

SUPPLEMENTAL TABLE 2

			Summary Difference				TMP vs. Vh				Dox/TMP vs. Vh					
			Dox vs. Vh		p-value		TMP vs. Vh		p-value		Dox/TMP vs. Vh		p-value			
ALLC_HUMAN	Average	St. Dev.	q-value	Stor Average	St. Dev.	q-value	Stor Average	St. Dev.	q-value	Stor Average	St. Dev.	q-value	Stor prioritized FI Secretary?			
P13667	PDI4A	Protein disulfide-isomerase A4 OS	-0.036583	0.2312397	0.6901033	0.2300344	1.1729845	0.2177922	7.47E-06	0.0003486	0.7876477	0.241629	0.0073319	0.116089	TRUE	TRUE
P14625	HSP90B1	Endoplasmic reticulum OS	-0.320475	0.2601488	0.0172645	0.0175842	1.0746761	0.2981099	7.581E-05	0.0017689	0.4284759	0.1933722	0.02135	0.0845103	TRUE	TRUE
P11021	HSPA5	78 kDa glucose-regulated protein OS	-0.357474	0.2646001	0.0117219	0.01535	1.351948	0.4356388	0.0001761	0.002739	0.9261142	0.3741535	0.0158181	0.1001812	TRUE	TRUE
Q9Y4L1	HYOU1	Hypoxia up-regulated protein 1 OS	-0.099746	0.2523979	0.336036	0.1257278	0.8593252	0.2969419	0.0002593	0.0032056	0.6620047	0.2383886	0.0115122	0.0911379	TRUE	TRUE
Q9UBS4	DNAJB11	Dnaj homolog subfamily B member 11 OS	-0.251335	0.2546744	0.0400637	0.0262322	0.677796	0.2631247	0.0004894	0.0045678	0.4641012	0.3001376	0.053605	0.1543174	TRUE	TRUE
P27797	CALR	Calreticulin OS	-0.550777	0.2325057	0.0007663	0.0028099	0.5125014	0.3499378	0.0082197	0.063931	-0.068928	0.3000627	0.6771748	0.579564	TRUE	TRUE
Q14257	RCN2	Reticulocalbin-2 OS	-0.426971	0.224564	0.0131422	0.0160627	0.3621158	0.3653084	0.090963	0.2653099	-0.024111	0.6925975	0.9686831	0.6668471	TRUE	TRUE
Q07955	SRSF1	Serine/arginine-rich splicing factor 1 OS	0.0723938	0.318354	0.6015291	0.2042228	0.2792407	0.3446242	0.1039434	0.2694828	0.0483091	0.1237426	0.4919668	0.5024446	TRUE	FALSE
P07237	PAHB	Protein disulfide-isomerase OS	-0.331388	0.3104067	0.0301716	0.0221258	0.2782163	0.3224652	0.0625675	0.2246013	0.1946408	0.2695867	0.2444659	0.386639	TRUE	TRUE
Q02809	PLD1	Phosphatidylcholine-specific phospholipase C1 OS	-0.676972	0.2075997	0.0018804	0.0043093	-0.288562	0.3073412	0.1037186	0.2847178	-0.798538	0.0808886	0.0034028	0.1077567	TRUE	TRUE
Q05707	COL14A1	Collagen alpha-1(XIV) chain OS	-0.557277	0.0123888	0.0001647	0.0015097	0.5747125	0.2726671	0.0675206	0.2250688	0.3576059				TRUE	TRUE
P30101	PDI3A	Protein disulfide-isomerase A3 OS	-0.299934	0.2381446	0.0517631	0.0169995	0.2678003	0.3022794	0.057531	0.2237316	-0.039505	0.2865636	0.8006693	0.6338632	TRUE	TRUE
Q13724	MOGS	Mannosyl-oligosaccharide glucosidase OS	-0.46823	0.2083968	0.194106	0.0751151	-0.447137	0.0548484	0.0550812	0.236777	-0.415095				TRUE	TRUE
Q71U36	TUBA1A	Tubulin alpha-1A chain OS	-0.003483	0.2882285	0.9775341	0.2937944	0.32613	0.3681215	0.082135	0.2555312	0.481697	0.3660861	0.1502996	0.3172991	TRUE	FALSE
Q8NBJ5	GLT25D1	Proteoglycan galactosyltransferase 1 OS	-0.753672	0.2218248	0.000106	0.0019431	-0.201574	0.2893205	0.1148461	0.2820783	-0.617783	0.2575846	0.01724	0.0909889	TRUE	TRUE
Q14697	GANAB	Neutral alpha-glucosidase AB OS	-0.742162	0.2410047	0.0001838	0.0011234	-0.193875	0.2802377	0.1169239	0.2728224	-0.593316	0.2641877	0.0205933	0.0931601	TRUE	TRUE
Q15084	PDI6A	Protein disulfide-isomerase A6 OS	-0.448138	0.2531954	0.0033861	0.0068976	0.3911737	0.3946671	0.0394627	0.1841594	0.068972	0.1604777	0.4532244	0.4784035	TRUE	TRUE
P30040	ERP29	Endoplasmic reticulum resident protein 29 OS	-0.794315	0.2476164	0.0076737	0.0108219	-0.514402	0.2393216	0.023157	0.15438	-0.826351	0.0925908	0.0503341	0.1593913	TRUE	TRUE
Q92896	GLG1	Golgi apparatus protein 1 OS	0.2050994	0.0588388	0.1274122	0.0556164	-0.114662	0.0410513	0.1578493	0.3507763	-0.531325	0.246929	0.2021296	0.3555983	TRUE	TRUE
Q43493	TGOLN2	Trans-Golgi network integral membrane protein OS	-0.236461	0.1884358	0.3266778	0.1247728	-0.510737	0.0442289	0.0389343	0.2018814	-0.344227				TRUE	TRUE
P34931	HSPA1L	Heat shock 70 kDa protein 1-like OS	0.0539744	0.3227324	0.7600343	0.2361688	0.4693401	0.5418938	0.1816587	0.3853366	0.4068648	0.9043029	0.4345231	0.4744793	TRUE	FALSE
Q9BS26	ERP44	Endoplasmic reticulum resident protein 44 OS	-0.075511	0.2106157	0.4200261	0.1509898	0.2758737	0.2358535	0.0351947	0.2053026	0.1766761	0.2318082	0.3176331	0.3868609	TRUE	TRUE
Q969V3	NCLN	Nicalin OS	-0.545481	0.0819438	0.0673714	0.0374286	-0.245894	0.1514497	0.2330805	0.4532122	-0.523398				TRUE	TRUE
P55145	MANF	Mesencephalic astrocyte-derived neurotrophic factor OS	0.1806661	0.066202	0.1614032	0.0657568	0.3749943	0.1978205	0.2272944	0.4611771	0.1702561	0.0902814	0.228376	0.3615953	TRUE	TRUE
Q14737	PDCD5	Programmed cell death protein 5 OS	-0.202758	0.586477	0.7104985	0.0256229	0.1650754	0.1011953	0.2603942	0.4860692	0.1049966	0.0617322	0.2508288	0.3610415	TRUE	FALSE
P13674	HAHA1	Prolyl 4-hydroxylase subunit alpha-1 OS	-0.642668	0.1736381	0.0234794	0.0195661	-0.330993	0.3716614	0.2629011	0.4718738	-0.476562				TRUE	TRUE
Q94905	ERLIN1	Erlin-1 OS	-0.498335	0.1396726	0.0056659	0.0103875	-0.245894	0.4174238	0.3236813	0.5035042	-0.915116				TRUE	TRUE
P57976	CPD	Carboxypeptidase D OS	-0.152682	0.1649275	0.0726545	0.0391765	-0.082315	0.1837276	0.3224685	0.5189148	-0.331232	0.1801077	0.0860279	0.2095552	TRUE	FALSE
P39656	DOST2	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 2 OS	-0.576607	0.3448842	0.0442632	0.0270497	-0.236186	0.3605243	0.2814066	0.4863818	-0.028398	0.0625654	0.9061892	0.6521816	TRUE	FALSE
P53396	ACLY	ATP-citrate lyase OS	0.0988213	0.043924	0.0204948	0.0187869	0.3046931	0.4967037	0.3073776	0.5122959	0.2171114	0.4415964	0.6132124	0.5548113	TRUE	FALSE
Q96AY3	FKBP10	Peptidyl-prolyl cis-trans isomerase FKBP10 OS	-0.565879	0.1558876	0.1224751	0.0561344	-0.536525	0.4800673	0.3591277	0.5237278	-0.894934				TRUE	TRUE
Q8NBJ7	GOLM1	Golgi membrane protein 1 OS	-0.101227	0.306984	0.72221	0.2282848	-0.153611	0.1370854	0.3583712	0.5394835					TRUE	TRUE
P46977	STT3A	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 3 OS	-0.354025	0.676391	0.3720973	0.1364357	-0.236726	0.4592397	0.3784033	0.5351158	0.440611	0.1856924	0.1843815	0.3434557	TRUE	TRUE
Q8WVK2	SNRNP27	U4/U6.U5 small nuclear ribonucleoprotein 27 OS	0.3377782	0.1252063	0.0428761	0.0271056	-0.158499	0.2769899	0.4260979	0.5681305	0.2379211	0.7507738	0.6381739	0.5613567	TRUE	FALSE
Q94905	ERLIN2	Erlin-2 OS	-0.648072	0.0931228	0.0068122	0.0104076	-0.302518	0.5251485	0.4235096	0.5812876	-0.257147	0.5204965	0.6117635	0.5697797	TRUE	TRUE
P27824	CANX	Calnexin OS	-0.306138	0.2758668	0.0260804	0.0199225	0.0713011	0.2607304	0.4965414	0.6436648	-0.250075	0.2475338	0.1369505	0.3089548	TRUE	TRUE
Q07065	CKAP4	Cytoskeleton-associated protein 4 OS	-0.278041	0.3036189	0.1099781	0.0544936	-0.087154	0.26444	0.5020336	0.6331955	-0.299279	0.1537663	0.2218472	0.3697454	TRUE	TRUE
P13162	PRDX4	Peroxisiredoxin-4 OS	-0.19441	0.5008915	0.0505744	0.0299096	-0.19181	0.6100552	0.520788	0.607586	-0.270437	0.7650221	0.6026885	0.5783374	TRUE	FALSE
P04844	RPND4	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 4 OS	-0.509792	0.2265955	0.1938657	0.0772653	-0.149155	0.2227772	0.5173742	0.6190803	-0.279244	0.1068062	0.1681552	0.3328071	TRUE	TRUE
O60568	PLOD3	Procollagen-lysine-2-oxoglutarate 5-dioxygenase OS	-0.52997	0.304802	0.017729	0.0171069	-0.087259	0.2748142	0.1569077	0.6347989	-0.620588	0.0125658	0.0091114	0.0962065	TRUE	TRUE
O43853	CALU	Calumenin OS	-0.166898	0.1684386	0.0910546	0.0463704	-0.094188	0.3189333	0.5451128	0.6204536	0.0510878	0.720803	0.9364014	0.6589491	TRUE	TRUE
Q96H7E	ERO1L	ERO1-like protein alpha OS	-0.431444	0.2691141	0.0230672	0.020138	-0.06635	0.2848699	0.583172	0.6479689	-0.365525	0.2642975	0.3008877	0.3811244	TRUE	TRUE
P04843	RPN1	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 1 OS	-0.25574	0.3294794	0.0858109	0.0449486	0.0727363	0.3462345	0.5984363	0.6494657	-0.247844	0.36298	0.2654388	0.3654592	TRUE	TRUE
P49257	LMAN1	Protein ERGIC-53 OS	-0.150026	0.2309473	0.1364683	0.0581842	-0.048891	0.278008	0.6575535	0.6974052	-0.110647	0.2339442	0.413972	0.4681827	TRUE	TRUE
Q08380	LGALS3BP	Galectin-3-binding protein OS	-0.370781	0.2433778	0.0135465	0.0155221	-0.096558	0.5368334	0.6779148	0.7030227	-0.218676	0.4632811	0.4995133	0.49431	TRUE	FALSE
Q8NBS9	TXNDC5	Thioredoxin domain-containing protein 5 OS	-0.11925	0.1666708	0.1400597	0.0583582	0.0594168	0.3870207	0.7223043	0.6879089	0.0901041	0.5798955	0.813045	0.6279616	TRUE	TRUE
P68371	TUBB2C	Tubulin beta-2C chain OS	-0.10774	0.2510816	0.5347906	0.1849905	0.0655654	0.2672895	0.7122773	0.7228002	0.0608299				TRUE	FALSE
P02786	TFRC	Transferrin receptor protein 1 OS	-0.051532	0.2855306	0.7071781	0.2315186	0.0648822	0.3801527	0.7221367	0.7020773	0.0548422	0.3622192	0.8176743	0.6165004	TRUE	FALSE
P23284	PF1B	Peptidyl-prolyl cis-trans isomerase B OS	-0.435472	0.1137261	0.0002322	0.0010644	0.0449634	0.2866445	0.7166006	0.7115237	-0.277035	0.1843749	0.057434	0.151562	TRUE	TRUE
Q9P035	PTPLAD1	3-hydroxyacyl-CoA dehydratase 3 OS	-0.677797	0.2378474	0.0386806	0.0262646	0.1195915	0.5338488	0.7354134	0.6863859	0.2132603				TRUE	TRUE
P50454	SERPINH1	Serpin H1 OS	-0.607615	0.2403058	0.0016009	0.0041928	-0.03693	0.3074625	0.78042	0.7141098	-0.268208	0.3116714	0.2745702	0.3622802	TRUE	TRUE
Q16706	MAN2A1	Alpha-mannosidase 2 OS	-0.014725	0.3966609	0.9666081	0.2953525	0.0249355	0.1078348	0.7987916	0.7168643	-0.576281				TRUE	TRUE
P14314	PRKCSH	Glucosylase 2 subunit beta OS	-0.533779	0.2271609	0.0007999	0.0024441	0.0204245	0.2698741	0.8478001	0.7326668	-0.229513	0.4532787	0.3857931	0.4524734	TRUE	TRUE
Q14204	DYNLC1H1	Cytosolic dynein 1 heavy chain 1 OS	-0.089881	0.1033445	0.4377161	0.154323	0.0202204	0.1164393	0.8466884	0.7455118					TRUE	FALSE
Q13217	DNAJC3	Dnaj homolog subfamily C member 3 OS	-0.159162	0.102165	0.1142752	0.0551328	0.0293891	0.2737593	0.8696408	0.737877	0.0193887				TRUE	TRUE
Q9NVU2	UGT1T1	UDP-glucosyltransferase 1 OS	-0.303694	0.2351689	0.0250032	0.0199301	0.0191161	0.2875121	0.8770052	0.7308377	-0.358412	0.1238417	0.0375683	0.1321848	TRUE	TRUE
Q9H3N1	TMX1	Thioredoxin-related transmembrane protein 1 OS	-0.463617	0.3047219	0.1188667	0.0558775	0.0195013	0.2737573	0.9130846	0.7475546	-0.120576	0.4038269	0.7456401	0.6213667	TRUE	TRUE
Q9Y2B0	CNPY2	Protein domain homolog 2 OS	-0.510078	0.1416489	0.1234385	0.0551961	-0.075046	0.822410								