

Resource Summary Report

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Bio Basic Inc

RRID:SCR_012675

Type: Tool

Proper Citation

Bio Basic Inc (RRID:SCR_012675)

Resource Information

URL: <http://www.scienceexchange.com/facilities/bio-basic-inc>

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Description: Bio Basic Inc. is a privately owned dynamic biotechnology company. The company was founded in 1990 in Toronto, Canada. From 1990 to 1995, Bio Basic Inc.'s primary focus has been in the field of biochemicals. Starting in 1995, Bio Basic Inc. has begun manufacturing various Life Science Products. Throughout the past years, the company has developed rapidly and served as a one-stop-shop to our customers in the life sciences field. To date, Bio Basic Inc. has approximately 600 employees, seven laboratories, three factory buildings, 40 international distributors and over 10,000 customers worldwide. Bio Basic Inc. is committed to innovation and excellence. In an effort to achieve this, Bio Basic Inc. has continuously directed efforts in R & D and improving the quality of our products. Bio Basic Inc. is proud to offer a breath of products and services to customers worldwide. Several of our product lines such as, Molecular Biology Kits, Biochemicals, Synthetic Genes and Synthetic Oligos have already shown dominant market presence and have been widely recognized by customers worldwide.

Abbreviations: BBI

Synonyms: Bio Basic, Bio Basic Inc., Bio Basic Canada Inc., Bio Basic Canada Inc, Bio Basic Canada

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

Resource Name: Bio Basic Inc

Resource ID: SCR_012675

Alternate IDs: SciEx_8824

Record Creation Time: 20220129T080311+0000

Ratings and Alerts

No rating or validation information has been found for Bio Basic Inc.

No alerts have been found for Bio Basic Inc.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Martínez-Armenta CA, et al. (2026) Evaluation of the CIB1R peptide derived from the cytoplasmic domain of neprilysin on cell migration in an in vitro model of lung cancer. *Frontiers in oncology*, 16, 1825515.

Tsuji S, et al. (2018) Real-time multiplex PCR for simultaneous detection of multiple species from environmental DNA: an application on two Japanese medaka species. *Scientific reports*, 8(1), 9138.

Zhao L, et al. (2017) Genetic variation of *Sporothrix globosa* isolates from diverse geographic and clinical origins in China. *Emerging microbes & infections*, 6(10), e88.

Hanapi UF, et al. (2017) Tracking the virus-like particles of *Macrobrachium rosenbergii* nodavirus in insect cells. *PeerJ*, 5, e2947.

Ceccoli RD, et al. (2017) Cloning and characterization of the Type I Baeyer-Villiger monooxygenase from *Leptospira biflexa*. *AMB Express*, 7(1), 87.

Law KB, et al. (2017) The peroxisomal AAA ATPase complex prevents pexophagy and development of peroxisome biogenesis disorders. *Autophagy*, 13(5), 868.

Breton S, et al. (2017) The extremely divergent maternally- and paternally-transmitted mitochondrial genomes are co-expressed in somatic tissues of two freshwater mussel species with doubly uniparental inheritance of mtDNA. *PloS one*, 12(8), e0183529.

Rocco BR, et al. (2017) Fluorescence-based cell-specific detection for laser-capture microdissection in human brain. *Scientific reports*, 7(1), 14213.

Aziz F, et al. (2017) Novel route synthesis of porous and solid gold nanoparticles for investigating their comparative performance as contrast agent in computed tomography scan and effect on liver and kidney function. *International journal of nanomedicine*, 12, 1555.

Hong Y, et al. (2017) Role of melatonin combined with exercise as a switch-like regulator for circadian behavior in advanced osteoarthritic knee. *Oncotarget*, 8(57), 97633.

Zeng W, et al. (2016) Improvement of *Bacillus subtilis* for poly- γ -glutamic acid production by genome shuffling. *Microbial biotechnology*, 9(6), 824.

Gusman A, et al. (2016) Pursuing the quest for better understanding the taxonomic distribution of the system of doubly uniparental inheritance of mtDNA. *PeerJ*, 4, e2760.