

Invitation to NGS Seminars

Wednesday Nov the 16th

Kontinkangas and Linnanmaa campuses



At 10:00 – 11:00 – RNA-seq and microbiome enrichment

Linnanmaa campus, room B235

Do you need increased sensitivity and specificity from your RNA-seq experiments? Do you have ever-decreasing amounts of input RNA? – We will present NEBNext Ultra II RNA library prep, the gold standard in transcriptome analysis and learn how to facilitate enrichment of microbial DNA from samples containing methylated host DNA to enable microbiome whole genome sequencing, even for samples with high levels of host DNA.

At 12:15 – 13:15 – Overcoming bottlenecks in NGS and discover EM-seq

Kontinkangas campus, room 289B

Are you challenged with trying to obtain higher library yields using ever-decreasing input amounts? - We will present how to overcome bottlenecks in NGS using an optimized library preparation workflow and how enzymatic methyl sequencing is a superior solution in comparison to traditional bisulfite conversion for epigenetic & methylation studies.

Presenters: **Björn Textor**, Senior Application Manager at New England Biolabs

Klaus Tangsgaard, Application specialist at New England Biolabs

Register for the seminars by email to pia.ojala@bionordika.fi. Please mention if you have diet restrictions.



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