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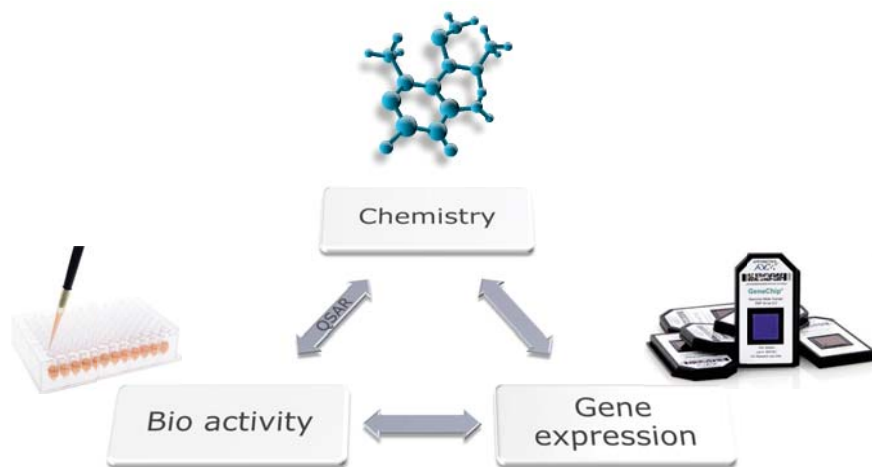
QSTAR Session - NCS2012 Potsdam

Quantitative Structure Transcriptional Activity Relationship



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Understanding New Molecules



QSTAR Session - NCS2012 Potsdam

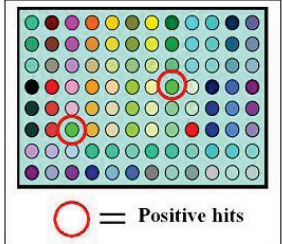
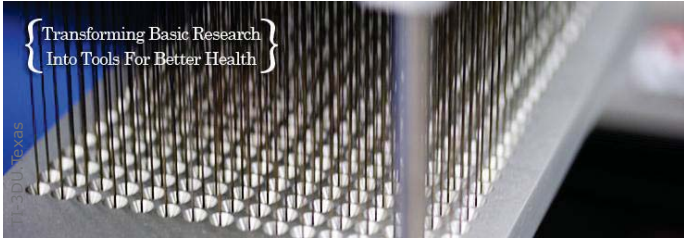
Quantitative Structure **Transcriptional** Activity Relationship

$$= \text{QSAR} + \text{T}$$

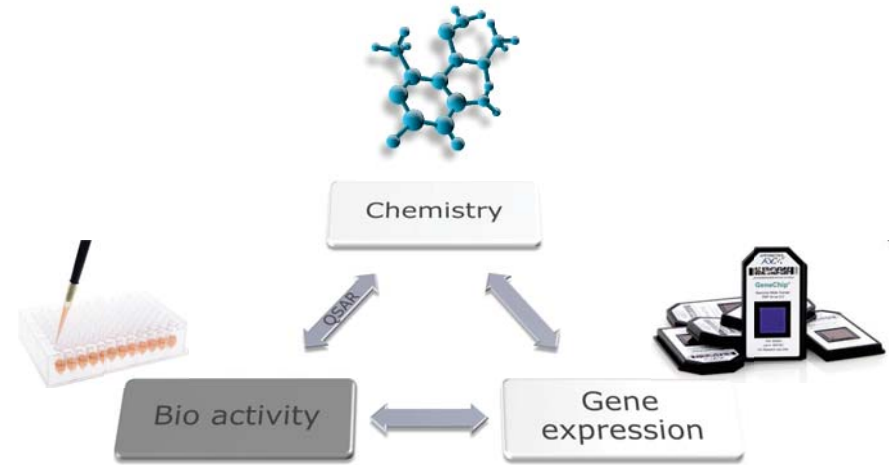


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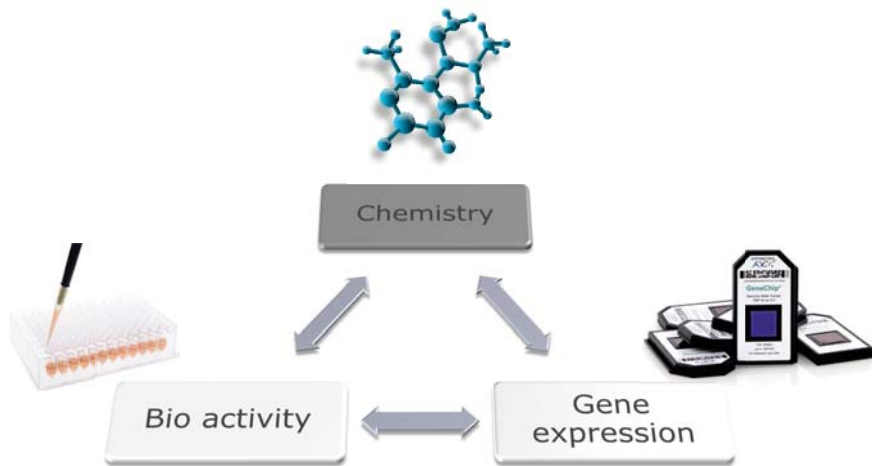
Bioassays – High Troughput Screening



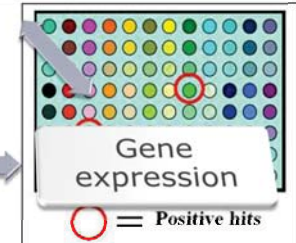
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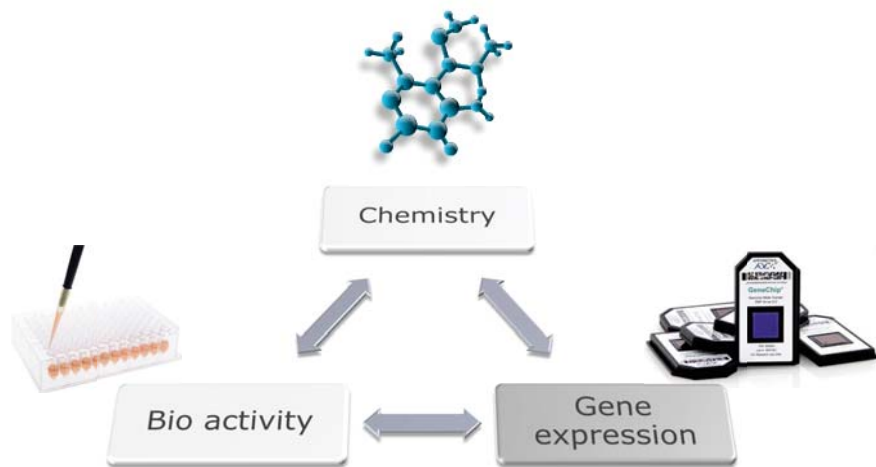
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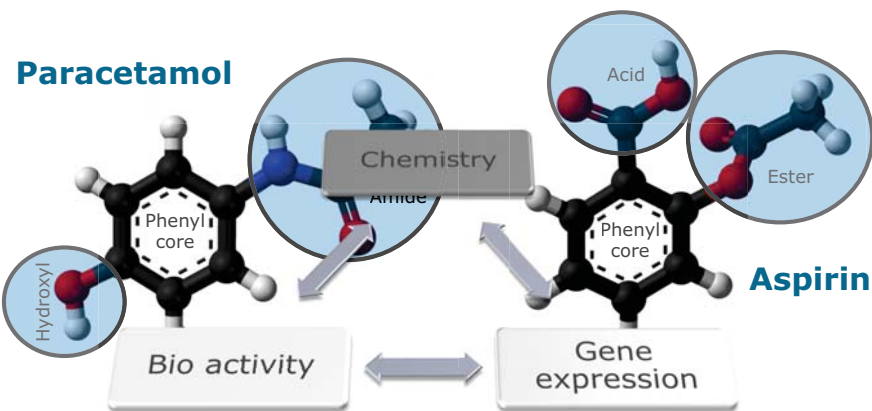
Bioassays – High Troughput Screening



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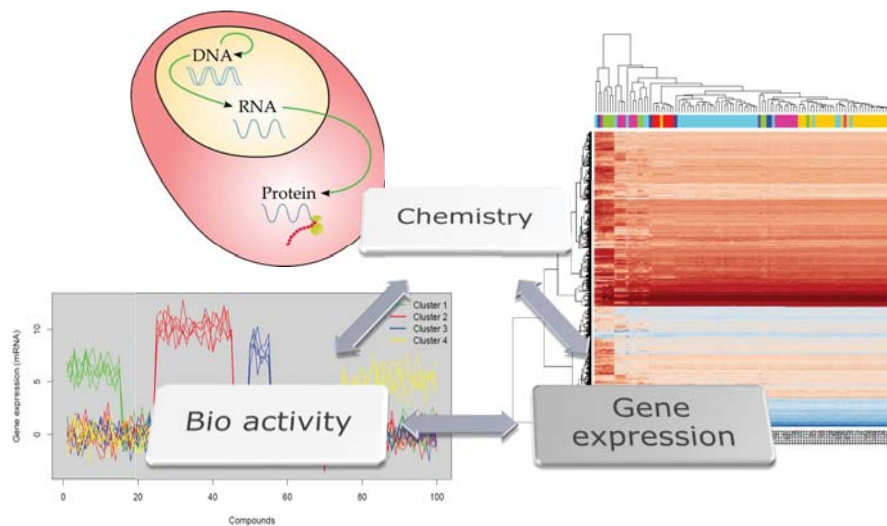


Chemical Structures

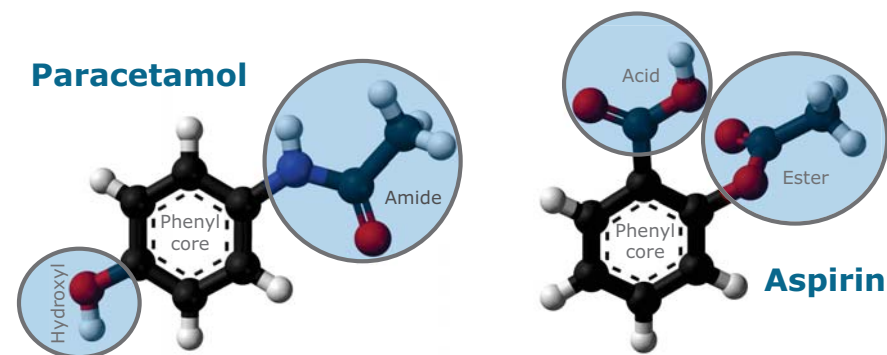


From Wikipedia

Gene Expression Profiling



Chemical Structures



From Wikipedia

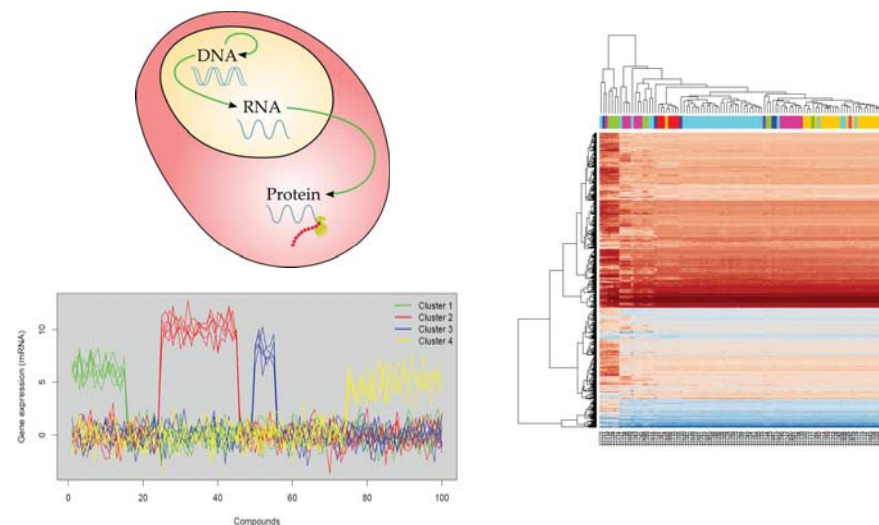
QSTAR Session Terminology

Gene Expression (GE)
gene 001, gene 002, ...

Bio-Activity (BA)

Chemistry (CH)
Compounds : cpd 001, cpd 002, ...
Finger print features : fpf 001, fpf 002, ...

Gene Expression Profiling



QSTAR Session Contents

Development of Predictive Models and Feature Selection Using LASSO and Elastic Net
Pushpike Thilakarathne

Analoging in large databases with structural fingerprint features
Martin Heusel

Integrated analysis of microarray data, biological data and chemical data in drug development
Nolen Joy Perualila

Biclustering in drug design
Djork-Arné Clevert

Integrated statistical analysis of three data sources for the detection of chemical fingerprint features that are simultaneously associated with bio-assay and transcriptomics data

Olivier Thas, Federico Mattiello

Understanding New Molecules

