

## Basic Application Training for Real Time PCR (LightCycler System) Agenda

| Time        | Friday (22-06-2012)   | Saturday (23-06-2012)  | Sunday (24-06-2012)  |   |
|-------------|---|--|--|---|
| 09:00-09:30 | <b>Welcome</b> <ul style="list-style-type: none"> <li>Introduction / Training Aims</li> <li>PCR Basics (if necessary)</li> <li>Real Time PCR Applications</li> </ul>  | <b>Short Review</b>  | <b>Short Review</b>  |   |
| 09:30-10:30 | <b>Presentation</b> <ul style="list-style-type: none"> <li>The New LC System for Real Time PCR</li> <li>Software 4.1</li> <li>General Overview</li> <li>LC2.0 Disposables , Kits (reagents)</li> </ul>  | <b>Experimental Work</b> <ul style="list-style-type: none"> <li>Detection of human (CYP2C9*2 and CYP2C9*3) SNPs</li> <li>Pipetting and Programming ( 3 Groupwork )</li> <li>Start Run</li> </ul>   | <b>Experimental Work</b> <ul style="list-style-type: none"> <li>Swine Inf A /H1N1 Detection Set</li> <li>Pipetting and Programming ( 3 Group work)</li> <li>Start Run</li> <li>Presentation and Qualitative Detection</li> </ul> |   |
| 10:30-10:45 | <b>Break</b>  |  |  |   |
| 10:45-12:00 | <b>LightCycler Basics</b> (Presentation) <ul style="list-style-type: none"> <li>Assay Formats &amp; Dyes</li> </ul> <b>Presentation / Demonstration</b> <ul style="list-style-type: none"> <li>Programming</li> <li>Templates &amp; Macros</li> </ul> | <b>Lab Work</b><br><b>Melting Curve Analysis</b> <ul style="list-style-type: none"> <li>Simulate T<sub>M</sub>-Calling Experiment</li> </ul> <b>SW Demo: Export of Files</b><br><b>SW Demo: Import of files</b>  | <b>Analysis of Results by Participants</b><br><br><b>Type of Analysis</b> <ul style="list-style-type: none"> <li>Absolute Quantification &amp; Qualitative detection.</li> </ul>   | • |
| 12:15-13:15 | <b>Lunch Break</b>  |  |  |   |
| 13:15-15:15 | <b>Experimental Work</b> <ul style="list-style-type: none"> <li>Apply Macro and Run LC Color Compensation Set for the Multiplex HybProbe Probe Technology</li> <li>Analysis and Create a CCC Object</li> </ul>  | <b>Experimental Work</b> <ul style="list-style-type: none"> <li>Analysis of Experiments (CYP2C9*2*3) SNPs</li> <li>Identify Genotypes</li> <li>Presentation of Results by Participants</li> </ul> <b>Type of Analysis</b> <ul style="list-style-type: none"> <li>Melting Curve Analysis (Genotyping -T<sub>M</sub>-Calling)</li> </ul> | <b>Lab Work</b> <ul style="list-style-type: none"> <li>Apply the Absolute Quantification, Qualitative Detection and T<sub>M</sub>-Calling analysis for the Run</li> <li>Apply the Analysis by Participants (each)</li> </ul>     |   |
| 15:15-15:30 | <b>Break</b>  |  |  |   |
| 15:30-16:45 | <b>Experimental Work</b><br><br><b>DNA Extraction</b>   | <b>Experimental Work</b> <ul style="list-style-type: none"> <li>Apply Macro and Run LC Color Compensation</li> <li>Set for the Multiplex TaqMan Probe Technology</li> <li>Analysis and Create a CCC Object</li> <li>Apply the H1N1 assay Compensation</li> </ul>   | <b>Real Time PCR Troubleshooting</b> <ul style="list-style-type: none"> <li>General troubleshooting for the LightCycler2.0 as a Real Time PCR System</li> <li>Interpretation of troubleshooting for the Assays done</li> </ul>   |   |

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| 16:45-17:00 | Summary - Open Questions | Summary - Open Questions | Summary - Open Questions |  |
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