D. melanogaster **Northern analysis**

cDNA ID	cDNA length, bp	Lengths of transcripts detected on Northern blots, kb					
		Schneider S2 cells	0-2h embryo	0-24h embryo	Late third instar	Pupa	Adult
AT08427	1808	Negative	Negative	Negative	Negative	Negative	Negative
AT11392	994	Negative	Negative	Negative	Negative	Negative	Negative
AT13310	606	Negative	Negative	Negative	Negative	Negative	Negative
AT14266	781	Negative	Negative	Negative	Negative	Negative	Negative
AT21289	1415	Negative	Negative	Negative	Negative	Negative	Negative
AT22150	1080	Negative	Negative	Negative	Negative	Negative	Negative
AT24650	578	Negative	Negative	Negative	Negative	Negative	0.6
AT27057	840	1.7	1.7	1.7	1.7	1.7	1.7
AT28783	1553	Negative	Negative	Negative	Negative	Negative	Negative
AT31442	1111	Negative	Negative	Negative	Negative	Negative	Negative
GH03576	931	Negative	1	1	1	1	1
GH05108	1094	1.8/2.8	Negative	1.8/2.8	1.8/2.8	1.8/2.8	1.8/2.8
GH05710	1211	Negative	Negative	Negative	Negative	Negative	Negative
GH06385	1190	Negative	Negative	Negative	1.2	1.2	1.2
GH08923	2462	6	5	6	6	6	5.0/6.0
GH14469	695	0.7	0.7/1.2	0.7/1.2	0.7	0.7	0.7/1.2
GH22170	743	Negative	Negative	Negative	Negative	Negative	Negative
GH25188	827	Negative	Negative	Negative	Negative	Negative	Negative
GH25505	2811	Negative	Negative	10	Negative	10	10
GH26692	1674	5.0/6.0	5	5	5	5	5
GH26828	2335	Negative	Negative	Negative	Negative	Negative	Negative
GH27201	4554	Negative	Negative	Negative	Negative	Negative	Negative
GM01028	1157	1.2/2.3/4.0	0.7/1.2	1.2/2.3/4.0	1.2/2.3/4.0	1.2/2.3/4.0	0.7/1.2/2.3/4.0
GM01206	1214	~9.0	6	~9.0	~9.0	~9.0	~9.0
GM02923	1035	9	9	9	9	9	9
GM03003	1884	~9.0	6	~9.0	~9.0	~9.0	~9.0
GM03661	1159	10	10	10	10	10	10
GM03914	1046	5	5	5	5	5	5
GM04921	1159	Negative	Negative	Negative	Negative	Negative	Negative
GM07040	3544	Negative	Negative	Negative	Negative	Negative	Negative
GM07077	1385	4	4	4.0/9.0 dbl	4.0/9.0 dbl	4.0/9.0 dbl	4.0/9.0 dbl
GM07658	2095	6	6	6	6	6	6
GM07660	853	4	4	4	4	4	4
GM07702	850	Negative	Negative	Negative	Negative	Negative	Negative
GM09444	653	10	10	10	10	10	10
GM09534	912	Negative	Negative	8	8	8	8
GM09668	834	3.8	3.3	3.8	3.8	3.8	3.8
GM10787	1695	1.5 to >10	1.5 to >10	1.5 to >10	1.5 to >10	1.5 to >10	1.5 to >10
GM12657	708	Negative	Negative	Negative	Negative	Negative	Negative
GM24362	450	8	8	8	6.0/8.0	8	8
HL05775	1890	Negative	~7.0 dblt.	~7.0 dblt.	~7.0 dblt./9.0	~7.0 dblt./9.0	6
LD11130	1187	6	5	6	6	6	5.0/6.0
LD11162	1549	Negative	1.5	1.5	1.5	1.5	1.5
LD13184	2222	Negative	Negative	Negative	2.2	2.2	2.2
LD16711	1672	Negative	Negative	Negative	Negative	Negative	Negative
LD23922	1721	Negative	Negative	Negative	Negative	Negative	Negative
LD28571	3302	Negative	Negative	Negative	Negative	Negative	Negative
LD30889	1069	Negative	Negative	Negative	Negative	Negative	Negative
LD38553	1317	Negative	Negative	Negative	Negative	Negative	Negative
LP01161	527	Negative	Negative	Negative	Negative	Negative	Negative
LP03188	553	Negative	Negative	Negative	0.6	0.6	Negative
LP06491	403	Negative	Negative	Negative	0.6	0.6	0.6
LP11739	1222	Negative	Negative	8	8	8	8
LP12023	313	Negative	Negative	Negative	0.3	0.3	Negative
RE21847	3191	5	Negative	5	Negative	Negative	Negative
RE28911	1005	Negative	Negative	1	1	1	1
RE45760	758	1.4	1.4	1.4	1.4	1.4	1.4
RE54004	562	Negative	Negative	0.6	0.6	0.6	0.6
RE63504	331	0.3	0.3	0.3	0.3	0.3	0.3
RE65113	1062	4	4	4	4	4	4
RE66017	3461	Negative	Negative	Negative	Negative	Negative	Negative
RH09485	518	Inconclusive	0.5	0.5	0.5	0.5	0.5
RH45340	407	0.4	0.4	0.4	0.4	0.4	0.4
RH57193	981	1	1	1	1	1	1
RH62702	2049	Negative	Inconclusive	>10.0	>10.0	>10.0	>10.0
RH62830	326	Negative	Negative	Negative	0.3	0.3	0.3
RH63361	424	Negative	Negative	Negative	Negative	0.4	0.4
SD03144	3956	Negative	Negative	Negative	Negative	Negative	Negative
SD04448	2301	Negative	6.5	6.5	6.5	6.5	6.5
SD05379	1391	4.5	4.5	4.5	4.5	4.5	4.5
SD10988	1415	1.4	Negative	Negative	Negative	Negative	Negative
SD25037	1560	Negative	Negative	Negative	Negative	Negative	Negative