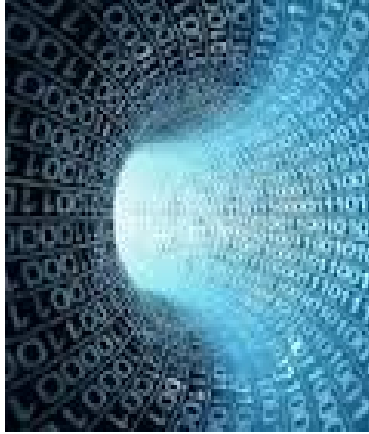# Dive deep into the Mouse Transcriptome

Genes, Exons, lincRNA, Isomers, microRNA and SNP's

Seminar by Ryan Estep, Field Application Scientist

Drowning in NGS data?
Delayed publishing?
Transcriptome assays
are your answer!

Go beyond the gene to see what you are missing…

- Differential gene expression from RNA to result in a week, at a low cost with high reproducibility.
- Using the right tools : Arrays vs RNA-Seq
- Isoforms matter, their specific expression too! Can you afford to miss these critical insights
- Equal to 480 RPM or 2 Full Lanes of RNA-Seq

Date: Wednesday, October 8, 2014

Time: 11:00 AM to 12:00 PM

Location: WRB 1413

Hosted by: Marti Veigl, Genomics Core

TAC software workshop: 12:30 to 2:00pm

Location: Wolstein Conference Room 3-136 (lunch served)

Please RSVP for TAC software workshop to Susan_Vigilante@affymetrix.com

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