

Supplemental Table S7

Site Table lncRNA Granulosa-Vinclozolin 1e-04

test_id	Chr	Start	mRNA_size	ORF_size	log2.fold_change.	p_value	q_value	Gene	Category
TCONS_00000352	1	24745599	4555	279	1.372	5.00E-05	1.05E-02	-	
TCONS_00004728	1	240939992	7401	276	1.386	5.00E-05	1.05E-02	-	
TCONS_00004962	1	257019716	848	171	2.829	5.00E-05	1.05E-02	-	
TCONS_00010933	1	1825190	614	123	Inf	5.00E-05	1.05E-02	-	
TCONS_00012305	1	48310460	821	126	1.184	5.00E-05	1.05E-02	-	
TCONS_00012351	1	50393212	10693	303	1.49	5.00E-05	1.05E-02	-	
TCONS_00013750	1	115080716	2830	222	2.085	5.00E-05	1.05E-02	-	
TCONS_00015105	1	174762178	670	147	Inf	5.00E-05	1.05E-02	-	
TCONS_00015170	1	176009693	2483	180	2.9	5.00E-05	1.05E-02	-	
TCONS_00015463	1	193045918	545	75	Inf	5.00E-05	1.05E-02	-	
TCONS_00016233	1	222293495	1241	135	2.237	5.00E-05	1.05E-02	-	
TCONS_00016335	1	226615363	3253	216	1.49	5.00E-05	1.05E-02	-	
TCONS_00017160	1	257056364	612	108	Inf	5.00E-05	1.05E-02	-	
TCONS_00021691	10	39116799	5574	309	1.312	5.00E-05	1.05E-02	-	
TCONS_00024792	10	28935709	958	216	Inf	5.00E-05	1.05E-02	-	
TCONS_00025407	10	47063559	2117	222	1.775	5.00E-05	1.05E-02	-	
TCONS_00026194	10	74070666	3910	258	1.243	5.00E-05	1.05E-02	-	
TCONS_00029682	11	83690227	4619	417	-1.146	5.00E-05	1.05E-02	-	
TCONS_00030978	11	53621867	2754	180	-1.009	5.00E-05	1.05E-02	-	
TCONS_00034580	12	47214615	4246	345	1.917	5.00E-05	1.05E-02	-	
TCONS_00039644	13	48937120	715	186	Inf	5.00E-05	1.05E-02	-	
TCONS_00043160	14	114756254	2817	216	1.463	5.00E-05	1.05E-02	U6	
TCONS_00043568	14	20102593	4081	255	-1.279	5.00E-05	1.05E-02	-	
TCONS_00043580	14	20953245	1999	282	2.13	5.00E-05	1.05E-02	-	
TCONS_00044783	14	3768966	2041	420	2.44	5.00E-05	1.05E-02	-	
TCONS_00046458	14	80901648	907	123	Inf	5.00E-05	1.05E-02	-	
TCONS_00046541	14	83497468	837	186	Inf	5.00E-05	1.05E-02	-	
TCONS_00047279	14	115450565	630	123		5.00E-05	1.05E-02	-	
TCONS_00048337	15	58774172	9067	405	0.999	5.00E-05	1.05E-02	-	
TCONS_00051870	15	59672522	946	78	Inf	5.00E-05	1.05E-02	-	
TCONS_00053583	16	27233712	1851	342	2.48	5.00E-05	1.05E-02	-	
TCONS_00056199	16	32563043	1269	141	1.895	5.00E-05	1.05E-02	-	
TCONS_00056536	16	49848305	683	60	Inf	5.00E-05	1.05E-02	-	
TCONS_00058299	17	43410024	7859	381	1.232	5.00E-05	1.05E-02	-	
TCONS_00058580	17	57207235	1726	144	-0.815	5.00E-05	1.05E-02	-	
TCONS_00059501	17	16726113	3190	264	1.223	5.00E-05	1.05E-02	-	
TCONS_00062125	17	43943905	2125	192	1.903	5.00E-05	1.05E-02	-	
TCONS_00065566	18	27651600	5598	327	1.063	5.00E-05	1.05E-02	-	
TCONS_00067146	18	31500550	1587	132	1.833	5.00E-05	1.05E-02	-	
TCONS_00068805	19	11037180	5219	222	1.662	5.00E-05	1.05E-02	-	
TCONS_00070384	19	25770960	1473	303	1.382	5.00E-05	1.05E-02	AC120246.2	
TCONS_00070642	19	37176721	3033	294	1.528	5.00E-05	1.05E-02	-	
TCONS_00071728	19	26011317	707	180	Inf	5.00E-05	1.05E-02	-	
TCONS_00072575	19	56354147	652	48	Inf	5.00E-05	1.05E-02	-	
TCONS_00078405	2	211761933	4309	135	-0.873	5.00E-05	1.05E-02	-	
TCONS_00078496	2	225503345	6959	192	1.123	5.00E-05	1.05E-02	-	
TCONS_00080110	2	47396892	554	24	Inf	5.00E-05	1.05E-02	-	
TCONS_00081459	2	118780886	4393	216	1.462	5.00E-05	1.05E-02	-	

TCONS_00081521	2	121969115	895	171	Inf		5.00E-05	1.05E-02	-	
TCONS_00081554	2	123197337	5380	402		2.257	5.00E-05	1.05E-02	-	
TCONS_00081622	2	126767447	640	99	Inf		5.00E-05	1.05E-02	-	
TCONS_00082894	2	193594576	921	168			5.00E-05	1.05E-02	-	
TCONS_00083124	2	198611760	1137	168	Inf		5.00E-05	1.05E-02	-	
TCONS_00086902	20	17024092	1382	210		-1.133	5.00E-05	1.05E-02	-	
TCONS_00086997	20	24494382	1621	159		-1.91	5.00E-05	1.05E-02	-	
TCONS_00087422	20	1254514	1950	315		-1.123	5.00E-05	1.05E-02	-	
TCONS_00093440	3	74226406	1856	360		-3.325	5.00E-05	1.05E-02	AABR07052758.1	
TCONS_00095063	3	164366919	5222	309		1.206	5.00E-05	1.05E-02	-	
TCONS_00095808	3	20242429	778	192			5.00E-05	1.05E-02	-	
TCONS_00096591	3	45498519	628	48	Inf		5.00E-05	1.05E-02	-	
TCONS_00097141	3	66641751	855	117	Inf		5.00E-05	1.05E-02	-	
TCONS_00097336	3	80015344	1107	285	Inf		5.00E-05	1.05E-02	-	
TCONS_00097380	3	82369699	711	63	Inf		5.00E-05	1.05E-02	-	
TCONS_00097707	3	101374114	5869	447		2.53	5.00E-05	1.05E-02	-	
TCONS_00097937	3	111301230	1723	123		2.433	5.00E-05	1.05E-02	-	
TCONS_00098252	3	124046187	686	123	Inf		5.00E-05	1.05E-02	-	
TCONS_00098426	3	127684176	562	96			5.00E-05	1.05E-02	-	
TCONS_00099934	3	164494270	2434	219		1.67	5.00E-05	1.05E-02	-	
TCONS_00101374	4	78404074	5115	234		1.62	5.00E-05	1.05E-02	-	
TCONS_00103456	4	45561146	5859	282		1.59	5.00E-05	1.05E-02	-	
TCONS_00104453	4	119783973	2722	159		2.075	5.00E-05	1.05E-02	-	
TCONS_00104569	4	127330752	3631	381		-1.553	5.00E-05	1.05E-02	-	
TCONS_00105621	4	7114007	768	210		2.413	5.00E-05	1.05E-02	-	
TCONS_00106507	4	51726363	1304	186		1.369	5.00E-05	1.05E-02	-	
TCONS_00107711	4	99132048	788	87	Inf		5.00E-05	1.05E-02	-	
TCONS_00108035	4	118384253	1587	252		2.921	5.00E-05	1.05E-02	-	
TCONS_00109077	4	152580822	656	48	Inf		5.00E-05	1.05E-02	-	
TCONS_00109555	4	170339540	4275	258		-2.045	5.00E-05	1.05E-02	-	
TCONS_00111747	5	150453439	2380	315		1.691	5.00E-05	1.05E-02	U11	
TCONS_00112828	5	34123313	11222	285		1.655	5.00E-05	1.05E-02	-	
TCONS_00115858	5	29615530	6451	174		1.547	5.00E-05	1.05E-02	-	
TCONS_00117565	5	120817142	4805	480		2.286	5.00E-05	1.05E-02	-	
TCONS_00119103	5	170877110	692	189			5.00E-05	1.05E-02	-	
TCONS_00121198	6	134627277	1861	510		-1.442	5.00E-05	1.05E-02	AABR07065559.1	
TCONS_00121470	6	2184106	781	135	Inf		5.00E-05	1.05E-02	-	
TCONS_00121472	6	2193035	7745	243		2.228	5.00E-05	1.05E-02	-	
TCONS_00121795	6	26497445	1208	369		1.9	5.00E-05	1.05E-02	-	
TCONS_00122293	6	72608369	5247	201		2.325	5.00E-05	1.05E-02	-	
TCONS_00122969	6	124746082	6034	219		2.053	5.00E-05	1.05E-02	-	
TCONS_00123445	6	2158215	664	105	Inf		5.00E-05	1.05E-02	-	
TCONS_00124674	6	40994355	719	114	Inf		5.00E-05	1.05E-02	-	
TCONS_00124886	6	44441812	688	102	Inf		5.00E-05	1.05E-02	-	
TCONS_00125842	6	93365463	3915	291		2.384	5.00E-05	1.05E-02	-	
TCONS_00125904	6	94763839	3955	204		2.3	5.00E-05	1.05E-02	-	
TCONS_00126115	6	101277521	3060	222		-1.482	5.00E-05	1.05E-02	-	
TCONS_00128996	7	117191898	3221	570		1.788	5.00E-05	1.05E-02	-	
TCONS_00133530	7	33585150	709	138	Inf		5.00E-05	1.05E-02	-	
TCONS_00134610	7	73993687	570	45	Inf		5.00E-05	1.05E-02	-	
TCONS_00135857	7	115484611	677	120			5.00E-05	1.05E-02	-	

TCONS_00136210	7	124940735	2696	393	2.329	5.00E-05	1.05E-02	-	
TCONS_00136450	7	135850715	3427	168	1.788	5.00E-05	1.05E-02	-	
TCONS_00136486	7	137875401	1127	144	2.46	5.00E-05	1.05E-02	-	
TCONS_00136499	7	137944316	700	81	Inf	5.00E-05	1.05E-02	-	
TCONS_00136882	8	6493323	7903	549	-4.13	5.00E-05	1.05E-02	-	
TCONS_00136883	8	6562365	6538	471	-2.607	5.00E-05	1.05E-02	-	
TCONS_00138634	8	114943678	4023	492	2.067	5.00E-05	1.05E-02	-	
TCONS_00138795	8	117564649	5118	303	1.267	5.00E-05	1.05E-02	-	
TCONS_00139049	8	128555940	2276	297	-1.741	5.00E-05	1.05E-02	-	
TCONS_00141081	8	107614723	11083	333	-0.932	5.00E-05	1.05E-02	-	
TCONS_00141552	8	129450300	2511	213	-2.366	5.00E-05	1.05E-02	-	
TCONS_00141692	8	5214271	672	147		5.00E-05	1.05E-02	-	
TCONS_00141752	8	6476872	2063	219	-3.134	5.00E-05	1.05E-02	-	
TCONS_00143956	8	83936931	765	207	Inf	5.00E-05	1.05E-02	-	
TCONS_00144419	8	102004542	637	57	Inf	5.00E-05	1.05E-02	-	
TCONS_00144853	8	118708715	756	135	Inf	5.00E-05	1.05E-02	-	
TCONS_00145047	8	127681266	647	84	Inf	5.00E-05	1.05E-02	-	
TCONS_00145783	9	42396618	2354	351	1.768	5.00E-05	1.05E-02	-	
TCONS_00149530	9	19988334	4997	213	1.743	5.00E-05	1.05E-02	-	
TCONS_00150389	9	60430237	3785	258	1.888	5.00E-05	1.05E-02	-	
TCONS_00150877	9	80904959	622	66		5.00E-05	1.05E-02	-	
TCONS_00154848	X	33583322	1406	231	-0.854	5.00E-05	1.05E-02	-	
TCONS_00159499	X	139003967	875	102	Inf	5.00E-05	1.05E-02	-	
TCONS_00159615	X	146006894	1117	156	2.753	5.00E-05	1.05E-02	-	