

Identify Biomarkers for Susceptibility to Both Type 1 Diabetes and Mental Disorders

Agata Jurczyk, Ph.D.

University of Massachusetts

Recent studies show that patients with diabetes have a much higher likelihood of depression than the general population, and young people with type 1 diabetes (T1D) had 11 times the suicide rate. Our goal is to identify genetic signatures in white blood cells that distinguish non-progressor T1D patients and T1D patients that do progress to psychiatric illness.

With the funding received from DRC, we conducted an unbiased search for changes in gene expression in white blood cells from patients with T1D, T1D with depression, and healthy control patients. From UMass Medical Center Biorepository we have obtained blood samples from 60 male patients with T1D (20), T1D with depression (20) and healthy controls (20). We have isolated RNA from all of these samples and constructed and sequenced libraries. Libraries with sequenced depth over 8 million reads were analyzed and successfully mapped to human genome (about 20% efficiency). Out of the 20 libraries each, the sequence libraries with over 8 millions reads included: 17 T1D, 12 T1D with depression and 11 healthy patients. The biometrics for each of these patients is summarized in Table 1. Patients were matched for body mass index (BMI). One of the patients sets: healthy and T1D had statistically significant age difference but not the others. Age of onset of T1D was not different between T1D and T1D with depression.

Table 1.

Health Status	Subject ID	Gender	AGE	BMI [kg/m2]	History	T1D Onset
Healthy	392	M	19		-	NA
Healthy	775	M	24	20.3	Depression	NA
Healthy	613	M	26	23.7	-	NA
Healthy	607	M	27	25.5	-	NA
Healthy	1716	M	40	23.0	Alcohol abuse	NA
Healthy	505	M	40		-	NA
Healthy	1378	M	40		-	NA
Healthy	5115	M	50	28.4	-	NA
Healthy	2906	M	50	24.7	-	NA
Healthy	3475	M	50	29.9	-	NA
Healthy	541	M	50	30.5	-	NA
		Average	37.82	25.75		
		STD	11.87	3.57		
		St Err	3.58	1.26		
T1D	6740	M	26	-	T1D	3
T1D	6525	M	26	24.3	T1D	9
T1D	6526	M	39	22.9	T1D	22
T1D	10737	M	46	37.6	T1D	19
T1D	1025	M	48	25.4	T1D	21
T1D	1209	M	51	26.4	T1D	14

T1D	1229	M	51	16.5	T1D	18
T1D	6123	M	53	18.8	T1D	40
T1D	6723	M	53	27.4	T1D	29
T1D	1630	M	56	22.9	T1D	2
T1D	2799	M	56	22.8	T1D	14
T1D	9727	M	56	24.8	T1D	10
T1D	8952	M	56	27.7	T1D	49
T1D	5354	M	59	-	T1D	8
T1D	3082	M	61	31.5	T1D	-
T1D	10586	M	62	22.0	T1D	17
T1D	6232	M	63	26.1	T1D	25
T1D vs Healthy		Average	50.71	25.12		18.75
		STD	11.08	5.02		12.63
		St Err	2.69	1.30		3.37
		Ttest	0.01	-		-
T1D & Dep	4828	M	20	19.5	Anxiety Disorder	20
T1D & Dep	6556	M	23	23.7	Anxiety/Depression	6
T1D & Dep	5010	M	35	20.5	Depression	15
T1D & Dep	6602	M	42	31.7	Chronic Depression	26
T1D & Dep	6048	M	46	24.2	Depression	25
T1D & Dep	1681	M	50	24.4	Depression	13
T1D & Dep	1993	M	50	27.6	Depression, Suicide ideation	35
T1D & Dep	1626	M	51	29.5	Anxiety/Depression	15
T1D & Dep	8020	M	53	24.6	Anxiety disorder, depression	20
T1D & Dep	8135	M	55	29.9	Anxiety and Depression	25
T1D & Dep	512	M	57	27.0	Bipolar, Major Depressive Disorder	38
T1D & Dep	2333	M	61	39.0	Anxiety/Depression	43
T1D Dep vs Healthy T1D vs T1D Dep		Average	45.25	26.79		23.42
		STD	13.03	5.32		10.95
		St Err	3.76	1.53		3.16
		Ttest	0.17	-		-
		Ttest	0.24	-		0.32

We were able to detect over 21,000 differentially expressed genes from each of the analyzed set: healthy controls vs T1D patients; healthy controls vs T1D patients with depression and T1D vs T1D patients with depression. Unfortunately, there were no significant differentially expressed genes between T1D and T1D with depression. We are currently reanalyzing this data set to make sure this result is accurate. There are 10 significant down regulated genes in common between T1D vs Healthy and T1D with depression vs Healthy (Table 2).

Table 2.

Gene ID	T1D vs Healthy Fold	P value	T1D Dep vs Healthy Fold	P value
RSRP1	0.625	0.001	0.649	0.037
S100A8	0.169	0.007	0.141	0.008
RPL34	0.255	0.010	0.161	2.83 E-08
SNHG10	0.503	0.022	0.431	0.002
TXN	0.435	0.023	0.397	0.031
HMGB2	0.501	0.028	0.443	0.019
RPL36A	0.268	0.032	0.216	0.0004
RPL27	0.506	0.032	0.390	1.57 E-06
S100A12	0.252	0.033	0.184	0.021
EEF1B2	0.496	0.041	0.437	0.025

There were 28 significantly up regulated genes and 77 significantly down regulated genes between T1D vs. Healthy (Table 3).

Table 3.

Gene ID	T1D vs Healthy Fold Down	P value	Gene ID	T1D vs Healthy Fold Up	P value
CHRM3	0.594	0.001	RBBP7	1.477	0.004
ITGB2-AS1	0.564	0.002	SF3A3	1.403	0.004
FAM157C	0.524	0.007	PABPC1	1.352	0.002
LOC254896	0.464	0.002	SLC6A19	6.202	0.027
LOC441081	0.446	0.008	HRASLS5	4.018	0.049
CSNK1A1L	0.390	0.001	COL6A3	2.515	0.048
RPS24	0.369	0.001	FOSB	2.050	0.033
C9orf84	0.367	0.008	MYADM	1.699	0.013
CASP5	0.367	0.008	FAM115A	1.623	0.049
C14orf2	0.358	0.007	TAF15	1.527	0.011
LINC00998	0.241	0.007	HTATSF1	1.509	0.032
CHCHD2	0.665	0.046	DDX24	1.508	0.011
HIST1H2AC	0.663	0.050	WBP11	1.490	0.028
PIGH	0.646	0.038	PARP1	1.467	0.011
RMRP	0.633	0.023	PYGB	1.447	0.041
ATP6V1G1	0.618	0.049	TAGAP	1.439	0.013
BLOC1S2	0.612	0.034	GOLGB1	1.422	0.010
RPS6	0.601	0.041	EZR	1.398	0.048
FAU	0.598	0.028	PDCD4	1.390	0.040
SLC8A1	0.597	0.050	PBXIP1	1.350	0.041

GTF2B	0.591	0.027	TFIP11	1.341	0.046
PSMA6	0.584	0.048	LUZP1	1.319	0.048
B2M	0.577	0.033	HNRNPD	1.312	0.022
GLRX	0.556	0.034	CSNK2A1	1.312	0.046
RPS18	0.552	0.028	CLINT1	1.311	0.029
ATP5L	0.546	0.036	SMARCC1	1.288	0.016
RPS27A	0.542	0.011	XRCC6	1.288	0.022
ATP5J	0.540	0.039	USP7	1.237	0.038
HIST1H4F	0.537	0.038			
SNORA49	0.532	0.040			
CLEC4A	0.530	0.022			
FAM157A	0.528	0.049			
AIF1	0.520	0.021			
RPS29	0.514	0.036			
GMFG	0.512	0.023			
TUBD1	0.510	0.046			
PTENP1	0.510	0.027			
LINC01094	0.495	0.032			
COX7C	0.481	0.036			
MGAM	0.478	0.027			
NIPSNAP3A	0.458	0.011			
RPS15A	0.454	0.022			
TOMM7	0.437	0.027			
SLC8A1-AS1	0.434	0.016			
SCARNA7	0.420	0.028			
NDUFB1	0.410	0.029			
TMA7	0.404	0.036			
RPS17	0.393	0.029			
TP53TG1	0.366	0.024			
RPL23	0.347	0.048			
RPL39	0.345	0.035			
CEP57L1	0.337	0.023			
RPS7	0.335	0.018			
COX7B	0.311	0.033			
RPL31	0.306	0.011			
RPL26	0.259	0.022			
AP3B2	0.100	0.044			
CIAO1	0.824	0.032			
UTY	0.783	0.029			
PHYKPL	0.745	0.041			

TRIM13	0.723	0.023
ARGLU1	0.717	0.028
METTL22	0.709	0.046
SLC35F5	0.701	0.011
YIPF4	0.699	0.011
SMA5	0.688	0.021
CD48	0.669	0.041

There were 12 significantly down regulated and 6 significantly up regulated genes between T1D with depression vs Healthy (Table 4).

Table 4.

Gene ID	T1D Dep vs Healthy Fold Down	P Value	Gene ID	T1D Dep vs Healthy Fold Up	P Value
RPS20	0.514	0.0086	PRAMEF6	35.840	0.0247
RPL35A	0.450	0.0003	ZNF334	17.060	0.0374
RPS21	0.441	0.0002	RGMB-AS1	12.273	0.0317
RPL9	0.302	0.0000	CDC42EP1	3.675	0.0374
NEIL3	0.218	0.0002	SIDT2	1.557	0.0490
ANKRD36BP2	0.171	0.0069	RBBP7	1.458	0.0374
TMBIM4	0.615	0.0401			
SCARNA10	0.477	0.0490			
CLEC4D	0.353	0.0312			
VNN1	0.328	0.0317			
SLC35F5	0.695	0.0401			
YIPF4	0.675	0.0480			

Our goal of this study was to identify cellular and molecular biomarkers in white blood cells (wbc's) that associate with T1D and major depression in hope to identify at-risk individuals for preventative treatments. We did identify significantly up regulated and down regulated genes between T1D vs healthy and T1D with Depression vs healthy sets and we are currently working to analyze these genes patterns for gene signature that could be used in the clinic. This study preliminary work should be followed with increased sample size that is beyond the scope of this grant. We will continue analyzing these data sets in order to discover genetic biomarkers that identify T1D patients with susceptibility to depression, and vice versa, so that appropriate preventative measures can be taken. Our hope is to find a genetic association linking T1D and depression which will uncover possible novel mechanisms to improve therapies, even preventative ones, to alleviate suffering caused by both illnesses which are very common and extraordinarily costly—both in relation to healthcare, but also in regards to individual quality of life.