GENOMENON

For Immediate Release
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Genomenon Attends Two Key Genetic & Pathology Industry Events

A Recap of Genomenon at the American Society for Human Genetics 2016 Show and Looking Forward to the Association for Molecular Pathology Show in November.

Ann Arbor, Mich. - October 28, 2016 - The Genomenon team recently attended the American Society for Human Genetics 2016 show (ASHG) and connected with over 8,000 attendees from around the world. The event, held in Vancouver this year, is the largest gathering of geneticists in the world and offers an extensive agenda from training sessions, networking opportunities, press gatherings and an exposition area.

Next up for the Genomenon team is the annual meeting for the Association for Molecular Pathology (AMP) in November. The three day event in Charlotte features a variety of activities including a corporate workshop day that invites conference attendees and the general public to learn about the latest in scientific advancements, as well as break-throughs technology and equipment. AMP also hosts several networking opportunities, as well as a science educator workshop and a product showcase within their exhibition hall.

Genomenon will be onsite at booth #1028, where we will be highlighting our product Mastermind, a comprehensive knowledge-base that takes the painstaking process of searching medical literature for disease-causing variants from hours to minutes by automatically sorting through the extensive genomic literature, which will be released in December.

If you would like to schedule a meeting with the Genomenon team, email sales@genomenon.com or visit www.genomenon.com for more information.

About Genomenon:
Genomenon's genome interpretation software is used for clinical diagnostics in cancer and genetic diseases. Genomenon reduces the time Pathologists and Geneticists spend researching and interpreting genetic variants from hours to minutes. Mastermind, our comprehensive knowledge-base, is built on mining millions of medical publications that automatically finds correlations between genes, variants and diseases tied to the primary scientific literature required for clinical diagnosis.

For more information, visit www.genomenon.com or email sales@genomenon.com.

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