البحث الثاني

Diagnostic potential of metastasis-associated-lungadenocarcinomatranscript- 1 (MALAT-1) and TNFa and hnRNPL related immunoregulatory long non-coding RNA (THRIL) in systemic lupus erythematosus patients: Relation to disease activity

ABSTRACT

Aim of the work: To determine expression levels and diagnostic value of metastasis-associated-lung-adenocarcinoma-transcript-1 (MALAT-1) and TNFa and hnRNPL related immunoregulatory long non-coding RNA (THRIL) in systemic lupus erythematosus (SLE), and to assess their role in the clinical characteristics of SLE and disease activity.

Patients and methods: Study included 40 patients with SLE and 30 matched controls. SLE Disease Activity Index (SLEDAI) score was assessed. Expression levels of MALAT-1 and THRIL were detected in the serum by using Real-time polymerase chain reaction and 2_DDCT method.

Results: Mean age of patients was 40.1 ± 9 years (25–55 years), they were 38 females and 2males and disease duration was 16.5 ± 3.9 years. Their mean SLEDAI was 5.8 ± 5.3 . Expression levels of MALAT-1 and THRIL were found to be significantly upregulated in the serum of SLE patients compared with controls (set as 1). MALAT-1 fold change = 3.7 ± 3.8 (p = 0.009), and THRIL fold change = 3.6 ± 3.4 (p = 0.026). There were significant correlations between MALAT-1 with THRILL (r = 0.44, p = 0.005), proteinuria (r = 0.45, p = 0.006), erythrocyte sedimentation rate (r = 0.43, p = 0.006) and SLEDAI (r = 0.36, p = 0.024). No significant correlations were found between THRIL and study parameters. Sensitivity and specificity of MALAT-1 and THRIL were determined (sensitivity 67.5% and 65% respectively), (specificity 100% for both, total accuracy 80% and 81.4% respectively), and the combined effect of both increased sensitivity and total

Conclusion: MALAT-1 and THRIL may be potential diagnostic biomarkers for SLE and only MALAT-1 may be valuable in detecting disease activity.

accuracy to 70% and 82.9% respectively. THRIL was a significant predictor for SLE

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disease (p = 0.02).

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