

Table S1. The baseline demographics and clinical characteristics of enrolled subjects.

Variables	Testing set			Validation set		
	HT (n=10)	Normal (n=10)	P value	HT (n=30)	Normal (n=30)	P value
Age (years)	47.5 ± 11.6	47.1 ± 11.9	0.532	47.1 ± 11.2	46.4 ± 12.1	0.127
Sex (male/female)	4/6	4/6	-	9/21	9/21	-
BMI (kg/m ²)	23.1 ± 2.8	22.9 ± 2.1	0.216	23.4 ± 3.4	23.1 ± 2.7	0.178
TSH (mIU/L)	2.8 (0.7-3.8)	2.4 (1.2-3.6)	0.187	2.2 (0.1-6.6)	2.0 (0.2-5.4)	0.467
FT3 (pmol/L)	4.2 (3.7-4.5)	4.3 (3.5-4.3)	0.623	4.7 (3.4-7.5)	4.5 (3.2-5.3)	0.381
FT4 (pmol/L)	15.1 (14.2-17.1)	15.4 (13.9-18.4)	0.327	15.2 (12.8-33.0)	15.2 (10.9-23.4)	0.169
TgAb (IU/mL)	209.5 (58.1-4000)	38.9 (11.2-112)	<0.001	253 (115-2298)	13.5 (10.1-73.4)	<0.001
TPOAb (IU/mL)	187.5 (60.2-600)	17.6 (9.0-32)	<0.001	204 (44-600)	9 (9-16.2)	<0.001

HT: Hashimoto thyroiditis; BMI: body mass index; TSH: thyroid stimulating hormone; FT3: free T3; FT4: free T4; TgAb: thyroglobulin antibody; TPOAb: thyroperoxidase antibody.

Table S2. Univariate and multiple analysis for difference miRNAs.

miRNAs	Univariate		Multivariate	
	OR (95%CI)	P value	OR (95%CI)	P value
miR-142-3p	1.613 (1.185-2.195)	0.002*	1.437 (1.016-2.032)	0.040*
miR-146a-5p	1.649 (1.176-2.311)	0.004*	1.372 (0.965-1.951)	0.078

CI: confidence interval; OR: Odds ratio; *statistically significant difference.

Figure S1. The uncropped original western blots from main figures.

Fig 1C

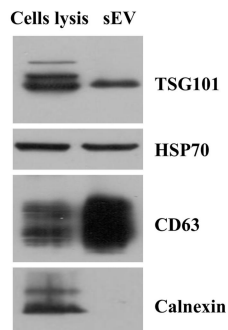


Fig 2E

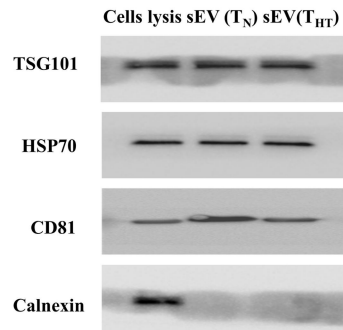


Fig 3B

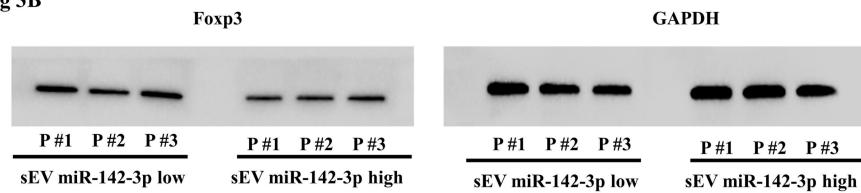


Fig 5D

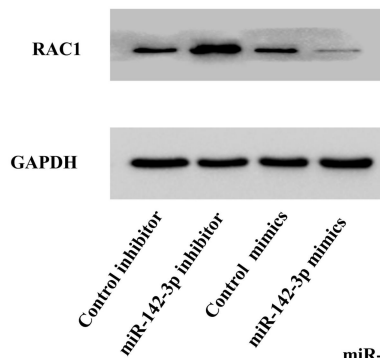


Fig 5I

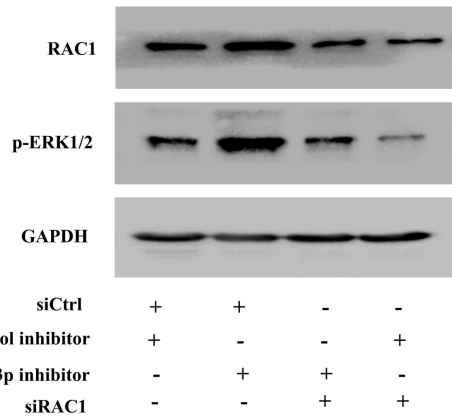


Fig 6F

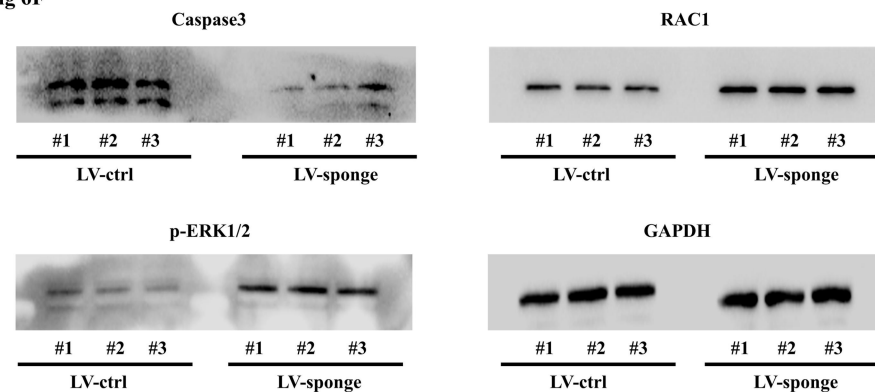


Figure S2. The lowest RNA concentration of tissue sEV total RNA and levels of tissue sEV miR-103a-3p

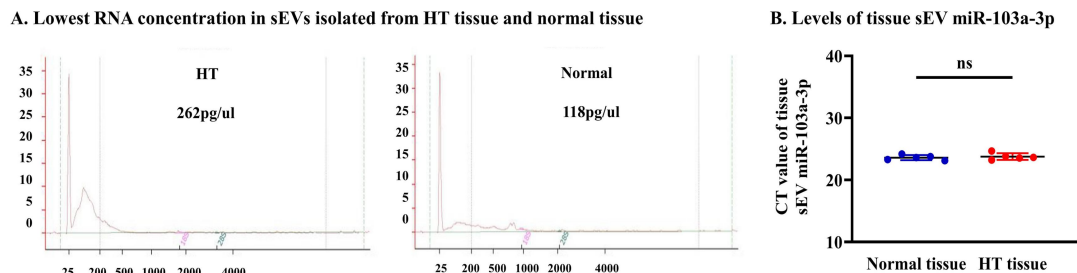


Figure S3. Tissue sEVs isolated from HT patients promotes apoptosis of thyrocyte cells which can be prevented by miR-142-3p inhibitor

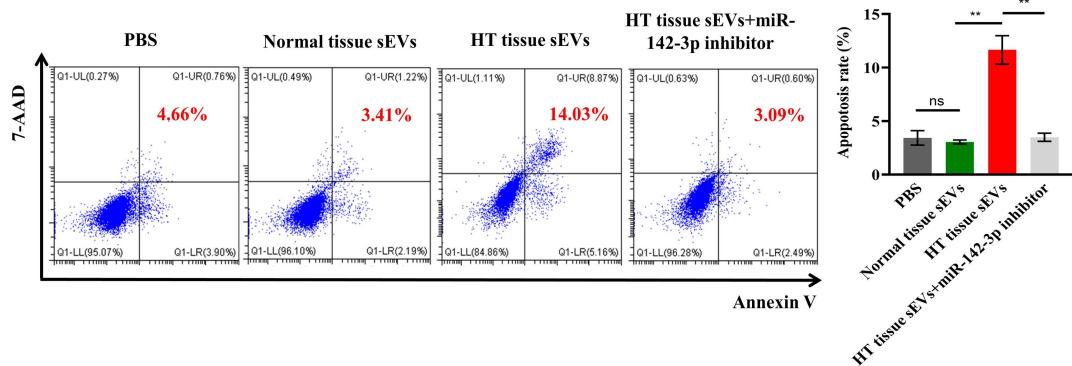
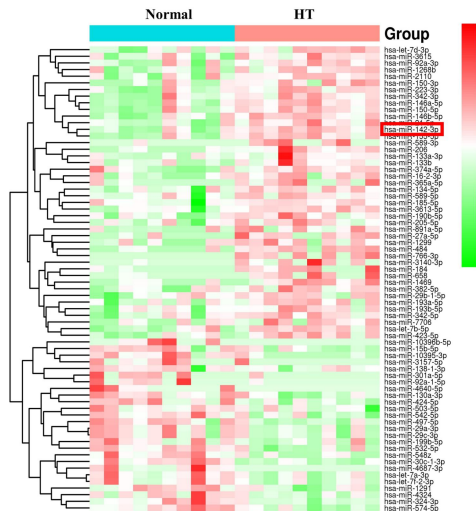
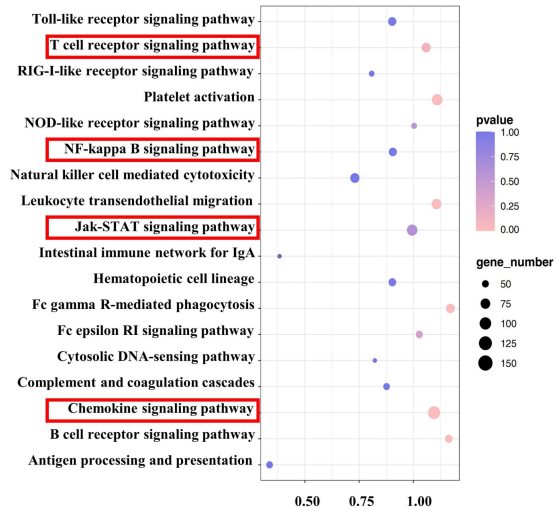


Figure S4. Biological function enrichment analysis and validation of tissue sEV miRNAs

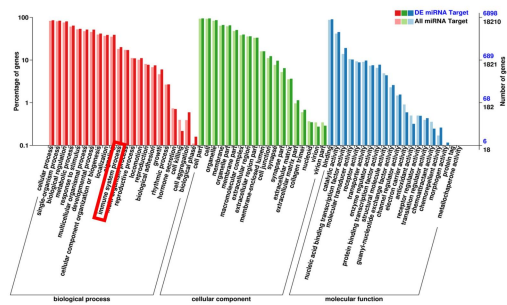
A. Heatmap across 20 tissue samples



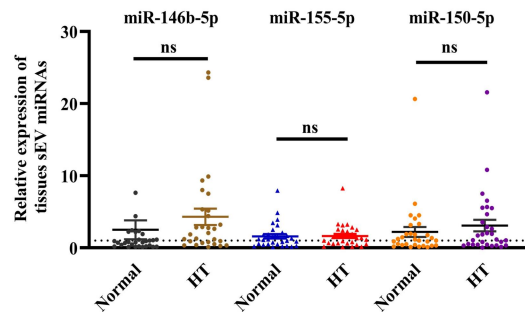
B. KEGG enrichment



C. GO enrichment



D. Validation of tissue sEV miRNAs



E. Association of tissue sEV miR-142-3p with thyroid function

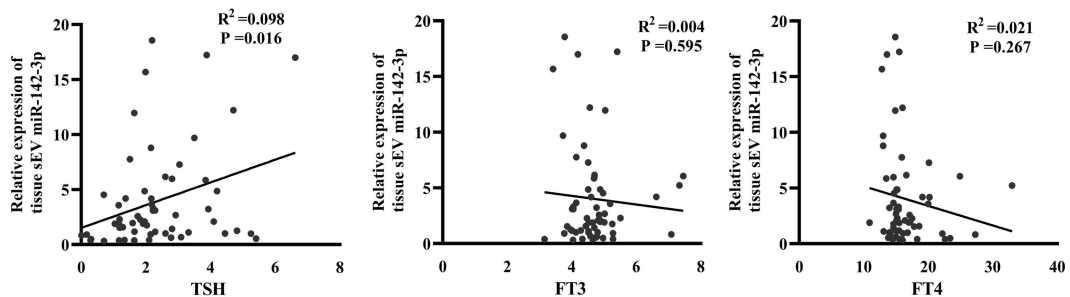


Figure S5. The levels of miR-142-3p in specific immune cell types and T lymphocyte-derived sEV

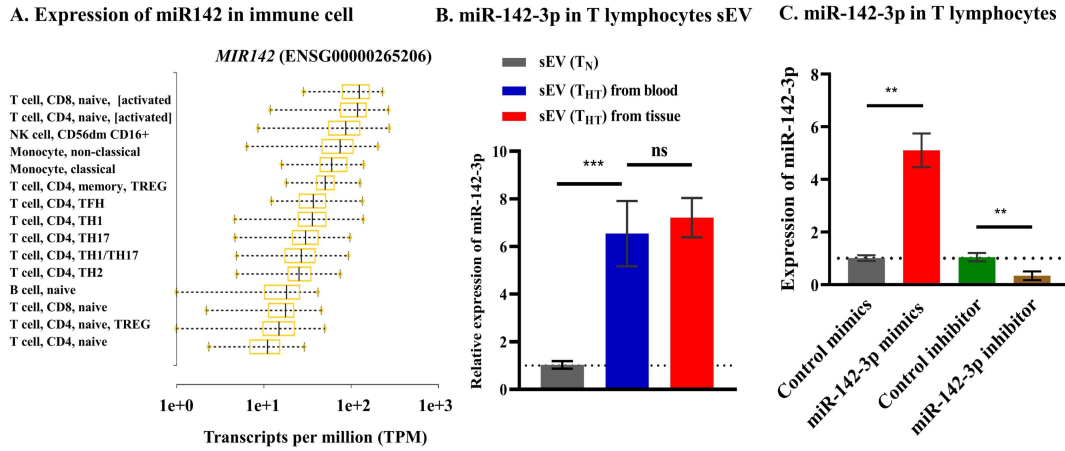


Figure S6. The miR-142-3p levels after transfection with mimics or inhibitor in thyrocyte cells, T lymphocyte migration and the effect of miR-142-3p on CXCL10 secretion

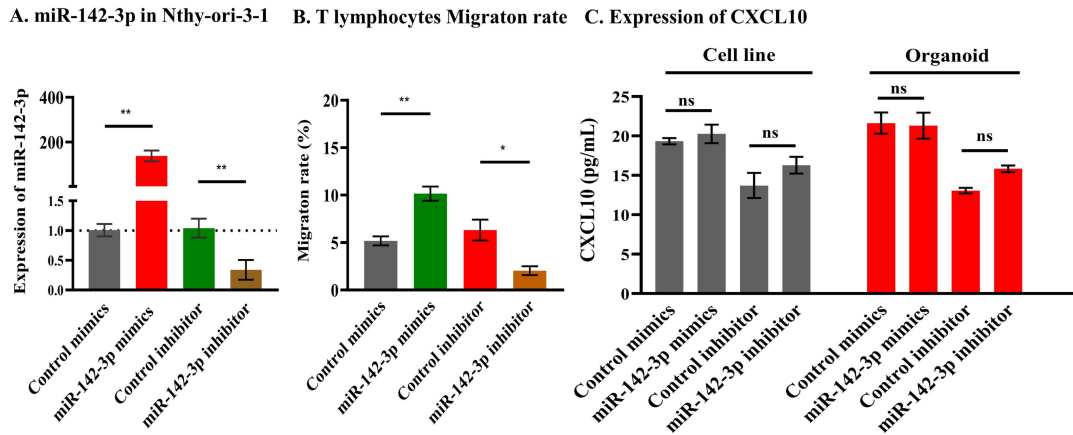


Figure S7. Predicted target and inhibition efficiency of siRNA for RAC1

