

Resource Summary Report

Generated by SPARC SAWG on Sep 14, 2026

ContigExpress

RRID:SCR_012276

Type: Tool

Proper Citation

ContigExpress (RRID:SCR_012276)

Resource Information

URL: https://tools.thermofisher.com/content/sfs/manuals/VectorNTI_UserManual.pdf

Proper Citation: ContigExpress (RRID:SCR_012276)

Description: A genomics data service provider offering genomic project management, data analysis, and bio-IT consulting services.

Abbreviations: ContigExpress

Resource Type: commercial organization, software application, software resource

Resource Name: ContigExpress

Resource ID: SCR_012276

Alternate IDs: SciEx_11277

Old URLs: <http://www.scienceexchange.com/facilities/contigexpress>

Record Creation Time: 20220129T080309+0000

Record Last Update: 20260912T195747+0000

Ratings and Alerts

No rating or validation information has been found for ContigExpress.

No alerts have been found for ContigExpress.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 198 mentions in open access literature.

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

Liu R, et al. (2025) A special short-wing petal faba genome and genetic dissection of floral and yield-related traits accelerate breeding and improvement of faba bean. *Genome biology*, 26(1), 62.

Zhang W, et al. (2025) Evaluation of different ecological regions for cultivation of best quality *Bupleurum*: a case study from Shanxi, China. *BMC plant biology*, 25(1), 542.

Meng Q, et al. (2025) ?*Lijiangomyces laojunensis* gen. et sp. nov. (Mytiliniaceae), and *Sclerococcum stictiae* (Dactylosporaceae), a new lichenicolous species from Yunnan, China. *MycKeys*, 114, 277.

Jiang X, et al. (2024) Gamma-glutamyl transferase secreted by *Helicobacter pylori* promotes the development of gastric cancer by affecting the energy metabolism and histone methylation status of gastric epithelial cells. *Cell communication and signaling : CCS*, 22(1), 402.

Yang Z, et al. (2024) Morphological and molecular characterization of *Heterodera ripae*, a new record cyst nematode in the rhizosphere soil of *Fagopyrum esculentum*. *Scientific reports*, 14(1), 9958.

Cheng Z, et al. (2024) Partial Correspondence between Host Plant-Related Differentiation and Symbiotic Bacterial Community in a Polyphagous Insect. *Animals : an open access journal from MDPI*, 14(2).

Yu J, et al. (2024) Whole-Genome Sequencing of the Entomopathogenic Fungus *Fusarium solani* KMZW-1 and Its Efficacy Against *Bactrocera dorsalis*. *Current issues in molecular biology*, 46(10), 11593.

Shang LH, et al. (2024) Role of Gut Bacteria in Enhancing Host Adaptation of *Tuta absoluta* to Different Host Plants. *Insects*, 15(10).

Tedesco P, et al. (2024) Morphological and molecular data on acuariid nematodes in European great cormorants (*Phalacrocorax carbo sinensis*) and pygmy cormorants (*Microcarbo pygmaeus*). *Scientific reports*, 14(1), 13712.

Li CJ, et al. (2024) ?Additional four species of *Tatraea* (Leotiomyces, Helotiales) in Yunnan Province, China. *MycKeys*, 102, 127.

Li T, et al. (2024) Characterization and genomic analysis of *Bacillus megaterium* with the ability to degrade aflatoxin B1. *Frontiers in microbiology*, 15, 1407270.

Zhou B, et al. (2024) "Point by point" source: The Chinese pine plantations in North China by evidence from mtDNA. *Ecology and evolution*, 14(6), e11570.

Promrug D, et al. (2023) Cocultures of *Enterococcus faecium* and *Aeromonas veronii* Induce the Secretion of Bacteriocin-like Substances against *Aeromonas*. *Journal of*

agricultural and food chemistry, 71(43), 16194.

Kim S, et al. (2023) Application of a Quantitative PCR to Investigate the Distribution and Dynamics of Two Morphologically Similar Species, *Karenia mikimotoi* and *K. papilionacea* (Dinophyceae) in Korean Coastal Waters. *Toxins*, 15(7).

Romey A, et al. (2023) Field Evaluation of a Safe, Easy, and Low-Cost Protocol for Shipment of Samples from Suspected Cases of Foot-and-Mouth Disease to Diagnostic Laboratories. *Transboundary and emerging diseases*, 2023, 9555213.

Motta G, et al. (2023) Parasitic infection in the scyphozoan *Rhizostoma pulmo* (Macri, 1778). *Scientific reports*, 13(1), 5549.

Feng Y, et al. (2023) CRISPR/Cas9-mediated SNAC9 mutants reveal the positive regulation of tomato ripening by SNAC9 and the mechanism of carotenoid metabolism regulation. *Horticulture research*, 10(4), uhad019.

Zhong J, et al. (2023) A point mutation in MC06g1112 encoding FLOWERING LOCUS T decreases the first flower node in bitter melon (*Momordica charantia* L.). *Frontiers in plant science*, 14, 1153208.

Krivitskaya V, et al. (2023) Characterization of a Panel of Monoclonal Antibodies Targeting the F-Protein of the Respiratory Syncytial Virus (RSV) for the Typing of Contemporary Circulating Strains. *Tropical medicine and infectious disease*, 9(1).

Yan WL, et al. (2023) The Study of Forensically Important Insects Recovered from Human Corpses in Taiwan. *Insects*, 14(4).