

Table S2 List of all proteins (n =1167) identified in the study

Accession	Description	113/114	113/115	113/116	113/117	113/118	113/119	113/121	Log2(113/113)	Log2(114/113)	Log2(115/113)	Log2(116/113)	Log2(117/113)	Log2(118/113)	Log2(119/113)	Log2(121/113)	Log	Ratio	P-value
F1RT25	Uncharacterized protein OS=Sus scrofa GN=SNX3 PE=4 SV=2 - [F1RT25_PIG]	1.233	0.916	1.117	0.983	1.076	1.064	1.121	0	-0.30	0.13	-0.16	0.02	-0.11	-0.09	-0.16	-1.08E-06	1.0000	
Q06A94	ROA1 OS=Sus scrofa GN=HNRNPA1 PE=2 SV=1 - [Q06A94_PIG]	1.055	1.124	0.467	1.465	0.466	0.905	0.896	0	-0.08	-0.17	1.10	-0.55	1.10	0.14	0.16	1.29E-04	0.9998	
F2Z5D4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=SYNCRIP PE=4 SV=2 - [F2Z5D4_PIG]	1.253	1.234	1.041	1.169	1.009	1.207	1.131	0	-0.33	-0.30	-0.23	-0.01	-0.27	-0.18	-1.32E-04	0.9990		
Q3HUX1	Fatty acid translocase/CD36 OS=Sus scrofa GN=CD36 PE=2 SV=1 - [Q3HUX1_PIG]	0.954	0.913	0.826	0.89	0.976	0.814	1.018	0	0.07	0.13	0.28	0.17	0.04	0.30	-0.03	-1.77E-04	0.9985	
F1SEF7	Serine/threonine-protein phosphatase (Fragment) OS=Sus scrofa GN=PPP1CB PE=3 SV=1 - [F1SEF7_PIG]	0.902	1.068	0.96	1.075	0.911	0.969	0.974	0	0.15	-0.09	0.06	-0.10	0.13	0.05	0.04	1.99E-04	0.9979	
P80276	Aldose reductase OS=Sus scrofa GN=AKR1B1 PE=1 SV=2 - [ALDR_PIG]	1.122	1.065	1.183	1.05	1.136	1.053	1.126	0	-0.17	-0.05	-0.24	-0.07	-0.18	-0.07	-0.17	-1.72E-04	0.9978	
F1SSD1	Uncharacterized protein OS=Sus scrofa GN=INVS PE=4 SV=2 - [F1SSD1_PIG]	1.509	0.973	0.917	1.109	1.271	0.972	0.986	0	-0.59	0.04	0.13	-0.15	-0.35	0.04	0.02	-1.20E-03	0.9951	
F1S596	Uncharacterized protein OS=Sus scrofa GN=PRKCSH PE=4 SV=1 - [F1S596_PIG]	1.194	1.11	1.065	1.201	0.941	1.074	1.165	0	-0.26	-0.15	-0.09	-0.26	-0.09	-0.10	-0.22	-6.53E-04	0.9948	
P08059	Glucose-6-phosphate isomerase OS=Sus scrofa GN=GPI PE=1 SV=3 - [G6PI_PIG]	0.893	1.133	1.203	1.053	1.132	1.001	1.018	0	0.16	-0.18	-0.27	-0.07	-0.18	0.00	-0.03	7.39E-04	0.9945	
F1S6Q1	Uncharacterized protein OS=Sus scrofa GN=NDUFA13 PE=4 SV=2 - [F1S6Q1_PIG]	0.97	1.008	1.397	1.288	1.349	0.885	0.884	0	0.04	-0.48	-0.37	-0.43	0.18	0.18	1.75E-03	0.9936		
F2Z5G5	Uncharacterized protein OS=Sus scrofa GN=ACTR1A PE=3 SV=1 - [F2Z5G5_PIG]	0.836	0.877	0.904	0.94	0.889	0.914	0.869	0	0.26	0.19	0.15	0.09	0.17	0.13	0.20	-5.16E-04	0.9934	
P63317	Troponin C, slow skeletal and cardiac muscles OS=Sus scrofa GN=TNNC1 PE=1 SV=1 - [TNNC1_PIG]	0.645	0.975	0.741	0.824	0.745	1.101	0.693	0	0.63	0.04	0.43	0.28	0.42	-0.14	0.53	-1.84E-03	0.9934	
K7GP58	Uncharacterized protein OS=Sus scrofa GN=AIFM1 PE=4 SV=1 - [K7GP58_PIG]	1.011	0.937	0.948	1.003	1.002	0.966	0.924	0	-0.02	0.09	0.08	0.00	0.00	0.05	0.11	4.00E-04	0.9922	
F1SMB6	Uncharacterized protein OS=Sus scrofa GN=LOC100520143 PE=3 SV=2 - [F1SMB6_PIG]	1.059	0.898	0.857	0.965	0.844	1.005	0.993	0	-0.08	0.16	0.22	0.05	0.24	-0.01	0.01	9.70E-04	0.9918	
F1RX99	Uncharacterized protein OS=Sus scrofa GN=TPD52 PE=4 SV=2 - [F1RX99_PIG]	1.131	0.942	0.62	1.038	0.732	1.157	0.757	0	-0.18	0.09	0.69	-0.05	0.45	-0.21	0.40	-2.68E-03	0.9918	
Q64L94	Proteasome activator complex subunit 1 OS=Sus scrofa GN=PSME1 PE=2 SV=1 - [PSME1_PIG]	1.003	0.906	0.584	0.95	0.813	0.933	0.741	0	0.00	0.14	0.78	0.07	0.30	0.10	0.43	-2.22E-03	0.9917	
F1SB09	Uncharacterized protein OS=Sus scrofa GN=SEC22B PE=4 SV=1 - [F1SB09_PIG]	1.018	1.026	1.08	1.069	0.978	1.058	1.021	0	-0.03	-0.04	-0.11	-0.10	0.03	-0.08	-0.03	-4.23E-04	0.9914	
F1SMN5	Uncharacterized protein OS=Sus scrofa GN=FLNC PE=4 SV=1 - [F1SMN5_PIG]	1.025	1.029	1.028	0.982	1.045	0.993	1.063	0	-0.04	-0.04	-0.04	0.03	-0.06	0.01	-0.09	3.51E-04	0.9909	
K7GLM2	Uncharacterized protein OS=Sus scrofa GN=NDUFB11 PE=4 SV=1 - [K7GLM2_PIG]	0.942	0.744	0.892	0.951	0.889	0.823	0.895	0	0.09	0.43	0.16	0.07	0.17	0.28	0.16	1.40E-03	0.9895	
F1RR69	Uncharacterized protein OS=Sus scrofa GN=SET PE=3 SV=1 - [F1RR69_PIG]	1.377	0.983	0.833	0.906	0.977	1.15	1.115	0	-0.46	0.02	0.26	0.14	0.03	-0.20	-0.16	-2.38E-03	0.9894	
I3LJ88	Uncharacterized protein OS=Sus scrofa GN=TMOD1 PE=4 SV=1 - [I3LJ88_PIG]	0.992	1.112	1.023	0.938	1.098	1.05	1.046	0	0.01	-0.15	-0.03	0.09	-0.13	-0.07	-0.06	-8.99E-04	0.9893	
I3LFC4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=AIMP1 PE=4 SV=1 - [I3LFC4_PIG]	0.93	0.829	0.853	0.897	0.89	0.916	0.897	0	0.10	0.27	0.23	0.16	0.17	0.13	0.16	9.27E-04	0.9886	
P21753	Thymosin beta-10 OS=Sus scrofa GN=TMSB10 PE=1 SV=2 - [TYB10_PIG]	1.113	1.208	1.221	1.229	0.966	1.289	1.079	0	-0.15	-0.27	-0.29	-0.30	0.05	-0.37	-0.11	-2.10E-03	0.9861	
F1RNA6	Uncharacterized protein (Fragment) OS=Sus scrofa GN=FLOT2 PE=4 SV=1 - [F1RNA6_PIG]	1.67	1.394	1.816	1.405	1.47	1.358	1.523	0	-0.74	-0.48	-0.86	-0.49	-0.56	-0.44	-0.61	-3.74E-03	0.9853	
Q551U1	Heat shock protein beta-1 OS=Sus scrofa GN=HSPB1 PE=2 SV=1 - [HSPB1_PIG]	1.014	0.901	0.915	0.736	1.081	0.916	1.158	0	-0.02	0.15	0.13	0.44	-0.11	0.13	-0.21	-3.42E-03	0.9827	
F1SLG9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100738341 PE=4 SV=1 - [F1SLG9_PIG]	1.023	0.839	0.991	0.782	1.053	1.032	1.092	0	-0.03	0.25	-0.13	0.35	-0.07	-0.05	-0.13	3.26E-03	0.9818	
D3K5K2	FK506 binding protein 4 OS=Sus scrofa GN=FKBP4 PE=2 SV=1 - [D3K5K2_PIG]	0.874	1.496	1.024	1.13	0.712	1.332	1.227	0	0.19	-0.58	-0.03	-0.18	0.49	-0.41	-0.30	6.51E-03	0.9810	
I3LLC8	Uncharacterized protein OS=Sus scrofa GN=EIF5 PE=4 SV=1 - [I3LLC8_PIG]	0.924	0.957	1.022	1.048	0.927	0.916	1.02	0	0.11	0.06	-0.03	-0.07	0.11	0.13	-0.03	-1.58E-03	0.9794	
F1SJC2	Uncharacterized protein OS=Sus scrofa GN=PPPIR3A PE=4 SV=2 - [F1SJC2_PIG]	0.915	0.963	1.116	0.956	0.984	1.032	1.008	0	0.13	0.05	-0.16	0.06	0.02	-0.05	-0.01	1.76E-03	0.9793	
F1RFY2	Beta-enolase OS=Sus scrofa GN=ENO3 PE=3 SV=1 - [F1RFY2_PIG]	0.826	0.954	0.965	0.947	0.96	0.893	0.932	0	0.28	0.07	0.05	0.08	0.06	0.16	0.10	1.80E-03	0.9788	
Q29214	60S acidic ribosomal protein P0 OS=Sus scrofa GN=RPLP0 PE=2 SV=2 - [RLA0_PIG]	1.027	1.195	1.068	1.038	1.218	1.003	1.026	0	-0.04	-0.26	-0.09	-0.05	-0.28	0.00	-0.04	2.67E-03	0.9761	
F1STY1	NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Sus scrofa GN=NDUFC2 PE=3 SV=2 - [F1STY1_PIG]	0.701	0.755	0.661	0.993	0.73	0.804	0.589	0	0.51	0.41	0.60	0.01	0.45	0.31	0.76	6.83E-03	0.9745	
F2Z5N8	Ras-related protein Rab-14 OS=Sus scrofa GN=RAB14 PE=3 SV=1 - [F2Z5N8_PIG]	0.8	0.782	0.757	0.945	0.799	0.829	0.748	0	0.32	0.35	0.40	0.08	0.32	0.27	0.42	4.12E-03	0.9728	
F1SKL2	Uncharacterized protein OS=Sus scrofa GN=TST PE=4 SV=1 - [F1SKL2_PIG]	0.94	0.966	0.852	0.975	0.911	0.921	0.951	0	0.09	0.05	0.23	0.04	0.13	0.12	0.07	-2.01E-03	0.9718	
Q19KE2	Fatty acid binding protein 5 (Fragment) OS=Sus scrofa GN=FABP5 PE=2 SV=1 - [Q19KE2_PIG]	0.998	0.725	0.822	0.849	0.767	0.946	0.952	0	0.00	0.46	0.28	0.24	0.38	0.08	0.07	5.07E-03	0.9713	
M3UZ37	Heterogeneous nuclear ribonucleoprotein A2/B1 OS=Sus scrofa GN=HNRNPA2B1 PE=2 SV=1 - [M3UZ37_PIG]	1.152	0.784	0.735	0.91	0.809	0.847	1.045	0	-0.20	0.35	0.44	0.14	0.31	0.24	-0.06	6.70E-03	0.9701	
I3LD45	Uncharacterized protein OS=Sus scrofa GN=SIRT5 PE=4 SV=1 - [I3LD45_PIG]	1.04	0.743	0.978	0.925	0.797	0.989	1.054	0	-0.06	0.43	0.03	0.11	0.33	0.02	-0.08	-6.04E-03	0.9671	
F1S288	Uncharacterized protein OS=Sus scrofa GN=SNTB1 PE=4 SV=2 - [F1S288_PIG]	1.119	0.842	1.036	1.097	0.828	1.052	1.006	0	-0.16	0.25	-0.05	-0.13	0.27	-0.07	-0.01	5.52E-03	0.9661	
F1RSG9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100152491 PE=4 SV=2 - [F1RSG9_PIG]	0.905	0.956	0.79	1.014	0.778	0.923	0.926	0	0.14	0.06	0.34	-0.02	0.36	0.12	0.11	4.90E-03	0.9655	
F1RG99	Uncharacterized protein OS=Sus scrofa GN=CTF1 PE=4 SV=1 - [F1RG99_PIG]	6.158	1.874	2.362	1.596	3.001	2.191	2.794	0	-2.62	-0.91	-1.24	-0.67	-1.13	-1.48	-2.63E-02	0.9654		
I3L9N1	Uncharacterized protein OS=Sus scrofa GN=ADPRHL1 PE=4 SV=1 - [I3L9N1_PIG]	0.747	0.909	0.991	0.86	0.902	0.908	0.943	0	0.42	0.14	0.01	0.22	0.15	0.14	0.08	4.70E-03	0.9646	
F1RVN2	Double C2-like domain-containing protein (Fragment) OS=Sus scrofa GN=LOC100525027 PE=4 SV=1 - [F1RVN2_PIG]	1.437	1.417	1.024	1.266	1.135	1.288	1.149	0	-0.52	-0.50	-0.03	-0.34	-0.18	-0.37	-0.20	-7.09E-03	0.9640	
F1RL20	Uncharacterized protein OS=Sus scrofa GN=CHCHD10 PE=4 SV=1 - [F1RL20_PIG]	1.533	1.125	1.07	1.183	1.037	1.04	1.414	0	-0.62	-0.17	-0.10	-0.24	-0.05	-0.06	-0.50	8.17E-03	0.9637	
I3L675	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYH15 PE=4 SV=1 - [I3L675_PIG]	0.768	1.049	0.795	1.027	0.735	0.829	1.048	0	0.38	-0.07	0.33	-0.04	0.44	0.27	-0.07	-8.53E-03	0.9612	
Q3ZDQ5	Caveolin-3 OS=Sus scrofa GN=CAV3 PE=2 SV=1 - [CAV3_PIG]	1.027	1.134	1.064	1.229	0.937	1.153	0.948	0	-0.04	-0.18	-0.09	-0.30	0.09	-0.21	0.08	-5.65E-03	0.9594	
B2ZF47	Mitochondrial aldehyde dehydrogenase 2 OS=Sus scrofa GN=ALDH2 PE=2 SV=1 - [B2ZF47_PIG]	0.996	1.073	1.188	1.153	1.155	1	0.94	0	0.01	-0.10	-0.25	-0.21	-0.21	0.00	0.09	5.10E-03	0.9592	
Q9MYT8	ATP synthase subunit e, mitochondrial OS=Sus scrofa GN=ATP5I PE=3 SV=4 - [ATP5I_PIG]	0.865	0.883	0.666	1.016	0.795	1.105	0.591	0	0.21	0.18	0.59	-0.02	0.33	-0.14	0.76	-1.31E-02	0.9578	
F1RF45	Uncharacterized protein (Fragment) OS=Sus scrofa GN=HOMER1 PE=4 SV=2 - [F1RF45_PIG]	1.029	1.064	0.877	1.057	0.883	1.066	0.952	0	-0.04	-0.09	0.19	-0.08	0.18	-0.09	0.07	4.92E-03	0.9578	
F1SU59	Annexin OS=Sus scrofa GN=ANXA7 PE=3 SV=2 - [F1SU59_PIG]	0.915	1.036	1.174	0.991	1.12	0.968	1.05	0	0.13	-0.05	-0.23	0.01	-0.16	0.05	-0.07	-4.91E-03	0.9574	
Q75NH0	Troponin T fast skeletal muscle type OS=Sus scrofa GN=tmt3 PE=2 SV=1 - [Q75NH0_PIG]	0.639	0.751	1.17	0.767	0.592	1.025	1.267	0	0.65	0.41	-0.23	0.38	0.76	-0.04	-0.34	-1.77E-02	0.9564	
Q29105	Ryanodine receptor OS=Sus scrofa GN=RYR1 PE=2 SV=1 - [Q29105_PIG]	0.944	1.129	1.124	1.083	1.05	1.02	1.044	0	0.08	-0.18	-0.17	-0.12	-0.07	-0.03	-0.06	-3.89E-03	0.9551	
K7ZRK0	IgA heavy chain constant region (Fragment) OS=Sus scrofa GN=IGHA PE=4 SV=1 - [K7ZRK0_PIG]	0.615	1.408	0.84	0.62	1.423	0.982	0.89	0	0.70	-0.49	0.25	0.69	-0.51	0.03	0.17	-2.10E-02	0.9541	
F1SSA6	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYH10 PE=4 SV=2 - [F1SSA6_PIG]	0.986	0.685	0.883	0.801	0.889	0.96	0.893	0	0.02	0.55	0.18	0.32	0.17	0.06	0.16	-8.41E-03	0.9531	
F1RQQ7	Phosphorylase (Fragment) OS=Sus scrofa GN=PYGB PE=3 SV=2 - [F1RQQ7_PIG]	0.845	1.034	1.082	0.97	1.048	0.998	0.918	0	0.24	-0.05	-0.11	0.04	-0.07	0.00	0.12	5.40E-03	0.9528	
F1SCD0	Uncharacterized protein OS=Sus scrofa GN=SERPINA3 PE=3 SV=1 - [F1SCD0_PIG]	0.946	0.839	0.893	0.761	1.614	0.734	0.825	0	0.08	0.25	0.16	0.39	-0.69	0.45	0.28	-1.74E-02	0.9514	
F1SGD7	Uncharacterized protein OS=Sus scrofa GN=NAP1L1 PE=3 SV=1 - [F1SGD7_PIG]	0.923	0.969	0.907	0.952	0.833	1.021	0.989	0	0.12	0.05	0.14	0.07	0.26	-0.03	0.02	4.67E-03	0.9504	
P02067	Hemoglobin subunit beta OS=Sus scrofa GN=HBB PE=1 SV=3 - [HBB_PIG]	0.75	0.599	0.975	0.962	0.875	1.182	0.411	0	0.42									

L8B149	IgG heavy chain OS=Sus scrofa GN=IGHG PE=2 SV=1 - [L8B149_PIG]	0.932	0.688	0.648	1.026	0.455	1.025	0.939	0	0.10	0.54	0.63	-0.04	1.14	-0.04	0.09	-2.82E-02	0.9335
F1SSC7	Uncharacterized protein OS=Sus scrofa GN=MURC PE=4 SV=2 - [F1SSC7_PIG]	0.971	1.005	0.806	0.945	0.876	0.974	0.956	0	0.04	-0.01	0.31	0.08	0.19	0.04	0.06	7.28E-03	0.9328
F1SFR3	Uncharacterized protein OS=Sus scrofa GN=ESPL1 PE=4 SV=1 - [F1SFR3_PIG]	0.94	1.392	1.291	1.212	1.201	1.08	1.113	0	0.09	-0.48	-0.37	-0.28	-0.26	-0.11	-0.15	-1.27E-02	0.9326
F1SMX5	Uncharacterized protein OS=Sus scrofa GN=CAB39 PE=4 SV=1 - [F1SMX5_PIG]	0.803	1.104	1.005	1.005	0.826	1.127	0.921	0	0.32	-0.14	-0.01	-0.01	0.28	-0.17	0.12	1.21E-02	0.9323
F2Z421	Uncharacterized protein OS=Sus scrofa GN=YWHAG PE=3 SV=1 - [F2Z421_PIG]	0.791	0.961	0.998	0.94	0.946	0.991	0.841	0	0.34	0.06	0.00	0.09	0.08	0.01	0.25	8.42E-03	0.9322
I3L5C0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=VAPA PE=4 SV=1 - [I3L5C0_PIG]	0.98	1.116	1.147	1.07	1.22	0.97	1.014	0	0.03	-0.16	-0.20	-0.10	-0.29	0.04	-0.02	-8.39E-03	0.9298
F1SED4	Uncharacterized protein OS=Sus scrofa GN=PREB PE=4 SV=2 - [F1SED4_PIG]	1.596	1.476	1.55	1.254	1.176	2.041	1.292	0	-0.67	-0.56	-0.63	-0.33	-0.23	-1.03	-0.37	-2.27E-02	0.9279
F6Q5P0	40S ribosomal protein S13 OS=Sus scrofa GN=RP513 PE=3 SV=2 - [F6Q5P0_PIG]	1.302	1.067	1.317	1.243	1.167	1.206	1.077	0	-0.38	-0.09	-0.40	-0.31	-0.22	-0.27	-0.11	-1.06E-02	0.9265
F1SJS8	Uncharacterized protein OS=Sus scrofa GN=TAGL9 PE=4 SV=2 - [F1SJS8_PIG]	1.058	0.961	0.971	0.872	1.133	1.031	0.947	0	-0.08	0.06	0.04	0.20	-0.18	-0.04	0.08	8.37E-03	0.9264
F1ST02	Proteasome subunit beta type OS=Sus scrofa GN=PSMB4 PE=3 SV=1 - [F1ST02_PIG]	0.839	0.877	1.069	0.894	0.987	1.005	0.909	0	0.25	0.19	-0.10	0.16	0.02	-0.01	0.14	-8.84E-03	0.9262
Q29388	Phosphoglucutase (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29388_PIG]	0.892	0.825	1.005	0.893	0.839	0.952	1.012	0	0.16	0.28	-0.01	0.16	0.25	0.07	-0.02	8.76E-03	0.9258
F1S155	Basic leucine zipper and W2 domain-containing protein 1 OS=Sus scrofa GN=BZW1 PE=2 SV=2 - [F1S155_PIG]	1.243	1.933	1.685	1.518	1.626	1.385	1.26	0	-0.31	-0.95	-0.75	-0.60	-0.70	-0.47	-0.33	-2.23E-02	0.9254
F1RH20	Uncharacterized protein OS=Sus scrofa GN=MYBPC2 PE=4 SV=2 - [F1RH20_PIG]	0.882	1.156	1.157	1.081	1.025	1.037	1	0	0.18	-0.21	-0.21	-0.11	-0.04	-0.05	0.00	9.49E-03	0.9253
I3LEI8	1,2-dihydroxy-3-keto-5-methylthiopentene dioxigenase OS=Sus scrofa GN=AD11 PE=3 SV=1 - [I3LEI8_PIG]	0.844	0.939	0.78	0.963	0.766	0.871	0.993	0	0.24	0.09	0.36	0.05	0.38	0.20	0.01	-1.14E-02	0.9250
I3LK43	Uncharacterized protein OS=Sus scrofa GN=NDUF57 PE=3 SV=1 - [I3LK43_PIG]	0.952	0.782	0.861	1.048	0.796	0.852	0.931	0	0.07	0.35	0.22	-0.07	0.33	0.23	0.10	-1.15E-02	0.9247
P00339	L-lactate dehydrogenase A chain OS=Sus scrofa GN=LDHA PE=1 SV=3 - [LDHA_PIG]	0.968	1.197	1.292	1.134	1.126	1.048	1.087	0	0.05	-0.26	-0.37	-0.18	-0.17	-0.07	-0.12	1.04E-02	0.9238
F1RH92	Uncharacterized protein OS=Sus scrofa GN=MYOT PE=4 SV=1 - [F1RH92_PIG]	1.102	0.9	0.905	0.876	0.99	0.939	1.074	0	-0.14	0.15	0.14	0.19	0.01	0.09	-0.10	9.35E-03	0.9231
F1S193	Uncharacterized protein OS=Sus scrofa GN=RCN2 PE=4 SV=2 - [F1S193_PIG]	1.048	1.072	1.105	1.07	0.992	1.134	1.046	0	-0.07	-0.10	-0.14	-0.10	0.01	-0.18	-0.06	-5.08E-03	0.9226
F1S110	Uncharacterized protein OS=Sus scrofa GN=RPL10L PE=4 SV=1 - [F1S110_PIG]	1.322	1.14	0.822	1.345	1.088	1.058	0.847	0	-0.40	-0.19	0.28	-0.43	-0.12	-0.08	0.24	-2.05E-02	0.9214
F2Z4Y1	Uncharacterized protein OS=Sus scrofa GN=YWHAH PE=3 SV=1 - [F2Z4Y1_PIG]	0.791	0.984	0.772	0.97	0.795	1.058	0.702	0	0.34	0.02	0.37	0.04	0.33	-0.08	0.51	1.73E-02	0.9211
I3LCX6	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LCX6_PIG]	0.887	1.313	1.029	1.11	1.051	1.027	1.036	0	0.17	-0.39	-0.04	-0.15	-0.07	-0.04	-0.05	-1.27E-02	0.9202
K7GP05	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100153260 PE=4 SV=1 - [K7GP05_PIG]	1.128	0.896	0.806	1.055	0.812	1.098	0.904	0	-0.17	0.16	0.31	-0.08	0.30	-0.13	0.15	-1.55E-02	0.9185
F1S7V6	Uncharacterized protein OS=Sus scrofa GN=PPP1R14C PE=4 SV=1 - [F1S7V6_PIG]	1.085	0.97	1.273	1.246	1.073	0.969	0.998	0	-0.12	0.04	-0.35	-0.32	-0.10	0.05	0.00	1.28E-02	0.9179
I3LEW5	Uncharacterized protein OS=Sus scrofa GN=PSMD2 PE=4 SV=1 - [I3LEW5_PIG]	0.999	0.835	1.059	0.901	0.911	1.026	1.021	0	0.00	0.26	-0.08	0.15	0.13	-0.04	-0.03	9.74E-03	0.9175
F1SGC6	Uncharacterized protein OS=Sus scrofa GN=NDUF85 PE=4 SV=1 - [F1SGC6_PIG]	0.944	0.853	1.053	1.127	0.962	0.936	0.862	0	0.08	0.23	-0.07	-0.17	0.06	0.10	0.21	-1.14E-02	0.9174
F1SB33	Uncharacterized protein OS=Sus scrofa GN=LOC100518030 PE=4 SV=1 - [F1SB33_PIG]	0.944	1.316	0.964	0.961	1.009	1.178	1.09	0	0.08	-0.40	0.05	0.06	-0.01	-0.24	-0.12	-1.40E-02	0.9170
ASA758	Keratin 1 (Fragment) OS=Sus scrofa GN=KRT1 PE=2 SV=1 - [ASA758_PIG]	1.289	1.227	1.423	1.237	1.208	1.283	1.134	0	-0.37	-0.20	-0.51	-0.37	-0.36	-0.18	1.25E-02	0.9161	
F1ST73	Uncharacterized protein (Fragment) OS=Sus scrofa GN=GRHRP PE=3 SV=2 - [F1ST73_PIG]	0.932	0.933	1.015	0.96	0.996	0.946	0.965	0	0.10	0.10	-0.02	0.06	0.01	0.08	0.05	4.00E-03	0.9154
F1RJH2	Short-chain-specific acyl-CoA dehydrogenase, mitochondrial OS=Sus scrofa GN=ACADS PE=3 SV=2 - [F1RJH2_PIG]	1.163	1.055	1.119	1.062	1.052	1.055	1.188	0	-0.22	-0.08	-0.16	-0.09	-0.07	-0.08	-0.25	-7.10E-03	0.9153
B2ZFN7	Actin (Fragment) OS=Sus scrofa PE=2 SV=1 - [B2ZFN7_PIG]	0.832	1.115	1.185	1.039	1.012	1.11	0.977	0	0.27	-0.16	-0.24	-0.06	-0.02	-0.15	0.03	-1.32E-02	0.9149
F1SNF6	Uncharacterized protein OS=Sus scrofa GN=CHCHD3 PE=4 SV=2 - [F1SNF6_PIG]	1.01	0.881	0.926	1.026	0.903	1.017	0.853	0	-0.01	0.18	0.11	-0.04	0.15	-0.02	0.23	8.97E-03	0.9149
P42639	Tropomyosin alpha-1 chain OS=Sus scrofa GN=TPM1 PE=1 SV=2 - [TPM1_PIG]	0.942	0.951	1.176	0.902	1.251	1.036	0.938	0	0.09	0.07	-0.23	0.15	-0.32	-0.05	0.09	-1.44E-02	0.9144
Q6UAQ8	Electron transfer flavoprotein subunit beta OS=Sus scrofa GN=ETFB PE=2 SV=3 - [ETFB_PIG]	0.97	0.819	0.851	0.914	0.936	0.913	0.845	0	0.04	0.29	0.23	0.13	0.10	0.13	0.24	8.67E-03	0.9144
F1S9D6	Myosin-7 OS=Sus scrofa GN=MYH7 PE=4 SV=1 - [F1S9D6_PIG]	0.937	1.317	1.426	1.219	1.139	1.094	1.105	0	0.09	-0.40	-0.51	-0.29	-0.19	-0.13	-0.14	1.71E-02	0.9143
F1SQP4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=NDUFA12 PE=4 SV=1 - [F1SQP4_PIG]	0.941	0.81	0.94	1.047	0.959	0.821	0.841	0	0.09	0.30	0.09	-0.07	0.06	0.28	0.25	1.19E-02	0.9135
F1SC07	Uncharacterized protein (Fragment) OS=Sus scrofa GN=DNAH11 PE=4 SV=2 - [F1SC07_PIG]	1.089	1.01	0.999	0.959	0.944	1.102	1.127	0	-0.12	-0.01	0.00	0.06	0.08	-0.14	-0.17	-8.29E-03	0.9130
I3LUE8	Uncharacterized protein (Fragment) OS=Sus scrofa GN=IARS2 PE=4 SV=1 - [I3LUE8_PIG]	1.508	1.055	1.44	1.205	1.293	1.211	1.155	0	-0.59	-0.08	-0.53	-0.27	-0.37	-0.28	-0.21	1.80E-02	0.9115
I3L594	Uncharacterized protein OS=Sus scrofa GN=EIF4H PE=4 SV=1 - [I3L594_PIG]	1.053	0.726	0.923	0.942	0.866	0.933	0.891	0	-0.07	0.46	0.12	0.09	0.21	0.10	0.17	1.43E-02	0.9105
G0Z3A1	Phosphoglucutase 1 OS=Sus scrofa GN=PGM1 PE=2 SV=1 - [G0Z3A1_PIG]	0.86	1.342	1.36	1.051	1.386	1.07	1.071	0	0.22	-0.42	-0.44	-0.07	-0.47	-0.10	-0.10	-2.22E-02	0.9102
F1SM98	Uncharacterized protein OS=Sus scrofa GN=NDUFV2 PE=4 SV=1 - [F1SM98_PIG]	1.048	0.746	0.918	1.003	1.052	0.786	0.824	0	-0.07	0.42	0.12	0.00	-0.07	0.35	-0.28	1.77E-02	0.9101
F1SE26	Uncharacterized protein OS=Sus scrofa GN=VDAC1P5 PE=4 SV=2 - [F1SE26_PIG]	1.051	1.032	0.965	0.996	1.048	0.988	1.004	0	-0.07	-0.05	0.05	0.01	-0.07	0.02	-0.01	3.90E-03	0.9098
F6PSL2	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MLC2V PE=4 SV=1 - [F6PSL2_PIG]	0.937	0.966	0.947	0.903	1.087	1.193	0.865	0	0.09	0.05	0.08	0.15	0.17	-0.25	0.21	1.31E-02	0.9093
K9IVW4	Myotrophin OS=Sus scrofa GN=MTPN PE=4 SV=1 - [K9IVW4_PIG]	0.873	0.932	0.96	1.03	0.933	0.918	0.903	0	0.20	0.10	0.06	-0.04	0.10	0.12	0.15	-7.10E-03	0.9087
F1RYY6	Transaldolase (Fragment) OS=Sus scrofa GN=TALDO1 PE=3 SV=2 - [F1RYY6_PIG]	0.803	0.681	0.919	0.812	1.015	0.978	0.579	0	0.32	0.55	0.12	0.30	-0.02	0.03	0.79	2.67E-02	0.9079
I3LEL7	Uncharacterized protein OS=Sus scrofa GN=LOC10739335 PE=4 SV=1 - [I3LEL7_PIG]	0.848	0.806	1.237	0.748	0.953	1.134	0.982	0	0.24	0.31	-0.31	0.42	0.07	-0.18	0.03	2.27E-02	0.9071
I3LD72	Uncharacterized protein OS=Sus scrofa GN=EHD2 PE=4 SV=1 - [I3LD72_PIG]	1.028	0.979	0.965	0.974	0.916	1.02	1.05	0	-0.04	0.03	0.05	0.04	0.13	-0.03	-0.07	5.86E-03	0.9055
Q2EN81	ATP synthase subunit O, mitochondrial OS=Sus scrofa GN=ATP5O PE=1 SV=1 - [ATP5O_PIG]	1.122	1.01	1.005	1.221	0.99	1.033	0.941	0	-0.17	-0.01	-0.01	-0.29	0.01	-0.05	0.09	-1.13E-02	0.9052
I3LCX3	Uncharacterized protein OS=Sus scrofa GN=DUSP3 PE=4 SV=1 - [I3LCX3_PIG]	0.978	0.947	1.017	1.012	0.977	0.985	0.958	0	0.03	0.08	-0.02	-0.02	0.03	0.02	0.06	3.43E-03	0.9052
F1RNN0	Uncharacterized protein OS=Sus scrofa GN=VPS29 PE=4 SV=2 - [F1RNN0_PIG]	1.205	1.312	1.174	1.288	1.19	1.067	1.095	0	-0.27	-0.39	-0.23	-0.37	-0.25	-0.09	-0.13	1.29E-02	0.9038
F1SQY0	Uncharacterized protein OS=Sus scrofa GN=LOC100156975 PE=4 SV=1 - [F1SQY0_PIG]	1.008	0.714	0.932	0.962	0.597	1.083	0.996	0	-0.01	0.49	0.10	0.06	0.74	-0.12	0.01	2.87E-02	0.9033
F1RZR6	Uncharacterized protein OS=Sus scrofa GN=CUTA PE=4 SV=1 - [F1RZR6_PIG]	1.02	0.927	0.985	1.023	0.91	0.996	0.988	0	-0.03	0.11	0.02	-0.03	0.14	0.01	0.02	5.97E-03	0.9032
I3LBK9	Uncharacterized protein OS=Sus scrofa GN=DNAJC3 PE=4 SV=1 - [I3LBK9_PIG]	0.564	0.655	0.263	0.805	0.303	0.636	0.523	0	0.83	0.61	1.93	0.31	1.72	0.65	0.94	6.50E-02	0.9012
I3LPW0	Uncharacterized protein OS=Sus scrofa GN=NDUFB4 PE=4 SV=1 - [I3LPW0_PIG]	0.959	0.696	0.755	1.033	0.785	0.826	0.707	0	0.06	0.52	0.41	-0.05	0.35	0.28	0.50	2.24E-02	0.9008
F2Z5G3	Uncharacterized protein OS=Sus scrofa GN=LOC100522926 PE=4 SV=1 - [F2Z5G3_PIG]	0.622	1.408	1.316	0.809	1.148	1.343	0.822	0	0.69	-0.49	-0.40	0.31	-0.20	-0.43	0.28	4.22E-02	0.9004
F1RYS9	Uncharacterized protein OS=Sus scrofa GN=FAM47E-STBD1 PE=4 SV=2 - [F1RYS9_PIG]	0.981	1.242	1.216	1.129	1.008	1.205	1.039	0	0.03	-0.31	-0.28	-0.18	-0.01	-0.27	-0.06	1.41E-02	0.9000
E7BXW9	Musculoskeletal embryonic nuclear protein 1 OS=Sus scrofa GN=MUSTN1 PE=4 SV=1 - [E7BXW9_PIG]	1.344	0.798	0.88	1.049	1.017	0.927	1.015	0	-0.43	0.33	0.18	-0.07	-0.02	0.11	-0.02	-2.22E-02	0.8990
I3LG74	Uncharacterized protein OS=Sus scrofa GN=KPN4A PE=4 SV=1 - [I3LG74_PIG]	1.091	1.807	1.626	1.319	1.711	1.345	1.154	0	-0.13	-0.85	-0.70	-0.40	-0.77	-0.43	-0.21	-3.20E-02	0.8988
F1S7K4	Uncharacterized protein OS=Sus scrofa GN=PLIN4 PE=4 SV=2 - [F1S7K4_PIG]	1.015	0.958	0.982	1.046	0.991	0.992	0.941	0	-0.02	0.06	0.03	-0.06	0.01</				

Q6QAT1	40S ribosomal protein S28 OS=Sus scrofa GN=RP528 PE=3 SV=2 - [RS28_PIG]	1.004	0.863	0.721	0.893	0.87	0.861	0.984	0	-0.01	0.21	0.47	0.16	0.20	0.22	0.02	-1.88E-02	0.8814
F1RW61	Uncharacterized protein OS=Sus scrofa GN=PYURF PE=4 SV=1 - [F1RW61_PIG]	0.983	1.321	0.743	1.01	1.039	0.894	0.953	0	0.02	-0.40	0.43	-0.01	-0.06	0.16	0.07	2.75E-02	0.8812
I3LDC3	Uncharacterized protein OS=Sus scrofa GN=NDUF810 PE=4 SV=1 - [I3LDC3_PIG]	1.083	0.807	0.921	0.974	0.994	0.884	0.861	0	-0.05	0.31	0.12	0.04	0.01	0.18	0.22	1.48E-02	0.8805
Q9GJT2	S-formylglutathione hydrolase OS=Sus scrofa GN=ESD PE=2 SV=1 - [ESTD_PIG]	0.993	1.011	1.161	0.998	1.129	0.958	1.118	0	0.01	-0.02	-0.22	0.00	-0.18	0.06	-0.16	-1.25E-02	0.8804
Q9T5X9	Peroxisidoxin-6 OS=Sus scrofa GN=PRDX6 PE=2 SV=3 - [PRDX6_PIG]	0.937	0.995	0.973	0.949	1.049	0.927	1.003	0	0.09	0.01	0.04	0.08	-0.07	0.11	0.00	-7.26E-03	0.8783
Q29582	Pyruvate kinase M2 (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29582_PIG]	0.989	1.088	1.2	1.043	1.135	0.991	1.064	0	0.02	-0.12	-0.26	-0.06	-0.18	0.01	-0.09	1.22E-02	0.8781
A1EAJ7	Bridging integrator 1 OS=Sus scrofa PE=2 SV=1 - [A1EAJ7_PIG]	0.98	0.985	1.116	1.065	0.947	1.014	1.083	0	0.03	0.02	-0.16	-0.09	0.08	-0.02	-0.12	-1.00E-02	0.8772
E3VVJ2	Prosaposin variant 2 OS=Sus scrofa GN=PSAP PE=2 SV=1 - [E3VVJ2_PIG]	1.062	0.742	0.663	0.655	0.896	0.92	0.884	0	-0.09	0.43	0.59	0.61	0.16	0.12	0.18	3.26E-02	0.8764
F1S733	Uncharacterized protein OS=Sus scrofa GN=UTRN PE=4 SV=2 - [F1S733_PIG]	1.085	1.009	0.912	0.853	1.177	0.901	1.047	0	-0.12	-0.01	0.13	0.23	-0.24	0.15	-0.07	1.90E-02	0.8764
P79263	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Sus scrofa GN=ITIH4 PE=1 SV=1 - [ITIH4_PIG]	1.069	0.886	1.193	0.899	1.136	1.068	1.092	0	-0.10	0.17	-0.25	0.15	-0.18	-0.09	-0.13	-1.90E-02	0.8760
F1S970	Uncharacterized protein (Fragment) OS=Sus scrofa GN=DNAJB4 PE=4 SV=1 - [F1S970_PIG]	0.974	0.778	0.809	0.769	0.914	0.921	1.002	0	0.04	0.36	0.31	0.38	0.13	0.12	0.00	-2.04E-02	0.8729
F1SRG2	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 OS=Sus scrofa GN=NDUFA6 PE=3 SV=2 - [F1SRG2_PIG]	5.383	6.147	4.956	2.445	6.43	5.121	1.408	0	-2.43	-2.62	-2.31	-1.29	-2.68	-2.36	-0.49	1.33E-01	0.8725
P10173	Fumarate hydratase, mitochondrial OS=Sus scrofa GN=FH PE=1 SV=1 - [FUMH_PIG]	1.197	1.02	1.265	1.245	1.061	1.002	1.11	0	-0.26	-0.03	-0.34	-0.32	-0.09	0.00	-0.15	1.80E-02	0.8719
F1RX68	Serine/threonine-protein phosphatase (Fragment) OS=Sus scrofa GN=PPP2CA PE=3 SV=2 - [F1RX68_PIG]	0.853	0.767	0.718	0.858	0.912	0.766	0.828	0	0.23	0.38	0.48	0.22	0.13	0.38	0.27	-1.98E-02	0.8708
K9IVQ6	Plectin (Fragment) OS=Sus scrofa GN=PLEC PE=2 SV=1 - [K9IVQ6_PIG]	1.051	0.961	1.048	0.992	1.034	1.033	1.015	0	-0.07	0.06	-0.07	0.01	-0.05	-0.05	-0.02	-5.74E-03	0.8707
Q007T2	Cell division control protein 42 homolog OS=Sus scrofa GN=CDC42 PE=2 SV=2 - [CDC42_PIG]	0.961	1.02	1.026	1.216	1.036	0.807	0.932	0	0.06	-0.03	-0.04	-0.28	-0.05	0.31	0.10	2.15E-02	0.8707
Q9GIX2	Diazepam binding inhibitor (Fragment) OS=Sus scrofa GN=DBI PE=2 SV=1 - [Q9GIX2_PIG]	0.869	1.05	0.837	1.088	0.884	1.007	0.736	0	0.20	-0.07	0.26	-0.12	0.18	-0.01	0.44	2.49E-02	0.8705
E1CAJ6	Protein disulfide isomerase P5 OS=Sus scrofa GN=pidi-p5 PE=2 SV=1 - [E1CAJ6_PIG]	0.986	1.088	0.787	0.989	1.001	0.908	0.987	0	0.02	-0.12	0.35	0.02	0.00	0.14	0.02	-1.79E-02	0.8702
F1S9Q3	Uncharacterized protein OS=Sus scrofa PE=3 SV=2 - [F1S9Q3_PIG]	0.953	1.045	1.02	0.965	1.091	0.961	1.029	0	0.07	-0.06	-0.03	0.05	-0.13	0.06	-0.04	-8.87E-03	0.8695
F2ZSE6	Uncharacterized protein OS=Sus scrofa GN=RP55 PE=3 SV=1 - [F2ZSE6_PIG]	1.008	1.248	0.775	1.223	0.787	0.984	1.131	0	-0.01	-0.32	0.37	-0.29	0.35	0.02	-0.18	-3.40E-02	0.8695
F1RXA7	Uncharacterized protein OS=Sus scrofa GN=LOC100512253 PE=4 SV=2 - [F1RXA7_PIG]	0.9	0.889	0.862	1.133	0.692	1.068	0.892	0	0.15	0.17	0.21	-0.18	0.53	-0.09	0.16	-2.88E-02	0.8685
F1S8Q4	Uncharacterized protein OS=Sus scrofa GN=ADCK3 PE=4 SV=2 - [F1S8Q4_PIG]	1.122	0.988	0.902	1.098	0.868	1	1.002	0	-0.17	0.02	0.15	-0.13	0.20	0.00	0.00	1.66E-02	0.8675
P00355	Glyceraldehyde-3-phosphate dehydrogenase OS=Sus scrofa GN=GAPDH PE=1 SV=4 - [G3P_PIG]	0.918	0.919	1.017	0.937	0.975	0.953	0.966	0	0.12	0.12	-0.02	0.09	0.04	0.07	0.05	7.20E-03	0.8670
I3LQN4	Uncharacterized protein OS=Sus scrofa GN=PHB2 PE=4 SV=1 - [I3LQN4_PIG]	0.997	1.032	0.97	1.197	0.967	0.976	0.921	0	0.00	-0.05	0.04	-0.26	0.05	0.04	0.12	-1.50E-02	0.8667
A5GF43	ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit OS=Sus scrofa GN=ATP5E PE=4 SV=1 - [A5GF43_PIG]	1.589	1.043	1.189	1.27	1.162	1.049	1.177	0	-0.67	-0.06	-0.25	-0.34	-0.22	-0.07	-0.24	2.83E-02	0.8666
Q3YB86	Plasma membrane calcium ATPase 4 variant b (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q3YB86_PIG]	1.112	1.219	1.112	1.164	0.926	1.231	1.199	0	-0.15	-0.29	-0.15	-0.22	0.11	-0.30	-0.26	-1.95E-02	0.8665
F1SECO	Uncharacterized protein OS=Sus scrofa GN=DCNT3 PE=4 SV=1 - [F1SECO_PIG]	0.741	0.993	0.927	0.887	0.921	1.019	0.866	0	0.43	0.01	0.11	0.17	0.12	-0.03	0.21	-2.00E-02	0.8664
K7GLH9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=IGBP1 PE=4 SV=1 - [K7GLH9_PIG]	0.826	0.865	0.668	1.016	0.586	0.934	0.955	0	0.28	0.21	0.58	-0.02	0.77	0.10	0.07	-3.85E-02	0.8661
Q5S3G4	Cytochrome c oxidase subunit 5B, mitochondrial OS=Sus scrofa GN=COX5B PE=2 SV=1 - [COX5B_PIG]	1.276	0.96	1.047	0.996	1.105	1.072	1.037	0	-0.35	0.06	-0.07	0.01	-0.14	-0.10	-0.05	1.70E-02	0.8659
I3LUC7	Uncharacterized protein OS=Sus scrofa GN=LOC100517662 PE=4 SV=1 - [I3LUC7_PIG]	0.702	0.548	0.841	0.807	0.825	0.639	0.842	0	0.51	0.87	0.25	0.31	0.28	0.65	0.25	-3.67E-02	0.8653
ASGHK3	Prefoldin 4 OS=Sus scrofa GN=PFND4 PE=4 SV=1 - [ASGHK3_PIG]	0.882	0.898	0.74	1.107	0.682	1.037	0.755	0	0.03	0.16	0.43	-0.15	0.55	-0.05	0.41	3.57E-02	0.8626
B5SYT7	Adenosine monophosphate deaminase 1 isoform M OS=Sus scrofa GN=AMPD1 PE=2 SV=1 - [B5SYT7_PIG]	0.875	1.228	1.138	1.14	1.116	1.03	0.99	0	0.19	-0.30	-0.19	-0.19	-0.16	-0.04	-0.01	-2.13E-02	0.8621
I3L815	Uncharacterized protein (Fragment) OS=Sus scrofa GN=EIF6 PE=3 SV=1 - [I3L815_PIG]	0.847	0.869	0.623	0.744	0.764	0.726	1.013	0	0.24	0.20	0.68	0.43	0.39	0.46	-0.02	3.34E-02	0.8608
F1RLH7	Uncharacterized protein OS=Sus scrofa GN=COX6B PE=4 SV=2 - [F1RLH7_PIG]	1.332	0.998	0.953	1.034	1.031	1.062	1.057	0	-0.41	0.00	0.07	-0.05	-0.04	-0.09	-0.08	2.06E-02	0.8594
F1SBN7	Uncharacterized protein OS=Sus scrofa GN=ATP5F1 PE=4 SV=1 - [F1SBN7_PIG]	0.871	0.996	0.876	1.232	0.882	0.98	0.772	0	0.20	0.01	0.19	-0.30	0.18	0.03	0.37	-2.84E-02	0.8590
Q95249	Myosin heavy chain (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q95249_PIG]	0.998	1.582	1.585	1.17	1.418	1.197	1.134	0	0.00	-0.66	-0.66	-0.23	-0.50	-0.26	-0.18	3.80E-02	0.8589
F1SHL9	Pyruvate kinase (Fragment) OS=Sus scrofa GN=PKM PE=3 SV=2 - [F1SHL9_PIG]	0.969	1.037	1.135	1.027	1.069	0.981	1.029	0	0.05	-0.05	-0.18	-0.04	-0.10	0.03	-0.04	1.04E-02	0.8581
A0PFK6	F-actin capping protein subunit alpha 2 OS=Sus scrofa GN=CP alpha2 PE=2 SV=1 - [A0PFK6_PIG]	0.952	1.087	1.114	1.013	1.01	1.04	1.053	0	0.07	-0.12	-0.16	-0.02	-0.01	-0.06	-0.07	1.03E-02	0.8572
F1SV23	Uncharacterized protein OS=Sus scrofa GN=NDUF85 PE=4 SV=1 - [F1SV23_PIG]	0.963	0.915	0.94	1.135	1.101	0.919	0.773	0	0.05	-0.13	0.09	-0.18	-0.14	0.12	0.37	-2.50E-02	0.8553
G9F6X6	Mitochondrial heat shock 60 kDa protein 1 OS=Sus scrofa PE=2 SV=1 - [G9F6X6_PIG]	0.939	1.1	1.08	1.026	1.188	1.021	0.963	0	0.09	0.14	-0.15	-0.04	-0.25	-0.03	0.05	-1.66E-02	0.8537
C3URJ6	Mitochondrial carnitine palmitoyltransferase II OS=Sus scrofa GN=CPT2 PE=2 SV=1 - [C3URJ6_PIG]	1.09	0.844	1.011	0.92	0.982	0.983	1.102	0	-0.12	0.24	-0.02	0.12	0.03	0.02	-0.14	-1.84E-02	0.8529
K7GPC2	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LICAM PE=4 SV=1 - [K7GPC2_PIG]	0.95	1.15	1.361	1.036	1.219	0.942	1.157	0	-0.20	-0.44	-0.05	-0.29	0.09	-0.21	2.78E-02	0.8518	
F1RKM1	Uncharacterized protein (Fragment) OS=Sus scrofa GN=ALDH7A1 PE=3 SV=2 - [F1RKM1_PIG]	0.945	0.819	0.919	1.027	0.908	0.871	0.915	0	0.08	0.29	0.12	-0.04	0.14	0.20	0.13	-1.58E-02	0.8481
F1RZQ6	Uncharacterized protein OS=Sus scrofa GN=SLC25A4 PE=3 SV=1 - [F1RZQ6_PIG]	0.97	1.314	1.121	1.488	1.051	1.078	0.934	0	0.04	-0.39	-0.16	-0.57	-0.34	-0.11	0.10	-3.50E-02	0.8476
D2WKD8	Sodium/potassium-transporting ATPase subunit alpha-2 OS=Sus scrofa GN=ATP1A2 PE=1 SV=1 - [AT1A2_PIG]	0.992	1.048	0.732	0.99	0.86	0.942	1.021	0	0.01	-0.07	0.45	0.01	0.22	0.09	-0.03	-2.64E-02	0.8459
F1S1A9	Uncharacterized protein OS=Sus scrofa GN=APOA2 PE=4 SV=1 - [F1S1A9_PIG]	2.461	2.047	1.311	1.035	2.505	1.73	1.165	0	-1.30	-1.03	-0.39	-0.05	-1.32	-0.79	-0.22	8.45E-02	0.8453
I3LIE7	Uncharacterized protein OS=Sus scrofa GN=MYBPH PE=4 SV=1 - [I3LIE7_PIG]	0.873	1.193	0.945	0.909	1.012	1.086	1.049	0	0.20	-0.25	0.08	0.14	-0.02	-0.12	-0.07	-2.26E-02	0.8446
I3L8H4	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3L8H4_PIG]	1.087	0.692	0.813	0.893	0.839	0.866	0.866	0	-0.12	0.53	0.30	0.16	0.25	0.21	0.21	3.05E-02	0.8437
F1S814	Uncharacterized protein OS=Sus scrofa GN=PGM1 PE=3 SV=2 - [F1S814_PIG]	0.827	1.048	1.122	0.917	1.065	0.93	1.007	0	0.27	-0.07	-0.17	0.13	-0.09	0.10	-0.01	2.21E-02	0.8432
M3TYC8	DEAD (Asp-Glu-Ala-Asp) box helicase 1 OS=Sus scrofa GN=DDX1 PE=2 SV=1 - [M3TYC8_PIG]	0.781	0.776	0.624	0.843	0.679	0.919	0.791	0	0.36	0.37	0.68	0.25	0.56	0.12	0.34	-3.45E-02	0.8431
Q69DL4	Complement C1qB (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q69DL4_PIG]	1.236	0.921	1.246	1.068	0.977	1.226	1.197	0	-0.31	0.12	-0.32	-0.09	0.03	-0.29	-0.26	-2.76E-02	0.8430
F1RV23	cAMP-dependent protein kinase type I-alpha regulatory subunit OS=Sus scrofa GN=PRKARIA PE=4 SV=1 - [F1RV23_PIG]	0.958	1.287	1.207	1.043	1.129	1.115	1.066	0	0.06	-0.36	-0.27	-0.06	-0.18	-0.16	-0.09	2.21E-02	0.8425
Q29387	Elongation factor 1-gamma (Fragment) OS=Sus scrofa GN=EEF1G PE=2 SV=2 - [EF1G_PIG]	1.076	1.192	1.037	1.1	1.174	1.045	1.026	0	-0.11	-0.25	-0.05	-0.14	-0.23	-0.06	-0.04	-1.45E-02	0.8425
Q71UN0	Glyceraldehyde-3-phosphate dehydrogenase (Fragment) OS=Sus scrofa GN=GAPDH PE=2 SV=1 - [Q71UN0_PIG]	0.85	0.745	0.791	0.647	1.064	0.988	0.819	0	0.23	0.42	0.34	0.63	-0.09	0.02	0.29	-3.83E-02	0.8424
I3L660	Profilin (Fragment) OS=Sus scrofa GN=PFN2 PE=3 SV=1 - [I3L660_PIG]	1.061	0.753	0.934	0.767	0.909	1.023	0.964	0	-0.09	0.41	0.10	0.38	0.14	-0.03	0.05	2.95E-02	0.8403
F1STY7	Uncharacterized protein OS=Sus scrofa GN=AAMDCE PE=4 SV=2 - [F1STY7_PIG]	1.012	0.892	1.036	0.967	1.025	0.99	0.982	0	-0.02	0.16	-0.05	0.05	-0.04	0.01	0.03	-1.08E-02	0.8403
I3LC47	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LC47_PIG]	1.004	1.13	1.123	1.071	1.132	0.994	1.097	0	-0.01	-0.18	-0.17	-0.10	-0.18	0.01	-0.13	-1.33E-02	0.8398
Q9TV62	Myosin-4 OS=Sus scrofa GN=MYH4 PE=2 SV=1 - [MYH4_PIG]	0.965	1.275	1.497	1.449	0.981	1.206											

I3LT9	Protein phosphatase methyltransferase 1 (Fragment) OS=Sus scrofa GN=PPME1 PE=3 SV=1 - [I3LT9_PIG]	0.97	1.104	1.136	1.1	1.074	0.989	0.998	0	0.04	-0.14	-0.18	-0.14	-0.10	0.02	0.00	1.53E-02	0.8269
F1SBY1	Ribosomal protein S6 kinase (Fragment) OS=Sus scrofa GN=RPSEK2 PE=3 SV=2 - [F1SBY1_PIG]	0.916	1.129	1.343	1.022	1.154	1.12	0.966	0	0.13	-0.18	-0.43	-0.03	-0.21	-0.16	0.05	3.06E-02	0.8263
A1XQT5	HUMMLC2B OS=Sus scrofa GN=HUMMLC2B PE=3 SV=1 - [A1XQT5_PIG]	0.864	0.832	0.885	0.842	0.885	0.932	0.879	0	0.21	0.27	0.18	0.25	0.18	0.10	0.19	1.49E-02	0.8259
P00506	Aspartate aminotransferase, mitochondrial OS=Sus scrofa GN=GOT2 PE=1 SV=2 - [AATM_PIG]	1.108	0.885	0.881	0.954	1.006	0.91	0.937	0	-0.15	0.18	0.18	0.07	-0.01	0.14	0.09	1.95E-02	0.8254
I3LKF3	Fascin OS=Sus scrofa GN=LOC100625519 PE=3 SV=1 - [I3LKF3_PIG]	0.993	0.867	0.968	0.964	0.94	0.987	0.962	0	0.01	0.21	0.05	0.05	0.09	0.02	0.06	-1.15E-02	0.8254
F1SV22	Uncharacterized protein OS=Sus scrofa GN=MACF1 PE=4 SV=2 - [F1SV22_PIG]	1.107	1.12	1.427	1.029	1.275	1.113	1.117	0	-0.15	-0.16	-0.51	-0.04	-0.35	-0.15	-0.16	2.93E-02	0.8241
I3LCN1	Uncharacterized protein OS=Sus scrofa GN=ENO2 PE=3 SV=1 - [I3LCN1_PIG]	0.917	0.796	0.916	0.921	0.981	0.963	0.815	0	0.13	0.33	0.13	0.12	0.03	0.05	0.30	-2.12E-02	0.8233
F1SGR8	Uncharacterized protein OS=Sus scrofa GN=SDR39U1 PE=4 SV=1 - [F1SGR8_PIG]	0.973	0.995	0.853	1.103	0.968	0.9	0.909	0	0.04	0.01	0.23	-0.14	0.05	0.15	0.14	-2.02E-02	0.8229
F1RMN7	Hemopexin OS=Sus scrofa GN=HPX PE=4 SV=2 - [F1RMN7_PIG]	0.913	0.996	0.944	0.845	1.157	1.012	0.927	0	0.13	0.01	0.08	0.24	-0.21	-0.02	0.11	-2.39E-02	0.8219
F1RL74	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=2 - [F1RL74_PIG]	0.95	1.005	0.895	0.983	0.921	0.95	1.028	0	0.07	-0.01	0.16	0.02	0.12	0.07	-0.04	-1.23E-02	0.8200
A4Z6H0	Adenylosuccinate synthetase isozyme 1 OS=Sus scrofa GN=ADSSL1 PE=2 SV=1 - [PURA1_PIG]	0.964	0.863	0.873	0.834	0.959	0.937	0.926	0	0.05	0.21	0.20	0.26	0.06	0.09	0.11	1.64E-02	0.8197
Q9GJV4	Four and a half LIM domains 1 protein, isoform C OS=Sus scrofa GN=fhl1C PE=2 SV=1 - [Q9GJV4_PIG]	1.111	1.09	1.218	1.24	1.044	0.882	1.189	0	-0.15	-0.12	-0.28	-0.31	-0.06	0.18	-0.25	2.99E-02	0.8189
I3LIJ3	Uncharacterized protein OS=Sus scrofa GN=EEF2 PE=4 SV=1 - [I3LIJ3_PIG]	0.965	1.154	1.061	1.077	1.084	1.004	1.051	0	0.05	-0.21	-0.09	-0.11	-0.12	-0.01	-0.07	-1.51E-02	0.8153
F2Z545	Uncharacterized protein OS=Sus scrofa GN=TCEB1 PE=4 SV=1 - [F2Z545_PIG]	0.851	0.898	1.017	0.956	0.915	1.004	0.927	0	0.23	0.16	-0.02	0.06	0.13	-0.01	0.11	-1.67E-02	0.8152
I3LCE9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100738863 PE=4 SV=1 - [I3LCE9_PIG]	0.608	0.828	0.542	0.792	0.571	0.726	0.712	0	0.72	0.27	0.88	0.34	0.81	0.46	0.49	5.58E-02	0.8134
F1SSZ2	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CHD5 PE=4 SV=2 - [F1SSZ2_PIG]	1.104	1.748	1.251	1.11	1.302	1.197	1.23	0	-0.14	-0.81	-0.32	-0.15	-0.38	-0.26	-0.30	4.55E-02	0.8107
F2Z558	Uncharacterized protein (Fragment) OS=Sus scrofa GN=YWHAZ PE=3 SV=2 - [F2Z558_PIG]	0.943	1.043	0.919	1	1.003	1.099	0.868	0	0.08	-0.06	0.12	0.00	0.00	-0.14	0.20	-2.05E-02	0.8096
F1SR5Y	Uncharacterized protein OS=Sus scrofa GN=PGAM1 PE=3 SV=2 - [F1SR5Y_PIG]	0.874	0.937	0.973	1.064	0.817	1.023	0.831	0	0.19	0.09	0.04	-0.09	0.29	-0.03	0.27	2.72E-02	0.8092
F1RW7H	Uncharacterized protein (Fragment) OS=Sus scrofa GN=PNPO PE=4 SV=2 - [F1RW7H_PIG]	0.786	0.789	1.103	0.907	0.886	0.612	1.182	0	0.35	0.34	-0.14	0.14	0.17	0.71	-0.24	5.87E-02	0.8079
K9IW73	Dihydrolipoyllysine-residue succinyltransferase OS=Sus scrofa GN=DLST-v1 PE=2 SV=1 - [K9IW73_PIG]	1.191	0.895	0.911	0.906	1.028	1.033	1.091	0	-0.25	0.16	0.13	0.14	-0.04	-0.05	-0.13	-2.81E-02	0.8073
P62197	26S protease regulatory subunit 8 OS=Sus scrofa GN=PSMCS PE=2 SV=1 - [PKS8_PIG]	0.963	1.039	0.883	1.066	0.919	0.982	0.867	0	0.05	-0.06	0.18	-0.09	0.12	0.03	0.21	2.08E-02	0.8072
Q9GL01	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase subunit 2 OS=Sus scrofa GN=RPN2 PE=2 SV=1 - [RPN2_PIG]	1.085	1.295	1.214	1.185	1.016	1.093	1.207	0	-0.12	-0.37	-0.28	-0.24	-0.02	-0.13	-0.27	2.57E-02	0.8072
F1SRB9	Uncharacterized protein OS=Sus scrofa GN=ST13 PE=4 SV=1 - [F1SRB9_PIG]	1.015	1.005	1.031	0.885	1.058	0.969	1.102	0	-0.02	-0.01	-0.04	0.18	-0.08	0.05	-0.14	1.82E-02	0.8059
O77773	Uncharacterized protein OS=Sus scrofa GN=CACNA2D1 PE=2 SV=1 - [O77773_PIG]	0.934	1.05	1.112	1.058	1.027	1.039	1.005	0	0.10	-0.07	-0.15	-0.08	-0.04	-0.06	-0.01	-1.43E-02	0.8059
F1SJP6	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 (Fragment) OS=Sus scrofa GN=NDUFA6 PE=3 SV=1 - [F1SJP6_PIG]	0.288	0.199	0.301	0.487	0.31	0.217	0.353	0	1.80	2.33	1.73	1.04	-1.69	2.20	1.50	1.44E-01	0.8056
I3L7X3	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3L7X3_PIG]	0.981	1.182	1.441	1.124	1.105	0.934	1.285	0	0.03	-0.24	-0.53	-0.17	-0.14	0.10	-0.36	4.12E-02	0.8053
Q2VTP6	Peptidyl-prolyl cis-trans isomerase OS=Sus scrofa GN=FKBP1A PE=4 SV=1 - [Q2VTP6_PIG]	0.984	0.857	0.946	0.945	0.872	0.907	1.016	0	0.02	0.22	0.08	0.08	0.20	0.14	-0.02	1.78E-02	0.8043
I3L893	Uncharacterized protein OS=Sus scrofa GN=GDH1 PE=4 SV=1 - [I3L893_PIG]	0.835	0.675	0.63	0.834	0.645	1.096	0.709	0	0.26	0.57	0.67	0.26	0.63	-0.13	0.50	-5.88E-02	0.8032
I3LDC2	Uncharacterized protein OS=Sus scrofa GN=REM1 PE=4 SV=1 - [I3LDC2_PIG]	0.901	0.742	0.883	0.715	1.061	0.938	0.924	0	0.15	0.43	0.18	0.48	-0.09	0.09	0.11	-3.89E-02	0.8032
F2Z5K2	Proteasome subunit alpha type OS=Sus scrofa GN=PSMA5 PE=3 SV=1 - [F2Z5K2_PIG]	0.908	1.214	1.218	1.026	1.137	1.169	1.069	0	0.14	-0.28	-0.28	-0.04	-0.19	-0.23	-0.10	-2.97E-02	0.8028
F1RG85	Uncharacterized protein OS=Sus scrofa GN=MYO18B PE=4 SV=2 - [F1RG85_PIG]	0.828	3.301	0.728	1.174	1.124	0.897	1.159	0	0.27	-1.72	0.46	-0.23	-0.17	0.16	-0.21	1.34E-01	0.8009
Q0QEV2	60S ribosomal protein L18 (Fragment) OS=Sus scrofa GN=RPL18 PE=2 SV=1 - [Q0QEV2_PIG]	0.948	1.174	0.895	1.369	0.884	0.982	0.947	0	0.08	-0.23	0.16	-0.45	0.18	0.03	0.08	-4.40E-02	0.7970
I3LCX2	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3LCX2_PIG]	1.096	0.816	1.089	0.943	1.049	1.141	0.766	0	-0.13	0.29	-0.12	0.08	-0.07	-0.19	0.38	4.30E-02	0.7961
F1SSS1	Uncharacterized protein OS=Sus scrofa GN=LOC100153701 PE=4 SV=2 - [F1SSS1_PIG]	0.901	0.97	0.83	1.092	0.832	0.876	0.839	0	0.15	0.04	0.27	-0.13	0.27	0.19	0.25	2.99E-02	0.7953
I3LJ05	Uncharacterized protein OS=Sus scrofa GN=C21orf33 PE=4 SV=1 - [I3LJ05_PIG]	1.055	0.944	0.991	0.957	1.068	0.982	0.946	0	-0.08	0.08	0.01	0.06	-0.09	0.03	0.08	1.40E-02	0.7950
F1SPG0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=RAB7A PE=4 SV=2 - [F1SPG0_PIG]	1.391	1.39	0.819	0.977	0.981	1.442	0.959	0	-0.48	-0.48	0.29	0.03	0.03	-0.53	0.06	6.42E-02	0.7948
F1RWK8	Uncharacterized protein OS=Sus scrofa GN=CACNB1 PE=4 SV=1 - [F1RWK8_PIG]	0.99	1.124	1.074	1.013	1.146	0.971	1.007	0	0.01	-0.17	-0.10	-0.02	-0.20	0.04	-0.01	1.86E-02	0.7935
Q1W0Y0	Cytochrome c oxidase subunit VIIc (Fragment) OS=Sus scrofa PE=4 SV=1 - [Q1W0Y0_PIG]	1.083	0.563	0.826	0.977	0.935	1.154	0.597	0	-0.12	0.83	0.28	0.03	0.10	-0.21	0.74	-8.04E-02	0.7927
F1RKI3	Uncharacterized protein OS=Sus scrofa GN=HINT1 PE=4 SV=1 - [F1RKI3_PIG]	1.015	0.94	1.018	0.891	1	1.0786	1.141	0	-0.02	0.09	-0.03	0.17	-0.14	0.35	-0.19	3.60E-02	0.7914
I3LQ51	Proteasome subunit beta type OS=Sus scrofa GN=PSMB1 PE=3 SV=1 - [I3LQ51_PIG]	0.877	0.965	0.748	0.875	0.933	0.88	0.815	0	0.19	0.05	0.42	0.19	0.10	0.18	0.30	2.82E-02	0.7914
K7GRY0	Uncharacterized protein OS=Sus scrofa GN=UBA1 PE=3 SV=1 - [K7GRY0_PIG]	1.04	1.069	1	1.064	1.081	1.027	1.062	0	-0.06	-0.10	-0.14	-0.09	-0.11	-0.04	-0.09	-9.18E-03	0.7911
F1RIU5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=PIHGK PE=4 SV=2 - [F1RIU5_PIG]	0.937	1.121	1.173	1.046	1.12	1.001	0.984	0	0.06	-0.12	-0.23	-0.06	-0.16	0.00	0.02	2.36E-02	0.7907
I3LLD5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=EIF4A1 PE=4 SV=1 - [I3LLD5_PIG]	0.999	1.209	0.937	1.03	0.895	1.042	1.089	0	0.00	-0.27	0.09	-0.04	0.16	-0.06	-0.12	2.84E-02	0.7868
F1ST01	Uncharacterized protein OS=Sus scrofa GN=SELENBP1 PE=4 SV=1 - [F1ST01_PIG]	0.955	0.895	0.943	0.92	0.854	1.048	0.927	0	0.07	0.16	0.08	0.12	0.23	-0.07	0.11	1.96E-02	0.7868
F1SGT3	Uncharacterized protein OS=Sus scrofa GN=PDHX PE=3 SV=1 - [F1SGT3_PIG]	1.101	0.884	0.954	0.873	1.041	0.996	0.959	0	-0.14	0.18	0.07	0.20	-0.06	0.01	0.06	2.43E-02	0.7859
I3LEK3	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CALML6 PE=4 SV=1 - [I3LEK3_PIG]	0.571	0.923	0.282	0.921	0.377	0.805	0.797	0	0.81	0.12	1.83	0.12	1.41	0.31	0.33	-1.46E-01	0.7849
I3LFU4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100625259 PE=4 SV=1 - [I3LFU4_PIG]	0.996	1.375	1.362	0.978	1.241	1.099	1.237	0	0.01	-0.46	-0.45	0.03	-0.31	-0.14	-0.31	4.42E-02	0.7848
I3LER5	Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Sus scrofa GN=COX4I1 PE=4 SV=1 - [I3LER5_PIG]	0.991	0.943	0.883	1.08	0.914	0.996	0.891	0	0.01	0.08	0.18	-0.11	0.13	0.01	0.17	-2.16E-02	0.7844
I3LRR4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=NDUFV3 PE=4 SV=1 - [I3LRR4_PIG]	1.053	0.678	0.991	0.915	0.954	0.975	0.935	0	-0.07	0.56	0.01	0.13	0.07	0.04	0.10	-4.24E-02	0.7840
A1XQV4	Tropomyosin alpha-3 chain OS=Sus scrofa GN=TPM3 PE=2 SV=1 - [TPM3_PIG]	1.277	1.115	1.127	0.96	1.006	1.207	1.252	0	-0.35	-0.16	-0.17	0.06	-0.01	-0.27	-0.32	3.42E-02	0.7837
F1S1G8	Amine oxidase OS=Sus scrofa GN=AOC3 PE=3 SV=2 - [F1S1G8_PIG]	0.995	0.963	0.952	0.934	0.991	0.995	0.969	0	0.01	0.05	0.07	0.10	0.01	0.01	0.05	7.91E-03	0.7814
F1RS36	78 kDa glucose-regulated protein OS=Sus scrofa GN=HSPA5 PE=3 SV=2 - [F1RS36_PIG]	1.013	0.942	1.081	0.976	1.033	1.019	1.042	0	-0.02	0.09	-0.11	0.04	-0.05	-0.03	-0.06	-1.34E-02	0.7801
F1RVC9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=HNRNPD PE=4 SV=1 - [F1RVC9_PIG]	1.141	0.916	1.079	0.96	1.182	0.894	1.017	0	-0.19	0.13	-0.11	0.06	-0.24	0.16	-0.02	3.21E-02	0.7798
F1SCC9	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [F1SCC9_PIG]	0.764	0.924	0.577	0.741	0.83	0.959	0.808	0	0.39	0.11	0.79	0.43	0.27	0.06	0.31	-5.66E-02	0.7787
I3LVJ7	Uncharacterized protein (Fragment) OS=Sus scrofa GN=PSMA7 PE=4 SV=1 - [I3LVJ7_PIG]	0.799	0.859	0.88	0.945	0.896	0.871	0.757	0	0.32	0.22	0.18	0.08	0.16	0.20	0.40	2.84E-02	0.7775
Q8SPR7	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase delta-4 OS=Sus scrofa GN=PLCD4 PE=2 SV=1 - [PLCD4_PIG]	0.868	1.013	1.137	0.944	1.146	0.964	1.043	0	0.20	-0.02	-0.19	0.08	-0.20	0.05	-0.06	-3.04E-02	0.7755
F1S073	Annexin OS=Sus scrofa GN=ANXA2 PE=3 SV=2 - [F1S073_PIG]	0.876	0.848	0.828	0.892	0.784	1.051	0.755	0	0.19	0.24	0.27	0.16	0.35	-0.07	0.41	3.71E-02	0.7743
P13618	ATP synthase-cou																	

F1SRCS5	Aconitate hydratase, mitochondrial (Fragment) OS=Sus scrofa GN=ACO2 PE=4 SV=2 - [F1SRCS5_PIG]	1.114	0.909	0.897	0.935	1.048	0.916	0.938	0	-0.16	0.14	0.16	0.10	-0.07	0.13	0.09	2.74E-02	0.7575
I3LQ57	Uncharacterized protein OS=Sus scrofa GN=LOC100739181 PE=4 SV=1 - [I3LQ57_PIG]	1.156	0.978	1.005	0.936	1.078	1.053	1.003	0	-0.21	0.03	-0.01	0.10	-0.11	-0.07	0.00	2.31E-02	0.7561
F1RM74	Glyceraldehyde-3-phosphate dehydrogenase OS=Sus scrofa GN=GAPDH5 PE=3 SV=1 - [F1RM74_PIG]	1.001	0.965	1.006	1.017	0.955	1.025	1.082	0	-0.01	0.05	-0.08	-0.02	0.07	-0.04	-0.11	-1.50E-02	0.7560
ASA779	Geranylgeranyl transferase type-2 subunit alpha OS=Sus scrofa GN=RABGGTA PE=2 SV=1 - [PGTA_PIG]	1.097	1.153	1.113	1.066	1.053	1.176	1.123	0	-0.13	-0.21	-0.15	-0.09	-0.07	-0.23	-0.17	-1.86E-02	0.7553
F1SPT5	Uncharacterized protein OS=Sus scrofa GN=LAMB2 PE=4 SV=2 - [F1SPT5_PIG]	1.028	0.922	1.093	1.259	0.854	1.174	0.934	0	-0.04	0.12	-0.13	-0.33	0.23	-0.23	0.10	-4.66E-02	0.7540
F1SHL3	Uncharacterized protein OS=Sus scrofa GN=SEC23A PE=4 SV=1 - [F1SHL3_PIG]	0.892	0.966	0.924	1.009	0.915	0.877	0.934	0	0.16	0.05	0.11	-0.01	0.13	0.19	0.10	1.86E-02	0.7501
F1SP54	Uncharacterized protein OS=Sus scrofa GN=NIPSNAP3B PE=4 SV=2 - [F1SP54_PIG]	1.172	1.295	1.078	1.106	1.213	1.015	1.099	0	-0.23	-0.37	-0.11	-0.15	-0.28	-0.02	-0.14	3.22E-02	0.7487
F2Z5N0	Proteasome subunit alpha type OS=Sus scrofa GN=PSMA6 PE=3 SV=1 - [F2Z5N0_PIG]	0.919	0.907	0.963	0.978	0.902	0.998	0.951	0	0.12	0.14	0.05	0.03	-0.15	0.00	0.07	-1.52E-02	0.7481
F1SAI8	Uncharacterized protein OS=Sus scrofa GN=HYOU1 PE=3 SV=1 - [F1SAI8_PIG]	1.319	1.212	1.027	1.04	1.043	1.224	1.115	0	-0.40	-0.28	-0.04	-0.06	-0.06	-0.29	-0.16	3.73E-02	0.7471
F1RWK0	Uncharacterized protein OS=Sus scrofa GN=JPH1 PE=4 SV=1 - [F1RWK0_PIG]	0.942	0.973	1.076	0.982	1.018	0.967	0.979	0	0.09	0.04	-0.11	0.03	-0.03	0.05	0.03	1.49E-02	0.7462
I3LN25	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LN25_PIG]	1.191	0.994	1.356	1.026	1.202	1.14	1.289	0	-0.25	0.01	-0.44	-0.04	-0.27	-0.19	-0.37	-4.37E-02	0.7451
F1S880	Uncharacterized protein OS=Sus scrofa GN=NSFL1C PE=4 SV=1 - [F1S880_PIG]	0.898	0.853	0.733	0.922	0.754	0.959	0.947	0	0.16	0.23	0.45	0.12	0.41	0.06	0.08	-4.23E-02	0.7440
Q5S1S4	Carbonic anhydrase 3 OS=Sus scrofa GN=CA3 PE=2 SV=3 - [CAH3_PIG]	0.95	0.886	1.117	0.915	0.987	1.026	1.106	0	0.07	0.17	-0.16	0.13	0.02	-0.04	-0.15	-3.11E-02	0.7431
F1SCF9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=TECR PE=4 SV=2 - [F1SCF9_PIG]	0.912	1.453	1.298	1.571	1.185	1.082	1.04	0	0.13	-0.54	-0.38	-0.65	-0.24	-0.11	-0.06	-7.11E-02	0.7427
I3LS72	Uncharacterized protein OS=Sus scrofa GN=COL6A1 PE=4 SV=1 - [I3LS72_PIG]	0.991	0.744	0.966	1.099	0.884	0.915	0.903	0	0.01	0.43	0.05	-0.14	0.18	0.13	0.15	-4.31E-02	0.7421
F1S232	4-trimethylaminobutylaldehyde dehydrogenase OS=Sus scrofa GN=ALDH9A1 PE=3 SV=2 - [F1S232_PIG]	1.073	1.094	1.071	1.097	0.914	1.152	1.185	0	-0.10	-0.13	-0.10	-0.13	0.13	-0.20	-0.24	-3.07E-02	0.7418
D0G6Y0	Hydroxysteroid (17-beta) dehydrogenase 12 (Fragment) OS=Sus scrofa GN=HSD17B12 PE=2 SV=1 - [D0G6Y0_PIG]	1.048	1.181	1.13	1.397	0.954	1.139	1.043	0	-0.07	-0.24	-0.18	-0.48	0.07	-0.19	-0.06	-4.47E-02	0.7415
F1SS64	Myosin-1 OS=Sus scrofa GN=MYH1 PE=4 SV=1 - [F1SS64_PIG]	1.048	0.849	0.981	0.92	1.036	0.881	0.96	0	-0.07	0.24	0.03	0.12	-0.05	0.18	0.06	2.87E-02	0.7393
F1SAJ9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MAPRE2 PE=4 SV=2 - [F1SAJ9_PIG]	1.043	1.013	1.137	1.043	1.027	1.051	1.024	0	-0.06	-0.02	-0.19	-0.06	-0.04	-0.07	-0.03	1.49E-02	0.7393
F1RV28	Uncharacterized protein OS=Sus scrofa GN=LOC100625946 PE=4 SV=1 - [F1RV28_PIG]	1.017	1.012	0.958	1.073	0.899	1.017	0.949	0	-0.02	-0.02	0.06	-0.10	0.15	-0.02	0.08	2.07E-02	0.7389
F1S827	Uncharacterized protein OS=Sus scrofa GN=SERBP1 PE=4 SV=2 - [F1S827_PIG]	1.054	0.883	0.964	0.904	0.982	0.971	0.981	0	-0.08	0.18	0.05	0.15	0.03	0.04	0.03	2.14E-02	0.7378
F1RLB6	Uncharacterized protein OS=Sus scrofa GN=NAPA PE=4 SV=2 - [F1RLB6_PIG]	0.892	0.944	0.717	0.892	0.809	0.809	0.924	0	0.16	0.08	0.48	0.16	0.31	0.31	0.11	4.06E-02	0.7377
F1S1J5	60S ribosomal protein L4 OS=Sus scrofa GN=RPL4 PE=4 SV=2 - [F1S1J5_PIG]	1.051	1.491	1.211	1.416	1.229	1.098	1.153	0	-0.07	-0.58	-0.28	-0.50	-0.30	-0.13	-0.21	-5.38E-02	0.7348
I3L6D2	Ras-related protein Rab-1A (Fragment) OS=Sus scrofa GN=RAB1A PE=3 SV=1 - [I3L6D2_PIG]	1.097	1.494	1.321	1.325	1.194	1.188	1.319	0	-0.13	-0.58	-0.40	-0.41	-0.26	-0.25	-0.40	-4.89E-02	0.7345
B6CVD6	Thioredoxin domain-containing 4 OS=Sus scrofa GN=TXNDC4 PE=2 SV=1 - [B6CVD6_PIG]	0.786	0.822	0.769	0.912	0.739	0.859	0.769	0	0.35	0.28	0.38	0.13	0.44	0.22	0.38	3.96E-02	0.7343
Q6UAAQ9	Electron transfer flavoprotein alpha subunit (Fragment) OS=Sus scrofa GN=ETFA PE=2 SV=1 - [Q6UAAQ9_PIG]	1.076	0.839	0.936	0.895	1.053	0.89	0.917	0	-0.11	0.25	0.10	0.16	-0.07	0.17	0.13	3.39E-02	0.7340
F1RNR7	Uncharacterized protein OS=Sus scrofa GN=LOC100525555 PE=4 SV=1 - [F1RNR7_PIG]	0.837	0.87	0.635	0.722	0.989	0.95	0.809	0	0.26	0.20	0.66	0.47	0.02	0.07	0.31	-6.18E-02	0.7331
I3L7H0	DCN1-like protein OS=Sus scrofa GN=DCUN1D1 PE=4 SV=1 - [I3L7H0_PIG]	0.701	0.933	0.792	0.87	0.828	0.875	0.926	0	0.51	0.10	0.34	0.20	0.27	0.19	0.11	-4.31E-02	0.7328
F1RN91	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=2 - [F1RN91_PIG]	0.986	0.997	0.936	1.105	0.85	1.09	0.983	0	0.02	0.00	0.10	-0.14	0.23	-0.12	0.02	-3.23E-02	0.7325
I3LKZ1	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LKZ1_PIG]	1.268	1.587	1.765	1.096	1.302	1.292	1.563	0	-0.34	-0.67	-0.82	-0.13	-0.38	-0.37	-0.64	7.54E-02	0.7317
F1ST49	Uncharacterized protein (Fragment) OS=Sus scrofa GN=NCBP1 PE=4 SV=2 - [F1ST49_PIG]	0.88	1.161	1.129	1.094	1.1	1.019	1.035	0	0.18	-0.22	-0.18	-0.13	-0.03	-0.05	-3.45E-02	0.7308	
F1RGD0	Uncharacterized protein OS=Sus scrofa GN=RIC8A PE=4 SV=1 - [F1RGD0_PIG]	0.703	1.005	0.915	1.058	0.785	0.896	1.015	0	0.51	-0.01	0.13	-0.08	0.35	0.16	-0.02	-5.61E-02	0.7300
E7E1Z0	Rho GDP dissociation inhibitor alpha OS=Sus scrofa GN=ARHGDI4 PE=2 SV=1 - [E7E1Z0_PIG]	0.938	0.921	0.934	1.034	0.94	0.978	0.894	0	0.09	0.12	0.10	-0.05	0.09	0.03	0.16	-1.87E-02	0.7299
F1RZ28	Uncharacterized protein OS=Sus scrofa GN=RPS10 PE=4 SV=1 - [F1RZ28_PIG]	1.136	1.02	1.074	1.189	1.103	0.937	1.101	0	-0.18	-0.03	-0.10	-0.25	-0.14	0.09	-0.14	-3.01E-02	0.7299
F1SR85	Uncharacterized protein OS=Sus scrofa GN=ABCB6 PE=3 SV=1 - [F1SR85_PIG]	1.04	1.017	1.169	0.937	1.264	1.055	1.097	0	-0.06	-0.02	-0.23	0.09	-0.34	-0.08	-0.13	-3.72E-02	0.7294
F1RVET7	Integrin beta OS=Sus scrofa GN=ITGB1 PE=3 SV=1 - [F1RVET7_PIG]	0.941	0.819	0.91	0.985	0.766	0.836	0.995	0	0.09	0.29	0.14	0.02	0.38	0.26	0.01	4.00E-02	0.7290
P00761	Trypsin OS=Sus scrofa PE=1 SV=1 - [TRY_PIG]	1.087	1.095	1.056	0.983	1.126	1.09	0.981	0	-0.12	-0.13	-0.08	0.02	-0.17	-0.12	0.03	2.17E-02	0.7264
I3LLD8	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100626701 PE=4 SV=1 - [I3LLD8_PIG]	0.952	0.761	0.671	0.844	0.813	0.907	0.902	0	0.07	0.39	0.58	0.24	0.30	0.14	0.15	-5.19E-02	0.7254
J9JIL3	Sodium/potassium-transporting ATPase subunit beta-1 (Fragment) OS=Sus scrofa GN=ATP1B1 PE=3 SV=1 - [J9JIL3_PIG]	1.171	0.801	1.159	0.864	1.024	0.982	1.079	0	-0.23	0.32	-0.21	0.21	-0.03	0.03	-0.11	5.34E-02	0.7252
I3LDC7	Isocitrate dehydrogenase [NADP] (Fragment) OS=Sus scrofa GN=IDH1 PE=3 SV=1 - [I3LDC7_PIG]	0.923	0.924	1.029	1.023	0.87	0.935	0.987	0	0.12	0.11	-0.04	-0.03	-0.20	0.10	0.02	2.39E-02	0.7247
F1SKK1	Uncharacterized protein OS=Sus scrofa GN=SLC25A20 PE=3 SV=1 - [F1SKK1_PIG]	1.379	1.04	1.208	1.277	1.07	1.119	1.289	0	-0.46	-0.06	-0.27	-0.35	-0.10	-0.16	-0.37	-4.65E-02	0.7246
F1S2E2	Annexin OS=Sus scrofa GN=ANXA11 PE=2 SV=2 - [F1S2E2_PIG]	0.828	1.034	0.971	0.973	0.968	1.001	1.018	0	0.03	-0.05	0.04	0.04	0.05	0.00	-0.03	9.70E-03	0.7243
F1SIF2	Uncharacterized protein OS=Sus scrofa GN=NDUF53 PE=3 SV=1 - [F1SIF2_PIG]	0.784	0.769	0.865	0.935	0.841	0.823	0.714	0	0.35	0.38	0.21	0.10	0.25	0.28	0.49	4.36E-02	0.7241
F1S0P0	Cytochrome c oxidase subunit 6C OS=Sus scrofa GN=COX6C PE=4 SV=1 - [F1S0P0_PIG]	0.764	1.1	0.845	1.22	0.86	1.301	0.674	0	0.39	-0.14	0.24	-0.29	0.22	-0.38	0.57	-9.34E-02	0.7236
I3LEC2	Uncharacterized protein OS=Sus scrofa GN=PCBP1 PE=4 SV=1 - [I3LEC2_PIG]	1.018	0.963	0.985	0.97	0.994	0.996	0.986	0	-0.03	0.05	0.02	0.04	0.01	0.01	0.02	7.07E-03	0.7235
F1SA62	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100517634 PE=4 SV=1 - [F1SA62_PIG]	1.105	0.728	1.006	0.93	1.081	0.758	0.892	0	-0.14	0.46	-0.01	0.10	-0.11	0.40	0.16	6.29E-02	0.7216
F1SD94	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100739748 PE=3 SV=2 - [F1SD94_PIG]	1.121	1.365	1.496	0.954	1.059	1.099	1.646	0	-0.16	-0.45	-0.58	0.07	-0.08	-0.14	-0.72	8.12E-02	0.7216
I3LSI3	Uncharacterized protein OS=Sus scrofa GN=AK2 PE=3 SV=1 - [I3LSI3_PIG]	1.042	0.829	0.978	0.966	1.02	0.881	0.888	0	-0.06	0.27	0.03	0.05	-0.03	0.18	0.17	3.30E-02	0.7214
I3LAR5	Uncharacterized protein OS=Sus scrofa GN=NACA PE=4 SV=1 - [I3LAR5_PIG]	1.019	0.844	0.978	0.859	1.08	1.018	0.982	0	-0.03	0.24	0.03	0.22	-0.11	-0.03	0.03	-3.52E-02	0.7192
F1SBU7	Lon protease homolog, mitochondrial OS=Sus scrofa GN=LONP1 PE=3 SV=1 - [F1SBU7_PIG]	1.091	1.022	0.982	1.071	0.941	0.932	1.086	0	-0.13	-0.03	0.03	-0.10	0.09	0.10	-0.12	2.55E-02	0.7188
B8LFE3	Cardiomyopathy associated 1 OS=Sus scrofa GN=CMYA1 PE=4 SV=1 - [B8LFE3_PIG]	1.124	0.899	0.957	0.982	0.918	0.992	1.214	0	-0.17	0.15	0.06	0.03	0.12	0.01	-0.28	-4.17E-02	0.7185
F1SA60	Uncharacterized protein (Fragment) OS=Sus scrofa GN=SPTB PE=4 SV=1 - [F1SA60_PIG]	0.935	1.046	0.913	1.121	0.808	0.903	0.974	0	0.10	-0.06	0.13	-0.16	0.31	0.15	0.04	4.12E-02	0.7184
F1RVD4	Uncharacterized protein OS=Sus scrofa GN=SEC31A PE=4 SV=2 - [F1RVD4_PIG]	0.994	1.642	2.171	1.339	1.71	1.509	1.39	0	0.01	-0.72	-1.12	-0.42	-0.77	-0.59	-0.48	-1.10E-01	0.7175
P62831	60S ribosomal protein L23 OS=Sus scrofa GN=RPL23 PE=2 SV=1 - [RL23_PIG]	1.119	1.132	1.1	1.236	1.121	0.969	1.133	0	-0.16	-0.18	-0.14	-0.31	-0.16	0.05	-0.18	-3.16E-02	0.7175
F1S3U9	Uncharacterized protein OS=Sus scrofa GN=PRDX1 PE=4 SV=2 - [F1S3U9_PIG]	0.809	0.803	0.83	0.689	1.011	0.927	0.975	0	0.31	0.32	0.27	0.54	-0.02	0.11	0.04	-5.59E-02	0.7164
F1SGI7	Uncharacterized protein (Fragment) OS=Sus scrofa GN=KRT75 PE=3 SV=2 - [F1SGI7_PIG]	1.753	1.821	2.23	1.773	1.389	1.744	1.234	0	-0.81	-0.86	-1.16	-0.83	-0.47	-0.80	-0.30	1.06E-01	0.7160
F1S765	Uncharacterized protein OS=Sus scrofa GN=C1orf123 PE=4 SV=2 - [F1S765_PIG]	0.961	0.747	0.904	0.908	0.8	1.069	0.957	0	0.06	0.42	0.15	0.14	0.32	-0.10	0.06	-4.89E-02	0.7145
C4MRC5	ADP-ribosylation factor 2 OS=Sus scrofa GN=ARF3 PE=3 SV=1 - [C4MRC5_PIG]	0.945	1.178	0.956	1.052	1.158	1.099	0.										

F1SDC7	Uncharacterized protein OS=Sus scrofa GN=ALDH3A2 PE=3 SV=2 - [F1SDC7_PIG]	1.037	1.095	1.087	1.111	1.018	1.048	1.092	0	-0.05	-0.13	-0.12	-0.15	-0.03	-0.07	-0.13	-1.71E-02	0.6971
I3L5L1	Uncharacterized protein OS=Sus scrofa GN=RAP1B PE=4 SV=1 - [I3L5L1_PIG]	1.002	1.146	1.157	1.013	1.226	1.11	1.059	0	0.00	-0.20	-0.21	-0.02	-0.29	-0.15	-0.08	-3.40E-02	0.6964
F1RFQ7	GTP-binding nuclear protein Ran OS=Sus scrofa GN=KAN PE=2 SV=2 - [F1RFQ7_PIG]	0.958	1.112	1.138	1.062	1.131	1.05	1.036	0	0.06	-0.15	-0.19	-0.09	-0.18	-0.07	-0.05	-2.70E-02	0.6964
Q68S99	Fatty acid binding protein 3 (Fragment) OS=Sus scrofa GN=FABP3 PE=2 SV=1 - [Q68S99_PIG]	1.018	0.879	1.078	0.954	1.09	1.057	0.96	0	-0.03	0.19	-0.11	0.07	-0.12	-0.08	0.06	-3.24E-02	0.6961
F1RW31	Uncharacterized protein OS=Sus scrofa GN=NUDT9 PE=4 SV=2 - [F1RW31_PIG]	0.776	0.863	0.633	0.548	1.015	0.836	0.698	0	0.37	0.21	0.66	0.87	-0.02	0.26	0.52	9.63E-02	0.6959
I3LEQ7	Uncharacterized protein OS=Sus scrofa GN=STRAP PE=4 SV=1 - [I3LEQ7_PIG]	1.154	1.079	1.126	1.076	0.986	1.172	1.244	0	-0.21	-0.11	-0.17	-0.11	0.02	-0.23	-0.31	-3.54E-02	0.6948
F1SSX4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100620549 PE=4 SV=2 - [F1SSX4_PIG]	1.013	0.756	0.848	0.992	0.693	1.002	0.78	0	-0.02	0.40	0.24	0.01	0.53	0.00	0.36	6.84E-02	0.6947
F1SCF0	Alpha-1-antitrypsin OS=Sus scrofa GN=SERPINA1 PE=3 SV=1 - [F1SCF0_PIG]	1.012	0.948	0.964	0.908	1.127	0.883	0.931	0	-0.02	0.08	0.05	0.14	-0.17	0.18	0.10	3.42E-02	0.6942
F1RW89	Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondrial OS=Sus scrofa GN=ETFDH PE=4 SV=2 - [F1RW89_PIG]	1.027	0.881	0.882	0.926	0.985	0.889	0.912	0	-0.04	0.18	0.18	0.11	0.02	0.17	0.13	2.75E-02	0.6940
I3L5M4	Uncharacterized protein OS=Sus scrofa GN=CA14 PE=4 SV=1 - [I3L5M4_PIG]	1.054	1.079	0.802	1.413	0.872	0.978	0.935	0	-0.08	-0.11	0.32	-0.50	0.20	0.03	0.10	-7.62E-02	0.6929
I3LEU7	Uncharacterized protein OS=Sus scrofa GN=CIT PE=4 SV=1 - [I3LEU7_PIG]	0.818	0.58	1.334	0.699	0.797	1.717	1.041	0	0.29	0.79	-0.42	0.52	0.33	-0.78	-0.06	-1.63E-01	0.6837
F1RQM4	Malic enzyme OS=Sus scrofa GN=ME1 PE=3 SV=2 - [F1RQM4_PIG]	0.98	0.844	0.852	1.205	0.758	1.005	0.921	0	0.03	0.24	0.23	-0.27	0.40	-0.01	0.12	-6.57E-02	0.6833
I3L6E7	Uncharacterized protein OS=Sus scrofa GN=PHOSPHO1 PE=4 SV=1 - [I3L6E7_PIG]	1.221	1.053	0.967	1.213	0.869	1.101	0.911	0	-0.29	-0.07	0.05	-0.28	0.20	-0.14	0.13	5.85E-02	0.6814
A8YQT9	Ribosomal protein L7 OS=Sus scrofa PE=2 SV=1 - [A8YQT9_PIG]	0.985	1.292	0.966	1.369	0.898	1.126	1.075	0	0.02	-0.37	0.05	-0.45	0.16	-0.17	-0.10	-6.89E-02	0.6807
F1S3J0	Uncharacterized protein OS=Sus scrofa GN=ACOT6 PE=4 SV=2 - [F1S3J0_PIG]	1.081	0.953	0.835	0.959	0.883	0.847	1.057	0	-0.11	0.07	0.26	0.06	0.18	0.24	-0.08	4.56E-02	0.6804
F1S8G6	Uncharacterized protein OS=Sus scrofa GN=SGTA PE=4 SV=1 - [F1S8G6_PIG]	1.058	0.558	0.928	1.021	0.523	0.922	0.776	0	-0.08	0.84	0.11	-0.03	0.94	0.12	0.37	1.30E-01	0.6798
P01965	Hemoglobin subunit alpha OS=Sus scrofa GN=HBA PE=1 SV=1 - [HBA_PIG]	0.752	1.012	1.176	1.084	1.012	1.26	0.401	0	0.41	-0.02	-0.23	-0.12	-0.02	-0.33	1.32	1.73E-01	0.6792
I3VKE6	Ceruloplasmin OS=Sus scrofa PE=2 SV=1 - [I3VKE6_PIG]	0.886	1.021	0.977	0.937	1.132	1.052	0.885	0	0.17	-0.03	0.03	0.09	-0.18	-0.07	0.18	-4.00E-02	0.6790
I3LH87	Uncharacterized protein (Fragment) OS=Sus scrofa GN=COQ9 PE=4 SV=1 - [I3LH87_PIG]	1.102	1.242	1.116	1.1	1.094	1.061	1.105	0	-0.14	-0.31	-0.16	-0.14	-0.13	-0.09	-0.14	2.86E-02	0.6765
F1SAK6	Uncharacterized protein OS=Sus scrofa GN=ATP5L PE=4 SV=1 - [F1SAK6_PIG]	1.07	1.221	1.089	1.331	1.032	1.123	1.046	0	-0.10	-0.29	-0.12	-0.41	-0.05	-0.17	-0.06	-4.54E-02	0.6761
Q9MZ15	Voltage-dependent anion-selective channel protein 2 OS=Sus scrofa GN=VDAC2 PE=2 SV=1 - [VDAC2_PIG]	1.089	0.931	0.89	0.853	0.967	0.992	0.997	0	-0.12	0.10	0.17	0.23	0.05	0.01	0.00	3.64E-02	0.6756
Q0MVN8	Quinone oxidoreductase OS=Sus scrofa GN=CRYZ PE=2 SV=1 - [QOR_PIG]	0.813	0.975	1.169	0.922	1.15	0.958	1.069	0	0.30	0.04	-0.23	0.12	-0.20	0.06	-0.10	-5.72E-02	0.6750
F1RJ93	Uncharacterized protein (Fragment) OS=Sus scrofa GN=TAGLN2 PE=4 SV=1 - [F1RJ93_PIG]	1.045	1.072	0.764	0.993	1.004	1.097	0.912	0	-0.06	-0.10	0.39	0.01	-0.01	-0.13	0.13	-5.52E-02	0.6745
F1SU00	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [F1SU00_PIG]	1.06	0.847	0.736	0.987	0.768	0.838	0.875	0	-0.08	0.24	0.44	0.02	0.38	0.25	0.19	6.24E-02	0.6737
F1SFP9	Uncharacterized protein OS=Sus scrofa GN=LMOD3 PE=4 SV=2 - [F1SFP9_PIG]	1.064	1.178	1.211	1.217	1.046	1.204	1.097	0	-0.13	-0.24	-0.28	-0.06	-0.27	-0.19	-0.33	-3.69E-02	0.6731
F1SNT8	Uncharacterized protein OS=Sus scrofa GN=DCTN1 PE=4 SV=2 - [F1SNT8_PIG]	1.03	1.239	1.103	1.057	1.042	1.148	1.01	0	-0.04	-0.31	-0.14	-0.08	-0.06	-0.20	-0.01	3.51E-02	0.6730
F1SLI3	Microtubule-associated protein OS=Sus scrofa GN=MAP4 PE=4 SV=2 - [F1SLI3_PIG]	1.166	0.999	1.175	1.189	1.013	1.031	1.246	0	-0.22	0.00	-0.23	-0.25	-0.02	-0.04	-0.32	-4.42E-02	0.6714
F1SO61	Uncharacterized protein OS=Sus scrofa GN=LOC100511049 PE=4 SV=2 - [F1SO61_PIG]	0.945	1.029	1.012	0.955	0.921	0.913	1.119	0	0.08	-0.04	-0.02	0.07	0.12	0.13	-0.16	3.28E-02	0.6709
F1SDX6	Uncharacterized protein OS=Sus scrofa GN=TGM2 PE=3 SV=1 - [F1SDX6_PIG]	0.927	1.211	1.012	1.087	0.904	1.023	0.999	0	0.11	-0.28	-0.02	-0.12	0.15	-0.03	0.00	4.45E-02	0.6679
ASA759	Keratin 2A OS=Sus scrofa GN=KRT2A PE=2 SV=1 - [ASA759_PIG]	2.196	1.639	2.485	2.116	1.355	1.617	1.267	0	-1.13	-0.71	-1.31	-1.08	-0.44	-0.69	-0.34	1.52E-01	0.6671
I3L829	Uncharacterized protein (Fragment) OS=Sus scrofa GN=NPLOC4 PE=4 SV=1 - [I3L829_PIG]	1.014	1.384	0.955	1.029	0.931	1.069	1.106	0	-0.02	-0.47	0.07	-0.02	0.10	-0.10	-0.15	6.07E-02	0.6659
F1S3B4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100626669 PE=4 SV=2 - [F1S3B4_PIG]	1.064	0.988	1.144	1.115	1.091	0.847	1.026	0	-0.09	0.02	-0.19	-0.16	-0.13	0.24	-0.04	4.65E-02	0.6658
F1SJ30	Mannose-6-phosphate isomerase OS=Sus scrofa GN=MPI PE=3 SV=1 - [F1SJ30_PIG]	1.111	1.032	1.08	0.992	0.962	0.944	1.223	0	-0.15	-0.05	-0.11	0.01	0.06	0.08	-0.29	4.21E-02	0.6657
F1RW55	Uncharacterized protein OS=Sus scrofa PE=4 SV=2 - [F1RW55_PIG]	1.012	1.011	0.834	1.209	0.922	0.981	0.905	0	-0.02	-0.02	0.26	-0.27	0.12	0.03	0.14	-5.35E-02	0.6656
A5D9P1	Estradiol 17-beta-dehydrogenase 8 OS=Sus scrofa GN=HSD17B8 PE=2 SV=1 - [A5D9P1_PIG]	1.078	1.009	1.21	1.238	0.727	1.183	1.727	0	-0.11	-0.01	-0.28	-0.31	0.46	-0.24	-0.79	-1.21E-01	0.6648
D0G781	Acyl carrier protein OS=Sus scrofa GN=NDUFAB1 PE=2 SV=1 - [D0G781_PIG]	0.639	0.99	0.865	1.154	1.005	1.127	0.591	0	0.65	0.01	0.21	-0.21	-0.01	-0.17	0.76	-1.24E-01	0.6633
I3LDK3	Uncharacterized protein OS=Sus scrofa GN=CCDC175 PE=4 SV=1 - [I3LDK3_PIG]	1.015	1.216	1.242	1.024	1.244	1.012	1.035	0	-0.02	-0.28	-0.31	-0.03	-0.31	-0.02	-0.05	5.01E-02	0.6624
P79384	Propionyl-CoA carboxylase beta chain, mitochondrial OS=Sus scrofa GN=PCCB PE=2 SV=1 - [PCCB_PIG]	0.835	0.977	0.831	1.068	0.936	0.667	0.827	0	0.26	0.03	0.27	-0.09	0.10	0.58	0.27	7.45E-02	0.6602
F1S9C9	Proteinase subunit beta type OS=Sus scrofa GN=LOC100155139 PE=3 SV=1 - [F1S9C9_PIG]	0.99	1.417	1.128	1.386	1.105	1.297	0.985	0	0.01	-0.50	-0.17	-0.47	-0.14	-0.38	0.02	-7.66E-02	0.6572
P54612	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Sus scrofa GN=PPP2R1A PE=2 SV=2 - [2AAA_PIG]	0.788	0.871	0.77	0.925	0.789	0.905	0.912	0	0.34	0.20	0.38	0.11	0.34	0.14	0.13	-4.72E-02	0.6571
Q19PY1	Phosphorylase (Fragment) OS=Sus scrofa GN=PYGM PE=2 SV=1 - [Q19PY1_PIG]	0.806	1.127	1.293	0.97	1.2	0.895	0.898	0	0.31	-0.17	-0.37	0.04	-0.26	0.16	0.16	8.21E-02	0.6564
F1SHX0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=NEB PE=4 SV=2 - [F1SHX0_PIG]	0.943	1.166	1.14	1.119	1.025	0.98	0.998	0	0.08	-0.22	-0.19	-0.16	-0.04	0.03	0.00	4.00E-02	0.6548
D2D0E1	Muscle type carnitine palmitoyltransferase 1B OS=Sus scrofa GN=CPT1B PE=3 SV=1 - [D2D0E1_PIG]	0.97	0.943	0.926	0.868	1.059	0.985	1.021	0	0.04	0.08	0.11	0.20	-0.08	0.02	-0.03	-3.15E-02	0.6541
Q6IM70	Reticulon OS=Sus scrofa GN=RTN4 PE=2 SV=1 - [Q6IM70_PIG]	0.77	1.098	0.818	1.091	0.83	0.935	0.62	0	0.38	-0.13	0.29	-0.13	0.27	0.10	0.69	9.94E-02	0.6530
F1RIU9	Uncharacterized protein OS=Sus scrofa GN=CHCHD2 PE=4 SV=2 - [F1RIU9_PIG]	1.221	0.932	1.088	0.989	1.091	0.997	1.023	0	-0.29	0.10	-0.12	0.02	-0.13	0.00	-0.03	4.25E-02	0.6527
F1S1P1	Uncharacterized protein OS=Sus scrofa GN=AKR7A3 PE=4 SV=2 - [F1S1P1_PIG]	0.961	0.848	0.947	0.96	0.866	0.977	0.868	0	0.06	0.24	0.08	0.06	0.21	0.03	0.20	3.26E-02	0.6526
Q29320	Lactate dehydrogenase-B (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29320_PIG]	1.249	1.247	1.539	1.131	1.332	1.251	1.046	0	-0.32	-0.32	-0.62	-0.18	-0.41	-0.32	-0.06	7.05E-02	0.6519
F1S538	Uncharacterized protein OS=Sus scrofa GN=GCLM PE=4 SV=1 - [F1S538_PIG]	1.399	1.29	1.469	1.33	1.172	1.371	1.415	0	-0.42	-0.37	-0.55	-0.41	-0.23	-0.46	-0.50	-6.33E-02	0.6506
F1S3W0	Cytochrome b-c1 complex subunit 6 OS=Sus scrofa GN=UQCRRH PE=3 SV=1 - [F1S3W0_PIG]	0.989	0.93	0.788	0.986	0.945	1.13	0.813	0	0.02	0.10	0.34	0.02	0.08	-0.18	0.30	-6.00E-02	0.6504
F1SQN4	Uncharacterized protein OS=Sus scrofa GN=RP56KA3 PE=4 SV=3 - [F1SQN4_PIG]	0.979	0.985	1.12	1.007	0.839	0.904	1.203	0	0.03	0.02	-0.16	-0.01	0.25	0.15	-0.27	5.83E-02	0.6489
F1SP81	Uncharacterized protein (Fragment) OS=Sus scrofa GN=PLAA PE=4 SV=2 - [F1SP81_PIG]	1.166	1.166	1.257	0.948	1.356	0.971	1.139	0	-0.22	-0.22	-0.33	0.08	-0.44	0.04	-0.19	6.64E-02	0.6481
P10668	Cofilin-1 OS=Sus scrofa GN=CFL1 PE=1 SV=3 - [COF1_PIG]	0.744	0.72	0.706	0.837	0.786	0.838	0.805	0	0.43	0.47	0.50	0.26	0.35	0.25	0.31	-5.77E-02	0.6479
P28491	Calreticulin OS=Sus scrofa GN=CALR PE=1 SV=3 - [CALR_PIG]	0.939	1.081	1.17	1.034	1.024	1.12	1.109	0	0.09	-0.11	-0.23	-0.05	-0.03	-0.16	-0.15	-3.68E-02	0.6477
A5GFS8	Vesicle-associated membrane protein-associated protein B OS=Sus scrofa GN=VAPB PE=3 SV=1 - [VAPB_PIG]	1.001	1.052	1.109	1.014	1.025	1.076	1.112	0	0.00	-0.07	-0.15	-0.02	-0.04	-0.11	-0.15	-2.27E-02	0.6476
I3L9X0	Coronin OS=Sus scrofa PE=3 SV=1 - [I3L9X0_PIG]	0.927	0.955	1.078	1.019	1.051	0.951	1.01	0	0.11	0.07	-0.11	-0.03	-0.07	0.07	-0.01	-2.71E-02	0.6473
Q99028	Catechol O-methyltransferase (Fragment) OS=Sus scrofa GN=COMT PE=1 SV=1 - [COMT_PIG]	0.822	0.814	0.817	0.922	0.733	0.891	0.787	0	0.28	0.30	0.29	0.12	0.45	0.17	0.35	5.15E-02	0.6443
F1SUE1	Uncharacterized protein OS=Sus scrofa PE=4 SV=2 - [F1SUE1_PIG]	1.156	0.567	0.876	1.187	0.731	0.944	1.005	0	-0.21	0.82	0.19	-0.25	0.45	0.08	-0.01	-1.30E-01	0.6414
F2ZSB1	Leukocyte elastase inhibitor OS=Sus scrofa GN=SERPINB1 PE=2 SV=1 - [F2ZSB1_PIG]	1.011	1.374	1.205	1.182	1.092	1.215	1.255	0	-0.02	-0.46	-0.27						

K7GSE5	Uncharacterized protein OS=Sus scrofa GN=NDUFB9 PE=4 SV=1 - [K7GSE5_PIG]	0.762	0.679	0.889	1.134	0.801	0.897	0.734	0	0.39	0.56	0.17	-0.18	0.32	0.16	0.45	-9.47E-02	0.6235
F1SQ89	Uncharacterized protein OS=Sus scrofa GN=CANDP PE=4 SV=1 - [F1SQ89_PIG]	0.944	1.093	1.152	1.111	0.982	1.018	0.954	0	0.08	-0.13	-0.20	-0.15	0.03	-0.03	0.07	4.15E-02	0.6228
M1FV56	Leucine-rich repeat interacting protein-1 OS=Sus scrofa PE=2 SV=1 - [M1FV56_PIG]	0.879	0.852	0.806	1.067	0.696	0.887	0.732	0	0.19	0.23	0.31	-0.09	0.52	0.17	0.45	8.10E-02	0.6209
FZ25S8	Uncharacterized protein OS=Sus scrofa GN=TUBA4A PE=3 SV=1 - [FZ25S8_PIG]	0.986	1.196	1.037	1.081	1.043	1.074	1.112	0	0.02	-0.26	-0.05	-0.11	-0.06	-0.10	-0.15	-3.47E-02	0.6202
F1SAE9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LAMB1 PE=4 SV=2 - [F1SAE9_PIG]	0.935	0.83	0.866	0.905	0.956	0.896	0.952	0	0.10	-0.27	0.21	0.14	0.06	0.16	0.07	-3.38E-02	0.6185
P02540	Desmin OS=Sus scrofa GN=DES PE=1 SV=4 - [DESM_PIG]	1.029	1.066	1.1	0.965	1.107	1.086	1.135	0	-0.04	-0.09	-0.14	0.05	-0.15	-0.12	-0.18	-3.15E-02	0.6178
F1SVD5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=SMYD1 PE=4 SV=2 - [F1SVD5_PIG]	0.78	0.879	0.854	0.871	0.844	0.923	0.983	0	0.36	0.19	0.23	0.20	0.24	0.12	0.02	-4.70E-02	0.6147
F1RPD4	Uncharacterized protein OS=Sus scrofa GN=LOC100620271 PE=4 SV=2 - [F1RPD4_PIG]	0.716	0.523	0.709	0.91	0.634	0.541	0.585	0	0.48	0.94	0.50	0.14	0.66	0.89	0.77	1.35E-01	0.6127
I3L197	Uncharacterized protein OS=Sus scrofa GN=CSRPI PE=4 SV=1 - [I3L197_PIG]	1.005	1.064	0.855	0.894	0.965	1.077	0.847	0	-0.01	-0.09	0.23	0.16	0.05	-0.11	0.24	5.41E-02	0.6121
F1RFH9	Uncharacterized protein OS=Sus scrofa GN=ATP2A1 PE=3 SV=1 - [F1RFH9_PIG]	0.99	1.343	1.306	1.116	1.212	1.045	1.008	0	0.01	-0.43	-0.39	-0.16	-0.28	-0.06	-0.01	7.13E-02	0.6108
Q1HL06	6-phosphofructokinase (Fragment) OS=Sus scrofa GN=PFMK PE=3 SV=1 - [Q1HL06_PIG]	0.887	0.976	1.1	1.016	1.047	1.04	0.956	0	0.17	0.04	-0.14	-0.02	-0.07	-0.06	0.06	-3.78E-02	0.6104
A0S0C8	ATP synthase subunit a OS=Sus scrofa GN=ATP6 PE=3 SV=1 - [A0S0C8_PIG]	1.422	2.056	1.532	1.825	1.505	1.64	1.402	0	-0.51	-1.04	-0.62	-0.87	-0.59	-0.71	-0.49	-1.24E-01	0.6078
I3LEV7	Uncharacterized protein OS=Sus scrofa GN=SH3BGR PE=4 SV=1 - [I3LEV7_PIG]	0.902	1.191	1.131	0.946	1.039	1.084	0.981	0	0.15	-0.25	-0.18	0.08	-0.06	-0.12	0.03	5.43E-02	0.6072
A9X3M3	CMYA4 OS=Sus scrofa PE=2 SV=1 - [A9X3M3_PIG]	0.968	1.023	1.057	0.987	1.143	1.018	0.992	0	0.05	-0.03	-0.08	0.02	-0.19	-0.03	0.01	-3.06E-02	0.6070
F1SL07	Uncharacterized protein OS=Sus scrofa GN=LOC100524622 PE=4 SV=2 - [F1SL07_PIG]	1.006	0.977	1.219	1.081	1.168	0.879	0.893	0	-0.01	0.03	-0.29	-0.11	-0.22	0.19	0.16	6.84E-02	0.6061
Q95KQ9	Cystatin B (Fragment) OS=Sus scrofa GN=ctsb PE=4 SV=1 - [Q95KQ9_PIG]	0.944	0.858	0.887	0.991	0.894	0.917	0.968	0	0.08	0.22	0.17	0.01	0.16	0.13	0.05	-3.26E-02	0.6049
F1SSH8	Uncharacterized protein OS=Sus scrofa GN=LOC100515636 PE=4 SV=2 - [F1SSH8_PIG]	1.067	1.047	0.987	1.017	0.991	1.003	1.041	0	-0.09	-0.07	0.02	-0.02	0.01	0.00	-0.06	1.68E-02	0.6032
I3LAT6	Uncharacterized protein OS=Sus scrofa GN=LOC100739149 PE=4 SV=1 - [I3LAT6_PIG]	0.665	0.763	0.626	1.029	0.629	1.008	0.697	0	0.59	0.39	0.68	-0.04	0.67	-0.01	0.52	-1.29E-01	0.6032
P00819	Acylphosphatase 2 OS=Sus scrofa GN=ACYP2 PE=1 SV=2 - [ACYP2_PIG]	1.129	1.084	1.152	0.948	1.161	1.051	1.086	0	-0.18	-0.12	-0.20	0.08	-0.22	-0.07	-0.12	4.16E-02	0.6030
F1S4X7	Uncharacterized protein OS=Sus scrofa GN=MYH7B PE=4 SV=2 - [F1S4X7_PIG]	1.088	1.101	0.992	1.064	1.003	1.084	1.109	0	-0.12	-0.14	0.01	-0.09	0.00	-0.12	-0.15	-2.76E-02	0.6020
P02189	Myoglobin OS=Sus scrofa GN=MB PE=1 SV=2 - [MYG_PIG]	0.926	0.937	0.986	1.042	0.889	1.015	0.782	0	0.11	0.09	0.02	-0.06	0.17	-0.02	0.35	5.46E-02	0.6020
F1S134	Uncharacterized protein OS=Sus scrofa GN=COX5A PE=4 SV=1 - [F1S134_PIG]	1.134	1.03	0.978	1.182	1.096	1.283	0.848	0	-0.18	-0.04	0.03	-0.24	-0.13	-0.36	0.24	-7.58E-02	0.6013
Q1AG08	Calscarin 1 OS=Sus scrofa PE=2 SV=1 - [Q1AG08_PIG]	0.997	0.886	1.031	0.963	0.91	1.04	1.138	0	0.00	0.17	-0.04	0.05	0.14	-0.06	-0.19	-4.69E-02	0.6011
FZ25V0	tRNA-splicing ligase RtcB homolog OS=Sus scrofa GN=RTCB PE=3 SV=1 - [FZ25V0_PIG]	1.057	1.012	1.09	1.076	1.029	0.994	0.998	0	-0.08	-0.02	-0.12	-0.11	-0.04	0.01	0.00	2.15E-02	0.6007
FZ25B6	Tropomyosin alpha-1 chain OS=Sus scrofa GN=TPM1 PE=3 SV=1 - [FZ25B6_PIG]	0.999	1	1.125	0.856	1.074	1.056	1.014	0	0.00	0.00	-0.17	0.22	-0.10	-0.08	-0.02	4.78E-02	0.5994
F1SS86	Uncharacterized protein OS=Sus scrofa GN=DHR57C PE=3 SV=2 - [F1SS86_PIG]	0.806	1.171	1.036	1.179	0.898	0.928	0.765	0	0.31	-0.23	-0.05	-0.24	0.16	0.11	0.39	9.49E-02	0.5986
F1SS57	Uncharacterized protein OS=Sus scrofa GN=AGL PE=4 SV=1 - [F1SS57_PIG]	1	1.066	1.186	1.052	1.124	0.976	0.974	0	0.00	-0.09	-0.25	-0.07	-0.17	0.04	0.04	4.24E-02	0.5984
F1RZR0	Uncharacterized protein OS=Sus scrofa GN=ITPR3 PE=4 SV=1 - [F1RZR0_PIG]	1.167	1.166	1.314	1.05	1.254	1.066	1.091	0	-0.22	-0.22	-0.39	-0.07	-0.33	-0.09	-0.13	5.59E-02	0.5959
F1SAC1	Uncharacterized protein OS=Sus scrofa GN=ECHS1 PE=3 SV=2 - [F1SAC1_PIG]	1.068	1.268	1.073	1.237	1.056	1.036	0.888	0	-0.09	-0.34	-0.10	-0.31	-0.08	-0.05	0.17	6.85E-02	0.5949
F1S269	Uncharacterized protein OS=Sus scrofa GN=RCSDI PE=4 SV=2 - [F1S269_PIG]	0.913	1.149	0.919	0.983	0.835	0.9	1.093	0	0.13	-0.20	0.12	0.02	0.26	0.15	-0.13	6.40E-02	0.5947
F1S430	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100525578 PE=4 SV=2 - [F1S430_PIG]	1.255	1.199	1.291	1.744	1.047	1.248	1.132	0	-0.33	-0.26	-0.37	-0.80	-0.07	-0.32	-0.18	-1.02E-01	0.5944
I3LJT1	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LJT1_PIG]	0.876	0.84	0.887	0.909	0.856	0.945	0.99	0	0.19	0.25	0.17	0.14	0.22	0.08	0.01	-3.94E-02	0.5938
F1RK10	Succinyl-CoA ligase [ADP-forming] subunit beta, mitochondrial OS=Sus scrofa GN=SUCLA2 PE=3 SV=2 - [F1RK10_PIG]	1.024	0.978	1.022	1.039	1.074	0.93	0.888	0	-0.03	0.03	-0.03	-0.06	-0.10	0.10	0.17	3.78E-02	0.5911
B0LY42	CD147 OS=Sus scrofa PE=2 SV=1 - [B0LY42_PIG]	1.044	1.045	0.987	1.04	1.06	1.003	1.017	0	-0.06	-0.06	0.02	-0.06	-0.08	0.00	-0.02	-1.56E-02	0.5910
F1SM78	Uncharacterized protein OS=Sus scrofa GN=MYL12B PE=4 SV=2 - [F1SM78_PIG]	0.961	1.103	0.89	1.039	0.9	1.081	1.075	0	0.06	-0.14	0.17	-0.06	0.15	-0.11	-0.10	-5.10E-02	0.5890
D6QST6	2,4-dienoyl-CoA reductase 1 OS=Sus scrofa GN=DECR1 PE=2 SV=1 - [D6QST6_PIG]	1.103	1.138	1.023	0.944	1.201	0.985	0.999	0	-0.14	-0.19	-0.03	0.08	-0.26	0.02	0.00	5.07E-02	0.5880
F1RN71	Uncharacterized protein OS=Sus scrofa GN=LOC100521221 PE=4 SV=2 - [F1RN71_PIG]	0.786	0.915	0.919	0.906	0.913	0.833	0.844	0	0.35	0.13	0.12	0.14	0.13	0.26	0.24	4.62E-02	0.5848
Q29573	Fructose-biphosphate aldolase (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29573_PIG]	0.847	0.83	0.772	0.964	0.73	0.983	0.967	0	0.24	0.27	0.37	0.05	0.45	0.02	0.05	-7.54E-02	0.5827
Q2EN76	Nucleoside diphosphate kinase B OS=Sus scrofa GN=NME2 PE=2 SV=1 - [NDKB_PIG]	1.107	1.157	1.168	1.132	1.066	1.063	1.066	0	-0.15	-0.21	-0.22	-0.18	-0.09	-0.09	-0.09	3.24E-02	0.5825
I3LLY8	Uncharacterized protein OS=Sus scrofa GN=KRT79 PE=3 SV=1 - [I3LLY8_PIG]	1.505	1.369	1.832	1.564	1.161	1.35	1.086	0	-0.59	-0.45	-0.87	-0.65	-0.22	-0.43	-0.12	1.26E-01	0.5823
D0G0B3	Acetyl-Coenzyme A acyltransferase 2 OS=Sus scrofa GN=ACAA2 PE=2 SV=1 - [D0G0B3_PIG]	1.112	1.007	0.937	0.896	1.042	0.968	1.034	0	-0.15	-0.01	0.09	0.16	-0.06	0.05	-0.05	4.18E-02	0.5821
I3LDM3	Uncharacterized protein (Fragment) OS=Sus scrofa GN=IP07 PE=4 SV=1 - [I3LDM3_PIG]	0.946	1.016	0.873	1.02	0.905	1.011	0.996	0	0.08	-0.02	0.20	-0.03	0.14	-0.02	0.01	-3.69E-02	0.5820
Q2YGT9	60S ribosomal protein L6 OS=Sus scrofa GN=RPL6 PE=2 SV=3 - [RL6_PIG]	0.916	1.418	1.069	1.365	1.168	1.059	1.066	0	0.13	-0.50	-0.10	-0.45	-0.22	-0.08	-0.09	-9.36E-02	0.5819
F1S4Y7	Adenosylhomocysteinase (Fragment) OS=Sus scrofa GN=AHCY PE=3 SV=1 - [F1S4Y7_PIG]	0.9	0.908	0.929	0.951	0.993	0.901	0.781	0	0.15	0.14	0.11	0.07	0.01	0.15	0.06	4.85E-02	0.5817
Q6WSP6	Heat shock 10kD protein OS=Sus scrofa PE=3 SV=1 - [Q6WSP6_PIG]	2.222	1.265	0.865	1.331	1.437	1.373	1.512	0	-1.15	-0.34	0.21	-0.41	-0.52	-0.46	-0.60	-1.17E-01	0.5793
F1S935	Uncharacterized protein OS=Sus scrofa GN=RP1.8A PE=4 SV=2 - [F1S935_PIG]	1.166	1.442	1.224	1.764	1.218	1.106	1.175	0	-0.22	-0.53	-0.29	-0.82	-0.28	-0.15	-0.23	-1.10E-01	0.5779
F1SAX3	Sodium/potassium-transporting ATPase subunit alpha-1 (Fragment) OS=Sus scrofa GN=ATP1A1 PE=3 SV=2 - [F1SAX3_PIG]	1.482	1.262	1.844	1.638	1.223	1.488	1.632	0	-0.57	-0.34	-0.88	-0.71	-0.29	-0.57	-0.17	-1.24E-01	0.5778
I3LV53	Uncharacterized protein OS=Sus scrofa GN=LOC100627810 PE=4 SV=1 - [I3LV53_PIG]	1.115	1.25	1.169	1.335	1.057	1.239	1.102	0	-0.16	-0.32	-0.23	-0.42	-0.08	-0.31	-0.14	-6.05E-02	0.5776
F1SSL6	Proteasome subunit alpha type OS=Sus scrofa GN=LOC100154408 PE=3 SV=1 - [F1SSL6_PIG]	0.899	0.973	0.951	0.949	1.151	0.897	0.974	0	0.15	0.04	0.07	0.08	-0.20	0.16	0.04	-4.95E-02	0.5767
F1RP44	Uncharacterized protein (Fragment) OS=Sus scrofa GN=IP05 PE=4 SV=1 - [F1RP44_PIG]	0.92	1.132	1.079	0.979	1.014	0.945	1.06	0	0.12	-0.18	-0.11	0.03	-0.02	0.08	-0.08	4.41E-02	0.5752
F1RSZ3	Uncharacterized protein OS=Sus scrofa GN=ACSL1 PE=4 SV=2 - [F1RSZ3_PIG]	1.159	1.025	1.083	1.082	1.05	1.092	1.131	0	-0.21	-0.04	-0.12	-0.11	-0.07	-0.13	-0.18	-3.13E-02	0.5705
I3LES8	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3LES8_PIG]	0.962	1.248	1.133	1.113	0.974	1.055	1.015	0	0.06	-0.32	-0.18	-0.15	0.04	-0.08	-0.02	5.72E-02	0.5699
F1SPR6	Uncharacterized protein OS=Sus scrofa GN=RNFI23 PE=4 SV=1 - [F1SPR6_PIG]	1.122	1.226	1.121	1.528	1.102	1.237	0.958	0	-0.17	-0.29	-0.16	-0.61	-0.14	-0.31	0.06	-9.30E-02	0.5697
F1SLF0	Uncharacterized protein OS=Sus scrofa GN=SPR PE=4 SV=2 - [F1SLF0_PIG]	0.832	0.835	0.835	0.959	0.823	0.746	0.835	0	0.27	0.26	0.26	0.06	0.28	0.42	0.26	5.97E-02	0.5696
Q000H8	Glutathione S-transferase mu 2 (Fragment) OS=Sus scrofa PE=3 SV=1 - [Q000H8_PIG]	0.727	0.813	0.711	0.818	0.828	0.922	0.822	0	0.46	0.30	0.49	0.29	0.27	0.12	0.28	-7.22E-02	0.5691
FZ25J8	Eukaryotic translation initiation factor 2 subunit 1 OS=Sus scrofa GN=EIF2S1 PE=4 SV=1 - [FZ25J8_PIG]	0.824	1.191	0.985	1.036	0.895	0.961	0.887	0	0.28	-0.25	0.02	-0.05	0.16	0.06	0.17	7.26E-02	0.5686
A8D737	T-cadherin OS=Sus scrofa PE=2 SV=1 - [A8D737_PIG]	1.322	1.079	1.193	1.104	1.04	1.093	1.16	0	-0.40	-0.11	-0.25	-0.14	-0.06	-0.13	-0.21	5.63E-02	0.5682
F1RQ31	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=2 - [F1RQ31_PIG]	1.034	0.906	1.255	1.076	1.135	1.007	1.144	0	-0.05	0.14	-0.33</						

F1RUK8	Rab GDP dissociation inhibitor beta OS= Sus scrofa GN=GD12 PE=4 SV=1 - [F1RUK8_PIG]	0.896	1.017	0.993	0.959	0.959	1.036	1.045	0	0.16	-0.02	0.01	0.06	0.06	-0.05	-0.06	-3.45E-02	0.5439
I3LQH7	Uncharacterized protein OS= Sus scrofa GN=BLVRB PE=4 SV=1 - [I3LQH7_PIG]	0.902	0.824	0.93	0.925	1.042	1.052	0.819	0	0.15	0.28	0.10	0.11	-0.06	-0.07	0.29	-6.62E-02	0.5433
F1SF15	Uncharacterized protein OS= Sus scrofa GN=HRG PE=4 SV=2 - [F1SF15_PIG]	1.044	1.116	0.862	1.343	1.367	0.898	0.865	0	-0.06	-0.16	0.21	-0.43	-0.45	0.16	0.21	-1.26E-01	0.5429
F1SK15	Inosine 5'-monophosphate dehydrogenase OS= Sus scrofa GN=IMPDH2 PE=3 SV=2 - [F1SK15_PIG]	0.867	1.148	1.041	1.028	0.955	1.151	1.102	0	0.21	-0.20	-0.06	-0.04	0.07	-0.20	-0.14	-6.63E-02	0.5423
Q75NG6	Tropoin T fast skeletal muscle type OS= Sus scrofa GN=tmt3 PE=2 SV=1 - [Q75NG6_PIG]	1.313	1.005	0.964	1.028	1.087	0.941	0.981	0	-0.39	-0.01	0.05	-0.05	-0.12	0.09	0.03	7.28E-02	0.5415
I3LG67	Uncharacterized protein OS= Sus scrofa GN=MARCKS PE=4 SV=1 - [I3LG67_PIG]	1.348	1.173	1.051	1.145	1.083	1.34	1.233	0	-0.43	-0.23	-0.07	-0.20	-0.12	-0.42	-0.30	-7.55E-02	0.5408
I3LKP0	Uncharacterized protein (Fragment) OS= Sus scrofa GN=SYNM PE=3 SV=1 - [I3LKP0_PIG]	0.999	1.073	1.098	1.069	1.013	1.063	1.106	0	0.00	-0.10	-0.13	-0.10	-0.02	-0.09	-0.15	-2.83E-02	0.5405
P82460	Thioredoxin OS= Sus scrofa GN=TXN PE=1 SV=3 - [THIO_PIG]	0.822	0.775	0.922	0.851	1.036	0.961	0.844	0	0.28	0.37	0.12	0.23	-0.05	0.06	0.24	-7.10E-02	0.5397
F1RCG9	Uncharacterized protein OS= Sus scrofa GN=PSMD13 PE=4 SV=1 - [F1RCG9_PIG]	0.889	0.907	1	0.984	0.881	0.941	0.889	0	0.17	0.14	0.00	0.02	0.18	0.09	0.17	3.82E-02	0.5386
P43368	Calpain-3 OS= Sus scrofa GN=CAPN3 PE=2 SV=2 - [CAN3_PIG]	0.931	1.101	1.081	1.151	0.965	1.114	1.034	0	0.10	-0.14	-0.11	-0.20	0.05	-0.16	-0.05	-5.19E-02	0.5384
F1SAQ4	Uncharacterized protein OS= Sus scrofa GN=SETD3 PE=4 SV=2 - [F1SAQ4_PIG]	1.155	0.927	1.126	1.017	1.066	0.967	0.995	0	-0.21	0.11	-0.17	-0.02	-0.09	0.05	0.01	5.22E-02	0.5383
I3LFR2	Uncharacterized protein OS= Sus scrofa GN=CUL5 PE=3 SV=1 - [I3LFR2_PIG]	0.892	0.996	0.805	0.982	0.886	0.955	1.003	0	0.16	0.01	0.31	0.03	0.17	0.07	0.00	-5.52E-02	0.5365
F1SED0	Uncharacterized protein (Fragment) OS= Sus scrofa GN=ACADSB PE=3 SV=2 - [F1SED0_PIG]	1.036	1.036	0.835	0.994	0.853	1.011	0.876	0	-0.05	-0.05	0.26	0.01	0.23	-0.02	0.19	6.38E-02	0.5359
F2Z5J1	Uncharacterized protein OS= Sus scrofa GN=PSMC1 PE=4 SV=1 - [F2Z5J1_PIG]	0.984	0.911	0.856	0.907	1.022	1.017	0.923	0	0.02	0.13	0.22	0.14	-0.03	-0.02	0.12	-4.53E-02	0.5356
B9W5V0	Elongation factor 1-alpha OS= Sus scrofa GN=EEF1A2 PE=3 SV=1 - [B9W5V0_PIG]	1.038	1.091	1.053	1.043	1.074	0.947	1.032	0	-0.05	-0.13	-0.07	-0.06	-0.10	0.08	-0.05	3.08E-02	0.5351
F1SIW8	Uncharacterized protein OS= Sus scrofa PE=4 SV=2 - [F1SIW8_PIG]	1.118	1.593	1.245	1.28	1.221	1.414	1.32	0	-0.16	-0.67	-0.32	-0.36	-0.29	-0.50	-0.40	-9.89E-02	0.5346
F1RQK8	Phosphorylase OS= Sus scrofa GN=PYGM PE=3 SV=2 - [F1RQK8_PIG]	0.846	1.058	1.101	0.959	1.039	0.946	0.874	0	0.24	-0.08	-0.14	0.06	-0.06	0.08	0.19	6.46E-02	0.5342
F1REW9	Uncharacterized protein OS= Sus scrofa GN=DENR PE=4 SV=2 - [F1REW9_PIG]	0.734	0.876	0.979	0.724	0.942	0.791	0.911	0	0.45	0.19	0.03	0.47	0.09	0.34	0.13	8.93E-02	0.5336
Q9TV69	Trans-1,2-dihydrobenzene-1,2-diol dehydrogenase OS= Sus scrofa GN=DHDH PE=1 SV=1 - [DHDH_PIG]	0.98	1.031	0.998	0.989	1.041	1.061	0.979	0	0.03	-0.04	0.00	0.02	-0.06	-0.09	0.03	-2.12E-02	0.5319
F1RVN1	Uncharacterized protein OS= Sus scrofa GN=NDUFV1 PE=4 SV=1 - [F1RVN1_PIG]	0.911	0.781	0.868	0.904	0.909	0.794	0.799	0	0.13	0.36	0.20	0.15	0.14	0.33	0.32	6.11E-02	0.5301
M3VH45	Annexin OS= Sus scrofa GN=ANXA6 PE=2 SV=1 - [M3VH45_PIG]	0.813	1.078	1.021	0.861	1.027	0.971	0.848	0	0.30	-0.11	-0.03	0.22	-0.04	0.04	0.24	7.44E-02	0.5300
F1SLZ5	Uncharacterized protein OS= Sus scrofa GN=COQ10A PE=4 SV=1 - [F1SLZ5_PIG]	1.058	0.974	1.189	1.073	1.011	0.923	1.055	0	-0.08	0.04	-0.25	-0.10	-0.02	0.12	-0.08	5.35E-02	0.5298
F1RFM8	Uncharacterized protein (Fragment) OS= Sus scrofa GN=DNAH10 PE=4 SV=2 - [F1RFM8_PIG]	0.998	1.298	1.335	0.954	1.202	1.015	1.149	0	0.00	-0.38	-0.42	0.07	-0.27	-0.02	-0.20	9.27E-02	0.5288
A6M930	Eukaryotic translation initiation factor 4A isoform 2 OS= Sus scrofa GN=EIF4A2 PE=2 SV=1 - [A6M930_PIG]	0.991	1.075	1.089	0.995	0.983	1.047	1.047	0	0.01	-0.10	-0.12	0.01	0.02	-0.07	-0.07	2.84E-02	0.5279
A0S0D2	NADH-ubiquinone oxidoreductase chain 4 OS= Sus scrofa GN=ND4 PE=3 SV=1 - [A0S0D2_PIG]	1.597	1.956	1.644	1.695	3.029	1.338	1.411	0	-0.68	-0.97	-0.72	-0.76	-1.60	-0.42	-0.50	-2.29E-01	0.5257
Q29092	Endoplasmic OS= Sus scrofa GN=HSP90B1 PE=2 SV=3 - [ENPL_PIG]	1.074	1.073	0.932	1.08	0.977	1.082	1.052	0	-0.10	-0.10	0.10	-0.11	0.03	-0.11	-0.07	-4.03E-02	0.5252
Q7M364	Antithrombin III OS= Sus scrofa domestica PE=1 SV=1 - [Q7M364_PIG]	1.131	1.035	1.061	0.971	1.054	1.057	1.05	0	-0.18	-0.05	-0.09	0.04	-0.08	-0.08	-0.07	3.22E-02	0.5248
I3L7W3	Uncharacterized protein (Fragment) OS= Sus scrofa GN=PIEZ02 PE=4 SV=1 - [I3L7W3_PIG]	1.641	1.524	1.662	0.982	1.363	1.388	1.485	0	-0.71	-0.61	-0.73	0.03	-0.45	-0.47	-0.57	1.48E-01	0.5245
Q5PXT2	LIM and cysteine-rich domains protein 1 OS= Sus scrofa GN=LMCD1 PE=2 SV=1 - [LMCD1_PIG]	1.179	0.951	1.039	0.923	1.039	0.994	1.054	0	-0.24	0.07	-0.06	0.12	-0.06	0.01	-0.08	5.34E-02	0.5243
F1RQI6	Uncharacterized protein (Fragment) OS= Sus scrofa GN=FLIP1 PE=4 SV=2 - [F1RQI6_PIG]	1.448	1.193	1.303	1.029	1.186	1.132	1.276	0	-0.53	-0.25	-0.38	-0.04	-0.25	-0.18	-0.35	8.82E-02	0.5241
I3LKS6	Dihydropyridine reductase OS= Sus scrofa GN=QDR PE=4 SV=1 - [I3LKS6_PIG]	1.003	1.779	1.18	0.973	1.412	1.004	0.983	0	0.00	-0.83	-0.24	0.04	-0.98	-0.01	0.02	1.59E-01	0.5240
F1SFF8	Phosphorylase OS= Sus scrofa GN=PYGL PE=3 SV=2 - [F1SFF8_PIG]	1.04	0.843	1.058	0.884	1.212	1.276	0.91	0	-0.06	0.25	-0.08	0.18	-0.28	-0.35	0.14	-1.06E-01	0.5236
F1RU49	Uncharacterized protein OS= Sus scrofa GN=LOC100519531 PE=4 SV=1 - [F1RU49_PIG]	0.853	1.094	1.097	0.99	0.991	0.964	0.915	0	0.23	-0.13	-0.13	0.01	0.01	0.05	0.13	6.06E-02	0.5227
I3LUZ8	Uncharacterized protein OS= Sus scrofa GN=LOC100516656 PE=4 SV=1 - [I3LUZ8_PIG]	1.036	1.265	1.025	1.047	1.078	1.069	0.938	0	-0.05	-0.34	-0.04	-0.07	-0.11	-0.10	0.09	6.18E-02	0.5226
I3LK59	Enolase OS= Sus scrofa GN=ENO1 PE=3 SV=1 - [I3LK59_PIG]	0.919	0.923	0.861	0.934	0.899	0.971	0.992	0	0.12	0.12	0.22	0.10	0.15	0.04	0.01	-3.68E-02	0.5224
Q863I2	Serine/threonine-protein kinase OSR1 OS= Sus scrofa GN=OSR1 PE=2 SV=1 - [OSXR1_PIG]	0.854	0.779	0.657	0.989	0.724	0.883	0.935	0	0.23	0.36	0.61	0.02	0.47	0.18	0.10	-1.09E-01	0.5218
E7EBY5	MACRO domain containing protein 1 OS= Sus scrofa GN=MACROD1 PE=2 SV=1 - [E7EBY5_PIG]	0.844	0.895	1.013	0.977	0.846	0.909	0.88	0	0.24	0.16	-0.02	0.03	0.24	0.14	0.18	5.27E-02	0.5210
F1SQ01	Uncharacterized protein OS= Sus scrofa GN=PRDX4 PE=4 SV=1 - [F1SQ01_PIG]	0.901	1.033	1.131	0.983	1.084	0.967	0.845	0	0.15	-0.05	-0.18	0.02	-0.12	0.05	0.24	6.84E-02	0.5201
B3VK29	Cyclic AMP-dependent protein kinase A catalytic subunit (Fragment) OS= Sus scrofa PE=2 SV=1 - [B3VK29_PIG]	0.991	1.228	1.057	1.03	1.132	0.969	0.96	0	0.01	-0.30	-0.08	-0.04	-0.18	0.05	0.06	6.15E-02	0.5200
F1RUJ4	GMP reductase OS= Sus scrofa GN=GMFR PE=3 SV=1 - [F1RUJ4_PIG]	0.95	1.051	1.107	1.021	0.979	0.977	1.027	0	0.07	-0.17	-0.15	-0.03	0.03	0.03	-0.04	3.50E-02	0.5186
K9IVR7	WD repeat-containing protein 1 OS= Sus scrofa GN=WDRI PE=2 SV=1 - [K9IVR7_PIG]	0.96	0.934	0.972	0.932	1.026	0.939	1.077	0	0.06	0.10	0.04	0.10	-0.04	0.09	-0.11	-3.75E-02	0.5181
F1RJW7	Calsequestrin OS= Sus scrofa GN=CASQ1 PE=2 SV=2 - [F1RJW7_PIG]	1.053	1.288	1.395	1.042	1.236	1.064	1.083	0	-0.07	-0.37	-0.48	-0.06	-0.31	-0.09	-0.12	8.76E-02	0.5178
F1ISF4	Uncharacterized protein OS= Sus scrofa GN=NEK7 PE=4 SV=2 - [F1ISF4_PIG]	0.915	1.099	1.122	1.036	1.016	0.977	0.955	0	0.13	-0.14	-0.17	-0.05	-0.02	0.03	0.07	5.00E-02	0.5171
F1S146	Uncharacterized protein OS= Sus scrofa GN=ANK2 PE=4 SV=2 - [F1S146_PIG]	1.401	1.165	1.192	1.198	1.223	1.001	1.048	0	-0.49	-0.22	-0.25	-0.29	0.00	-0.07	8.50E-02	0.5131	
Q8MKF3	Beta-tropomyosin (Fragment) OS= Sus scrofa GN=TPM2 PE=2 SV=1 - [Q8MKF3_PIG]	0.984	0.931	1.085	0.785	1.015	0.999	1.024	0	0.02	0.10	-0.12	0.35	-0.02	0.00	-0.03	7.16E-02	0.5125
F1SU55	Uncharacterized protein OS= Sus scrofa GN=SYNP02L PE=4 SV=2 - [F1SU55_PIG]	1.112	0.979	1.043	1.04	0.998	1.072	1.14	0	-0.15	0.03	-0.06	-0.06	0.00	-0.10	-0.19	-3.99E-02	0.5112
F1RT87	Uncharacterized protein OS= Sus scrofa PE=4 SV=2 - [F1RT87_PIG]	0.891	0.96	0.75	1.009	0.756	1.078	1.041	0	0.17	0.06	0.42	-0.01	0.40	-0.11	-0.06	-1.04E-01	0.5112
F2Z547	cAMP-dependent protein kinase inhibitor alpha OS= Sus scrofa GN=PKIA PE=4 SV=1 - [F2Z547_PIG]	1.092	1.24	0.773	1.216	1.156	0.69	0.615	0	-0.13	-0.31	0.37	-0.28	-0.21	0.54	0.70	2.03E-01	0.5111
F1SCH1	Uncharacterized protein (Fragment) OS= Sus scrofa GN=NDUFB7 PE=4 SV=2 - [F1SCH1_PIG]	0.915	0.965	0.913	0.986	1.007	0.897	0.731	0	0.13	0.05	0.13	0.02	-0.01	0.16	0.45	7.71E-02	0.5106
P63246	Guanine nucleotide-binding protein subunit beta-2-like 1 OS= Sus scrofa GN=GNB2L1 PE=2 SV=3 - [GBLP_PIG]	1.066	1.043	0.994	0.919	1.004	1.026	1.059	0	-0.09	-0.06	0.01	0.12	-0.01	-0.04	-0.08	3.52E-02	0.5094
F1SGT2	Methylthioribulose-1-phosphate dehydratase OS= Sus scrofa GN=APIP PE=3 SV=1 - [F1SGT2_PIG]	1.045	0.879	0.956	1.126	1.125	0.828	1.053	0	-0.06	0.19	0.06	-0.17	-0.17	0.27	-0.07	-8.27E-02	0.5093
P04178	Superoxide dismutase [Cu-Zn] OS= Sus scrofa GN=SOD1 PE=1 SV=2 - [SODC_PIG]	1.124	0.981	1.159	0.968	0.994	1.008	1.132	0	-0.17	0.03	-0.21	0.05	0.01	-0.01	-0.18	5.48E-02	0.5091
Q9MZ16	Voltage-dependent anion-selective channel protein 1 OS= Sus scrofa GN=VDAC1 PE=2 SV=3 - [VDAC1_PIG]	1.067	0.95	0.879	0.916	0.969	0.95	0.937	0	-0.09	0.07	0.19	0.13	0.05	0.07	0.09	4.33E-02	0.5075
F1RZQ7	Uncharacterized protein OS= Sus scrofa GN=VBP1 PE=4 SV=2 - [F1RZQ7_PIG]	0.932	0.879	0.792	0.889	0.808	0.903	0.856	0	0.10	0.19	0.34	0.17	0.31	0.15	0.22	5.62E-02	0.5067
F1RII5	Uncharacterized protein OS= Sus scrofa GN=LOC100515788 PE=3 SV=2 - [F1RII5_PIG]	0.795	0.779	0.501	0.821	0.454	0.846	0.541	0	0.33	0.36	1.00	0.28	1.14	0.24	0.89	2.16E-01	0.5059
F1RW98	Uncharacterized protein OS= Sus scrofa GN=LOC100621992 PE=4 SV=2 - [F1RW98_PIG]	1.063	0.971	0.923	0.994	0.996	1.022	1.027	0	-0.09	0.04	0.12	0.01	-0.03	-0.04	-3.13E-02	0.5058	
F1SF28	Zona pellucida-binding protein 1 OS= Sus scrofa GN=ZPB PE=4 SV=2 - [F1SF28_PIG]	0.966	1.227	1.34	1.021	1.208	0.955	1.035	0	-0.05	-0.30	-0.42	-0.03	-0.27	0.07	-0.05	9.54E-02	0.5057
F1RWW4	Uncharacterized protein OS= Sus scrofa GN=PDLIM5 PE=4 SV=2 - [F1RWW4_PIG]	1.181	1.085	1.141	1.033	1.062												



I3LDY1	Uncharacterized protein (Fragment) OS=Sus scrofa GN=EML1 PE=4 SV=1 - [I3LDY1_PIG]	0.933	0.967	1.289	0.993	1.022	1.299	1.179	0	0.10	0.05	-0.37	0.01	-0.03	-0.38	-0.24	-1.05E-01	0.4816
EOX6R2	Glycogen synthase 1 OS=Sus scrofa GN=GYS1 PE=2 SV=1 - [EOX6R2_PIG]	1.083	1.185	1.28	1.036	1.224	1.04	1.012	0	-0.12	-0.24	-0.36	-0.05	-0.29	-0.06	-0.02	7.49E-02	0.4815
Q6S4N2	Heat shock 70 kDa protein 1B OS=Sus scrofa GN=HSPA1B PE=2 SV=1 - [HS71B_PIG]	0.98	0.872	0.963	0.863	1.087	0.966	1.106	0	0.03	0.20	0.05	0.21	-0.12	0.05	-0.15	-7.11E-02	0.4788
F1RFK3	Uncharacterized protein (Fragment) OS=Sus scrofa GN=TRRAP PE=4 SV=2 - [F1RFK3_PIG]	1.04	1.081	1.28	1.021	1.059	1.075	1.047	0	-0.06	-0.11	-0.36	-0.03	-0.08	-0.10	-0.07	6.05E-02	0.4782
F1SHQ8	Uncharacterized protein OS=Sus scrofa GN=LOC100518253 PE=3 SV=1 - [F1SHQ8_PIG]	1.132	1.05	1.008	0.997	1.044	1.081	0.938	0	-0.18	-0.07	-0.01	0.00	-0.06	-0.11	0.09	4.57E-02	0.4767
F1SSE7	Uncharacterized protein (Fragment) OS=Sus scrofa GN=COL15A1 PE=4 SV=2 - [F1SSE7_PIG]	0.955	0.841	0.834	0.96	0.904	0.908	0.992	0	0.07	0.25	0.26	0.06	0.15	0.14	0.01	-5.57E-02	0.4761
F1SIS9	Uncharacterized protein OS=Sus scrofa GN=NDUFA10 PE=4 SV=1 - [F1SIS9_PIG]	0.94	0.825	0.984	0.918	1.035	0.803	0.791	0	0.09	0.28	0.02	0.12	-0.05	0.32	0.34	8.46E-02	0.4736
I3LEJ8	Uncharacterized protein OS=Sus scrofa GN=LOC100516293 PE=4 SV=1 - [I3LEJ8_PIG]	1.144	0.951	0.999	0.845	1.034	0.985	1.042	0	-0.19	0.07	0.00	0.24	-0.05	0.02	-0.06	6.93E-02	0.4731
Q29568	Phosphopyruvate hydratase (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29568_PIG]	0.889	0.924	0.934	0.928	0.926	1.039	0.961	0	0.17	0.11	0.10	0.11	0.11	-0.06	0.06	-4.03E-02	0.4712
K7GL43	Uncharacterized protein (Fragment) OS=Sus scrofa GN=THY1 PE=4 SV=1 - [K7GL43_PIG]	1.191	1.181	1.25	1.355	1.298	0.986	1.337	0	-0.25	-0.24	-0.32	-0.44	-0.38	0.02	-0.42	-9.98E-02	0.4698
F1RPG1	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [F1RPG1_PIG]	1.026	1.313	1.391	1.037	1.231	1.007	1.086	0	-0.04	-0.39	-0.48	-0.05	-0.30	-0.01	-0.12	1.06E-01	0.4687
F1RKG0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYL10 PE=4 SV=2 - [F1RKG0_PIG]	0.853	0.677	0.651	0.799	0.788	0.867	0.975	0	0.23	0.56	0.67	0.32	0.34	0.21	0.04	-1.25E-01	0.4680
F1SFR6	Succinyl-CoA ligase [GDP-forming] subunit beta, mitochondrial OS=Sus scrofa GN=SUCLG2 PE=3 SV=2 - [F1SFR6_PIG]	1.033	0.979	0.985	0.889	0.739	1.166	0.967	0	-0.05	0.03	0.02	0.17	0.44	-0.22	0.05	1.07E-01	0.4672
F1SLU6	Ubiquitin carboxyl-terminal hydrolase OS=Sus scrofa GN=USP5 PE=3 SV=2 - [F1SLU6_PIG]	0.887	0.877	0.891	0.967	0.904	0.905	0.986	0	0.17	0.19	0.17	0.05	0.15	0.14	0.02	-4.26E-02	0.4671
Q6IM77	Reticulon OS=Sus scrofa GN=RTN2 PE=2 SV=1 - [Q6IM77_PIG]	0.839	1.223	1.21	1.184	1.01	0.941	0.721	0	0.25	-0.29	-0.28	-0.24	-0.01	0.09	0.47	1.53E-01	0.4662
Q29024	Spliceosome RNA helicase DDX39B OS=Sus scrofa GN=DDX39B PE=3 SV=2 - [DX39B_PIG]	0.946	1.124	0.752	0.882	0.973	1.332	1.023	0	0.08	-0.17	0.41	0.18	0.04	-0.41	-0.03	-1.37E-01	0.4656
P56729	Sucrase-isomaltase, intestinal (Fragments) OS=Sus scrofa GN=SI PE=1 SV=2 - [SUI5_PIG]	0.974	1.283	1.323	1.003	1.191	0.967	1.072	0	0.04	-0.36	-0.40	0.00	-0.25	0.05	-0.10	1.04E-01	0.4652
F1RIA7	Uncharacterized protein OS=Sus scrofa GN=TMEM109 PE=4 SV=1 - [F1RIA7_PIG]	1.021	0.856	1.037	1.19	0.968	1.16	0.883	0	-0.03	0.22	-0.05	-0.25	0.05	-0.21	0.18	-9.51E-02	0.4643
F1SZ24	Uncharacterized protein OS=Sus scrofa GN=LAMA2 PE=4 SV=2 - [F1SZ24_PIG]	0.904	0.917	0.822	0.941	0.889	0.94	1.005	0	0.15	0.13	0.28	0.09	0.17	0.09	-0.01	-5.35E-02	0.4634
F1SCB0	Uncharacterized protein OS=Sus scrofa GN=FXR1 PE=4 SV=2 - [F1SCB0_PIG]	1.126	1.027	1.021	1.126	1.02	0.932	0.938	0	-0.17	-0.04	-0.03	-0.17	-0.03	0.10	0.09	5.84E-02	0.4616
F1S192	Uncharacterized protein OS=Sus scrofa GN=PFND2 PE=4 SV=1 - [F1S192_PIG]	1.137	1.072	0.932	0.996	0.951	0.93	1.085	0	-0.19	-0.10	0.10	0.01	0.07	0.10	-0.12	6.23E-02	0.4614
F1RSR4	Uncharacterized protein OS=Sus scrofa GN=RFC3 PE=4 SV=1 - [F1RSR4_PIG]	1.205	1.078	1.259	0.878	1.083	1.297	0.971	0	-0.27	-0.11	-0.33	0.19	-0.12	-0.38	0.04	1.12E-01	0.4594
F1RP45	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CISD1 PE=4 SV=1 - [F1RP45_PIG]	1.033	1.032	1.026	1.066	1.012	1.011	0.874	0	-0.05	-0.05	-0.04	-0.09	-0.02	-0.02	0.19	4.96E-02	0.4585
F1RFA5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=PGP PE=4 SV=1 - [F1RFA5_PIG]	0.99	0.912	0.851	0.996	0.863	1.019	1.055	0	0.01	0.13	0.23	0.01	0.21	-0.03	-0.08	-6.66E-02	0.4582
F1SNA8	Uncharacterized protein OS=Sus scrofa GN=HSDL2 PE=4 SV=1 - [F1SNA8_PIG]	1.282	1.539	1.377	1.221	1.283	1.141	1.118	0	-0.36	-0.62	-0.46	-0.29	-0.36	-0.19	-0.16	1.11E-01	0.4574
Q32YV9	Proteasome 26S subunit non-ATPase 4 OS=Sus scrofa GN=PSMD4 PE=2 SV=1 - [Q32YV9_PIG]	1.035	0.96	1.069	0.856	1.103	0.942	0.994	0	-0.05	0.06	-0.10	0.22	-0.14	0.09	0.01	6.62E-02	0.4570
K7GLE1	Annexin OS=Sus scrofa GN=ANXA1 PE=3 SV=1 - [K7GLE1_PIG]	0.867	0.939	0.886	1.092	0.959	0.911	0.9	0	0.21	0.09	0.17	-0.39	0.06	0.13	0.15	-6.29E-02	0.4553
G9B792	Ankyrin repeat domain-containing protein 2 OS=Sus scrofa GN=ANKRD2 PE=2 SV=1 - [G9B792_PIG]	1.08	0.917	0.986	0.999	0.941	1.05	1.213	0	-0.11	0.13	0.02	0.00	0.09	-0.07	-0.28	-7.35E-02	0.4544
F1RLG5	Uncharacterized protein OS=Sus scrofa GN=REEP5 PE=4 SV=1 - [F1RLG5_PIG]	0.762	0.991	0.878	1.025	0.847	0.808	0.666	0	0.39	0.01	0.19	-0.04	0.24	0.31	0.59	1.26E-01	0.4529
F1S4J7	Uncharacterized protein OS=Sus scrofa GN=FBP2 PE=3 SV=2 - [F1S4J7_PIG]	0.89	1.048	1.07	1.345	0.998	1.015	0.968	0	0.17	-0.07	-0.10	-0.43	0.00	-0.02	0.05	-1.01E-01	0.4525
F1SC98	Uncharacterized protein (Fragment) OS=Sus scrofa GN=IDE PE=3 SV=2 - [F1SC98_PIG]	1.288	1.164	1.094	1.053	1.155	1.24	1.418	0	-0.37	-0.22	-0.13	-0.07	-0.21	-0.31	-0.50	-9.57E-02	0.4509
F1S587	Uncharacterized protein OS=Sus scrofa GN=AP1M2 PE=4 SV=2 - [F1S587_PIG]	0.886	1.043	0.833	1.159	0.944	0.993	0.904	0	0.17	-0.06	0.26	-0.21	0.08	0.01	0.15	-8.79E-02	0.4491
F1S1U2	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase 48 kDa subunit OS=Sus scrofa GN=DDOST PE=4 SV=1 - [F1S1U2_PIG]	1.176	1.308	0.31	0.917	1.715	0.862	1.162	0	-0.23	-0.39	0.16	0.13	-0.78	0.21	-0.22	-4.31E-01	0.4478
F1RIH9	Uncharacterized protein OS=Sus scrofa GN=GLOD4 PE=4 SV=2 - [F1RIH9_PIG]	0.991	1.069	1.069	1.008	0.979	1.047	1.011	0	0.01	-0.10	-0.10	-0.01	0.03	-0.07	-0.02	2.91E-02	0.4466
F1RI18	Uncharacterized protein OS=Sus scrofa GN=UQCRCF PE=4 SV=1 - [F1RI18_PIG]	0.932	1.111	1.045	1.338	1.283	1.145	0.812	0	0.10	-0.15	-0.06	-0.42	-0.36	-0.20	0.30	-1.40E-01	0.4459
Q6QA55	60S ribosomal protein L12 (Fragment) OS=Sus scrofa GN=RPL12 PE=2 SV=1 - [Q6QA55_PIG]	1.095	0.946	0.985	1.169	0.947	1.012	1.09	0	-0.13	0.08	0.02	-0.23	0.08	-0.02	-0.12	-6.48E-02	0.4458
F1S9V9	Uncharacterized protein (Fragment) OS=Sus scrofa GN=TMEM38A PE=4 SV=2 - [F1S9V9_PIG]	1.031	1.358	1.187	1.485	1.139	1.289	1.063	0	-0.04	-0.44	-0.25	-0.57	-0.19	-0.37	-0.09	-1.20E-01	0.4452
Q8WNV7	Dehydrogenase/reductase SDR family member 4 OS=Sus scrofa GN=DHRS4 PE=1 SV=2 - [DHRS4_PIG]	1.455	2.006	1.406	1.147	1.593	2.345	1.89	0	-0.54	-1.00	-0.49	-0.20	-0.67	-1.23	-0.92	-2.41E-01	0.4440
F1RIHW4	Uncharacterized protein OS=Sus scrofa GN=RRAS PE=4 SV=1 - [F1RIHW4_PIG]	0.891	1	0.666	0.994	0.841	1.034	0.973	0	0.17	0.00	0.59	0.01	0.25	-0.05	0.04	-1.26E-01	0.4422
F1RNS8	Uncharacterized protein OS=Sus scrofa GN=PDAP1 PE=4 SV=1 - [F1RNS8_PIG]	1.074	0.726	0.716	0.81	0.846	1.184	1.096	0	-0.10	0.46	0.48	0.30	0.24	-0.24	-0.13	-1.68E-01	0.4422
I3LKY1	Uncharacterized protein OS=Sus scrofa GN=LOC100737979 PE=4 SV=1 - [I3LKY1_PIG]	1.099	1.024	0.555	0.719	0.668	0.941	0.777	0	-0.14	-0.03	0.85	0.48	0.58	0.09	0.36	2.08E-01	0.4413
F2Z578	Uncharacterized protein OS=Sus scrofa GN=LOC100512420 PE=4 SV=1 - [F2Z578_PIG]	1.053	0.898	1.021	0.971	1.128	1	1.031	0	-0.07	0.16	-0.03	0.04	-0.17	0.00	-0.04	-5.65E-02	0.4403
Q95KL4	Selenoprotein W OS=Sus scrofa GN=SEPW1 PE=3 SV=4 - [SELW_PIG]	1.095	0.91	1.039	1.04	1.057	0.831	0.899	0	-0.13	0.14	-0.06	-0.06	-0.08	0.27	0.15	8.35E-02	0.4398
I3LQ99	Uncharacterized protein OS=Sus scrofa GN=MRPS36 PE=4 SV=1 - [I3LQ99_PIG]	1.275	0.924	1.077	1.012	1.064	0.909	1.007	0	-0.35	0.11	-0.11	-0.02	-0.09	0.14	-0.01	9.11E-02	0.4391
Q9T774	Glyceraldehyde-3-phosphate dehydrogenase (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q9T774_PIG]	0.977	1.055	0.719	1.25	0.851	0.97	1.064	0	0.03	-0.08	0.48	-0.32	0.23	0.04	-0.09	-1.42E-01	0.4384
I3LLH4	Uncharacterized protein OS=Sus scrofa GN=STAC3 PE=4 SV=1 - [I3LLH4_PIG]	0.946	0.849	0.949	0.926	0.837	0.872	0.965	0	0.08	0.24	0.08	0.11	0.26	0.20	0.05	5.62E-02	0.4356
F1SNC6	Uncharacterized protein (Fragment) OS=Sus scrofa GN=KIAA0368 PE=4 SV=2 - [F1SNC6_PIG]	0.769	1.157	0.934	0.888	0.919	0.848	0.899	0	0.38	-0.21	0.10	0.17	0.12	0.24	0.15	1.04E-01	0.4349
I3L953	Uncharacterized protein (Fragment) OS=Sus scrofa GN=OBSCN PE=4 SV=1 - [I3L953_PIG]	0.965	1.018	1.083	0.965	1.06	0.965	0.962	0	0.05	-0.03	-0.12	0.05	-0.08	0.05	0.06	4.10E-02	0.4334
F1S031	Uncharacterized protein OS=Sus scrofa GN=NDUF56 PE=4 SV=1 - [F1S031_PIG]	0.944	0.707	0.907	1.007	1.008	0.877	0.904	0	0.08	0.50	0.14	-0.01	-0.01	0.19	0.15	-1.03E-01	0.4321
Q29117	Titin (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29117_PIG]	0.944	1.092	1.043	0.984	1.022	0.978	0.979	0	0.08	-0.13	-0.06	0.02	-0.03	0.03	0.03	3.98E-02	0.4316
B5KJG2	Phosphoglycerate mutase 2 OS=Sus scrofa GN=PGAM2 PE=2 SV=1 - [B5KJG2_PIG]	0.821	1	1.07	0.992	0.952	0.894	0.818	0	0.28	0.00	-0.10	0.01	0.07	0.16	0.29	8.68E-02	0.4291
I3LP41	Malate dehydrogenase OS=Sus scrofa GN=MDH2 PE=3 SV=1 - [I3LP41_PIG]	1.247	0.888	0.925	0.899	0.998	0.892	0.973	0	-0.32	0.17	0.11	0.15	0.00	0.16	0.04	9.89E-02	0.4287
F1SCC7	Uncharacterized protein OS=Sus scrofa GN=LOC100156325 PE=3 SV=1 - [F1SCC7_PIG]	0.438	1.045	0.758	0.599	1.053	1.073	1.22	0	1.19	-0.06	0.40	0.74	-0.07	-0.10	-0.29	-3.13E-01	0.4280
Q9N251	Titin (Fragment) OS=Sus scrofa GN=TTN PE=4 SV=1 - [Q9N251_PIG]	0.921	0.89	1.033	0.869	0.915	0.961	0.96	0	0.12	0.17	-0.05	0.20	0.13	0.06	0.06	5.18E-02	0.4276
P01025	Complement C3 OS=Sus scrofa GN=C3 PE=1 SV=2 - [C03_PIG]	1.054	1.098	1.175	0.945	1.14	1.089	0.949	0	-0.08	-0.13	-0.23	0.08	-0.19	-0.12	0.08	7.21E-02	0.4273
K7GMN8	Pyruvate dehydrogenase E1 component subunit alpha, somatic form, mitochondrial OS=Sus scrofa GN=PDHA1 PE=4 SV=1 - [K7GMN8_PIG]	1.148	0.894	0.855	0.943	0.924	0.889	0.901	0	-0.20	0.16	0.23	0.08	0.11	0.17	0.15	8.26E-02	0.4260
F1S1X3	Uncharacterized protein OS=Sus scrofa GN=NARS PE=3 SV=2 - [F1S1X3_PIG]	1.883	1.315	1.306	1.263	1.166	1.157	1.262	0	0.18	-0.40	-0.39	-0.34	-0.22	-0.21	-0.34	-1.26E-01	0.4258
F1SB81	Plasminogen OS=Sus scrofa GN=PLG PE=3 SV=1 - [F1																	

Q56P28	PRA1 family protein 3 OS=Sus scrofa GN=ARL6IP5 PE=2 SV=1 - [PRAF3_PIG]	0.621	1.118	0.888	1.254	0.959	1.153	0.796	0	0.69	-0.16	0.17	-0.33	0.06	-0.21	0.33	-2.10E-01	0.4055
G9F6X8	Prolyl 4-hydroxylase beta polypeptide OS=Sus scrofa PE=2 SV=1 - [G9F6X8_PIG]	0.945	1.059	1.17	0.95	1.13	0.96	0.89	0	0.08	-0.08	-0.23	0.07	-0.18	0.06	0.17	8.81E-02	0.4054
AT7X81	COP9 signalosome complex subunit 6 OS=Sus scrofa GN=COP56 PE=2 SV=1 - [CSN6_PIG]	0.81	0.918	0.64	1.044	0.785	0.908	0.963	0	0.30	0.12	0.64	-0.06	0.35	0.14	0.05	-1.48E-01	0.4047
F1SGH5	Uncharacterized protein OS=Sus scrofa GN=PDHB PE=4 SV=1 - [F1SGH5_PIG]	1.102	0.95	0.946	0.965	1.042	0.936	0.878	0	-0.14	0.07	0.08	0.05	-0.06	0.10	0.19	6.53E-02	0.4024
P59083	14 kDa phosphohistidine phosphatase (Fragments) OS=Sus scrofa GN=PHPT1 PE=1 SV=1 - [PHP14_PIG]	0.921	0.803	0.78	0.789	0.74	0.871	0.877	0	0.12	0.32	0.36	0.34	0.43	0.20	0.19	9.28E-02	0.4021
G8IFA6	Mitochondrial NADH dehydrogenase Fe-S protein 4 OS=Sus scrofa PE=2 SV=1 - [G8IFA6_PIG]	1.038	0.866	1.027	0.995	1.019	0.879	0.809	0	-0.05	0.21	-0.04	0.01	-0.03	0.19	0.31	8.92E-02	0.4021
F1S6B4	Uncharacterized protein OS=Sus scrofa GN=PRELP PE=4 SV=2 - [F1S6B4_PIG]	1.01	0.604	0.821	0.996	0.91	1.004	0.866	0	-0.01	0.73	0.28	0.01	0.14	-0.01	0.21	-1.63E-01	0.4015
F1S6Q7	ATP synthase subunit delta, mitochondrial OS=Sus scrofa GN=ATPSD PE=3 SV=1 - [F1S6Q7_PIG]	1.155	0.847	0.836	0.816	0.854	0.95	0.913	0	-0.21	0.24	0.26	0.29	0.23	0.07	0.13	1.09E-01	0.4013
F1RTY6	Uncharacterized protein OS=Sus scrofa GN=PPID PE=4 SV=1 - [F1RTY6_PIG]	1	1.094	1.089	0.984	1.068	1.129	1.199	0	0.00	-0.13	-0.12	0.02	-0.09	-0.18	-0.26	-6.40E-02	0.4007
I3LPH5	Uncharacterized protein OS=Sus scrofa GN=G3BP2 PE=4 SV=1 - [I3LPH5_PIG]	0.853	0.725	0.782	1.062	0.707	0.972	1.004	0	0.23	0.46	0.35	-0.09	0.504	0.04	-0.01	-1.50E-01	0.3998
P03974	Transitional endoplasmic reticulum ATPase OS=Sus scrofa GN=VCP PE=1 SV=5 - [TERA_PIG]	1.024	1.115	1.178	0.917	1.103	1.042	1.047	0	-0.03	-0.16	-0.24	0.13	-0.14	-0.06	-0.07	7.14E-02	0.3998
Q6QGC0	PDZ and LIM domain protein 3 OS=Sus scrofa GN=PDLM3 PE=2 SV=1 - [PDLI3_PIG]	1.039	0.971	1.041	1.004	1.029	1.017	1.099	0	-0.06	0.04	-0.06	-0.01	-0.04	-0.02	-0.14	-3.42E-02	0.3994
A5GFR5	Novel protein similar to vertebrate chromosome 6 open reading frame 153 (C6orf153) OS=Sus scrofa GN=C7H6orf153 PE=3 SV=1 - [A5GFR5_PIG]	0.98	1.178	1.334	1.081	1.055	0.958	1.063	0	0.03	-0.24	-0.42	-0.11	-0.08	0.06	-0.09	1.02E-01	0.3994
M3VK21	Solute carrier family 25 (Aspartate/glutamate carrier), member 12 OS=Sus scrofa GN=SLC25A12 PE=2 SV=1 - [M3VK21_PIG]	1.038	0.948	0.95	0.972	1.011	0.895	0.94	0	-0.05	0.08	0.07	0.04	-0.02	0.16	0.09	4.43E-02	0.3991
Q6QAS1	Glutaredoxin (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q6QAS1_PIG]	1.304	1.108	1.2	1.112	1.006	1.013	1.189	0	-0.38	-0.15	-0.26	-0.15	-0.01	-0.02	-0.25	9.09E-02	0.3984
F1S3H4	Uncharacterized protein (Fragment) OS=Sus scrofa GN=COQ6 PE=4 SV=1 - [F1S3H4_PIG]	1.073	1.043	1.087	1.028	1.035	0.905	1.082	0	-0.10	-0.06	-0.12	-0.04	-0.05	0.14	-0.11	5.59E-02	0.3983
F1SRF1	Uncharacterized protein OS=Sus scrofa GN=KLHL40 PE=4 SV=1 - [F1SRF1_PIG]	1.016	1.049	1.01	1.056	0.962	0.966	1.005	0	-0.02	-0.07	-0.01	-0.08	0.06	0.05	-0.01	3.16E-02	0.3976
ASASW8	Complement component 4A OS=Sus scrofa GN=C4A PE=4 SV=1 - [ASASW8_PIG]	1.202	0.911	1.028	1.053	1.141	1.068	1.093	0	-0.27	0.13	-0.04	-0.07	-0.19	-0.09	-0.13	-7.93E-02	0.3964
I3L8S3	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CMBL PE=4 SV=1 - [I3L8S3_PIG]	1.023	1.36	1.678	1.081	1.129	1.131	1.073	0	-0.03	-0.44	-0.75	-0.11	-0.18	-0.18	-0.10	1.64E-01	0.3956
F1RL80	Branched-chain-amino-acid aminotransferase OS=Sus scrofa GN=BCAT2 PE=2 SV=2 - [F1RL80_PIG]	0.858	0.812	0.934	0.845	0.884	0.779	0.908	0	0.22	0.30	0.10	0.24	0.18	0.36	0.14	7.51E-02	0.3953
P00348	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Sus scrofa GN=HADH PE=1 SV=2 - [HCDH_PIG]	1.136	1.035	1.073	1.136	0.978	1.036	0.879	0	-0.18	-0.05	-0.10	-0.18	0.03	-0.05	0.19	7.96E-02	0.3948
Q2HYU1	Mitochondrial creatine kinase 2 OS=Sus scrofa GN=CKMT2 PE=2 SV=1 - [Q2HYU1_PIG]	1.117	0.954	0.914	0.941	1.037	0.906	0.901	0	-0.16	0.07	0.13	0.09	-0.05	0.14	0.15	7.25E-02	0.3883
F1S0C1	Uncharacterized protein OS=Sus scrofa GN=ADH5 PE=3 SV=1 - [F1S0C1_PIG]	1.057	1.036	1.198	0.928	1.057	1.045	1.06	0	-0.08	-0.05	-0.26	0.11	-0.08	-0.06	-0.08	6.80E-02	0.3883
Q6QRN9	ADP/ATP translocase 3 OS=Sus scrofa GN=SLC25A6 PE=2 SV=3 - [ADT3_PIG]	1.29	1.302	1.284	1.162	0.988	1.269	1.086	0	-0.37	-0.38	-0.36	-0.22	0.02	-0.34	-0.12	1.12E-01	0.3876
D0G7F6	Triosephosphate isomerase OS=Sus scrofa GN=TIM1 PE=2 SV=1 - [D0G7F6_PIG]	0.996	0.908	1.063	0.964	0.975	0.919	0.975	0	0.01	0.14	-0.09	0.05	0.04	0.12	0.04	4.77E-02	0.3864
F1RP07	Uncharacterized protein OS=Sus scrofa GN=LOC100521794 PE=4 SV=2 - [F1RP07_PIG]	0.972	1.226	1.263	1.013	1.126	0.989	1.013	0	0.04	-0.29	-0.34	-0.02	-0.17	0.02	-0.02	9.93E-02	0.3861
F1RNZ1	Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Sus scrofa GN=UQCRCF1 PE=3 SV=1 - [F1RNZ1_PIG]	1.25	1.06	1.079	1.003	1.094	1.063	1.009	0	-0.32	-0.08	-0.11	0.00	-0.13	-0.09	-0.01	7.02E-02	0.3846
I3LLS0	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3LLS0_PIG]	1.079	0.98	1.073	0.967	0.982	1.004	1.058	0	-0.11	0.03	-0.10	0.05	0.03	-0.01	-0.08	4.24E-02	0.3845
F2Z5U5	Uncharacterized protein OS=Sus scrofa GN=LOC100516604 PE=4 SV=1 - [F2Z5U5_PIG]	0.848	1.064	1.16	0.968	0.854	1.044	0.899	0	0.24	-0.09	-0.21	0.05	0.23	-0.06	0.15	1.08E-01	0.3822
F1S1V8	Uncharacterized protein OS=Sus scrofa GN=LOC100156983 PE=4 SV=2 - [F1S1V8_PIG]	1.025	1.085	1.184	1.007	1.125	0.958	1.001	0	-0.04	-0.12	-0.24	-0.01	-0.17	0.06	0.00	6.94E-02	0.3800
Q29096	Proteasome subunit C9-like protein (Fragment) OS=Sus scrofa PE=2 SV=2 - [Q29096_PIG]	0.912	1.069	1.208	0.953	1.084	0.977	0.887	0	0.13	-0.10	-0.27	0.07	-0.12	0.03	0.17	9.89E-02	0.3793
F1S5Z3	Uncharacterized protein OS=Sus scrofa GN=SARS PE=4 SV=2 - [F1S5Z3_PIG]	0.791	1.048	0.782	0.972	0.912	1.027	0.968	0	0.34	-0.07	0.35	0.04	0.13	-0.04	0.05	-1.11E-01	0.3770
I3LS60	Uncharacterized protein OS=Sus scrofa GN=CARKD PE=3 SV=1 - [I3LS60_PIG]	1.045	0.849	0.921	0.877	0.916	0.827	0.983	0	-0.06	0.24	0.12	0.19	0.13	0.27	0.02	8.08E-02	0.3760
P79303	UTP-glucose-1-phosphate uridylyltransferase OS=Sus scrofa GN=UGP2 PE=2 SV=3 - [UGPA_PIG]	1.023	0.904	1.017	0.986	0.946	0.848	0.994	0	-0.03	0.15	-0.02	0.02	0.08	0.24	0.01	6.46E-02	0.3738
C3RZA5	SET and MYND domain-containing 1 OS=Sus scrofa GN=SMYD1 PE=2 SV=1 - [C3RZA5_PIG]	0.923	0.943	0.979	1.021	0.97	0.956	0.983	0	0.12	0.08	0.03	-0.03	0.04	0.06	0.02	-3.18E-02	0.3730
F1SUS0	Uncharacterized protein OS=Sus scrofa GN=MFAP2 PE=4 SV=1 - [F1SUS0_PIG]	0.914	0.826	0.979	0.917	1.015	1.066	0.926	0	0.13	0.28	0.03	0.13	-0.02	-0.09	0.11	-7.85E-02	0.3720
F1RYA4	Uncharacterized protein OS=Sus scrofa GN=NAP1L4 PE=3 SV=2 - [F1RYA4_PIG]	0.896	0.972	0.791	1.011	0.706	0.816	0.834	0	0.16	0.04	0.34	-0.02	0.50	0.29	0.26	1.26E-01	0.3720
I3LGF7	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100525988 PE=4 SV=1 - [I3LGF7_PIG]	0.982	1.145	1.25	1.048	1.003	0.978	1.078	0	0.03	-0.20	-0.32	-0.07	0.00	0.03	-0.11	8.57E-02	0.3704
I3LVD5	Actin, cytoplasmic 1 OS=Sus scrofa GN=ACTB PE=3 SV=1 - [I3LVD5_PIG]	1.042	0.885	0.926	0.943	0.939	0.89	0.931	0	-0.06	0.18	0.11	0.08	0.09	0.17	0.10	5.47E-02	0.3703
I3LAF6	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LAF6_PIG]	0.891	0.836	0.745	0.866	0.902	0.943	0.976	0	0.17	0.26	0.42	0.21	0.15	0.08	0.04	-9.34E-02	0.3698
F1SS61	Uncharacterized protein OS=Sus scrofa GN=MYH3 PE=4 SV=1 - [F1SS61_PIG]	1.195	1.3	1.298	1.031	1.026	1.163	1.222	0	-0.26	-0.38	-0.38	-0.04	-0.22	-0.29	1.06E-01	0.3693	
K7GPB9	Uncharacterized protein OS=Sus scrofa GN=SEPT6 PE=4 SV=1 - [K7GPB9_PIG]	0.803	0.736	0.559	0.853	0.773	0.806	1.077	0	0.32	0.44	0.84	0.23	0.37	0.31	-0.11	-1.98E-01	0.3683
F1SM07	Uncharacterized protein OS=Sus scrofa GN=COP58 PE=4 SV=1 - [F1SM07_PIG]	0.735	0.774	0.656	1.164	0.668	0.955	0.886	0	0.44	0.37	0.61	-0.22	0.58	0.07	0.17	-2.04E-01	0.3676
F1RFX9	Uncharacterized protein OS=Sus scrofa GN=OMC PE=3 SV=1 - [F1RFX9_PIG]	1.033	1.07	1.004	1.261	1.072	1.027	0.992	0	-0.05	-0.10	-0.01	-0.01	-0.04	0.01	-7.79E-02	0.3668	
K9IVR9	Importin subunit beta-1 OS=Sus scrofa GN=KPNB1 PE=2 SV=1 - [K9IVR9_PIG]	0.904	1.117	0.851	0.978	1.012	1.046	1.059	0	0.15	-0.16	0.23	0.03	-0.02	-0.06	-0.08	-8.79E-02	0.3659
Q29194	Ribosomal protein S2 (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29194_PIG]	0.875	1.205	1.112	1.247	1.136	1.07	1.054	0	0.19	-0.27	-0.15	-0.32	-0.18	-0.10	-0.08	-1.12E-01	0.3657
F1RIN7	Uncharacterized protein OS=Sus scrofa GN=HRC PE=4 SV=2 - [F1RIN7_PIG]	0.79	1.002	1.053	1.12	0.933	1.188	0.949	0	0.34	0.00	-0.07	-0.16	0.10	-0.25	0.08	-1.25E-01	0.3645
F1RM62	Uncharacterized protein OS=Sus scrofa GN=HSPB6 PE=3 SV=1 - [F1RM62_PIG]	0.986	0.753	0.925	0.929	0.996	0.934	1.054	0	0.02	0.41	0.11	0.11	0.01	0.10	-0.08	-1.02E-01	0.3645
G3CKJ2	Glyceraldehyde-3-phosphate dehydrogenase (Fragment) OS=Sus scrofa GN=GAPDH PE=2 SV=1 - [G3CKJ2_PIG]	1.012	1.03	0.95	0.957	0.95	0.968	1.023	0	-0.02	-0.04	0.07	0.06	0.07	0.05	-0.03	3.43E-02	0.3625
F1SEV8	Uncharacterized protein OS=Sus scrofa GN=LOC100737048 PE=3 SV=2 - [F1SEV8_PIG]	1.114	1.093	1.038	1.009	1.098	1.306	1.104	0	-0.16	-0.13	-0.05	-0.01	-0.13	-0.39	-0.14	-8.45E-02	0.3621
Q95319	Beta-actin, cytoplasmic (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q95319_PIG]	0.995	1.138	1.154	1.032	1.054	1.03	0.986	0	0.01	-0.19	-0.21	-0.05	-0.08	-0.04	0.02	6.06E-02	0.3618
F2Z5M3	Uncharacterized protein OS=Sus scrofa GN=EIF5AL1 PE=4 SV=1 - [F2Z5M3_PIG]	0.94	1.089	1.115	0.98	1.022	0.976	0.994	0	0.09	-0.12	-0.16	0.03	-0.03	0.04	0.01	5.81E-02	0.3609
P41367	Medium-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Sus scrofa GN=ACADM PE=1 SV=3 - [ACADM_PIG]	1.072	0.729	0.739	0.991	1.184	0.89	0.892	0	-0.10	0.46	0.44	0.01	-0.24	0.17	0.16	-1.72E-01	0.3604
L8B0S2	IgG heavy chain OS=Sus scrofa GN=IGHG PE=2 SV=1 - [L8B0S2_PIG]	0.76	1.024	0.976	0.901	0.888	0.924	0.718	0	0.40	-0.03	0.04	0.15	0.17	0.11	0.48	1.29E-01	0.3604
F1S3H7	Uncharacterized protein OS=Sus scrofa GN=LOC100156930 PE=4 SV=2 - [F1S3H7_PIG]	1.22	0.82	1.042	0.883	1.266	0.716	0.68	0	-0.29	0.29	-0.06	0.18	-0.34	0.48	0.56	2.34E-01	0.3576
D2SQP3	Cysteine and glycine-rich protein 3 OS=Sus scrofa GN=CSR3 PE=2 SV=1 - [D2SQP3_PIG]	0.823	0.521	0.765	0.733	0.927	0.773	1.376	0	0.28	0.94	0.39	0.45	0.11	0.37	-0.46	-2.85E-01	0.3560
B8Y4S4	Sarcoglycan beta OS=Sus scrofa PE=2 SV=1 - [B8Y4S4_PIG]	1.424	0.93	1.075	0.972	0.923	0.984	1.083	0	-0.51	0.10	-0.10	0.04	0.12	0.02	-0.12	1.44E-01	0.3535
F1S571	Uncharacterized protein (Fragment) OS=Sus scrofa GN=COL11A1 PE=4 SV=2 - [F1S571_PIG]	1.111	0.753	1.12	1.225	1.1												

F1SIH8	Uncharacterized protein OS=Sus scrofa GN=LOC100516776 PE=4 SV=2 - [F1SIH8_PIG]	0.859	0.832	0.795	0.863	0.98	0.922	0.921	0	0.22	0.27	0.33	0.21	0.03	0.12	0.12	-8.45E-02	0.3366
F1SB60	Superoxide dismutase OS=Sus scrofa GN=SOD2 PE=3 SV=1 - [F1SB60_PIG]	1.226	1.048	1.128	1.013	1.105	0.954	1.077	0	-0.29	-0.07	-0.17	-0.02	-0.14	0.07	-0.11	8.34E-02	0.3365
F1SMB2	Dihydrolipoylysine-residue acetyltransferase OS=Sus scrofa GN=DLAT PE=2 SV=2 - [F1SMB2_PIG]	1.073	0.972	1.001	0.963	1.02	0.928	1.009	0	-0.10	0.04	0.00	0.05	-0.03	0.11	-0.01	4.57E-02	0.3364
I3LDC1	Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondrial OS=Sus scrofa GN=SDHB PE=4 SV=1 - [I3LDC1_PIG]	1.041	0.839	0.777	0.802	0.831	0.898	0.833	0	-0.06	0.25	0.36	0.32	0.27	0.16	0.26	1.11E-01	0.3362
P50578	Alcohol dehydrogenase [NADP(+)] OS=Sus scrofa GN=AKR1A1 PE=1 SV=2 - [AK1A1_PIG]	1.639	1.285	1.098	0.971	0.915	1.294	1.149	0	-0.71	-0.36	-0.13	0.04	0.13	-0.37	-0.20	2.02E-01	0.3361
F1SEQ7	Uncharacterized protein OS=Sus scrofa GN=FAM213A PE=4 SV=1 - [F1SEQ7_PIG]	1.145	0.997	0.942	0.975	1.178	1.018	1.234	0	-0.20	0.00	0.09	0.04	-0.24	-0.03	-0.30	-1.06E-01	0.3350
P79293	Myosin-7 OS=Sus scrofa GN=MYH7 PE=2 SV=2 - [MYH7_PIG]	1.06	1.318	1.295	0.954	1.123	1.061	1.135	0	-0.08	-0.40	-0.37	0.07	-0.17	-0.09	-0.18	1.22E-01	0.3336
P61288	Transitionally-controlled tumor protein OS=Sus scrofa GN=TPST1 PE=2 SV=1 - [TCPT_PIG]	1.071	0.977	1.01	1.098	1.16	1.095	0.936	0	-0.10	0.03	-0.01	-0.13	-0.21	-0.13	-0.10	-7.62E-02	0.3330
F1RIF8	6-phosphogluconate dehydrogenase, decarboxylating (Fragment) OS=Sus scrofa GN=PGD PE=3 SV=2 - [F1RIF8_PIG]	1.309	1.009	1.03	1.064	0.915	1.139	0.794	0	-0.39	-0.01	-0.04	-0.09	0.13	-0.19	0.33	1.57E-01	0.3321
F1RI39	Uncharacterized protein OS=Sus scrofa GN=LOC100517284 PE=4 SV=2 - [F1RI39_PIG]	1.056	0.927	0.952	0.976	1.073	1.011	1.02	0	-0.08	0.01	0.07	0.04	-0.10	-0.02	-0.03	-5.32E-02	0.3295
F1S564	Uncharacterized protein OS=Sus scrofa GN=RTCA PE=4 SV=2 - [F1S564_PIG]	0.422	0.613	0.874	0.73	0.513	0.597	0.356	0	1.24	0.71	0.19	0.45	0.96	0.74	1.49	3.77E-01	0.3293
I3LDU9	Uncharacterized protein OS=Sus scrofa GN=SNTB2 PE=4 SV=1 - [I3LDU9_PIG]	1.093	1.182	1.28	1.179	1.075	1.301	1.396	0	-0.13	-0.24	-0.36	-0.24	-0.10	-0.38	-0.48	-1.19E-01	0.3289
F1RLM1	Uncharacterized protein (Fragment) OS=Sus scrofa GN=COL4A1 PE=4 SV=2 - [F1RLM1_PIG]	0.978	0.825	0.976	0.969	0.981	0.964	1.064	0	0.03	0.28	0.04	0.05	0.03	0.05	-0.09	-7.70E-02	0.3282
F2Z5V1	Uncharacterized protein OS=Sus scrofa GN=TMA7 PE=4 SV=1 - [F2Z5V1_PIG]	1.27	1.238	1.418	1.234	0.99	1.143	1.106	0	-0.34	-0.31	-0.50	-0.30	0.01	-0.19	-0.15	1.32E-01	0.3277
F1RKY2	Uncharacterized protein OS=Sus scrofa GN=SERPIND1 PE=3 SV=1 - [F1RKY2_PIG]	0.776	0.852	0.845	0.816	1.197	0.913	0.945	0	0.37	0.23	0.24	0.29	-0.26	0.13	0.08	-1.48E-01	0.3273
I3LS51	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3LS51_PIG]	1.081	1.16	1.086	1.013	0.978	1.025	1.121	0	-0.11	-0.21	-0.12	-0.02	0.03	-0.04	-0.16	6.46E-02	0.3272
M3TYR4	Mitogen-activated protein kinase kinase 3 OS=Sus scrofa GN=MAP2K3 PE=2 SV=1 - [M3TYR4_PIG]	1.053	1.038	0.929	1.668	0.845	1.107	1.199	0	-0.07	-0.05	0.11	-0.74	0.24	-0.15	-0.26	-2.20E-01	0.3259
K9IW80	Nicotinamide nucleotide transhydrogenase OS=Sus scrofa GN=NNT PE=2 SV=1 - [K9IW80_PIG]	1.139	0.96	0.992	1.033	0.99	0.909	0.955	0	-0.19	0.06	0.01	-0.05	0.01	0.14	0.07	7.23E-02	0.3220
I3LUM2	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LUM2_PIG]	0.957	0.359	1.051	1.131	1.316	0.888	0.895	0	0.06	1.48	-0.07	-0.18	-0.40	0.17	0.16	-4.28E-01	0.3214
I3LJ87	Uncharacterized protein OS=Sus scrofa GN=RP52E PE=3 SV=1 - [I3LJ87_PIG]	0.976	1.033	1.166	1.389	1.161	1.035	1.008	0	0.04	-0.05	-0.22	-0.47	-0.22	-0.05	-0.01	-1.29E-01	0.3214
I3LC07	Uncharacterized protein OS=Sus scrofa GN=RAB18 PE=3 SV=1 - [I3LC07_PIG]	0.929	0.984	0.959	1.314	0.883	0.998	1.096	0	0.11	0.02	0.06	-0.39	0.18	0.00	-0.13	-1.33E-01	0.3200
F1RZA1	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100157017 PE=3 SV=2 - [F1RZA1_PIG]	0.664	0.873	0.754	0.9	1.056	0.773	0.96	0	0.59	0.20	0.41	0.15	-0.08	0.37	0.06	-1.73E-01	0.3200
F1RIX8	Uncharacterized protein OS=Sus scrofa GN=CBR4 PE=3 SV=2 - [F1RIX8_PIG]	1.503	1.102	1.087	1.564	1.156	1.539	1.121	0	-0.59	-0.14	-0.12	-0.65	-0.59	-0.62	-0.16	-1.98E-01	0.3196
F1RS49	Uncharacterized protein OS=Sus scrofa GN=ABCE1 PE=3 SV=2 - [F1RS49_PIG]	0.884	1.149	0.701	1.224	0.935	1.074	0.979	0	0.18	-0.20	0.51	-0.29	0.10	-0.10	0.03	-1.89E-01	0.3180
F1SHU0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=HIBADH PE=4 SV=2 - [F1SHU0_PIG]	1.131	0.92	0.877	0.801	1.028	0.868	0.915	0	-0.18	0.12	0.19	0.32	-0.04	0.20	0.13	1.20E-01	0.3176
F1RP34	NAD(P)H-hydrate epimerase OS=Sus scrofa GN=APOA1BP PE=3 SV=2 - [F1RP34_PIG]	1	1.025	0.995	1.012	0.986	1.066	1.037	0	0.00	-0.04	0.01	-0.02	0.02	-0.09	-0.05	-2.83E-02	0.3175
Q9TV36	Fibrillin-1 OS=Sus scrofa GN=FBN1 PE=2 SV=1 - [FBN1_PIG]	1.074	0.918	0.948	0.979	0.992	0.83	0.919	0	-0.10	0.12	0.08	0.03	0.01	0.27	0.12	8.39E-02	0.3167
I3LC10	Uncharacterized protein OS=Sus scrofa GN=LOC100525869 PE=4 SV=1 - [I3LC10_PIG]	1.023	0.841	0.929	0.958	0.905	0.853	0.863	0	-0.03	0.25	0.11	0.06	0.14	0.23	0.21	8.11E-02	0.3161
F1SGV7	Uncharacterized protein OS=Sus scrofa GN=TTG3 PE=4 SV=2 - [F1SGV7_PIG]	0.966	1.098	1.027	1.157	1.065	1.036	1.015	0	0.05	-0.13	-0.04	-0.21	-0.09	-0.05	-0.02	-6.26E-02	0.3145
M3VK13	Eukaryotic translation initiation factor 5 (v1) OS=Sus scrofa GN=EIF5 PE=2 SV=1 - [M3VK13_PIG]	1.119	1.194	1.358	1.213	0.981	1.091	0.974	0	-0.16	-0.26	-0.44	-0.28	0.03	-0.13	0.04	1.30E-01	0.3144
I3LRW8	Serine/arginine-rich-splicing factor 2 (Fragment) OS=Sus scrofa GN=SRSF2 PE=4 SV=1 - [I3LRW8_PIG]	0.992	0.941	1.147	1.104	0.941	1.118	1.308	0	0.01	0.09	-0.20	-0.14	0.09	-0.16	-0.39	-1.26E-01	0.3130
A7TX80	COP9 constitutive photomorphogenic-like subunit 5 OS=Sus scrofa GN=COPS5 PE=2 SV=1 - [A7TX80_PIG]	0.989	1.104	1.187	1.076	0.94	1.005	1.016	0	0.02	-0.14	-0.25	-0.11	0.09	-0.01	-0.02	8.19E-02	0.3117
F1SAU3	Uncharacterized protein (Fragment) OS=Sus scrofa GN=ATRN1L1 PE=4 SV=2 - [F1SAU3_PIG]	2.013	1.351	1.676	1.109	1.565	1.073	1.133	0	-1.01	-0.43	-0.75	-0.15	-0.65	-0.10	-0.18	2.78E-01	0.3112
F1S4D5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CCBL2 PE=4 SV=2 - [F1S4D5_PIG]	1.127	1.221	1.205	1.045	1.117	1.068	1.077	0	-0.17	-0.29	-0.27	-0.06	-0.16	-0.09	-0.11	7.61E-02	0.3110
I3LTV6	Cytochrome c oxidase subunit 6A, mitochondrial OS=Sus scrofa GN=COX6A2 PE=3 SV=1 - [I3LTV6_PIG]	1.294	1.138	1.048	1.132	0.988	1.066	0.958	0	-0.37	-0.19	-0.07	-0.18	0.02	-0.09	0.06	1.09E-01	0.3101
I3LKD2	Uncharacterized protein OS=Sus scrofa GN=FAM162A PE=4 SV=1 - [I3LKD2_PIG]	1.097	1.604	1.08	1.143	0.982	1.057	0.97	0	-0.13	-0.68	-0.11	-0.19	0.03	-0.08	0.04	1.81E-01	0.3078
I3LD43	Uncharacterized protein OS=Sus scrofa GN=LOC100627689 PE=4 SV=1 - [I3LD43_PIG]	1.174	0.953	1.131	0.839	1.232	0.81	0.937	0	-0.23	0.07	-0.18	0.25	-0.30	0.50	0.09	1.72E-01	0.3071
F1RUN2	Serum albumin OS=Sus scrofa GN=ALB PE=4 SV=1 - [F1RUN2_PIG]	0.98	1.025	0.832	0.77	1.059	0.861	0.937	0	0.03	-0.04	0.27	0.38	-0.08	0.22	0.29	1.35E-01	0.3058
F1RVL5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYOM2 PE=4 SV=2 - [F1RVL5_PIG]	0.962	1.133	1.164	0.982	1.094	0.938	0.975	0	0.06	-0.18	-0.22	0.03	-0.13	0.09	0.04	9.22E-02	0.3054
F1SSR6	Uncharacterized protein OS=Sus scrofa PE=4 SV=2 - [F1SSR6_PIG]	0.922	1.048	0.959	0.944	0.974	0.937	0.945	0	0.12	0.07	0.06	0.08	0.04	0.09	0.08	4.67E-02	0.3046
F1SIF7	Alpha-2-HS-glycoprotein (Fragment) OS=Sus scrofa GN=AHSG PE=4 SV=2 - [F1SIF7_PIG]	0.974	0.815	0.738	0.893	1.035	0.962	0.94	0	0.04	0.30	0.44	0.16	-0.05	0.06	0.09	-1.28E-01	0.3030
I3LB92	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYOM3 PE=4 SV=1 - [I3LB92_PIG]	1.064	1.053	0.938	0.968	0.889	0.947	1.052	0	-0.09	-0.07	0.09	0.05	0.17	0.08	-0.07	7.34E-02	0.3030
Q0QF01	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Sus scrofa GN=SDHA PE=1 SV=2 - [DHSA_PIG]	1.009	0.865	0.855	0.864	0.875	0.87	0.918	0	-0.01	0.21	0.23	0.21	0.19	0.20	0.12	7.64E-02	0.3029
P18648	Apolipoprotein A1 OS=Sus scrofa GN=APOA1 PE=1 SV=4 - [APOA1_PIG]	1.12	0.986	0.92	0.895	1.268	1.347	1.052	0	-0.16	0.02	0.12	0.16	-0.34	-0.43	-0.07	-1.66E-01	0.3009
F1SDN2	Uncharacterized protein OS=Sus scrofa GN=HADHB PE=3 SV=2 - [F1SDN2_PIG]	1.135	0.979	1.054	1.016	1.028	0.93	1.006	0	-0.18	0.03	-0.08	-0.02	-0.04	0.10	-0.01	6.53E-02	0.3008
Q52NJ1	Ras-related protein Rab-11A OS=Sus scrofa GN=RAB11A PE=2 SV=3 - [RB11A_PIG]	0.775	0.934	0.678	0.9	0.826	1.081	0.987	0	0.37	0.10	0.56	0.15	0.28	-0.11	0.02	-1.73E-01	0.3001
F2Z553	Uncharacterized protein OS=Sus scrofa GN=EIF1 PE=4 SV=1 - [F2Z553_PIG]	0.865	0.657	0.946	1.025	0.91	0.936	0.955	0	0.21	0.61	0.08	-0.04	0.14	0.10	0.07	-1.58E-01	0.2995
Q5G6V9	Cofilin-2 OS=Sus scrofa GN=CFL2 PE=2 SV=4 - [COF2_PIG]	0.918	0.851	0.919	0.815	0.905	0.903	0.893	0	0.12	0.23	0.12	0.30	0.14	0.15	0.16	6.79E-02	0.2991
I3LP92	Uncharacterized protein OS=Sus scrofa GN=RANGRF PE=4 SV=1 - [I3LP92_PIG]	1.063	1.013	0.675	1.206	0.889	1.142	1.046	0	-0.09	-0.02	0.57	-0.27	0.17	-0.19	-0.06	-2.04E-01	0.2990
F1RJ11	Uncharacterized protein OS=Sus scrofa GN=MLEC PE=4 SV=1 - [F1RJ11_PIG]	0.875	1.096	1.067	1.176	1.028	1.126	1.001	0	0.19	-0.13	-0.09	-0.23	-0.04	-0.17	0.00	-1.03E-01	0.2985
I3LLU0	Uncharacterized protein OS=Sus scrofa GN=GPDI1 PE=3 SV=1 - [I3LLU0_PIG]	0.852	1.066	0.956	1.026	1.004	1.019	1.027	0	0.23	-0.09	0.06	-0.04	-0.01	-0.03	-0.04	-7.80E-02	0.2984
Q7M2W6	Alpha-crystallin B chain OS=Sus scrofa GN=CRYAB PE=2 SV=1 - [CRYAB_PIG]	1.07	0.848	0.98	0.887	1.16	1.021	1.362	0	-0.10	0.24	0.03	0.17	-0.21	-0.03	-0.45	-1.72E-01	0.2950
Q1HL08	Peroxisomal enoyl coenzyme A hydratase 1 (Fragment) OS=Sus scrofa GN=ECH1 PE=4 SV=1 - [Q1HL08_PIG]	1.282	0.537	0.996	0.999	1.168	1.275	1.114	0	-0.36	0.90	0.01	0.00	-0.22	-0.35	-0.16	-3.18E-01	0.2949
U3MG62	Carnosine synthase 1 OS=Sus scrofa GN=CARNS1 PE=2 SV=1 - [U3MG62_PIG]	1.061	1.017	0.929	1.024	1.015	0.873	0.859	0	-0.09	-0.02	0.11	-0.03	-0.02	0.20	0.22	9.08E-02	0.2946
F1SK00	Isocitrate dehydrogenase [NADP] (Fragment) OS=Sus scrofa GN=IDH2 PE=3 SV=2 - [F1SK00_PIG]	1.336	1.071	1.132	1.073	1.096	0.975	1.039	0	-0.42	-0.10	-0.18	-0.10	-0.13	0.04	-0.06	1.11E-01	0.2945
L8AXK3	IgG heavy chain OS=Sus scrofa GN=IGHG PE=2 SV=1 - [L8AXK3_PIG]	0.985	0.807	0.749	0.821	0.786	0.819	0.805	0	0.02	0.31	0.42	0.28	0.35	0.29	0.31	1.21E-01	0.2928
F1SFA7	Uncharacterized protein OS=Sus scrofa GN=COL1A2 PE=4 SV=1 - [F1SFA7_PIG]	0.969	0.421	0.927	1.188	1.095	0.865	0.931	0	0.05	1.25	0.11	-0.25	-0.13	0.21	0.10	-3.67E-02	0.2915
F1S1A8	Uncharacterized protein OS=Sus scrofa GN=NDUF52 PE=3 SV=1 - [F1S1A8_PIG]	0.938	0.882	0.912	0.942	0.938	0.774											

I3LEF8	Uncharacterized protein OS=Sus scrofa GN=HSD17B4 PE=3 SV=1 - [I3LEF8_PIG]	1.319	1.111	1.283	1.21	1.107	1.002	0.918	0	-0.40	-0.15	-0.36	-0.28	-0.15	0.00	0.12	1.52E-01	0.2760
FUSGS9	Catalase OS=Sus scrofa GN=CAT PE=3 SV=1 - [FUSGS9_PIG]	0.919	0.821	0.931	0.939	0.734	0.937	0.735	0	0.12	0.28	0.10	0.09	0.45	0.09	0.44	1.41E-01	0.2747
F1SV06	Uncharacterized protein OS=Sus scrofa GN=PABPC4 PE=4 SV=1 - [F1SV06_PIG]	0.968	0.848	0.947	1.007	0.89	1.055	1.073	0	0.05	0.24	0.08	-0.01	0.17	-0.08	-0.10	-9.60E-02	0.2743
F1SLE7	Uncharacterized protein OS=Sus scrofa GN=LOC100522732 PE=4 SV=2 - [F1SLE7_PIG]	0.927	1.057	1.052	1.09	1.026	1.027	1.056	0	0.11	-0.08	-0.07	-0.12	-0.04	-0.04	-0.08	-5.87E-02	0.2728
F1S156	Uncharacterized protein OS=Sus scrofa GN=SYNP02 PE=4 SV=2 - [F1S156_PIG]	1.028	0.945	1.061	0.905	1.013	0.918	1.013	0	-0.04	0.08	-0.09	0.14	-0.02	0.12	-0.02	6.85E-02	0.2722
F1SDT0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100155350 PE=4 SV=1 - [F1SDT0_PIG]	0.948	0.944	0.935	0.968	0.811	0.906	0.955	0	0.08	0.08	0.10	0.05	0.30	0.14	0.07	7.52E-02	0.2706
I3LSD3	60S ribosomal protein L13 OS=Sus scrofa GN=RPL13 PE=3 SV=1 - [I3LSD3_PIG]	1.113	1.249	1.102	1.227	1.009	0.941	0.848	0	-0.15	-0.32	-0.14	-0.30	-0.01	0.09	0.24	1.58E-01	0.2697
Q9GKX6	Aldose 1-epimerase OS=Sus scrofa GN=GALM PE=2 SV=1 - [GALM_PIG]	1.109	1.224	0.979	1.12	1.032	1.237	1.38	0	-0.15	-0.29	0.03	-0.16	-0.05	-0.31	-0.46	-1.43E-01	0.2694
Q6UV40	Fructose-bisphosphate aldolase A (Fragment) OS=Sus scrofa GN=ALDOA PE=4 SV=1 - [Q6UV40_PIG]	0.857	0.849	0.874	0.914	0.945	0.933	0.956	0	0.22	0.24	0.19	0.13	0.08	0.10	0.06	-6.92E-02	0.2692
Q29561	UMP-CMP kinase OS=Sus scrofa GN=CMPK1 PE=1 SV=1 - [KCY_PIG]	1.097	0.896	1.082	0.88	0.907	0.991	1.015	0	-0.13	0.16	-0.11	0.18	0.14	0.01	-0.02	1.01E-01	0.2692
F1RLZ7	Uncharacterized protein (Fragment) OS=Sus scrofa GN=DPYSL3 PE=4 SV=2 - [F1RLZ7_PIG]	1.236	0.992	0.884	1.212	1.12	1.032	1.128	0	-0.31	0.01	0.18	-0.28	-0.16	-0.05	-0.17	-1.36E-01	0.2679
F1S415	Uncharacterized protein OS=Sus scrofa GN=BAG3 PE=4 SV=1 - [F1S415_PIG]	1.089	0.968	1.004	1.009	1.013	1.1	1.147	0	-0.12	0.05	-0.01	-0.01	-0.02	-0.14	-0.20	-7.13E-02	0.2664
Q9XSD9	Decorin OS=Sus scrofa GN=DCN PE=2 SV=1 - [PGS2_PIG]	1.019	0.781	1.134	1.112	1.151	1.037	1.025	0	-0.03	0.36	-0.18	-0.15	-0.20	-0.05	-0.04	-1.48E-01	0.2658
F1RR89	Uncharacterized protein (Fragment) OS=Sus scrofa GN=RARS PE=4 SV=2 - [F1RR89_PIG]	1.098	1.081	1.246	0.915	1.078	1.09	1.021	0	-0.13	-0.11	-0.32	0.13	-0.11	-0.12	-0.03	1.08E-01	0.2651
I3LPB5	Creatine kinase B-type (Fragment) OS=Sus scrofa GN=CKB PE=3 SV=1 - [I3LPB5_PIG]	0.893	0.787	0.918	0.774	0.914	0.886	0.684	0	0.16	0.35	0.12	0.37	0.13	0.17	0.55	1.47E-01	0.2646
F1S4A6	Uncharacterized protein (Fragment) OS=Sus scrofa GN=SYNJ2BP PE=4 SV=1 - [F1S4A6_PIG]	1.053	1.079	1.129	1.111	0.988	1.226	1.473	0	-0.07	-0.11	-0.18	-0.15	0.02	-0.29	-0.56	-1.57E-01	0.2635
F1SH25	Uncharacterized protein OS=Sus scrofa GN=RAB21 PE=3 SV=1 - [F1SH25_PIG]	0.999	1.121	1.118	1.218	1.178	0.982	1.193	0	0.00	-0.16	-0.16	-0.28	-0.24	0.03	-0.25	-1.06E-01	0.2628
D2K6G0	Thioredoxin reductase 2 OS=Sus scrofa PE=2 SV=1 - [D2K6G0_PIG]	0.924	0.592	0.896	0.536	0.823	0.715	0.743	0	0.11	0.76	0.16	0.90	0.28	0.48	0.43	2.66E-01	0.2627
F1SNG6	Uncharacterized protein OS=Sus scrofa GN=EXOC4 PE=4 SV=2 - [F1SNG6_PIG]	1.117	1.164	1.212	1.027	1.082	0.97	1.127	0	-0.16	-0.22	-0.28	-0.04	-0.11	0.04	-0.17	9.39E-02	0.2627
P00889	Citrate synthase, mitochondrial OS=Sus scrofa GN=CS PE=1 SV=2 - [CISY_PIG]	1.072	1.027	1.047	1.065	1.027	0.961	0.898	0	-0.10	-0.04	-0.07	-0.09	-0.04	0.06	0.16	7.21E-02	0.2626
D0G773	Isovaleryl Coenzyme A dehydrogenase OS=Sus scrofa GN=IVD PE=2 SV=1 - [D0G773_PIG]	1.011	1.271	1.282	1.116	1.076	1.037	0.743	0	-0.02	-0.35	-0.36	-0.16	-0.11	-0.05	0.43	2.08E-01	0.2615
Q9TV61	Myosin-1 OS=Sus scrofa GN=MYH1 PE=2 SV=1 - [MYH1_PIG]	1.038	1.177	0.999	0.952	1.042	0.955	1.028	0	-0.05	-0.24	0.00	0.07	-0.06	0.07	-0.04	8.14E-02	0.2612
F1RXZ6	Uncharacterized protein OS=Sus scrofa GN=BIN1 PE=4 SV=2 - [F1RXZ6_PIG]	1.874	1.835	2.176	2.068	1.149	0.951	0.947	0	-0.91	-0.88	-1.12	-1.05	-0.02	0.07	0.08	4.52E-01	0.2608
F1SAD9	Uncharacterized protein OS=Sus scrofa GN=PDIA4 PE=3 SV=1 - [F1SAD9_PIG]	0.881	0.729	0.801	0.952	0.691	0.753	0.71	0	-0.11	0.46	0.32	0.07	0.53	0.41	0.49	2.11E-01	0.2603
F2Z5B9	60S ribosomal protein L34 OS=Sus scrofa GN=RPL34 PE=4 SV=1 - [F2Z5B9_PIG]	0.899	1.031	1.195	1.173	1.15	1.083	1.048	0	0.15	-0.04	-0.26	-0.23	-0.20	-0.12	-0.07	-1.17E-01	0.2548
I3LB90	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LB90_PIG]	1.022	0.784	0.848	0.856	1.012	1.109	1.105	0	-0.03	0.35	0.24	0.22	-0.02	-0.15	-0.14	-1.61E-01	0.2535
D0EW19	Calcineurin catalytic subunit alpha isoform (Fragment) OS=Sus scrofa PE=2 SV=1 - [D0EW19_PIG]	1.074	1.056	0.936	1.347	0.875	1.262	1.185	0	-0.10	-0.08	0.10	-0.43	0.19	-0.34	-0.24	-1.83E-01	0.2532
I3L9T6	Uncharacterized protein OS=Sus scrofa GN=LOC100739576 PE=4 SV=1 - [I3L9T6_PIG]	1.118	0.648	1.162	1.287	1.288	1.075	1.002	0	-0.16	0.63	-0.22	-0.36	-0.37	-0.10	0.00	-2.71E-01	0.2526
F1SEN8	Uncharacterized protein OS=Sus scrofa GN=LDB3 PE=4 SV=2 - [F1SEN8_PIG]	1.054	0.987	1.082	0.961	1.018	0.986	1.022	0	-0.08	0.02	-0.11	0.06	-0.03	0.02	-0.03	4.78E-02	0.2510
Q1AG05	Calsarcin 3 OS=Sus scrofa PE=2 SV=1 - [Q1AG05_PIG]	1.1	0.871	1.038	0.87	0.97	0.91	0.972	0	-0.14	0.20	-0.05	0.20	0.04	0.14	0.04	1.03E-01	0.2506
F1RXC2	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CA2 PE=4 SV=1 - [F1RXC2_PIG]	0.984	0.973	1.168	1.003	1.135	1.208	1.092	0	0.02	0.04	-0.22	0.00	-0.18	-0.27	-0.13	-1.06E-01	0.2498
D0G0C7	Antioxidant protein 1 homolog (Yeast) OS=Sus scrofa GN=ATOX1 PE=4 SV=1 - [D0G0C7_PIG]	1.372	1.525	1.279	1.871	1.201	1.545	1.479	0	-0.46	-0.61	-0.36	-0.90	-0.26	-0.63	-0.56	-2.35E-01	0.2492
F1SM75	Uncharacterized protein OS=Sus scrofa GN=MYOM1 PE=4 SV=2 - [F1SM75_PIG]	0.979	1.092	1.119	0.994	1.025	0.96	1.003	0	0.04	-0.13	-0.16	0.01	-0.04	0.06	0.00	6.82E-02	0.2471
I3LMB7	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3LMB7_PIG]	1.008	0.815	0.778	0.439	0.702	0.885	0.916	0	-0.01	0.30	0.36	1.19	0.51	0.18	0.13	3.39E-01	0.2455
I3LD74	60S ribosomal protein L11 OS=Sus scrofa GN=RPL11 PE=3 SV=1 - [I3LD74_PIG]	1.076	1.059	1.052	1.142	1.072	1.022	1.111	0	-0.11	-0.08	-0.07	-0.19	-0.10	-0.03	-0.15	-5.34E-02	0.2452
F2Z5P1	Histone H2A (Fragment) OS=Sus scrofa GN=H2AFV PE=3 SV=1 - [F2Z5P1_PIG]	1.086	1.149	1.154	1.044	1.297	1.413	1.117	0	-0.12	-0.20	-0.21	-0.06	-0.38	-0.50	-0.16	-1.42E-01	0.2450
A9JPD4	Carnitine acetyl transferase OS=Sus scrofa GN=CRAT PE=2 SV=1 - [A9JPD4_PIG]	1.172	1.1	1.185	1.038	1.13	0.972	1.035	0	-0.23	-0.14	-0.24	-0.05	-0.18	0.04	-0.05	9.31E-02	0.2418
A8D349	Keratin 18 (Fragment) OS=Sus scrofa GN=KRT18 PE=2 SV=1 - [A8D349_PIG]	1.048	0.703	0.831	1.119	0.864	1.119	1.004	0	-0.07	0.51	0.27	-0.16	0.21	-0.16	-0.01	-2.07E-01	0.2407
Q4JH15	Troponin I OS=Sus scrofa GN=TNNI2 PE=2 SV=1 - [Q4JH15_PIG]	0.891	0.905	0.903	0.871	0.894	0.932	0.834	0	0.17	0.14	0.15	0.20	0.16	0.10	0.26	6.67E-02	0.2396
A3RCG1	Vitamin D3 receptor OS=Sus scrofa GN=VDR PE=2 SV=1 - [VDR_PIG]	1.136	1.276	1.054	1.035	1.447	1.245	1.294	0	-0.18	-0.35	-0.08	-0.05	-0.53	-0.32	-0.37	-1.65E-01	0.2392
I3LQ84	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LQ84_PIG]	1.037	0.649	0.914	1.152	1.01	0.981	0.981	0	-0.05	0.62	0.13	-0.20	-0.01	0.03	0.03	-2.16E-01	0.2356
Q0PY11	Elongation factor 1-alpha OS=Sus scrofa GN=EEF1A PE=2 SV=1 - [Q0PY11_PIG]	1.169	1.009	0.916	1.041	1.201	1.097	1.076	0	-0.23	0.01	0.13	-0.06	-0.26	-0.13	-0.11	-1.12E-01	0.2352
P00571	Adenylate kinase isoenzyme 1 OS=Sus scrofa GN=AK1 PE=1 SV=1 - [KAD1_PIG]	0.928	0.852	1.009	0.941	0.906	0.886	0.804	0	0.11	0.23	-0.01	0.09	0.14	0.17	0.31	9.84E-02	0.2352
Q2XVP4	Tubulin alpha-1B chain OS=Sus scrofa GN=TUBA1B PE=2 SV=1 - [TBA1B_PIG]	1.124	0.982	1.029	1.084	1.082	1.151	1.028	0	-0.17	0.03	-0.04	-0.12	-0.11	-0.20	-0.04	-7.23E-02	0.2335
F1SAD1	Uncharacterized protein OS=Sus scrofa GN=TPK1 PE=4 SV=2 - [F1SAD1_PIG]	0.846	0.923	0.9	0.904	0.967	0.98	1.044	0	0.24	0.12	0.15	0.15	0.05	0.03	-0.06	-8.70E-02	0.2335
F1RKU0	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial OS=Sus scrofa GN=IDH3A PE=3 SV=1 - [F1RKU0_PIG]	1.088	0.949	0.959	0.921	0.991	0.932	0.965	0	-0.12	0.08	0.06	0.12	0.01	0.10	0.05	6.76E-02	0.2321
I3LJ48	Uncharacterized protein (Fragment) OS=Sus scrofa GN=EHHADH PE=3 SV=1 - [I3LJ48_PIG]	1.26	0.853	1.116	1.053	0.857	0.915	0.877	0	-0.33	0.23	-0.16	-0.07	0.22	0.13	0.19	1.82E-01	0.2320
P67985	60S ribosomal protein L22 OS=Sus scrofa GN=RPL22 PE=2 SV=2 - [RL22_PIG]	1.052	1.008	0.883	0.916	0.836	0.903	1.012	0	-0.07	-0.01	0.18	0.13	0.26	0.15	-0.02	1.05E-01	0.2293
Q8WN73	Carbamoyl-phosphate synthetase 1 (Fragment) OS=Sus scrofa domesticus PE=2 SV=1 - [Q8WN73_PIG]	1.256	1.209	1.237	1.579	0.512	0.888	0.713	0	-0.33	-0.27	-0.31	-0.66	0.97	0.17	0.49	4.69E-01	0.2291
F1SEN2	Glutamate dehydrogenase 1, mitochondrial OS=Sus scrofa GN=GLUD1 PE=3 SV=1 - [F1SEN2_PIG]	1.008	0.842	0.936	0.91	0.876	0.839	0.924	0	-0.01	0.25	0.10	0.14	0.19	0.25	0.11	9.06E-02	0.2288
I3LF10	Uncharacterized protein (Fragment) OS=Sus scrofa GN=RAC1 PE=3 SV=1 - [I3LF10_PIG]	0.966	1.17	0.918	1.067	0.784	0.889	0.898	0	0.05	-0.23	0.12	-0.09	0.35	0.17	0.16	1.59E-01	0.2283
I3LK07	Uncharacterized protein OS=Sus scrofa GN=LOC100623679 PE=3 SV=1 - [I3LK07_PIG]	0.968	1.173	1.237	1.081	0.944	1.006	0.952	0	0.05	-0.23	-0.31	-0.11	0.08	-0.01	0.07	1.31E-01	0.2275
I3LMH4	Uncharacterized protein OS=Sus scrofa GN=EIF4G1 PE=4 SV=1 - [I3LMH4_PIG]	1.1	1.105	1.13	0.993	1.028	0.971	1.114	0	-0.14	-0.14	-0.18	0.01	-0.04	0.04	-0.16	7.87E-02	0.2266
I3LQ17	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LQ17_PIG]	1.144	0.975	1.003	0.986	1.193	1.154	1.109	0	-0.19	0.04	0.00	0.02	-0.25	-0.21	-0.15	-1.07E-01	0.2261
P00336	L-lactate dehydrogenase B chain OS=Sus scrofa GN=LDHB PE=1 SV=3 - [LDHB_PIG]	1.124	1.16	1.389	1.003	1.03	1.051	1.128	0	-0.17	-0.21	-0.47	0.00	-0.04	-0.07	-0.17	1.41E-01	0.2261
I3LAY6	Uncharacterized protein OS=Sus scrofa GN=NEXN PE=4 SV=1 - [I3LAY6_PIG]	0.932	0.978	0.94	1.023	0.899	1.105	1.129	0	0.10	0.03	0.09	-0.03	0.15	-0.14	-0.18	-1.05E-01	0.2260
F1SJR3	Uncharacterized protein OS=Sus scrofa GN=CSRNPI PE=4 SV=1 - [F1SJR3_PIG]	1.015	0.873	0.929	0.733	0.897	0.855	0.992	0	-0.02	0.20	0.11	0.45	0.16	0.23	0.01	1.40E-01	0.2253
F1SFG7	Uncharacterized protein OS=Sus scrofa GN=OPA1 PE=3 SV=2 - [F1SFG7_PIG]	0.945	1.16	1.054	1.011	0.909	0.984	0.979	0	0.08	-0.21	-0.08	-0.02	0.14				

F1RIE2	Uncharacterized protein OS=Sus scrofa GN=DDB1 PE=4 SV=1 - [F1RIE2_PIG]	0.899	1.01	1.066	0.927	0.913	0.957	0.966	0	0.15	-0.01	-0.09	0.11	0.13	0.06	0.05	7.67E-02	0.2115	
P02543	Vimentin OS=Sus scrofa GN=VIM PE=1 SV=2 - [VIME_PIG]	1.132	0.863	1.016	1.01	1.101	1.199	1.079	0	-0.18	0.21	-0.02	-0.01	-0.14	-0.26	-0.11	-1.34E-01	0.2096	
I3L6N8	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3L6N8_PIG]	0.93	1.165	1.253	0.918	1.039	0.871	1.033	0	0.10	-0.22	-0.33	0.12	-0.06	0.20	-0.05	-1.65E-01	0.2077	
B7STY5	Chloride intracellular channel 5 OS=Sus scrofa GN=CLUC5 PE=2 SV=1 - [B7STY5_PIG]	0.884	1.129	1.037	1.108	1.168	1.016	1.112	0	0.02	-0.18	-0.05	-0.15	-0.22	-0.02	-0.15	-8.60E-02	0.2077	
F2Z5F2	Uncharacterized protein OS=Sus scrofa GN=RAB10 PE=3 SV=1 - [F2Z5F2_PIG]	1.166	1.279	1.159	1.199	0.983	1.005	0.982	0	-0.22	-0.36	-0.21	-0.26	0.02	-0.01	0.03	1.43E-01	0.2074	
F1SFQ4	Uncharacterized protein OS=Sus scrofa GN=PCBP2 PE=4 SV=2 - [F1SFQ4_PIG]	0.995	1.04	1.089	1.026	1.079	1.104	1.068	0	0.01	-0.06	-0.12	-0.04	-0.11	-0.14	-0.09	-5.30E-02	0.2065	
Q6RVA9	Caveolin-1 OS=Sus scrofa GN=CAV1 PE=2 SV=1 - [CAV1_PIG]	0.949	0.906	0.971	1.005	1.018	1.087	0.931	0	0.08	0.14	0.04	-0.01	-0.03	-0.12	0.10	-7.76E-02	0.2064	
F1RPS8	ATP synthase subunit alpha heart isoform, mitochondrial (Fragment) OS=Sus scrofa GN=ATP5A1 PE=3 SV=1 - [F1RPS8_PIG]	1.093	0.969	0.938	0.969	0.974	0.936	0.918	0	-0.13	0.05	0.09	0.05	0.04	0.10	0.12	7.32E-02	0.2063	
F1SA33	Uncharacterized protein OS=Sus scrofa GN=LOC100157249 PE=4 SV=2 - [F1SA33_PIG]	1.247	1.284	1.271	1.17	1.018	1.092	1.076	0	-0.32	-0.36	-0.35	-0.23	-0.03	-0.13	-0.11	1.35E-01	0.2062	
F1RZV8	Uncharacterized protein OS=Sus scrofa GN=KLHL31 PE=4 SV=1 - [F1RZV8_PIG]	0.936	0.724	0.708	0.714	0.777	0.654	0.765	0	0.10	0.47	0.50	0.49	0.36	0.61	0.39	1.97E-01	0.2058	
F1RL38	Uncharacterized protein OS=Sus scrofa GN=DDT PE=4 SV=2 - [F1RL38_PIG]	1.502	1.225	1.354	1.095	1.103	1.065	1.168	0	-0.59	-0.29	-0.44	-0.13	-0.14	-0.09	-0.22	1.82E-01	0.2046	
P11607	Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 OS=Sus scrofa GN=ATP2A2 PE=1 SV=1 - [AT2A2_PIG]	1.198	1.205	1.204	1.02	0.994	1.088	1.145	0	-0.26	-0.27	-0.27	-0.03	0.01	-0.12	-0.20	1.15E-01	0.2045	
F1S2E3	Peptidyl-prolyl cis-trans isomerase OS=Sus scrofa GN=LOC100152612 PE=3 SV=1 - [F1S2E3_PIG]	1.378	1.026	1.07	1.016	0.96	0.957	1.046	0	-0.46	-0.04	-0.10	-0.02	0.06	0.06	-0.06	1.58E-01	0.2043	
F1SN27	Sorbitol dehydrogenase OS=Sus scrofa GN=SORD PE=3 SV=1 - [F1SN27_PIG]	1.004	0.843	0.822	0.961	1.062	1.074	0.922	0	-0.01	0.25	0.28	0.06	-0.09	-0.10	0.12	-1.35E-01	0.2043	
I3LUR7	Uncharacterized protein OS=Sus scrofa GN=COL6A3 PE=4 SV=1 - [I3LUR7_PIG]	0.979	0.718	0.901	1.095	0.962	0.982	0.974	0	0.03	0.48	0.15	-0.13	0.06	0.03	0.04	-1.47E-01	0.2042	
F1SDX9	Peroxisiredoxin-2 OS=Sus scrofa GN=PRDX2 PE=4 SV=2 - [F1SDX9_PIG]	1.069	1.067	1.207	0.94	1.14	0.971	0.934	0	-0.10	-0.09	-0.27	0.09	-0.19	0.04	0.10	1.26E-01	0.2041	
D0G778	Macrophage migration inhibitory factor (Fragment) OS=Sus scrofa GN=MIF PE=2 SV=1 - [D0G778_PIG]	1.116	1.056	1.07	1.023	1.146	1.165	1.143	0	-0.16	-0.08	-0.10	-0.03	-0.20	-0.22	-0.19	-7.70E-02	0.2040	
F1SPJ0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=WIBG PE=4 SV=2 - [F1SPJ0_PIG]	0.707	0.279	0.281	0.874	0.788	0.755	0.696	0	0.50	1.84	1.83	0.19	0.34	0.41	0.52	-6.77E-01	0.2034	
A1XQU1	Proteasome subunit beta type-7 OS=Sus scrofa GN=PSMB7 PE=2 SV=2 - [PSB7_PIG]	0.918	1.011	0.998	0.914	1.008	0.931	0.821	0	0.12	-0.02	0.00	0.13	-0.01	0.10	0.28	9.88E-02	0.2016	
F1RXU6	Uncharacterized protein OS=Sus scrofa GN=OSGEPL1 PE=4 SV=1 - [F1RXU6_PIG]	1.135	1.114	1.336	0.97	1.036	1.068	1.085	0	-0.18	-0.16	-0.42	0.04	-0.05	-0.09	-0.12	1.34E-01	0.2007	
I3L9U0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYL5 PE=4 SV=1 - [I3L9U0_PIG]	1.102	0.951	1.026	0.79	1.087	0.976	0.649	0	-0.14	0.07	-0.04	0.34	-0.12	0.04	0.62	2.46E-01	0.2007	
F1SR61	Uncharacterized protein OS=Sus scrofa GN=LOC100511787 PE=3 SV=2 - [F1SR61_PIG]	0.995	0.956	1.039	0.903	0.985	0.945	0.998	0	0.01	0.06	-0.06	0.15	0.02	0.08	0.00	5.91E-02	0.1975	
F1RXQ9	Ornithine carbamoyltransferase, mitochondrial OS=Sus scrofa GN=OTC PE=3 SV=2 - [F1RXQ9_PIG]	0.512	0.355	0.505	0.47	0.398	0.372	0.537	0	0.97	1.49	0.99	1.09	1.33	1.43	1.49	4.71E-01	0.1960	
F1SFE6	Uncharacterized protein OS=Sus scrofa GN=PSMC6 PE=4 SV=1 - [F1SFE6_PIG]	0.976	0.853	0.848	0.938	0.798	0.835	0.811	0	0.04	0.23	0.24	0.09	0.33	0.26	0.30	1.19E-01	0.1952	
A3RKG5	Progesterone receptor membrane component 2 OS=Sus scrofa GN=PGRCM2 PE=2 SV=1 - [A3RKG5_PIG]	0.986	1.325	0.957	0.926	0.971	0.953	0.934	0	0.02	-0.41	0.06	0.11	0.04	0.07	0.10	1.61E-01	0.1951	
D0G7F3	SH3-domain GRB2-like endophilin B1 OS=Sus scrofa GN=SH3GLB1 PE=2 SV=1 - [D0G7F3_PIG]	0.905	0.984	0.779	1.147	0.989	1.038	0.911	0	0.14	0.02	0.36	-0.20	0.02	-0.05	0.13	-1.57E-01	0.1949	
F1RHF0	Ubiquitin carboxyl-terminal hydrolase isozyme L3 (Fragment) OS=Sus scrofa GN=UCHL3 PE=4 SV=2 - [F1RHF0_PIG]	0.953	0.82	0.894	0.878	0.793	0.914	0.788	0	0.07	0.29	0.19	0.33	0.13	0.34	1.20E-01	0.1940		
Q6R185	Phosphoglycerate kinase 2 OS=Sus scrofa GN=PGK2 PE=2 SV=1 - [PGK2_PIG]	0.93	0.827	0.988	0.847	0.979	0.794	0.774	0	0.10	0.27	0.02	0.24	0.03	0.33	0.37	1.44E-01	0.1936	
Q7JFN3	Calcium/calmodulin-dependent protein kinase II isoform gamma-C protein kinase II OS=Sus scrofa GN=CAMK2G PE=2 SV=1 - [Q7JFN3_PIG]	0.849	0.974	0.938	1.026	1.011	0.966	0.972	0	0.24	0.04	0.09	-0.04	-0.02	0.05	0.04	-8.21E-02	0.1927	
F1SP32	Uncharacterized protein OS=Sus scrofa GN=RAD23B PE=4 SV=1 - [F1SP32_PIG]	1.067	1.037	1.155	0.947	0.982	1.046	1.044	0	-0.09	-0.05	-0.21	0.08	0.03	-0.06	-0.06	8.29E-02	0.1922	
B8Y4S3	Sarcoglycan alpha OS=Sus scrofa PE=2 SV=1 - [B8Y4S3_PIG]	1.22	0.934	0.982	0.979	0.714	1.015	0.884	0	-0.29	0.10	0.03	0.03	0.49	-0.02	0.18	2.09E-01	0.1920	
I3LNG8	Uncharacterized protein OS=Sus scrofa GN=STIP1 PE=4 SV=1 - [I3LNG8_PIG]	1.035	1.046	0.946	1.109	1.03	0.995	1.253	0	-0.05	-0.06	0.08	-0.15	-0.04	0.01	-0.33	-1.19E-01	0.1901	
F2Z572	Uncharacterized protein OS=Sus scrofa GN=LOC100517744 PE=3 SV=1 - [F2Z572_PIG]	0.91	1.079	1.026	0.989	0.905	0.954	0.929	0	0.14	-0.11	-0.04	0.02	0.14	0.07	0.11	8.62E-02	0.1901	
F1SOC0	Methionine aminopeptidase OS=Sus scrofa GN=LOC100513747 PE=2 SV=1 - [F1SOC0_PIG]	1.028	0.873	0.852	0.912	0.793	0.904	0.829	0	-0.04	0.20	0.23	0.13	0.33	0.15	0.27	1.24E-01	0.1901	
Q6V9B4	Adiponectin (Fragment) OS=Sus scrofa GN=APM1 PE=2 SV=1 - [Q6V9B4_PIG]	0.9	0.848	0.948	0.959	1.084	0.972	0.946	0	0.15	0.24	0.08	0.06	-0.12	0.04	0.08	-1.00E-01	0.1896	
F1RX35	Uncharacterized protein OS=Sus scrofa GN=LOC100627396 PE=4 SV=2 - [F1RX35_PIG]	1.215	0.893	1.042	0.935	1.4	1.288	1.281	0	-0.28	0.16	-0.06	0.10	-0.49	-0.37	-0.36	-2.33E-01	0.1895	
F1RI15	Uncharacterized protein OS=Sus scrofa GN=HSPA4 PE=3 SV=1 - [F1RI15_PIG]	0.898	0.975	0.893	1.001	0.949	0.971	1.053	0	0.16	0.04	0.16	0.00	0.08	0.04	-0.07	-7.82E-02	0.1874	
I3L7S4	Uncharacterized protein OS=Sus scrofa GN=CALD1 PE=4 SV=1 - [I3L7S4_PIG]	1.015	0.893	0.973	1.219	0.906	1.18	1.046	0	-0.02	0.16	0.04	-0.29	0.14	-0.24	-0.06	-1.57E-01	0.1873	
F1SB63	T-complex protein 1 subunit alpha OS=Sus scrofa GN=TCP1 PE=3 SV=1 - [F1SB63_PIG]	0.906	1.193	1.015	1.12	1.126	1.112	1.1	0	0.14	-0.25	-0.02	-0.16	-0.17	-0.15	-0.14	-1.23E-01	0.1869	
I3L8B1	Uncharacterized protein OS=Sus scrofa GN=NOL3 PE=4 SV=1 - [I3L8B1_PIG]	1.002	0.929	1.197	0.937	0.792	1.017	0.941	0	0.00	0.11	-0.26	0.09	0.34	-0.02	0.09	1.62E-01	0.1858	
B5APU3	Actin-related protein 2-like protein OS=Sus scrofa GN=ACTR2 PE=2 SV=1 - [B5APU3_PIG]	0.743	0.546	0.764	0.817	0.904	0.897	1.037	0	0.43	0.87	0.39	0.29	0.15	0.16	-0.05	-2.87E-01	0.1857	
L8B0W9	IgG heavy chain OS=Sus scrofa GN=IGHG PE=2 SV=1 - [L8B0W9_PIG]	0.901	0.853	0.682	0.585	0.795	0.813	0.716	0.582	0	0.15	0.23	0.55	0.33	0.48	0.78	2.40E-01	0.1848	
I3LV37	Uncharacterized protein (Fragment) OS=Sus scrofa PE=4 SV=1 - [I3LV37_PIG]	0.9	1.355	1.086	1.547	1.008	1.363	1.32	0	0.15	-0.44	-0.12	-0.63	-0.01	-0.45	-0.40	-2.71E-01	0.1843	
F1S3H1	Uncharacterized protein OS=Sus scrofa GN=ALDH6A1 PE=3 SV=2 - [F1S3H1_PIG]	0.988	0.862	0.957	0.882	0.9	0.947	0.848	0	0.02	0.81	0.21	0.06	0.18	0.05	0.24	8.86E-02	0.1831	
F1S418	Uncharacterized protein OS=Sus scrofa GN=PRDX3 PE=4 SV=1 - [F1S418_PIG]	1.178	0.956	0.921	0.921	0.96	0.906	0.926	0	-0.24	0.06	0.12	0.12	0.06	0.14	0.11	1.21E-01	0.1824	
A0ENH4	Toll-like receptor 3 OS=Sus scrofa GN=TLR3 PE=2 SV=1 - [A0ENH4_PIG]	0.977	1.109	0.949	0.914	1.197	1.188	1.046	0	0.03	-0.15	0.08	0.13	-0.26	-0.25	-0.49	-2.07E-01	0.1824	
F1RHJ2	Uncharacterized protein OS=Sus scrofa GN=LOC100153512 PE=4 SV=1 - [F1RHJ2_PIG]	0.837	1.021	0.965	0.955	1.032	1.103	1.052	0	0.26	-0.03	0.05	0.07	-0.05	-0.14	-0.07	-1.18E-01	0.1802	
I3LR05	Uncharacterized protein (Fragment) OS=Sus scrofa GN=AKAP9 PE=4 SV=1 - [I3LR05_PIG]	0.811	0.877	0.923	0.829	1.091	1.148	1.057	0	0.30	0.19	0.12	0.27	-0.13	-0.20	-0.08	-1.85E-01	0.1798	
F1RMJ9	Oxysterol-binding protein OS=Sus scrofa GN=OSBP PE=3 SV=1 - [F1RMJ9_PIG]	0.897	1.039	1.121	1.154	1.05	1.144	1.044	0	0.16	-0.06	-0.16	-0.21	-0.07	-0.19	-0.06	-1.18E-01	0.1797	
Q5EEL9	Skeletal muscle myosin regulatory light chain 2 (Fragment) OS=Sus scrofa GN=HUMMLC2B PE=4 SV=1 - [Q5EEL9_PIG]	0.898	0.995	0.782	0.846	0.805	0.895	0.763	0	0.16	0.01	0.35	0.24	0.31	0.16	0.39	1.47E-01	0.1797	
L8B0W0	IgG heavy chain OS=Sus scrofa GN=IGHG PE=2 SV=1 - [L8B0W0_PIG]	0.967	1.053	0.705	1.006	0.983	1.107	1.245	0	0.05	-0.07	0.50	-0.01	0.02	-0.15	-0.32	-2.31E-01	0.1791	
I3LDM7	Uncharacterized protein OS=Sus scrofa GN=GLO1 PE=4 SV=1 - [I3LDM7_PIG]	1.006	0.934	1.294	0.82	1.03	0.905	0.943	0	-0.01	0.10	-0.37	0.29	-0.04	0.14	0.08	1.89E-01	0.1787	
F1RX36	Fibrinogen alpha chain OS=Sus scrofa GN=FGA PE=4 SV=2 - [F1RX36_PIG]	1.227	1.053	1.019	0.999	1.314	1.303	1.25	0	-0.30	-0.07	-0.03	0.00	-0.39	-0.38	-0.32	-1.75E-01	0.1786	
L8B0R9	IgG heavy chain OS=Sus scrofa GN=IGHG PE=2 SV=1 - [L8B0R9_PIG]	0.955	0.928	0.777	0.821	0.799	0.731	0.921	0	0.07	0.11	0.36	0.28	0.32	0.45	0.12	1.60E-01	0.1786	
Q1G1K9	Mitochondrial NAD+isocitrate dehydrogenase 3 beta variant 1 OS=Sus scrofa GN=IDH3B PE=2 SV=1 - [Q1G1K9_PIG]	1.069	1.102	0.884	0.921	0.962	0.936	0.926	0	-0.10	-0.14	0.18	0.12	0.06	0.10	0.11	1.10E-01	0.1774	
F1RFY1	Profilin OS=Sus scrofa GN=PFN1 PE=3 SV=1 - [F1RFY1_PIG]	1.07	1.084	1.126	1.08	1.13	1.101	1.167	0	0.10	-0.10	-0.12	-0.17	-0.11	-0.18	-0.14	-0.22	-6.59E-02	0.1772
F1SK45	Uncharacterized protein OS=Sus scrofa GN=AK3 PE=3 SV=1 - [F1SK45_PIG]	1.181	1.065	1.075	0.882	1.107	0.971	1	0	-0.24	-0.09	-0.10	0.18	-0.15	0.04	0.00	1.28E-01	0.1769	
K9IW14	Talin-1 OS=Sus scrofa GN=TLN1 PE=2 SV=1 - [K9IW14_PIG]	1.169	1.094	0.97	1.408	1.116													

F1RQU2	Uncharacterized protein OS=Sus scrofa GN=HSPCB PE=3 SV=1 - [F1RQU2_PIG]	0.85	1.125	0.896	1.055	1.048	1.051	1.115	0	0.23	-0.17	0.16	-0.08	-0.07	-0.07	-0.16	-1.49E-01	0.1567
I3LMH3	Uncharacterized protein OS=Sus scrofa GN=INTS5 PE=4 SV=1 - [I3LMH3_PIG]	0.308	0.583	0.381	0.862	0.62	0.831	0.906	0	1.70	0.78	1.39	0.21	0.69	0.27	0.14	-6.39E-01	0.1565
P53027	60S ribosomal protein L10a (Fragment) OS=Sus scrofa GN=RLP10A PE=2 SV=3 - [RL10A_PIG]	1.064	0.791	0.796	1.035	1	1	1.08	0	-0.09	0.34	0.33	-0.05	0.00	0.00	-0.11	-1.85E-01	0.1560
I3LAA4	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LAA4_PIG]	1.087	0.901	0.759	0.849	0.704	0.836	0.833	0	-0.12	0.15	0.40	0.24	0.51	0.26	0.26	2.09E-01	0.1548
F1S816	Uncharacterized protein OS=Sus scrofa GN=RAB28 PE=3 SV=1 - [F1S816_PIG]	1.076	1.205	0.979	0.994	0.957	1.005	0.97	0	-0.11	-0.27	0.03	0.01	0.06	-0.01	0.04	1.13E-01	0.1545
F1S4Z9	Uncharacterized protein OS=Sus scrofa GN=NTA1 PE=4 SV=1 - [F1S4Z9_PIG]	1.021	1.019	0.918	0.985	0.931	0.886	0.946	0	-0.03	-0.03	0.12	0.02	0.10	0.17	0.08	7.83E-02	0.1545
F1RIS3	Uncharacterized protein OS=Sus scrofa GN=TRIM72 PE=4 SV=2 - [F1RIS3_PIG]	1.04	1.027	1.167	1.021	1.031	0.952	0.959	0	-0.06	-0.04	-0.22	-0.03	-0.04	0.07	0.06	9.38E-02	0.1544
I3L6B9	Uncharacterized protein OS=Sus scrofa GN=PSMD7 PE=4 SV=1 - [I3L6B9_PIG]	1.213	0.994	1.116	0.987	0.984	0.963	1.036	0	-0.28	0.01	-0.16	0.02	0.02	0.05	-0.05	1.18E-01	0.1528
L7PBE6	T-complex protein 1 subunit epsilon OS=Sus scrofa GN=CCCT5 PE=2 SV=1 - [L7PBE6_PIG]	0.863	0.975	0.972	1.005	0.981	1.004	1.04	0	0.21	0.04	0.04	-0.01	0.03	-0.01	-0.06	-8.30E-02	0.1524
I3LQ50	Uncharacterized protein OS=Sus scrofa GN=HNRNPK PE=4 SV=1 - [I3LQ50_PIG]	0.992	0.92	0.854	0.945	1.072	1.072	0.987	0	0.01	0.12	0.23	0.08	-0.10	-0.10	0.02	-1.15E-01	0.1516
F2ZSP3	Uncharacterized protein OS=Sus scrofa GN=UBE2D4 PE=3 SV=1 - [F2ZSP3_PIG]	1.082	1.045	1.035	1.008	0.999	0.894	1.028	0	-0.11	-0.06	-0.05	-0.01	0.00	0.16	-0.04	8.46E-02	0.1485
P00503	Aspartate aminotransferase, cytoplasmic OS=Sus scrofa GN=GOT1 PE=1 SV=3 - [AATC_PIG]	0.967	0.904	0.979	0.848	0.989	0.803	0.911	0	0.05	0.15	0.03	0.24	0.02	0.32	0.13	1.20E-01	0.1480
P62936	Peptidyl-prolyl cis-trans isomerase A OS=Sus scrofa GN=PP1A PE=1 SV=2 - [PP1A_PIG]	0.914	1.164	0.867	1.188	0.975	1.208	1.128	0	0.13	-0.22	0.21	-0.25	0.04	-0.27	-0.17	-1.94E-01	0.1476
P79381	Epoxide hydrolase 1 OS=Sus scrofa GN=EPHX1 PE=2 SV=1 - [HYEP_PIG]	1.137	1.212	1.186	1.201	0.778	1.028	0.814	0	-0.19	-0.28	-0.25	-0.26	0.36	-0.04	0.30	2.66E-01	0.1466
F1RWH1	Uncharacterized protein OS=Sus scrofa GN=CBX1 PE=4 SV=2 - [F1RWH1_PIG]	0.991	0.9	1.124	0.86	0.957	1.422	1.356	0	0.01	0.15	-0.17	-0.12	0.06	-0.51	-0.44	-2.50E-01	0.1465
F1SUU5	Uncharacterized protein OS=Sus scrofa GN=DNAJC16 PE=4 SV=1 - [F1SUU5_PIG]	1.211	0.9	1.16	1.059	0.957	0.827	0.806	0	-0.28	0.15	-0.21	-0.08	0.06	0.27	0.31	2.26E-01	0.1455
I3LJX2	Uncharacterized protein OS=Sus scrofa PE=4 SV=1 - [I3LJX2_PIG]	0.982	0.465	1.081	1.436	1.51	1.022	1.008	0	0.03	1.10	-0.11	-0.52	-0.59	-0.03	-0.01	-5.45E-01	0.1446
F1RQ91	Uncharacterized protein OS=Sus scrofa GN=RP54 PE=3 SV=2 - [F1RQ91_PIG]	1.008	1.127	1.097	1.141	1.119	1.08	1.122	0	-0.01	-0.17	-0.13	-0.19	-0.16	-0.11	-0.17	-7.80E-02	0.1437
F1SCR1	Uncharacterized protein (Fragment) OS=Sus scrofa GN=ROCK2 PE=4 SV=2 - [F1SCR1_PIG]	1.297	1.244	1.257	0.948	1.01	1.118	1.148	0	-0.38	-0.31	-0.33	0.08	-0.01	-0.16	-0.20	1.81E-01	0.1433
F1SHD6	Uncharacterized protein OS=Sus scrofa GN=EEF1B2 PE=3 SV=1 - [F1SHD6_PIG]	1.104	1.13	1.021	0.901	1.018	1.076	0.89	0	-0.14	-0.18	-0.03	0.15	-0.03	-0.11	0.17	1.34E-01	0.1428
F1S4N2	Uncharacterized protein OS=Sus scrofa GN=CYP5B1 PE=4 SV=1 - [F1S4N2_PIG]	1.112	1.029	1.02	1.26	1.055	1.093	1.105	0	-0.16	-0.04	-0.03	-0.33	-0.08	-0.13	-0.14	-1.12E-01	0.1428
I3LID5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=IGFN1 PE=4 SV=1 - [I3LID5_PIG]	0.993	0.721	0.924	0.958	1.08	0.998	1.16	0	0.01	0.47	0.11	0.06	-0.11	0.00	-0.21	-2.14E-01	0.1415
F1REW5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=CLIP1 PE=4 SV=2 - [F1REW5_PIG]	1.053	0.977	0.911	1.207	0.87	1.371	1.453	0	-0.54	0.03	0.13	-0.27	0.00	-0.46	-0.54	-2.90E-01	0.1413
C0JW62	Sarcoglycan gamma (Fragment) OS=Sus scrofa PE=2 SV=1 - [C0JW62_PIG]	1.064	1.073	0.895	0.943	0.608	1.016	0.762	0	-0.09	-0.10	0.16	0.08	0.72	-0.02	0.39	3.01E-01	0.1408
F1SUM3	Uncharacterized protein (Fragment) OS=Sus scrofa GN=SERPINH1 PE=3 SV=1 - [F1SUM3_PIG]	0.962	0.814	0.863	0.939	1.034	0.982	1.008	0	0.06	0.30	0.21	0.09	-0.05	0.03	-0.01	-1.27E-01	0.1404
F1SOA2	Peptidyl-prolyl cis-trans isomerase OS=Sus scrofa GN=LOC100737887 PE=3 SV=1 - [F1SOA2_PIG]	1.107	0.914	1.1	1.122	1.223	1.173	1.029	0	-0.15	0.13	-0.14	-0.17	-0.29	-0.23	-0.04	-1.43E-01	0.1402
Q9GMA6	Alpha-1-antichymotrypsin 2 OS=Sus scrofa GN=SERPINA3-2 PE=3 SV=1 - [Q9GMA6_PIG]	1.127	1.169	0.942	0.907	1	0.962	0.98	0	-0.17	-0.23	0.09	0.14	0.00	0.06	0.03	1.34E-01	0.1395
Q28936	Fibrinogen A-alpha-chain (Fragment) OS=Sus scrofa PE=4 SV=1 - [Q28936_PIG]	1.215	1.113	1.028	1.031	1.405	1.318	1.255	0	-0.28	-0.15	-0.04	-0.04	-0.49	-0.40	-0.33	-1.96E-01	0.1392
Q0R678	DJ-1 protein OS=Sus scrofa GN=PARK7 PE=2 SV=1 - [Q0R678_PIG]	1.037	1.036	1.123	1.018	1.035	0.947	0.972	0	-0.05	-0.05	-0.17	-0.03	-0.05	0.08	0.04	7.87E-02	0.1386
B5LOY6	Calpastatin OS=Sus scrofa GN=CAST PE=4 SV=1 - [B5LOY6_PIG]	0.949	0.936	1.079	0.984	0.97	0.921	0.954	0	-0.07	0.10	-0.11	0.02	0.04	0.12	0.07	8.43E-02	0.1386
F1RTN3	Moesin OS=Sus scrofa GN=MSN PE=4 SV=1 - [F1RTN3_PIG]	1.114	1.008	1.115	0.965	0.953	0.969	1.068	0	-0.16	-0.01	-0.16	0.05	0.07	0.05	-0.09	8.99E-02	0.1376
F1RHM8	Uncharacterized protein OS=Sus scrofa GN=РАНBP1 PE=4 SV=1 - [F1RHM8_PIG]	1.17	1.075	0.942	0.87	1.02	0.952	0.949	0	-0.23	-0.10	0.09	0.20	-0.03	0.07	0.08	1.41E-01	0.1376
A2IBS6	Sara2 protein OS=Sus scrofa PE=2 SV=1 - [A2IBS6_PIG]	1.008	1.19	1.191	1.013	1.097	0.899	0.854	0	-0.01	-0.25	-0.25	-0.02	-0.13	0.15	0.23	1.86E-01	0.1371
F1SSY3	Tropomodulin 4 OS=Sus scrofa GN=TMOD4 PE=2 SV=1 - [F1SSY3_PIG]	0.939	0.975	0.975	0.897	0.981	0.944	0.912	0	0.09	0.04	0.04	0.16	0.03	0.08	0.13	5.92E-02	0.1342
Q29554	Trifunctional enzyme subunit alpha, mitochondrial OS=Sus scrofa GN=HADHA PE=1 SV=1 - [ECHA_PIG]	1.066	1.044	1.065	1.015	1.039	0.92	0.99	0	-0.09	-0.06	-0.09	-0.02	-0.06	0.12	0.01	7.58E-02	0.1335
O02705	Heat shock protein HSP 90-alpha OS=Sus scrofa GN=HSP90AA1 PE=2 SV=3 - [HS90A_PIG]	0.904	1.083	0.97	0.99	1.1	1.063	1.183	0	0.15	-0.12	0.04	0.01	-0.14	-0.09	-0.24	-1.32E-01	0.1328
F2ZSL7	Proteasome subunit alpha type OS=Sus scrofa GN=LOC100516779 PE=3 SV=1 - [F2ZSL7_PIG]	0.977	0.957	1.008	0.989	0.954	0.946	0.896	0	0.03	0.06	-0.01	0.02	0.07	0.08	0.16	5.92E-02	0.1319
Q6JLA8	Four and a half LIM domains 3 (Fragment) OS=Sus scrofa PE=4 SV=1 - [Q6JLA8_PIG]	0.99	0.859	0.937	0.857	0.805	0.879	0.942	0	0.01	0.22	0.09	0.22	0.31	0.19	0.09	1.20E-01	0.1312
Q00P27	Glyceralddehyde-3-phosphate dehydrogenase (Fragment) OS=Sus scrofa GN=GAPDH PE=2 SV=1 - [Q00P27_PIG]	0.986	0.937	0.841	0.906	0.92	0.797	0.764	0	0.02	0.09	0.25	0.14	0.12	0.33	0.39	1.54E-01	0.1298
B2XWS1	Guanylate binding protein 1 OS=Sus scrofa GN=GBP1 PE=2 SV=1 - [B2XWS1_PIG]	1.083	1.139	0.914	1.103	1.161	1.195	1.075	0	-0.12	-0.19	0.13	-0.14	-0.22	-0.26	-0.10	-1.36E-01	0.1295
F1S1R8	Uncharacterized protein OS=Sus scrofa GN=PGBD1 PE=4 SV=2 - [F1S1R8_PIG]	0.919	0.998	1.126	1.015	0.737	0.927	0.844	0	0.12	0.00	-0.17	-0.02	0.44	0.11	0.24	2.05E-01	0.1269
Q1TTA9	Type VI collagen alpha-1 chain (Fragment) OS=Sus scrofa GN=COL6A1 PE=2 SV=1 - [Q1TTA9_PIG]	0.978	0.697	0.837	1.045	1.047	0.992	0.966	0	0.03	-0.92	-0.26	-0.06	-0.07	0.01	0.05	-2.19E-01	0.1269
F1RR64	Uncharacterized protein OS=Sus scrofa GN=ENDOG PE=4 SV=1 - [F1RR64_PIG]	0.671	0.475	0.859	0.964	0.837	0.98	1.32	0	0.58	1.07	0.22	0.05	0.26	0.03	-0.40	-4.83E-01	0.1265
P23687	Prolyl endopeptidase OS=Sus scrofa GN=PREP PE=1 SV=1 - [PPCE_PIG]	0.953	0.976	1.007	0.992	0.996	1.03	1.042	0	0.07	0.04	-0.01	0.01	0.01	-0.04	-0.06	-4.48E-02	0.1264
F1SAW8	Calsequestrin (Fragment) OS=Sus scrofa GN=CASQ2 PE=3 SV=2 - [F1SAW8_PIG]	1.046	1.464	1.213	1.166	1.361	1.628	1.762	0	-0.06	-0.55	-0.28	-0.22	-0.44	-0.70	-0.82	-3.23E-01	0.1264
I3LHC8	Uncharacterized protein OS=Sus scrofa GN=LOC100522692 PE=4 SV=1 - [I3LHC8_PIG]	1.324	1.07	1.174	0.839	0.927	0.928	1.177	0	-0.40	-0.10	-0.23	0.25	0.11	0.11	-0.24	2.42E-01	0.1248
D2XN65	Glycerol-3-phosphate dehydrogenase 1 OS=Sus scrofa PE=3 SV=1 - [D2XN65_PIG]	0.978	1.133	1.194	0.981	1.038	0.925	0.955	0	0.03	-0.18	-0.26	0.03	-0.05	0.11	0.07	1.39E-01	0.1247
I3LNK5	Uncharacterized protein (Fragment) OS=Sus scrofa GN=GSPT1 PE=4 SV=1 - [I3LKN5_PIG]	1.076	0.983	1.12	0.992	0.965	0.923	1.021	0	-0.11	0.02	-0.16	0.01	0.05	0.12	-0.03	9.83E-02	0.1191
ASPFO0	B-factor, proteridin OS=Sus scrofa GN=BF PE=3 SV=1 - [ASPFO0_PIG]	0.986	1.061	1.002	1.066	1.065	1.112	1.006	0	0.02	-0.09	0.00	-0.09	-0.09	-0.15	-0.01	-6.92E-02	0.1166
F1SUE3	Uncharacterized protein OS=Sus scrofa GN=PPA1 PE=4 SV=2 - [F1SUE3_PIG]	1.154	1.061	0.991	1.095	1.082	1.194	1.343	0	-0.21	-0.09	0.01	-0.13	-0.11	-0.26	-0.43	-1.62E-01	0.1157
A1XQU9	40S ribosomal protein S20 OS=Sus scrofa GN=RP520 PE=3 SV=1 - [RS20_PIG]	0.996	0.939	0.914	1.087	0.978	0.986	1.042	0	0.01	0.09	0.13	-0.12	0.03	0.02	-0.06	-8.84E-02	0.1154
F1RQF0	Uncharacterized protein (Fragment) OS=Sus scrofa GN=NIPAL4 PE=4 SV=2 - [F1RQF0_PIG]	0.964	1.17	1.199	0.954	0.991	0.974	0.975	0	0.05	-0.23	-0.26	0.07	0.01	0.04	0.04	1.48E-01	0.1143
I3LP02	Uncharacterized protein OS=Sus scrofa GN=ACAT1 PE=3 SV=1 - [I3LP02_PIG]	1.259	1.008	1.043	0.954	1.046	0.913	0.912	0	-0.33	-0.01	-0.06	0.07	-0.06	0.13	0.13	1.68E-01	0.1142
J9JIK8	Protein-L-isaspartate O-methyltransferase (Fragment) OS=Sus scrofa GN=PCMT1 PE=3 SV=1 - [J9JIK8_PIG]	1.073	0.999	1.053	0.922	0.932	0.952	1.056	0	-0.10	0.00	-0.07	0.12	0.10	0.07	-0.08	9.65E-02	0.1127
Q29571	Leucine aminopeptidase (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29571_PIG]	1.013	0.792	0.96	0.809	0.956	0.707	0.725	0	-0.02	0.34	0.06	0.31	0.06	0.50	0.46	2.40E-01	0.1124
F1SL51	Uncharacterized protein (Fragment) OS=Sus scrofa GN=LOC100739453 PE=4 SV=1 - [F1SL51_PIG]	0.997	0.874	0.762	1.039	1.138	0.958	1.016	0	0.00	0.19	0.39	-0.06	-0.19	0.06	-0.02	-1.98E-01	0.1119
F1RUE3	Uncharacterized protein OS=Sus scrofa GN=ALDH5A1 PE=3 SV=2 - [F1RUE3_PIG]	1.324	1.233	1.318	1.096	0.988	1.024	1.129	0	-0.40	-0.30	-0.40	-0.12	-0.02	-0.03	-0.18	1.95E-01	0.1116
F1RP17	T-complex protein 1 subunit gamma OS=Sus scrofa GN=CCCT3 PE=3 SV=2 - [F1RP17_PIG]	0.913	1.005	0.882	1.042	0.974	1.051	1.003	0	0.13	-0.01	0.18	-0.06					

A1XQT6	MLC1f OS-Sus scrofa GN=MYL1 PE=2 SV=1 - [A1XQT6_PIG]	1.015	0.955	0.975	0.762	0.95	0.959	0.885	0	-0.02	0.07	0.04	0.39	0.07	0.06	0.18	1.55E-01	0.0971
F1S2K3	Uncharacterized protein OS-Sus scrofa GN=LOC100621711 PE=4 SV=1 - [F1S2K3_PIG]	0.926	0.953	0.992	0.87	0.904	0.976	0.896	0	0.11	0.07	0.01	0.20	0.15	0.04	0.16	8.70E-02	0.0943
Q5XLd3	Creatine kinase M-type OS-Sus scrofa GN=CKM PE=2 SV=1 - [KCRM_PIG]	1.001	0.952	1.1	0.869	1.03	0.854	0.906	0	0.00	0.07	-0.14	0.20	-0.04	0.23	0.14	1.50E-01	0.0933
F1S6B5	Fibromodulin (Fragment) OS-Sus scrofa GN=FMOD PE=4 SV=2 - [F1S6B5_PIG]	0.972	0.172	0.53	1.352	1.3	0.88	1.981	0	0.04	2.54	0.92	-0.44	-0.38	0.18	-0.99	-1.28E+00	0.0930
F1RK48	Uncharacterized protein (Fragment) OS-Sus scrofa GN=SRL PE=4 SV=2 - [F1RK48_PIG]	1.047	1.184	1.219	0.988	1.067	0.974	0.963	0	-0.07	-0.24	-0.29	0.02	-0.09	0.04	0.05	1.53E-01	0.0922
F1SU52	Uncharacterized protein OS-Sus scrofa GN=LOC100522130 PE=4 SV=2 - [F1SU52_PIG]	1.052	1.062	0.926	0.899	0.999	0.921	0.91	0	-0.07	-0.09	0.11	0.15	0.00	0.12	0.14	1.15E-01	0.0902
F1RPB7	Uncharacterized protein OS-Sus scrofa GN=DRG1 PE=4 SV=2 - [F1RPB7_PIG]	1.115	1.232	1.124	0.998	0.783	1.064	0.978	0	-0.16	-0.30	-0.17	0.00	0.35	-0.09	0.03	2.31E-01	0.0895
F1SFF4	Uncharacterized protein OS-Sus scrofa GN=C14orf166 PE=4 SV=1 - [F1SFF4_PIG]	0.878	0.782	0.83	0.905	0.96	1.061	1.049	0	0.19	0.35	0.27	0.14	0.06	-0.09	-0.07	-1.91E-01	0.0870
F1RR78	Uncharacterized protein OS-Sus scrofa GN=LOC100049693 PE=4 SV=2 - [F1RR78_PIG]	0.959	0.929	0.946	1.031	0.948	1.091	1.038	0	0.06	0.11	0.08	-0.04	0.08	-0.13	-0.05	-9.83E-02	0.0851
P49931	Antibacterial peptide PMAP-36 OS-Sus scrofa GN=PMAP36 PE=2 SV=1 - [PMP36_PIG]	1.036	0.691	0.785	0.987	0.98	1.239	1.245	0	-0.05	0.53	0.35	0.02	0.03	-0.31	-0.32	-3.52E-01	0.0846
I3L918	Uncharacterized protein OS-Sus scrofa PE=4 SV=1 - [I3L918_PIG]	0.932	0.912	0.859	0.986	0.96	1.015	1.005	0	0.10	0.13	0.22	0.02	0.06	-0.02	-0.01	-1.01E-01	0.0834
I3LD55	Uncharacterized protein (Fragment) OS-Sus scrofa GN=COQ3 PE=4 SV=1 - [I3LD55_PIG]	0.77	1.833	1.721	1.69	2.577	1.735	2.277	0	0.38	-0.87	-0.78	-0.76	-1.37	-0.79	-1.19	-7.06E-01	0.0822
B5APV1	Actin-related protein 2/3 complex subunit 5 OS-Sus scrofa PE=2 SV=1 - [B5APV1_PIG]	0.734	1.089	0.645	1.188	1.216	0.95	1.183	0	0.45	-0.12	0.63	-0.25	-0.28	-0.07	-0.24	-4.14E-01	0.0814
K9IVP8	Obg-like ATPase 1 isoform 2 OS-Sus scrofa GN=OLA1 PE=2 SV=1 - [K9IVP8_PIG]	1.099	1.113	1.111	0.979	1.064	1.01	0.992	0	-0.14	-0.15	-0.15	0.03	-0.09	-0.01	0.01	9.52E-02	0.0813
F1RGP8	Uncharacterized protein (Fragment) OS-Sus scrofa GN=PBXIP1 PE=4 SV=2 - [F1RGP8_PIG]	0.99	1.1	0.944	1.095	1.02	1.231	1.146	0	0.01	-0.14	0.08	-0.13	-0.03	-0.30	-0.20	-1.54E-01	0.0805
Q27HS3	Vascular smooth muscle alpha-actin (Fragment) OS-Sus scrofa PE=2 SV=1 - [Q27HS3_PIG]	1.038	1.147	1.201	0.966	1.022	0.94	1.029	0	-0.05	-0.20	-0.26	0.05	-0.03	0.09	-0.04	1.46E-01	0.0798
F2Z5K4	Uncharacterized protein OS-Sus scrofa GN=RHOC PE=3 SV=1 - [F2Z5K4_PIG]	1.048	1.012	1.2	1.258	1.209	1.134	1.123	0	-0.07	-0.02	-0.26	-0.33	-0.27	-0.18	-0.17	-1.51E-01	0.0796
I3LTB8	Uncharacterized protein OS-Sus scrofa PE=4 SV=1 - [I3LTB8_PIG]	1.235	0.924	1.108	0.882	0.983	0.88	0.931	0	-0.30	0.11	-0.15	0.18	0.02	0.18	0.10	2.08E-01	0.0790
Q5GN48	Dystrophin OS-Sus scrofa GN=DMD PE=1 SV=1 - [DMD_PIG]	1.009	0.978	0.98	1.023	0.988	1.038	1.054	0	-0.01	0.03	0.03	-0.03	0.02	-0.05	-0.08	-4.83E-02	0.0781
F1RVA0	Histone H3 (Fragment) OS-Sus scrofa GN=H3F3B PE=3 SV=1 - [F1RVA0_PIG]	1.142	0.927	1.235	1.152	1.263	1.363	1.231	0	-0.19	0.11	-0.30	-0.20	-0.34	-0.45	-0.30	-2.25E-01	0.0772
F1RIU3	Uncharacterized protein OS-Sus scrofa GN=CTF6A PE=3 SV=1 - [F1RIU3_PIG]	0.902	1.073	1.116	1.072	1.191	1.132	1.171	0	0.15	-0.10	-0.16	-0.10	-0.25	-0.18	-0.23	-1.62E-01	0.0748
F1SSN5	Uncharacterized protein OS-Sus scrofa GN=KTN1 PE=4 SV=1 - [F1SSN5_PIG]	1.039	1.124	1.065	1.169	1.184	1.146	1.06	0	-0.06	-0.17	-0.09	-0.23	-0.24	-0.20	-0.08	-1.09E-01	0.0739
I3LBK8	Uncharacterized protein (Fragment) OS-Sus scrofa GN=LOC100626333 PE=3 SV=1 - [I3LBK8_PIG]	0.886	0.986	0.918	1.059	1.233	1.124	0.946	0	0.17	0.02	0.12	-0.08	-0.30	-0.17	0.08	-1.98E-01	0.0708
F1SI13	Peptidyl-prolyl cis-trans isomerase (Fragment) OS-Sus scrofa GN=FKBP3 PE=4 SV=2 - [F1SI13_PIG]	1.116	1.191	1.067	0.918	1.022	0.942	1.047	0	-0.16	-0.25	-0.09	0.12	-0.03	0.09	-0.07	1.54E-01	0.0701
Q7M3B0	Actin beta (Fragments) OS-Sus scrofa domestica PE=1 SV=1 - [Q7M3B0_PIG]	1.023	1.049	1.084	0.872	1.066	0.897	0.905	0	-0.03	-0.07	-0.12	0.20	-0.09	0.16	0.14	1.56E-01	0.0678
Q684M6	CDC37 cell division cycle 37 protein OS-Sus scrofa GN=CDC37 PE=4 SV=1 - [Q684M6_PIG]	0.885	0.939	0.913	0.975	0.983	1.166	1.05	0	0.18	0.09	0.13	0.04	0.02	-0.22	-0.07	-1.57E-01	0.0671
P80031	Glutathione S-transferase P OS-Sus scrofa GN=GSTP1 PE=1 SV=2 - [GSTP1_PIG]	0.99	0.921	0.93	0.916	0.938	0.876	0.837	0	0.01	0.10	0.13	0.09	0.19	0.26	1.07E-01	0.0644	
Pe2802	Histone H4 OS-Sus scrofa PE=1 SV=2 - [H4_PIG]	1.059	0.95	0.996	1.064	1.024	1.216	1.125	0	-0.08	0.07	0.01	-0.09	-0.03	-0.28	-0.17	-1.43E-01	0.0623
I3LAB6	Proteasome subunit alpha type OS-Sus scrofa GN=PSMA2 PE=2 SV=1 - [I3LAB6_PIG]	0.971	0.932	0.879	0.906	0.846	0.9	0.85	0	0.04	0.10	0.19	0.14	0.24	0.15	0.23	1.10E-01	0.0623
F1RZM4	Uncharacterized protein (Fragment) OS-Sus scrofa GN=LAMA4 PE=4 SV=2 - [F1RZM4_PIG]	0.942	0.885	0.828	0.849	0.785	0.859	0.75	0	0.09	0.18	0.27	0.24	0.35	0.22	0.42	1.71E-01	0.0622
F1S4R2	Uncharacterized protein (Fragment) OS-Sus scrofa GN=ATL2 PE=4 SV=2 - [F1S4R2_PIG]	1.034	1.197	1.202	1.017	0.997	0.961	0.966	0	-0.05	-0.26	-0.27	-0.02	0.00	0.06	0.05	1.65E-01	0.0621
F1SLZ3	Uncharacterized protein (Fragment) OS-Sus scrofa GN=LOC100520244 PE=4 SV=1 - [F1SLZ3_PIG]	0.292	0.539	0.529	1.103	0.862	0.872	1.093	0	1.78	0.89	0.92	-0.14	0.21	0.20	-0.13	-8.61E-01	0.0618
Q8WNY0	Glyceraldehyde-3-phosphate dehydrogenase (Fragment) OS-Sus scrofa PE=2 SV=1 - [Q8WNY0_PIG]	0.795	0.879	0.534	1.009	1.194	1.089	1.036	0	0.33	0.19	0.91	-0.01	-0.26	-0.12	-0.05	-4.66E-01	0.0609
F1SEL2	Uncharacterized protein (Fragment) OS-Sus scrofa GN=FRMPD2 PE=4 SV=2 - [F1SEL2_PIG]	1.227	0.914	1.051	0.902	0.961	0.847	0.729	0	-0.30	0.13	-0.07	0.15	0.06	0.24	0.46	2.85E-01	0.0604
B3VMQ9	Adenylosuccinate lyase OS-Sus scrofa GN=ADSL PE=2 SV=1 - [B3VMQ9_PIG]	0.908	0.906	1.031	1.083	1.101	1.077	0.979	0	0.14	0.14	-0.04	-0.12	-0.14	-0.11	0.03	-1.42E-01	0.0598
F1S891	Small nuclear ribonucleoprotein-associated protein OS-Sus scrofa GN=SNRPB PE=3 SV=2 - [F1S891_PIG]	1.094	1.088	0.867	1.132	1.249	1.212	1.083	0	-0.13	-0.12	0.21	-0.18	-0.32	-0.28	-0.12	-2.12E-01	0.0592
Q7M326	Acylinmoacyl-peptidase (Fragment) OS-Sus scrofa domestica PE=1 SV=1 - [Q7M326_PIG]	1.123	1.069	0.86	0.934	0.788	0.857	0.87	0	-0.17	-0.10	0.22	0.10	0.34	0.22	0.20	2.28E-01	0.0583
I3LM03	Carbonyl reductase [NADPH] 1 OS-Sus scrofa GN=CBR1 PE=3 SV=1 - [I3LM03_PIG]	1.158	1.043	1.077	1.05	0.917	0.944	0.959	0	-0.21	-0.06	-0.11	-0.07	0.13	0.08	0.06	1.44E-01	0.0570
F1SBS3	Uncharacterized protein OS-Sus scrofa GN=SYCP1 PE=4 SV=1 - [F1SBS3_PIG]	1.3	1.379	1.187	1	0.938	1.121	0.932	0	-0.38	-0.46	-0.25	0.00	0.09	-0.16	0.10	2.80E-01	0.0563
O11780	Transforming growth factor-beta-induced protein ig-h3 OS-Sus scrofa GN=TGFBI PE=1 SV=1 - [BGH3_PIG]	1.008	1.073	1.08	1.055	1.149	1.154	1.105	0	-0.01	-0.10	-0.11	-0.08	-0.20	-0.21	-0.14	-1.01E-01	0.0526
F1SUR3	Uncharacterized protein (Fragment) OS-Sus scrofa GN=PGM2L1 PE=4 SV=2 - [F1SUR3_PIG]	1.099	0.92	1.04	1.4	1.058	1.125	1.408	0	-0.14	0.12	-0.06	-0.49	-0.08	-0.17	-0.49	-2.89E-01	0.0517
F1S786	5'-AMP-activated protein kinase catalytic subunit alpha-2 (Fragment) OS-Sus scrofa GN=PRKAA2 PE=4 SV=2 - [F1S786_PIG]	1.026	0.988	0.875	0.974	0.634	0.76	0.749	0	-0.04	0.02	0.19	0.04	0.66	0.40	0.42	3.34E-01	0.0513
F1RHL9	Uncharacterized protein OS-Sus scrofa GN=ACTN2 PE=4 SV=1 - [F1RHL9_PIG]	1.041	1.063	1.036	0.917	0.978	0.946	1.03	0	-0.06	-0.09	-0.05	0.13	0.03	0.08	-0.04	9.79E-02	0.0510
Q29551	Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondrial OS-Sus scrofa GN=OXCT1 PE=1 SV=2 - [SCOT1_PIG]	1.05	1.024	0.941	0.997	0.896	0.853	0.801	0	-0.07	-0.03	0.09	0.00	0.16	0.23	0.32	1.82E-01	0.0507
F1SUC9	Uncharacterized protein OS-Sus scrofa GN=LRRRC20 PE=4 SV=2 - [F1SUC9_PIG]	0.975	1	1.013	1.005	0.926	0.881	0.886	0	0.04	0.00	-0.02	-0.01	0.11	0.18	0.17	1.11E-01	0.0503
L8B180	IgG heavy chain OS-Sus scrofa GN=IGHG PE=2 SV=1 - [L8B180_PIG]	0.946	0.685	0.603	0.601	0.47	0.642	0.527	0	0.08	0.55	0.73	1.09	0.64	0.92	5.08E-01	0.0469	
F1SIH4	Uncharacterized protein OS-Sus scrofa GN=GARS PE=4 SV=1 - [F1SIH4_PIG]	1.157	1.035	1.568	1.417	1.676	1.519	1.534	0	-0.21	-0.05	-0.65	-0.50	-0.75	-0.60	-0.62	-3.90E-01	0.0463
Q29014	Alpha-1 acid glycoprotein (Fragment) OS-Sus scrofa PE=2 SV=1 - [Q29014_PIG]	1.109	0.934	1.135	1.136	1.265	1.162	1.178	0	-0.15	0.10	-0.18	-0.34	-0.22	-0.24	-1.86E-01	0.0454	
I3LV18	60S ribosomal protein L7a OS-Sus scrofa GN=RPL7A PE=4 SV=1 - [I3LV18_PIG]	1.178	1.176	1.276	1.461	1.381	1.416	1.208	0	-0.24	-0.23	-0.35	-0.55	-0.47	-0.50	-0.27	-2.41E-01	0.0446
F1SID9	Uncharacterized protein OS-Sus scrofa GN=DARS PE=3 SV=1 - [F1SID9_PIG]	0.993	1.054	1.02	1.088	1.044	1.041	1.09	0	0.01	-0.08	-0.03	-0.12	-0.06	-0.06	-0.12	-6.79E-02	0.0427
I3LGNS	Uncharacterized protein (Fragment) OS-Sus scrofa PE=4 SV=1 - [I3LGNS_PIG]	0.892	0.749	0.807	0.966	0.997	1.063	1.043	0	0.16	0.42	0.31	0.05	0.00	-0.09	-0.06	-2.46E-01	0.0420
F1SRI8	Uncharacterized protein (Fragment) OS-Sus scrofa GN=MYBPC1 PE=4 SV=2 - [F1SRI8_PIG]	1.08	1.086	1.12	0.939	1.042	0.934	0.997	0	-0.11	-0.12	-0.16	0.09	-0.06	0.10	0.00	1.32E-01	0.0420
F1STB2	Uncharacterized protein OS-Sus scrofa GN=HINT2 PE=4 SV=1 - [F1STB2_PIG]	1.031	0.97	0.939	0.917	0.934	0.76	0.849	0	-0.04	0.04	0.09	0.13	0.10	0.40	0.24	1.91E-01	0.0407
F1SUE4	Uncharacterized protein OS-Sus scrofa GN=ASPN PE=4 SV=1 - [F1SUE4_PIG]	0.945	0.746	0.839	1.335	1.221	1.012	0.994	0	0.08	0.42	0.25	-0.42	-0.29	-0.02	0.01	-3.68E-01	0.0394
K7GQ48	Uncharacterized protein OS-Sus scrofa GN=A2M PE=4 SV=1 - [K7GQ48_PIG]	0.96	1.145	1.007	1.037	0.554	0.487	0.348	0	0.06	-0.20	-0.01	-0.05	0.85	1.04	1.52	8.77E-01	0.0393
DOG0C8	Chaperonin containing TCP1, subunit 2 (Beta) OS-Sus scrofa GN=CCT2 PE=2 SV=1 - [DOG0C8_PIG]	0.799	0.918	0.838	1.002	0.988	1.033	1.026	0	0.32	0.12	0.25	0.00	0.02	-0.05	-0.04	-1.93E-01	0.0390
F1SDJ8	Uncharacterized protein OS-Sus scrofa GN=GLRX3 PE=4 SV=1 - [F1SDJ8_PIG]	0.94	0.959	0.946	0.929	0.875	0.918	0.927	0	0.09	0.06	0.08	0.11	0.19	0.12	0.11	7.55E-02	0.0382
I3JL14	Uncharacterized protein (Fragment) OS-Sus scrofa GN=MYO1C PE=4 SV=1 - [I3JL14_PIG]	0.765	1.103	1.283	1.286	1.442	1.446	1.37	0	0.39	-0.14	-0.36	-0.36	-0.53	-0.5			

F1SLY4	Uncharacterized protein OS=Sus scrofa GN=LMD02 PE=4 SV=1 - [F1SLY4_PIG]	0.796	0.805	1.046	0.765	0.5	0.41	0.638	0	0.33	0.31	-0.06	0.39	1.00	1.29	0.65	6.86E-01	0.0216
F1RP05	Uncharacterized protein OS=Sus scrofa GN=DNAJA2 PE=3 SV=1 - [F1RP05_PIG]	0.898	1.007	0.957	0.911	0.868	0.881	0.839	0	0.16	-0.01	0.06	0.13	0.20	0.18	0.25	1.42E-01	0.0205
K7GNZ3	Uncharacterized protein OS=Sus scrofa GN=NACA PE=4 SV=1 - [K7GNZ3_PIG]	0.856	0.93	0.85	1.166	0.966	1.133	1.077	0	0.22	0.10	0.23	-0.22	0.05	-0.18	-0.11	-2.56E-01	0.0202
O19069	Succinyl-CoA ligase [ADP:GDP-forming] subunit alpha, mitochondrial OS=Sus scrofa GN=SUCLG1 PE=1 SV=2 - [SUCA_PIG]	0.943	0.881	0.952	1.04	1.059	0.993	1.064	0	0.08	0.18	0.07	-0.06	-0.08	0.01	-0.09	-1.39E-01	0.0194
I3LUN7	Uncharacterized protein OS=Sus scrofa GN=KCNS3 PE=4 SV=1 - [I3LUN7_PIG]	1.109	0.266	0.841	1.042	17.897	15.428	13.982	0	-0.15	1.91	0.25	-0.06	-4.16	-3.95	-3.81	-3.50E+00	0.0184
K7GKE7	Haptoglobin OS=Sus scrofa GN=HP PE=3 SV=1 - [K7GKE7_PIG]	0.415	1.019	0.885	1.859	1.41	1.719	1.461	0	1.27	-0.03	0.18	-0.89	-0.50	-0.78	-0.55	-1.03E+00	0.0184
G9F6X9	Dihydropyrimidinase-like 2 OS=Sus scrofa PE=2 SV=1 - [G9F6X9_PIG]	0.922	0.78	0.921	1.219	1.092	1.086	1.033	0	0.12	0.36	0.12	-0.29	-0.13	-0.12	-0.05	-2.93E-01	0.0177
I6LNT8	Glycogen synthase kinase 3 alpha OS=Sus scrofa GN=GSK3A PE=2 SV=1 - [I6LNT8_PIG]	0.986	1.198	1.062	1.363	1.14	1.384	1.395	0	0.02	-0.26	-0.09	-0.45	-0.19	-0.47	-0.48	-3.14E-01	0.0158
P11708	Malate dehydrogenase, cytoplasmic OS=Sus scrofa GN=MDH1 PE=1 SV=4 - [MDHC_PIG]	1.021	1.03	1.074	0.955	0.932	0.922	0.998	0	-0.03	-0.04	-0.10	0.07	0.10	0.12	0.00	1.16E-01	0.0132
Q29549	Clusterin OS=Sus scrofa GN=CLU PE=1 SV=1 - [CLUS_PIG]	1.089	0.883	0.977	1.119	1.143	1.28	1.195	0	-0.12	0.18	0.03	-0.16	-0.19	-0.36	-0.26	-2.65E-01	0.0128
B3STX9	Prothrombin OS=Sus scrofa PE=2 SV=1 - [B3STX9_PIG]	0.977	1.01	1.002	0.83	0.841	0.868	0.973	0	0.03	-0.01	0.00	0.27	0.25	0.20	0.04	1.87E-01	0.0127
B6V8C8	40S ribosomal protein S3a OS=Sus scrofa GN=RPS3A PE=2 SV=1 - [B6V8C8_PIG]	0.937	1.014	0.905	1.337	1.136	1.188	1.091	0	0.09	-0.02	0.14	-0.42	-0.18	-0.25	-0.13	-2.99E-01	0.0070
Q9N1F5	Glutathione S-transferase omega-1 OS=Sus scrofa GN=GSTO1 PE=1 SV=2 - [GSTO1_PIG]	0.936	0.96	0.949	0.915	0.896	0.905	0.902	0	0.10	0.06	0.08	0.13	0.16	0.14	0.15	8.74E-02	0.0066
I3LD20	Dystroglycan OS=Sus scrofa GN=DAG1 PE=4 SV=1 - [I3LD20_PIG]	0.809	0.848	0.802	1.044	1.143	1.082	1.232	0	0.31	0.24	0.32	-0.06	-0.19	-0.11	-0.30	-3.83E-01	0.0055
Q53DY6	Histone H1.3-like protein OS=Sus scrofa PE=2 SV=1 - [Q53DY6_PIG]	1.049	1.001	1.338	2.12	1.549	3.258	2.403	0	-0.07	0.00	-0.42	-1.08	-0.63	-1.70	-1.26	-1.05E+00	0.0051
P49924	Fatty acid-binding protein, liver OS=Sus scrofa GN=FABP1 PE=2 SV=1 - [FABPL_PIG]	1.045	0.929	0.956	0.492	0.316	0.683	0.507	0	-0.06	0.11	0.06	1.02	1.66	0.55	0.98	1.03E+00	0.0045
F1SH92	Inter-alpha-trypsin inhibitor heavy chain H4 OS=Sus scrofa GN=ITIH4 PE=4 SV=2 - [F1SH92_PIG]	0.863	0.991	0.961	1.372	1.301	1.479	1.157	0	0.21	0.01	0.06	-0.46	-0.38	-0.56	-0.21	-4.73E-01	0.0018
I3LFE7	Uncharacterized protein (Fragment) OS=Sus scrofa GN=MYOZ3 PE=4 SV=1 - [I3LFE7_PIG]	0.901	0.738	1.133	0.218	0.398	0.357	0.415	0	0.15	0.44	-0.18	2.20	1.33	1.49	1.27	1.47E+00	0.0011
P37111	Aminoacylase-1 OS=Sus scrofa GN=ACY1 PE=1 SV=2 - [ACY1_PIG]	0.983	0.965	1.011	0.921	0.896	0.892	0.869	0	0.02	0.05	-0.02	0.12	0.16	0.16	0.20	1.46E-01	0.0007
F1SHR2	Uncharacterized protein OS=Sus scrofa GN=BCL2L13 PE=4 SV=1 - [F1SHR2_PIG]	0.955	0.916	0.938	1.161	1.104	1.143	1.141	0	0.07	0.13	0.09	-0.22	-0.14	-0.19	-0.19	-2.57E-01	0.0002
Q29314	60S ribosomal protein P1 (Fragment) OS=Sus scrofa PE=2 SV=1 - [Q29314_PIG]																	
K9J4W2	NACHT, LRR and PYD domains-containing protein 12-like protein OS=Sus scrofa GN=NLRP12L PE=2 SV=1 - [K9J4W2_PIG]																	
P79309	Tropomyosin TM30-pl (Fragment) OS=Sus scrofa PE=2 SV=1 - [P79309_PIG]																	
Q06AS8	GBAK OS=Sus scrofa GN=GNAI3 PE=2 SV=1 - [Q06AS8_PIG]																	
Q2EN79	Ubiquinol-cytochrome c reductase complex OS=Sus scrofa GN=UQCR10 PE=4 SV=1 - [Q2EN79_PIG]																	
K9IVG2	Solute carrier family 2 (Facilitated glucose/fructose transporter), member 5 OS=Sus scrofa GN=SLC2A5 PE=2 SV=1 - [K9IVG2_PIG]																	
F1SE30	Uncharacterized protein OS=Sus scrofa GN=ANK1 PE=4 SV=2 - [F1SE30_PIG]																	
Q3Y5G4	Nebulin-related anchoring protein OS=Sus scrofa GN=NRAP PE=2 SV=2 - [Q3Y5G4_PIG]																	
F1RIX9	Uncharacterized protein OS=Sus scrofa GN=SH3RF1 PE=4 SV=2 - [F1RIX9_PIG]																	

Note: highlighted proteins with red color in this form were identified proteins in the present experiment without quantitative information.