

planetmath.org

Math for the people, by the people.

Jan Łukasiewicz

Canonical name JanLukasiewicz
Date of creation 2013-03-22 16:10:12
Last modified on 2013-03-22 16:10:12
Owner Mravinci (12996)
Last modified by Mravinci (12996)

Numerical id 17

Author Mravinci (12996)

Entry type Definition
Classification msc 03G12
Classification msc 03G30
Classification msc 03G10
Classification msc 92B05
Classification msc 01A60
Classification msc 03G20

Synonym Jan Łukasiewicz Synonym Jan Łukasiewicz Synonym Jan Łukasiewicz

Related topic GeneticNetsOrNetworks

Related topic AlgebraicCategoryOfLMnLogicAlgebras
Related topic AnalyticsAndOntologyFormalLogics
Related topic FormalLogicsAndMetaMathematics

Jan Łukasiewicz (1878 - 1956) Polish mathematician and logician mainly concerned with logic in mathematics probably best known for the threevalued logics, the Polish notation, the law of excluded middle and the axiomatizations of much classical propositional logic. He studied at the University of Lwów, earning doctorates in mathematics and philosophy in 1902. In 1919 he was Minister of Education in Poland. After WWII, he was exiled to Belgium. Arguably, the 3-valued logics named after him was the first published report of a non-Boolean, or non-Chrysippean logic (with only two logic values, 'true' or 'false', in the Chrysippean case). Subsequently, Łukasiewicz logic algebras were constructed by Grigore Moisil in 1940-1945 to define 'nuances' in logics, or many-valued logics, as well as 3-state control logic (electronic) circuits. Lukasiewicz-Moisil (LM_n) logic algebras were defined axiomatically after 1969 as n-valued logic algebra representations and extensions of the Lukasiewcz (3-valued) logics; then, the universal properties of http://planetmath.org/AlgebraicCategoryOfLMnLogicAlgebrascategories of LM_n -logic algebras were also investigated and reported in a series of recent publications. Recently, several modifications of LM_n -logic algebras are under consideration as valid candidates for representations of quantum logics, as well as for modeling non-linear biodynamics in genetic 'nets' or networks, as well as in single-cell organisms, or in tumor growth.