potential variations of sensitivity between In the study population, the frequency of GA and AA genotypes, and the A 18 18 thromboplastins and between in vivo samples with similar levels of rivaroxaban. allele of XRCC1+399A/G are significantly lower in cases compared to controls. Mass spectrometry, the current validated standard, was not used to measure Genotypes were determined by polymerase chain reaction-restriction fragment 19 19 plasma rivaroxaban concentration. length polymorphism. The major allele frequency was calculated after genotyping 100 normal Anti-Xa assav versus NR-rivaroxaban correlation coefficients were 0.97-0.99. 20 20 individuals for each SNP. **REFLECTIVE LEARNING** Molecular genetics is set to play an increasing and important In certain circumstances, monitoring of plasma rivaroxaban concentration 01 01 might be needed. Discuss. role in cytopathology. Discuss. Explain in detail why this paper by Kim et al. represents an advance Explain in detail why this paper by Abbas et al. represents 02 02

an advance in biomedical science.