

Resource Summary Report

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Q Squared Solutions Expression Analysis

RRID:SCR_012497

Type: Tool

Proper Citation

Q Squared Solutions Expression Analysis (RRID:SCR_012497)

Resource Information

URL: <https://www.q2labsolutions.com/genomics-laboratories>

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Description: Core provides whole genome to focused set gene expression and genotyping assays along with DNA sequencings services, sequence enrichment technologies and bioinformatics support. Platforms utilized include Affymetrix GeneChip, Agilent Sure Select, Fluidigm Access Arrays, Illumina BeadChip, iScan, Genome Analyzer and Hi-Seq, RainDance Technologies RDT 1000 and, the Pacific Biosciences PacBio RS. Expression Analysis offers solutions for challenging specimens such as whole blood and FFPE tissues, as well as nucleic acid isolation and data analysis services.

Abbreviations: Q2 Solutions, EA, Q squared solutions, Q2, Q squared

Synonyms: Q2 Solutions Expression Analysis, Q 2 Solutions Expression Analysis, Q 2 Expression Analysis

Resource Type: access service resource, commercial organization, core facility, service resource

Keywords: genomic, genotyping, DNA sequencing,

Availability: Available to external user

Resource Name: Q Squared Solutions Expression Analysis

Resource ID: SCR_012497

Alternate IDs: SciEx_366

Alternate URLs: <http://www.scienceexchange.com/facilities/expression-analysis>

Record Creation Time: 20220129T080310+0000

Record Last Update: 20260905T132724+0000

Ratings and Alerts

No rating or validation information has been found for Q Squared Solutions Expression Analysis.

No alerts have been found for Q Squared Solutions Expression Analysis.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Iqbal M, et al. (2013) Expression of HMA4 cDNAs of the zinc hyperaccumulator *Noccaea caerulescens* from endogenous NcHMA4 promoters does not complement the zinc-deficiency phenotype of the *Arabidopsis thaliana* hma2hma4 double mutant. *Frontiers in plant science*, 4, 404.