

	+ IL-7		- IL-7				
	Fold Change	P-Value	Fold Change	P-Value	PicTar	miRanda	PITA
Trp53inp1	0.336	0.063	0.308	0.007	X	X	X
Arid3a	0.389	0.032	0.368	0.04			X
Abtb1	0.432	0.042	0.458	0.044	X	X	X
Lactb	0.469	0.045	0.497	0.066	X	X	X
Ptpn18	0.48	0.052	0.396	0.04	X	X	
Atg4d	0.507	0.067	0.45	0.038	X		X
Btg2	0.523	0.123	0.543	0.027			X
Tmem123	0.546	0.081	0.453	0.022		X	X
MAP3K11	0.583	0.056	0.566	0.008		X	X
LBH	0.614	0.002	0.697	0.171	X	X	X
Enpep	0.616	0.13	0.537	0.026	X	X	X
Usp12	0.622	0.084	0.549	0.01	X		X
E2F2	0.632	0.094	0.632	0.012			X
Blzf1	0.642	0.076	0.65	0.042	X	X	X
Baz2a	0.658	0.062	0.669	0.036		X	X
Homez	0.665	0.072	0.611	0.004		X	X
Sirt7	0.682	0.199	0.619	0.031	X	X	X
Rhot2	0.7	0.064	0.651	0.021		X	
Sertad3	0.701	0.192	0.62	0.027	X	X	X
Tnfaip3/A20	0.766	0.437	0.465	0.022		X	X
Bak1	0.803	0.161	0.655	0.003	X	X	X
Rps6ka1	0.842	0.161	0.652	0.081	X	X	X
eIF1ad	0.846	0.357	0.653	0.034	X	X	X

"X" indicates that the gene is predicted by the respective algorithm