



Giovedì 17 Maggio 2018 - ore 10:30 - 13:15

Aula 6 Torre Nord piano 0 – Centro Servizi BRIT

- 10:30-10:40 **Why and what to consider for optimizing RNA studies**
Speaker: **Cristina Valsecchi**, Sr.TSS GeneExpression
- 10:40-11:10 **Take the best of the transcriptome with Microarrays technology: miRNA, mRNA, alternative splicing and lncRNA. New Affymetrix solutions for GeneExpression**
Speaker: **Cristina Valsecchi**, Sr.TSS GeneExpression
- 11:10-11:20 **Data integration and Data Analysis with Transcriptome Analysis Console: demo**
Speaker: **Cristina Valsecchi**, Sr.TSS Gene Expression
- 11:20-11:40 *Coffee Break*
- 11:40-12:10 **RNA world and NGS from targeted RNA to WT**
Speaker: **Salvatore Conti**, NGS TSS
- 12:10-12:45 **Gene Expression advances on qPCR: increase throughput and sensitivity**
Speaker: **Cristina Valsecchi**, Sr.TSS GeneExpression
- 12:45-13:00 **Thermo Fisher Cloud: Free Gene Expression Data Analysis**
Speaker: **Serena Grimaldi**, Sr. qPCR/CE Field Application Scientist
- 13:00-13:15 Closing and remarks

The event is free of charge. Deadline for registration: within Friday 11th of May

For information: Dr. Pietro Geraci, mobile: 349 8323310

Registration & Logistics: pietro.geraci@thermofisher.com