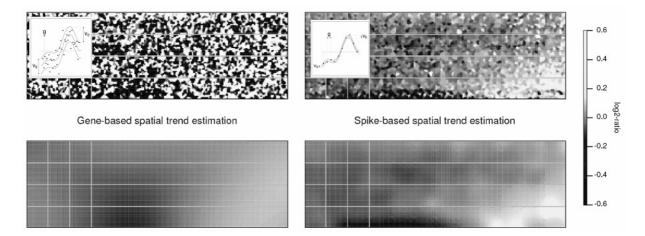
## **Tenfold improved performance of spatial normalization of microarrays**

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Spatial variance is typically 2-3 times stronger than residual noise in microarray experiments. Improved microarray designs facilitate an efficient removal of these signal distortions. While the use of dedicated probes for exogenous spikes is becoming more widespread, their exploitation for the removal of spatial artifacts has received less attention. In particular, their potential to systematically quanitfy spatial effect characteristics, has so far not been considered. Since better knowledge of the underlying processes will afford enhanced experimental and analytical approaches, this is a gap to be filled. We present a Gaussian process framework to determine dimensionless quantitative characteristics of spatial effects in microarrays and we have evaluated the performance of spatial normalization procedures exploiting normalization probes. While microarray real estate is costly we can demonstrate that using just 10% of probes for spikes a tenfold improved spatial normalization procedures relying on non-replicated gene probes.



<u>Acknowledgments:</u> The Boku Bioinformatics group acknowledges support by the Vienna Science and Technology Fund (WWTF), the Austrian Centre of Biopharmaceutical Technology (ACBT), Austrian Research Centres Seibersdorf (ARCS), and Baxter AG. Thomas Tüchler acknowledges partial support by the Austrian GEN-AU program. FlyChip acknowledges funding by the Biotechnology and Biological Science Research Council (BBSRC).