

Timothy F Havel

List of Publications by Year in descending order

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Version: 2024-02-01

50
papers

4,468
citations

270111

25
h-index

242451

47
g-index

55
all docs

55
docs citations

55
times ranked

2819
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Storing energy and powering small systems with mechanical springs made of carbon nanotube yarn. <i>Energy</i> , 2014, 76, 318-325. | 4.5 | 16 |
| 2 | Enhancing the Tensile Properties of Continuous Millimeter-Scale Carbon Nanotube Fibers by Densification. <i>ACS Applied Materials & Interfaces</i> , 2013, 5, 7198-7207. | 4.0 | 36 |
| 3 | Signatures of Incoherence in a Quantum Information Processor. <i>Quantum Information Processing</i> , 2007, 6, 431-444. | 1.0 | 4 |
| 4 | Quantum Control of Nuclear Spins. , 2006, , . | | 0 |
| 5 | Reflection symmetries for multiqubit density operators. <i>Journal of Mathematical Physics</i> , 2006, 47, 032104. | 0.5 | 4 |
| 6 | Solid-state NMR three-qubit homonuclear system for quantum-information processing: Control and characterization. <i>Physical Review A</i> , 2006, 73, . | 1.0 | 35 |
| 7 | Selective coherence transfers in homonuclear dipolar coupled spin systems. <i>Physical Review A</i> , 2005, 71, . | 1.0 | 14 |
| 8 | Geometric algebra and transition-selective implementations of the controlled-NOT gate. <i>Concepts in Magnetic Resonance</i> , 2004, 23A, 49-62. | 1.3 | 2 |
| 9 | Quantum process tomography of the quantum Fourier transform. <i>Journal of Chemical Physics</i> , 2004, 121, 6117-6133. | 1.2 | 131 |
| 10 | Experimental Simulation of Spin Squeezing by Nuclear Magnetic Resonance. <i>Quantum Information Processing</i> , 2003, 2, 433-448. | 1.0 | 14 |
| 11 | Metric matrix embedding in protein structure calculations, NMR spectra analysis, and relaxation theory. <i>Magnetic Resonance in Chemistry</i> , 2003, 41, S37-S50. | 1.1 | 10 |
| 12 | Robust control of quantum information. <i>Journal of Chemical Physics</i> , 2003, 119, 9993-10001. | 1.2 | 81 |
| 13 | Robust procedures for converting among Lindblad, Kraus and matrix representations of quantum dynamical semigroups. <i>Journal of Mathematical Physics</i> , 2003, 44, 534. | 0.5 | 113 |
| 14 | Exploring noiseless subsystems via nuclear magnetic resonance. <i>Physical Review A</i> , 2003, 67, . | 1.0 | 25 |
| 15 | The Role of Tensegrity in Distance Geometry. , 2002, , 55-68. | | 0 |
| 16 | Design of strongly modulating pulses to implement precise effective Hamiltonians for quantum information processing. <i>Journal of Chemical Physics</i> , 2002, 116, 7599-7606. | 1.2 | 204 |
| 17 | The Real Density Matrix. <i>Quantum Information Processing</i> , 2002, 1, 511-538. | 1.0 | 12 |
| 18 | Interaction and Entanglement in the Multiparticle Spacetime Algebra. , 2002, , 227-247. | | 2 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Hadamard products of product operators and the design of gradient-diffusion experiments for simulating decoherence by NMR spectroscopy. <i>Physics Letters, Section A: General, Atomic and Solid State Physics</i> , 2001, 280, 282-288. | 0.9 | 29 |
| 20 | Multiqubit logic gates in NMR quantum computing. <i>New Journal of Physics</i> , 2000, 2, 10-10. | 1.2 | 22 |
| 21 | Spatially encoded pseudopure states for NMR quantum-information processing. <i>Physical Review A</i> , 2000, 62, . | 1.0 | 19 |
| 22 | Quantum codes for controlling coherent evolution. <i>Journal of Chemical Physics</i> , 2000, 113, 10878-10885. | 1.2 | 2 |
| 23 | Generalized methods for the development of quantum logic gates for an NMR quantum information processor. <i>Physical Review A</i> , 1999, 60, 2777-2780. | 1.0 | 56 |
| 24 | Observations of quantum dynamics by solution-state NMR spectroscopy. <i>Concepts in Magnetic Resonance</i> , 1999, 11, 225-238. | 1.3 | 25 |
| 25 | Expressing the operations of quantum computing in multiparticle geometric algebra. <i>Physics Letters, Section A: General, Atomic and Solid State Physics</i> , 1998, 240, 1-7. | 0.9 | 66 |
| 26 | Nuclear magnetic resonance spectroscopy: An experimentally accessible paradigm for quantum computing. <i>Physica D: Nonlinear Phenomena</i> , 1998, 120, 82-101. | 1.3 | 339 |
| 27 | Embedding with a rigid substructure. <i>Journal of Mathematical Chemistry</i> , 1997, 21, 223-260. | 0.7 | 3 |
| 28 | A Robust Method for Estimating Cross-Relaxation Rates from Simultaneous Fits to Build-Up and Decay Curves. <i>Journal of Magnetic Resonance</i> , 1997, 124, 372-382. | 1.2 | 7 |
| 29 | Recent advances in molecular distance geometry. <i>Lecture Notes in Computer Science</i> , 1996, , 62-71. | 1.0 | 2 |
| 30 | Applications of geometric algebra to the theory of molecular conformation 2. The local deformation problem. <i>Computational and Theoretical Chemistry</i> , 1995, 336, 175-189. | 1.5 | 6 |
| 31 | Geometric Algebra and Möbius Sphere Geometry as a Basis for Euclidean Invariant Theory. , 1995, , 245-256. | | 14 |
| 32 | An evaluation of least-squares fits to COSY spectra as a means of estimating proton-proton coupling constants. I. Simulated test problems. <i>Journal of Biomolecular NMR</i> , 1994, 4, 807-826. | 1.6 | 10 |
| 33 | An evaluation of least-squares fits to COSY spectra as a means of estimating proton-proton coupling constants II. Applications to polypeptides. <i>Journal of Biomolecular NMR</i> , 1994, 4, 827-844. | 1.6 | 5 |
| 34 | Applications of geometric algebra to the theory of molecular conformation. <i>Computational and Theoretical Chemistry</i> , 1994, 308, 241-262. | 1.5 | 6 |
| 35 | Distance geometry and geometric algebra. <i>Foundations of Physics</i> , 1993, 23, 1357-1374. | 0.6 | 56 |
| 36 | The solution structure of eglin c based on measurements of many NOEs and coupling constants and its comparison with X-ray structures. <i>Protein Science</i> , 1992, 1, 736-751. | 3.1 | 411 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 37 | Bound Smoothing Under Chirality Constraints. <i>SIAM Journal on Discrete Mathematics</i> , 1991, 4, 535-549. | 0.4 | 9 |
| 38 | Alternating zinc fingers in the human male associated protein ZFY: 2D NMR structure of an even finger and implications for jumping-linker DNA recognition. <i>Biochemistry</i> , 1991, 30, 3371-3386. | 1.2 | 75 |
| 39 | A new method for building protein conformations from sequence alignments with homologues of known structure. <i>Journal of Molecular Biology</i> , 1991, 217, 1-7. | 2.0 | 132 |
| 40 | Some examples of the use of distances as coordinates for Euclidean geometry. <i>Journal of Symbolic Computation</i> , 1991, 11, 579-593. | 0.5 | 72 |
| 41 | The sampling properties of some distance geometry algorithms applied to unconstrained polypeptide chains: A study of 1830 independently computed conformations. <i>Biopolymers</i> , 1990, 29, 1565-1585. | 1.2 | 126 |
| 42 | Computational experience with an algorithm for tetrahedron inequality bound smoothing. <i>Bulletin of Mathematical Biology</i> , 1989, 51, 173-194. | 0.9 | 32 |
| 43 | Shortest-path problems and molecular conformation. <i>Discrete Applied Mathematics</i> , 1988, 19, 129-144. | 0.5 | 37 |
| 44 | Protein structures in solution by nuclear magnetic resonance and distance geometry. <i>Journal of Molecular Biology</i> , 1987, 196, 611-639. | 2.0 | 646 |
| 45 | An evaluation of the combined use of nuclear magnetic resonance and distance geometry for the determination of protein conformations in solution. <i>Journal of Molecular Biology</i> , 1985, 182, 281-294. | 2.0 | 383 |
| 46 | Solution conformation of proteinase inhibitor IIA from bull seminal plasma by ¹ H nuclear magnetic resonance and distance geometry. <i>Journal of Molecular Biology</i> , 1985, 182, 295-315. | 2.0 | 599 |
| 47 | The theory and practice of distance geometry. <i>Bulletin of Mathematical Biology</i> , 1983, 45, 665-720. | 0.9 | 324 |
| 48 | The combinatorial distance geometry method for the calculation of molecular conformation. I. A new approach to an old problem. <i>Journal of Theoretical Biology</i> , 1983, 104, 359-381. | 0.8 | 52 |
| 49 | The combinatorial distance geometry method for the calculation of molecular conformation II. Sample problems and computational statistics. <i>Journal of Theoretical Biology</i> , 1983, 104, 383-400. | 0.8 | 23 |
| 50 | Effects of distance constraints on macromolecular conformation. II. Simulation of experimental results and theoretical predictions. <i>Biopolymers</i> , 1979, 18, 73-81. | 1.2 | 155 |