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Apr 27, 2019 1. s. 1 2 3 Version 3.0.4. The following are some of the. Courselab 3 is much better than the prior version. a) has. Please wait while Courselab 3 loads. Courselab 3 is a Learning Management System. 2) Courselab 3 is a text-based. Shop Software & Browsers eBooks Learning Management Systems Software Licensing Antivirus & Malware Protection Other Miscellaneous LMS eLearning Free Software Courseware Additional Reading References External links Category:Information technology companies of the United States Category:Software companies of the United States Category:Educational computing Category:Companies based in Burlington, Vermont Category:Free educational software Category:Free learning support software Category:Textbook writers Category:Educational material sponsors Category:Software companies of the United StatesA new sequence variant of the amino-terminal domain of MLL protein. MLL protein, one of the major transcription factors essential for normal hematopoiesis and development of leukemia, belongs to the trithorax group proteins that possess amino-terminal methyltransferase activities. The coding region of human MLL gene, as of others, consists of 6 exons. The methyltransferase domain is encoded by exons 2-6, whereas the amino-terminal domains of MLL proteins are encoded by exons 1a, 1b and 1c. Recently, a new point mutation, G957A, in exon 1c was identified as a human leukemia-associated mutation and it was assumed that the mutation affected the amino-terminal domain. In the present study, we directly detected the expression of the amino-terminal domain of MLL protein with anti-peptide antibody. We also detected a novel mutation in exon 1c of MLL gene. The gene was amplified by polymerase chain reaction and directly sequenced. The mutated peptide in exon 1c was expressed as a recombinant protein and was analyzed by an enzyme-linked immunosorbent assay. The results showed that the novel peptide derived from MLL exon 1c, including G957A mutation, was not recognized by the anti-peptide antibody. These results showed that the mutant peptide does not have strong antigenicity and

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