

Resource Summary Report

Generated by T1D on Aug 9, 2026

Multi-Disciplinary Approach to the Study of Chronic Pelvic Pain

RRID:SCR_013754

Type: Tool

Proper Citation

Multi-Disciplinary Approach to the Study of Chronic Pelvic Pain (RRID:SCR_013754)

Resource Information

URL: <http://mappnetwork.org>

Proper Citation: Multi-Disciplinary Approach to the Study of Chronic Pelvic Pain (RRID:SCR_013754)

Description: Collaborative research network specializing in urological chronic pelvic pain disorders. Project involves conducting research primarily on interstitial cystitis/painful bladder syndrome and chronic prostatitis/chronic pelvic pain syndrome and involves researchers from multiple disciplines. MAPP Network includes researchers with clinical, epidemiological, and basic research expertise.

Abbreviations: MAPP

Synonyms: Multidisciplinary Approach to the Study of Chronic Pelvic Pain Research Network, MAPP Research Network

Resource Type: data or information resource, portal, disease-related portal, topical portal

Keywords: interstitial cystitis, painful bladder syndrome, chronic prostatitis, chronic pelvic pain syndrome, chronic pelvic pain

Related Condition: interstitial cystitis/painful bladder syndrome, IC/PBS, chronic prostatitis/chronic pelvic pain syndrome, CP/CPPS

Availability: Free, Freely available

Resource Name: Multi-Disciplinary Approach to the Study of Chronic Pelvic Pain

Resource ID: SCR_013754

Alternate IDs: SCR_014375

Record Creation Time: 20220129T080317+0000

Record Last Update: 20260808T190016+0000

Ratings and Alerts

No rating or validation information has been found for Multi-Disciplinary Approach to the Study of Chronic Pelvic Pain.

No alerts have been found for Multi-Disciplinary Approach to the Study of Chronic Pelvic Pain.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 189 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Priya M, et al. (2026) Computational Prediction of Deleterious SNPs in the GALNS Gene Implicated in Morquio A Syndrome (MPS IVA). ACS omega, 11(7), 11183.

Wolff DT, et al. (2026) Correlates of Positive Response to Therapeutic Hydrodistension in Interstitial Cystitis/Bladder Pain Syndrome. Neurourology and urodynamics, 45(1), 169.

Wu C, et al. (2026) Molecular Mechanisms of CLCN5 Missense Mutations in Dent Disease Type 1: A Comprehensive Computational Analysis and Clinical Correlations in a Chinese Cohort. Journal of cellular and molecular medicine, 30(6), e71108.

Dissanayake D, et al. (2026) Predicting pyrazinamide resistance in Mycobacterium tuberculosis using a graph convolutional network. BMC microbiology, 26(1).

Homer SR, et al. (2026) Designing a systemic intervention for student loneliness and social connectedness using a mixed-methods, co-creation approach. Npj mental health research, 5(1).

Rafi MO, et al. (2026) Predicting the Impact of Deleterious Single-Nucleotide Polymorphisms in the p47ING1a Isoform of Human ING1 Gene. Genetics research, 2026(1), e2859448.

Azmi MB, et al. (2026) Computational identification of rare pathogenic genomic variants in esophageal cancer markers: Transcript-level analysis, sequence-based insights, and structural-functional impacts of non-synonymous SNPs. Biochemistry and biophysics reports, 45, 102503.

Grigoriadis S, et al. (2026) A Brief Intervention for the Treatment of Anxiety in Pregnancy: A Pilot Randomized Controlled Trial (The TAP Study). Psychotherapy and

psychosomatics, 1.

Nila NN, et al. (2026) Comprehensive analysis of the deleterious nonsynonymous, non-coding SNPs and cancer variants of human aldo-keto reductase type 1 (AKR1C1) protein and their probable association with disease risk and progression: a computational study. *Biochemistry and biophysics reports*, 45, 102502.

Akçe?me B, et al. (2025) Identification of deleterious non-synonymous single nucleotide polymorphisms in the mRNA decay activator ZFP36L2. *RNA biology*, 22(1), 1.

Naliboff BD, et al. (2025) Relationship of Sex and Diagnosis With Symptoms and Illness Impact in Urologic Chronic Pelvic Pain; A Mapp Network Analysis. *Neurourology and urodynamics*, 44(2), 400.

Habib O, et al. (2025) Structural and immunological impacts of TOLLIP nsSNPs: A computational biology approach to drug discovery and immune system modulation. *PloS one*, 20(11), e0328573.

Jacinto J, et al. (2025) Exploring skeletal disorders in cattle and sheep: a WGS-based framework for diagnosis and classification. *Genetics, selection, evolution : GSE*, 57(1), 51.

Bayat S, et al. (2025) Expanding the Phenotype and Genotype Spectrum of a Novel Mutation in Hypomyelinating Leukodystrophy-5 With a Review of the Literature on 42 Cases. *Basic and clinical neuroscience*, 16(2), 505.

Guan Y, et al. (2025) Population frequency of Predicted pathogenic MisMatch Repair (MMR) gene variants in Lynch syndrome from bioinformatic analyses of the general population. *Scientific reports*, 15(1), 34545.

Rodrigues MLA, et al. (2025) Challenges and perceptions of dental undergraduate students regarding active learning in clinical practice: A qualitative study. *PloS one*, 20(9), e0328939.

Sultan R, et al. (2024) 3D printing of polypropylene reinforced with hemp fibers: Mechanical, water absorption and morphological properties. *Heliyon*, 10(4), e26617.

Bak M, et al. (2024) MAPP unravels frequent co-regulation of splicing and polyadenylation by RNA-binding proteins and their dysregulation in cancer. *Nature communications*, 15(1), 4110.

?astorálová M, et al. (2024) A myristoyl switch at the plasma membrane triggers cleavage and oligomerization of Mason-Pfizer monkey virus matrix protein. *eLife*, 13.

Nila NN, et al. (2024) Investigating the structural and functional consequences of germline single nucleotide polymorphisms located in the genes of the alternative lengthening of telomere (ALT) pathway. *Heliyon*, 10(12), e33110.