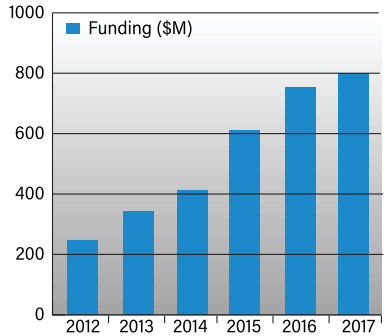
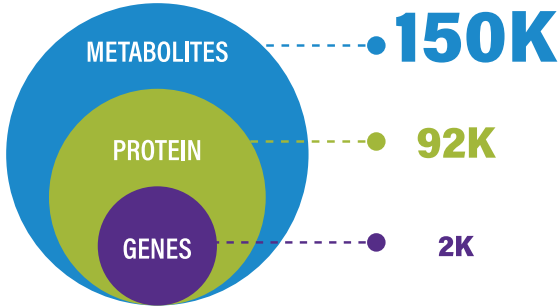


One Lab. So Many Answers.

Targeted Omics Method Library

The metabolomics space is **6 times** bigger than the genomics space



The amount of Metabolomics **funding is growing rapidly** (5YR CAGR = 25%)¹

Typical method development for a single targeted panel takes

8 months

and costs

\$100,000

+ opportunity costs²

The increased interest in targeted metabolomics studies means **method development will be a major bottleneck** for core labs!

11.7 projects per core FTE (2015) **up from 7.6** per FTE (2014)

54% of labs saw an increase in the number of customers

Waters Targeted Omics Method Library

contains downloadable method files and app notes for multi-analyte omics panels



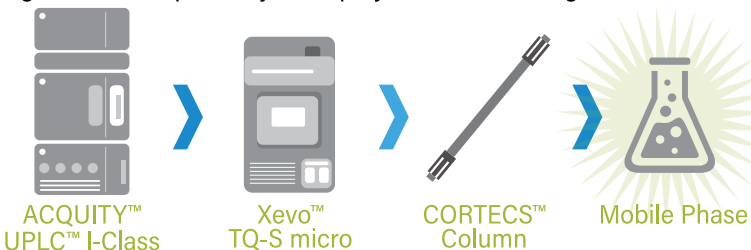
We are **empowered!**



It empowers labs by providing wide analyte coverage and fast method turnaround. Check it out at www.waters.com/targetedomics

Fast Forward to Results

Within the Targeted Omics Method Library we offer rapid (-R) method packages. All methods with an (-R) designation are rapid, easy-to-deploy methods sharing a common workflow:



That means each system can run multiple different analyses without changing columns and mobile phases resulting in less downtime and more coverage for your lab!

DOWNLOAD > IMPORT > GO!

Sources: 1: <https://projectreporter.nih.gov/reporter.cfm> 2: Internal poll of method development scientists

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