Supplemental material

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Supplementary table 1. Summary of transcriptome datasets of peripheral blood of patients with SjS

GEO ID	Participants	Samples	Techniques	Platforms	Outcomes
GSE48378	11 female SjS patients and age- and gender-matched 16 healthy controls	PBMCs	Array	GPL5175	Expression profiling
GSE51092	190 SjS patients and 32 controls	Whole blood	Array	GPL6884	Expression profiling
GSE66795	131 SjS patients and 29 age-matched non-SjS controls	Whole blood	Array	GPL10558	Expression profiling
GSE84844	30 SjS patients (Mean age: 61.1 years; Male, 3.3%) and 30 healthy controls (Mean age:39.3 years; Male, 0%)	Whole blood	Array	GPL570	Expression profiling
GSE145065	5 SjS patients and 5 age- and gender-matched healthy controls	PBMCs	RNA-seq	GPL16791	Expression profiling

(PBMCs, peripheral blood mononuclear cells; RNA-seq, RNA-sequencing.)