Scott P Narus

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11959271/publications.pdf

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| | | 1040056 | 940533 |
|----------|----------------|--------------|----------------|
| 19 | 266 | 9 | 16 |
| papers | citations | h-index | g-index |
| | | | |
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| | | | |
| 19 | 19 | 19 | 425 |
| all docs | docs citations | times ranked | citing authors |
| | | | |

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Comprehensive methodology to monitor longitudinal change patterns during EHR implementations: a case study at a large health care delivery network. Journal of Biomedical Informatics, 2018, 83, 40-53. | 4.3 | 11 |
| 2 | Development and classification of a robust inventory of near real-time outcome measurements for assessing information technology interventions in health care. Journal of Biomedical Informatics, 2017, 73, 62-75. | 4.3 | 6 |
| 3 | Health information technology adoption: Understanding research protocols and outcome measurements for IT interventions in health care. Journal of Biomedical Informatics, 2016, 63, 33-44. | 4.3 | 19 |
| 4 | Longitudinal Analysis of Computerized Alerts for Laboratory Monitoring of Post-liver Transplant Immunosuppressive Care. AMIA Annual Symposium proceedings, 2015, 2015, 1918-26. | 0.2 | 1 |
| 5 | A domain analysis model for eIRB systems: Addressing the weak link in clinical research informatics. Journal of Biomedical Informatics, 2014, 52, 121-129. | 4.3 | 8 |
| 6 | A Focus Area Maturity Model for a Statewide Master Person Index. Online Journal of Public Health Informatics, 2013, 5, 210. | 0.7 | 4 |
| 7 | Utility of gene-specific algorithms for predicting pathogenicity of uncertain gene variants. Journal of the American Medical Informatics Association: JAMIA, 2012, 19, 207-211. | 4.4 | 25 |
| 8 | Consensus: a framework for evaluation of uncertain gene variants in laboratory test reporting. Genome Medicine, 2012, 4, 48. | 8.2 | 10 |
| 9 | Federating clinical data from six pediatric hospitals: process and initial results for microbiology from the PHIS+ consortium. AMIA Annual Symposium proceedings, 2012, 2012, 281-90. | 0.2 | 8 |
| 10 | Federated Querying Architecture with Clinical & Translational Health IT Application. Journal of Medical Systems, 2011, 35, 1211-1224. | 3.6 | 19 |
| 11 | Federated querying architecture for clinical & translational health IT. , 2010, , . | | 3 |
| 12 | Computational Feature Selection and Classification of RET Phenotypic Severity. Journal of Data Mining in Genomics $\&$ Proteomics, 2010, 01, . | 0.5 | 5 |
| 13 | Integrating a Federated Healthcare Data Query Platform With Electronic IRB Information Systems. AMIA Annual Symposium proceedings, 2010, 2010, 291-5. | 0.2 | 2 |
| 14 | Architecture of a federated query engine for heterogeneous resources. AMIA Annual Symposium proceedings, 2009, 2009, 70-4. | 0.2 | 11 |
| 15 | Computerized Alerts Improve Outpatient Laboratory Monitoring of Transplant Patients. Journal of the American Medical Informatics Association: JAMIA, 2008, 15, 324-332. | 4.4 | 29 |
| 16 | Effectiveness of Topic-specific Infobuttons: A Randomized Controlled Trial. Journal of the American Medical Informatics Association: JAMIA, 2008, 15, 752-759. | 4.4 | 62 |
| 17 | System analysis and improvement in the process of transplant patient care. Studies in Health Technology and Informatics, 2007, 129, 915-9. | 0.3 | 5 |
| 18 | A Case for Manual Entry of Structured, Coded Laboratory Data from Multiple Sources into an Ambulatory Electronic Health Record. Journal of the American Medical Informatics Association: JAMIA, 2006, 13, 12-15. | 4.4 | 19 |

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| # | # | Article | IF | CITATIONS |
|---|----|---|-----|-----------|
| 1 | 19 | Using LOINC to link an EMR to the pertinent paragraph in a structured reference knowledge base. Proceedings, 2002, , 652-6. | 0.6 | 19 |