Lower the Barrier to Adopt Large Cancer Genomic Profiling in Diagnostic Labs: Break the Data Bottleneck Summit

Expert tips and tricks to improve the accuracy and efficiency of reporting relevant variants from large cancer genomic profiling

Date: Thursday, May 19, 2022 - All times are shown in EDT Registration Link: <u>https://event.on24.com/wcc/r/3760943/FF03E173FB7423E0CD6FC42A0BF09452/3738303</u>

Agenda

1:00 p.m.–1:30 p.m. EDT	Simplify your secondary analysis to 5 steps
	 Ajay Athavale, Senior Bioinformatics Engineer, QCI Services, QIAGEN Digital Insights Variant calling, filtering and visualization: Structural variants, indels, point mutations Quality control reports Get started quickly with configurable pre-made workflows
1:30 p.m.–2:15 p.m. EDT	Generate oncologist-ready clinical reports from comprehensive cancer panels within minutes
	 Beate Litzenburger, Ph.D., Associate Director, Global Product Management, QIAGEN Digital Insights Why high-quality, deeply curated knowledge is necessary to boost your molecular tumor boards? How does QCI interpret the effect of co-occurring variants on prognosis and diagnosis? How can you identify incidental findings in QCI? How does QCI display and interpret structural variants? How does QCI handle multi-sample analysis?
2:15 p.m2:45 p.m. EDT	Going beyond: Molecular evidence for off-label therapy use
	 Sheryl Elkin, Ph.D., Chief Scientific Officer, QIAGEN Digital Insights Using molecular pathways to expand therapy options Identifying evidence-based therapy options for cancer patients for whom there is no on-label therapy Providing confidence with expert-curated, oncologist-reviewed treatment strategies

