P#ST37



Use of Intra-Gross Examination Touch Prep Slides to Facilitate Turnaround Time (TAT) Advantage Using the Biocartis Idylla





Introduction EGFR A 24 Idylla KRAS Idylla NGS or direct Concordance MSI Idvlla MSI Concordance 22 NRAS/BRAF/EGEE L861Q As the role of molecular biomarkers for clinical management of cancer patients expands, the demand for sequencing MSS MSS Concordant 20 G12A WT KRAS G12A Concordant MSS MSS rapid molecular results is increasing. We sought to reduce TATs for specific actionable genetic alterations by Concordant 18 utilizing touch preparation (TP) samples as substrate material for the Biocartis Idylla system, a fully MSI-H MSI-H Concordant C2 G12V KRAS G12V Concordant WT MSS MSS Concordant automated molecular assay platform 14 MSI-H MSS Discordant NRAS/EGFR - WT WT No mutations Concordant/partia 12 MSI-H MSI-H Concordant BRAF - Invalio detected invalid Methods 10 MSS MSS Concordant WT WT No mutations Concordant MSS MSS Concordant Diff-Quik (DQ) stained TPs were prospectively collected from non-small cell lung carcinoma (NSCLC) and detected MSS MSS Concordant WT NRAS/EGFR - WT Concordant/partia colorectal adenocarcinoma (CRC) resection specimens at the time of gross examination. If tumor cells were C5 No mutations MSI-H MSI-H Concordant BRAF - Invalid detected invalid present microscopically, they were tested using the Biocartis Idylla system for EGFR, KRAS, BRAF, NRAS MSS Invalid Invalid and microsatellite instability (MSI) depending on tumor type. The accuracy of this approach was evaluated C6 WT NRAS/EGFR - WT BRAF V600F Concordant MSI-H MSS Discordant with subsequent corresponding formalin fixed paraffin embedded (FFPE) material. BRAF - V600E 20 Invalid MSI-H Invalid 20 40 10 50 MSS MSS Concordant G12A NRAS/BRAF-Invalid KRAS G12A Concordant/partial Figure 1: Case L10, Diff-Quik TP slide, chamber Invalid MSS Invalid EGER - WT invalid A. EGFR total CQ = 21.82, dCQ = 3.26 Results MSS MSS Concordan WT NRAS/EGFR - WT KRAS N116H Concordant/partial EGFR 16 CRCs and 13 NSCLCs were tested. BRAF - Invalid invalid* No mutation detected Idvlla Results Among CRCs, 14, 13, 6 and 13 had valid KRAS, NRAS, BRAF and MSI results, respectively Days compared to final C9 WT NRAS/EGER - WT No mutations Concordant/partia 4 KRAS mutations and 1 BRAF mutation were identified in CRCs (Table 1). Available pathology sign out date BRAF - Invalid detected invalid 2/5 MSI-High cases demonstrated discordant results (C5 and C12, Table 3), called MSI-H by Idylla but Partial results 7 days before C10 G12D WT KRAS G12D Concordant microsatellite stable by confirmatory testing. Partial results 37 days after Among NSCLCs, 13, 12 and 13 had valid EGFR, KRAS and BRAF results, respectively. C11 WT NRAS/EGFR - WT No mutations Concordant/partia Complete results 2 days before BRAF - Invalid detected invalid 2 EGFR mutations and 2 KRAS mutations were identified in NSCLCs (Table 2). Complete results Same day All calls with valid mutational results were concordant with orthogonal analysis except one NSCLC case WT WT KRAS G10dup Concordant' (L10) in which both a false positive (FP) EGFR and false negative (FN) BRAF mutation was evident on Complete results 1 day after WT NRAS/EGFR - WT BRAF V600E Concordant/partia Idylla L6 Complete results 11 days after BRAF - Invalid invalid Further evaluation of discordant cases was performed by testing unstained slides, which resulted in C14 WT NRAS/EGFR - WT No mutations Concordant/partia 17 Complete results 1 day before correct findings (Example case presented in Figures 1-3). BRAF - Invalio detected invalid Select cases with invalid results were re-run on the Idylla utilizing FFPE material or unstained TPs and Partial results 7 days before KRAS G13D Invalid Invalid Invalid 30 40 10 20 50 subsequently vielded valid results. Partial results 8 days before Figure 2: Case L10, Unstained TP slide, A subset of samples (n=22) tested with rapid intent were used for analysis of TATs. 17/22 had partial C16 Invalid Invalid No mutations Invalid chamber A. CQ = 17.83 C3 Partial results 5 days before results, defined as at least one result, or complete results available on the same day as the pathology detected report release (Table 4) Complete results 6 days before Table 2 M BRAF в 20 Complete results 4 days before Idvlla Idvlla Idvlla NGS or direct sequencing Concordance V600E/E2/D Discussion 18 EGFR KRAS BRAF Complete results Same day 16 EGFR DEL746 750 Exon 19 WT WT Concordant The aim of our study was to investigate the use of TPs on the Idylla system to reduce TATs such that results C7 Complete results 1 day before 14 deletion would be available with routine surgical pathology reports. Our approach of collecting TP samples directly WT G12C WT KRAS G12C Concordant C8 Complete results 4 days before 12 from tissue and testing them via the Idylla system significantly reduced potential TATs. Samples processed WT WT WT KRAS G13C Concordant* C9 10 Complete results 4 days after with rapid intent often provided results the same day as or before the pathology report release. Our study BRAF G466 Concordant/invalid suggests that DQ stain may cause interference, leading to invalid and FP results, thus use of a similar WT Invalid WT No mutations C10 Complete results 4 days before No mutations WT WT WT Concordant approach to accelerate TAT would require alternate methods to confirm tumor presence Complete results 11 days after WT WT WT No mutations Concordant W/T WT W/T EGFR A767 V769dup Concordant* C12 Complete results Same day References WT WT WT No mutations Concordant C13 Complete results Same day No mutations WT WT WT Concordant De Luca C, Conticelli F, Leone A et al. Is the Idylla EGFR Mutation Assay feasible on archival stained cytological smears? L8610 WT WT BRAF V600E Discordant Partial results 10 days before A pilot study, J Clin Pathol, 2019;72(9):609-614 G12V KRAS G12V WT WT Concordant 20 30 2. De Luca C, Gragnano G, Pisapia P et al. EGFR mutation detection on lung cancer cytological specimens by the novel fully WT WT WT No mutations Concordant Figure 3: Case L10, Unstained TP slide, Tables 1-4 Legend: WT = wild-type MSS= microsatellite stable MSI-H = atellite instability high, N/A = none applicable, * = mutation detected by automated PCR-based Idylla EGFR Mutation Assay. J Clin Pathol. 2017;70(4):295-300. WT WT W No mutations Concordant chamber B. SPC CQ = 28.70, dCQ = 2.14 orthogonal assay is not evaluated by Idvlla

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