

Table S3 Overlap of transcripts regulated by HP1, H1 and CHD1.

SYMBOL/ TRANSCRIPT	FLYBASE ID	FOLD CHANGE		
		HP1 KD	H1 KD	<i>Chd1</i>
HP1 depletion effect: UP				
<i>CR31451</i>	FBgn0051451	181.09	8.26	
<i>CG31140</i>	FBgn0051140	79.80	6.31	
<i>G6</i>	FBte0000958	67.66		
<i>CG31642</i>	FBgn0051642	51.14		0.435
<i>AT07338</i>	FBcl0481740	50.00		
<i>TART</i>		35.15		
<i>CG3635</i>	FBgn0032981	34.71	6.77	
<i>CG18605</i>	FBgn0034411	24.98		
<i>Def</i>	FBgn0010385	22.33		
<i>CG10126</i>	FBgn0038088	21.46		
<i>springer</i>	FBte0000333	20.62		
<i>CG17124</i>	FBgn0032297	20.00	2.46	
<i>gypsy6</i>	FBte0001175	19.16		
<i>CG11714</i>	FBgn0036170	19.04	3.88	
<i>CG40295</i>	FBgn0058295	18.09	11.97	
<i>HDC20537</i>		16.14	2.13	
<i>CG5773</i>	FBgn0034290	15.22		
<i>CG13795</i>	FBgn0031937	13.84		
<i>CG5171</i>	FBgn0031907	12.85		0.149
<i>accord</i>	FBte0000956	11.76		
<i>CG5399</i>	FBgn0038353	11.64	28.01	2.06
<i>CG32091</i>	FBgn0052091	10.80		
<i>edin</i>	FBgn0052185	10.48		
<i>CG32032</i>	FBgn0043806	10.10		
<i>gypsy2</i>	FBte0001040	9.84		
<i>tok</i>	FBgn0004885	9.72	2.11	
<i>CG9691</i>	FBgn0030160	9.48		
<i>Yp3</i>	FBgn0004047	9.00		0.180
<i>invader3</i>	FBte0000619	8.99		
<i>CG13315</i>	FBgn0040827	8.90		
<i>CG3961</i>	FBgn0036821	8.86		
<i>CAH1</i>	FBgn0027844	8.65	2.25	0.493
<i>AttC</i>	FBgn0041579	8.50		
<i>gypsy</i>	FBte0000021	8.45		
<i>Cyp6a13</i>	FBgn0033304	8.40	5.64	
<i>Drs</i>	FBgn0010381	8.33	4.28	0.451
<i>ldgf1</i>	FBgn0020416	8.29		
<i>Cyp12e1</i>	FBgn0037817	8.15		
<i>HMS-Beagle</i>	FBte0000726	7.86		
<i>CG33775</i>	FBgn0053775	7.68		
<i>KCNQ</i>	FBgn0033494	7.66		
<i>Gpb5</i>	FBgn0063368	7.47		
<i>Gr94a</i>	FBgn0041225	7.33	5.19	
<i>CG33468</i>	FBgn0053468	7.30	4.91	2.46
<i>CG4752</i>	FBgn0034733	7.25		
<i>CG9981</i>	FBgn0030746	7.06		2.77
<i>aret</i>	FBgn0000114	6.89		

CG45057	FBgn0266417	6.74	2.20	
<i>Cyp6d4</i>	FBgn0039006	6.66		
<i>CG40115</i>	FBgn0058115	6.47	10.26	
<i>HDC20531</i>		6.34		
<i>Sid</i>	FBgn0039593	6.31		0.261
<i>CT39784</i>		6.11	14.84	
<i>HDC20523</i>		6.01	2.81	
<i>CG11672</i>	FBgn0037563	5.98		
<i>Tsf3</i>	FBgn0034094	5.89		
<i>Rh4</i>	FBgn0003250	5.82		
<i>CG10877</i>	FBgn0038804	5.73	0.157	
<i>CG9717</i>	FBgn0039789	5.71		
<i>CG18473</i>	FBgn0037683	5.69	3.14	
<i>bru-3</i>	FBgn0264001	5.67		
<i>rau</i>	FBgn0031745	5.63		
<i>CG15661</i>	FBgn0034605	5.55	12.72	
<i>Gbp</i>	FBgn0034199	5.52		0.429
<i>CG3940</i>	FBgn0037788	5.43		0.356
<i>CG10799</i>	FBgn0033821	5.41	3.26	
<i>PGRP-SD</i>	FBgn0035806	5.40		
<i>PGRP-SC2</i>	FBgn0043575	5.39	2.66	0.222
<i>Cyp12e1</i>	FBgn0037817	5.34		
<i>CG14164</i>	FBgn0036057	5.31		
<i>AttA</i>	FBgn0012042	5.30		0.337
<i>CG4928</i>	FBgn0027556	5.30	0.315	
<i>CG14695</i>	FBgn0037850	5.29		
<i>Zasp52</i>	FBgn0265991	5.26	0.492	
<i>CG5367</i>	FBgn0032228	5.24		
<i>CG18446</i>	FBgn0033458	5.20	2.05	0.395
<i>Mct1</i>	FBgn0023549	5.19	0.395	
<i>invader4</i>	FBte0000292	5.17		
<i>Pka-C3</i>	FBgn0000489	5.16		
<i>CG17124</i>	FBgn0032297	5.14	4.46	
<i>CG9919</i>	FBgn0030742	5.13		
<i>SPH93</i>	FBgn0032638	5.12	11.67	
<i>Corp</i>	FBgn0030028	5.08	4.57	
<i>Apoltp</i>	FBgn0032136	5.06	2.89	5.70
<i>spir</i>	FBgn0003475	5.06		
<i>CG8586</i>	FBgn0033320	5.03		0.438
<i>CG11052</i>	FBgn0040524	5.03		
<i>yin</i>	FBgn0265575	5.00	0.449	0.398
<i>CG43110</i>	FBgn0262570	4.99	0.432	
<i>Cyp9c1</i>	FBgn0015040	4.93		3.41
<i>CG5381</i>	FBgn0032218	4.85		
<i>CG32687</i>	FBgn0052687	4.82	2.09	
<i>Cad96Ca</i>	FBgn0022800	4.81		
<i>Tsp2A</i>	FBgn0024361	4.77		
<i>Cyp28d1</i>	FBgn0031689	4.76		
<i>Mctp</i>	FBgn0034389	4.72	0.300	2.95
<i>Ama</i>	FBgn0000071	4.72	2.27	0.371
<i>Jheh3</i>	FBgn0034406	4.69		2.68
<i>alphaTub67C</i>	FBgn0087040	4.63		

CG5731	FBgn0032192	4.60	2.72	
GstE3	FBgn0063497	4.59		0.432
CG40124	FBgn0058124	4.58	11.10	
CG5853	FBgn0032167	4.57		
CT39116		4.56	103.1	
CG14606	FBgn0037485	4.54		
shf	FBgn0003390	4.52	2.97	
CG1294	FBgn0033030	4.49	4.55	
ItgalphaPS4	FBgn0034005	4.49	3.91	
PGRP-SA	FBgn0030310	4.46	2.20	
Cyp4p3	FBgn0033397	4.43	9.73	8.75
CG6709	FBgn0036056	4.43	2.66	
HDC20116		4.41		
Cyp12a5	FBgn0038680	4.30		
CG12698	FBgn0030721	4.26		
CG9170	FBgn0030716	4.26	3.83	
qbert		4.26		
RpS5b	FBgn0038277	4.25		
AttB	FBgn0041581	4.22		
Hf	FBgn0014000	4.22		
l(3)neo38	FBgn0265276	4.17		
HDC20280		4.17		
CG6330	FBgn0039464	4.15		
beat-IIIc	FBgn0032629	4.13	2.39	
alphaTub85E	FBgn0003886	4.12		
Swim	FBgn0034709	4.11	9.06	
Mal-A5	FBgn0050359	4.11		
GstE11	FBgn0034354	4.06	0.341	
CG2930	FBgn0028491	4.06		
Eip78C	FBgn0004865	4.04	0.136	0.144
Epac	FBgn0085421	4.01	9.43	
CG3246	FBgn0031538	4.00		
TpnC4	FBgn0033027	4.00	3.70	
TART		3.97		
Myo28B1	FBgn0040299	3.95	2.21	
CG14451	FBgn0037183	3.94		
kek1	FBgn0015399	3.90	5.43	
Esyt2	FBgn0266758	3.88		
CG13117	FBgn0032140	3.85	2.71	
CG40274	FBgn0058274	3.85	5.22	
micropia	FBte0000042	3.83		
CG5322	FBgn0032253	3.80		
CG11400	FBgn0034198	3.78	19.24	
CG16947	FBgn0031816	3.78	0.285	
GATE	FBte0000359	3.78		
NT5E-2	FBgn0050104	3.76	0.196	
CG9150	FBgn0031775	3.72		
spir	FBgn0003475	3.67		
lr41a	FBgn0040849	3.67		
Tsp42Ed	FBgn0029507	3.66		
l(3)neo38	FBgn0265276	3.65		
l(3)neo38	FBgn0265276	3.65		

CG12868	FBgn0033945	3.64	2.74	
CG17032	FBgn0036547	3.63		
CG31272	FBgn0051272	3.62		
Hsp68	FBgn0001230	3.60		
loh	FBgn0032252	3.59	2.15	
Tl	FBgn0262473	3.58	2.30	
sdk	FBgn0021764	3.58		
CG4301	FBgn0030747	3.58		
aret	FBgn0000114	3.58		
rdgB	FBgn0003218	3.58		
bgm	FBgn0027348	3.58	4.62	
vir-1	FBgn0043841	3.56	8.76	
CG43658	FBgn0263706	3.53		
LysX	FBgn0004431	3.51	33.21	0.344
eIF4E-6	FBgn0039622	3.50		
fend	FBgn0030090	3.47	3.52	
CG3679	FBgn0027521	3.43	4.21	
wrapper	FBgn0025878	3.42		
Glut1	FBgn0264574	3.36	0.312	0.395
CG34232	FBgn0085261	3.35		
CG12926	FBgn0033437	3.34	0.415	
HDC20112		3.32		
HMS-Beagle	FBte0000726	3.32		
Pepck	FBgn0003067	3.29		2.30
Mtk	FBgn0014865	3.29		0.438
Listericin	FBgn0033593	3.28		
Ugt35a	FBgn0026315	3.27	3.35	2.53
Ndae1	FBgn0259111	3.26		2.10
rdgB	FBgn0003218	3.26		
CG4269	FBgn0034741	3.25		
sda	FBgn0015541	3.24		
CG17207	FBgn0038051	3.24		
mthl14	FBgn0052476	3.23	15.70	
Spn31A	FBgn0032178	3.22		
CG5096	FBgn0032235	3.19	3.96	
nerfin-2	FBgn0041105	3.18		
CG10405	FBgn0038431	3.16		
hiw	FBgn0030600	3.15		
Arc1	FBgn0033926	3.15		
CG6357	FBgn0033875	3.14	10.96	
Nep2	FBgn0027570	3.13		0.283
d	FBgn0262029	3.13		
AttD	FBgn0038530	3.13	72.76	
Fie	FBgn0026592	3.12	2.46	
CG40040	FBgn0058040	3.10		
l(3)neo38	FBgn0265276	3.10		
CG32625	FBgn0052625	3.09	39.17	
CG9360	FBgn0030332	3.06	10.35	
CG3568	FBgn0029710	3.03		
CG31626	FBgn0051626	3.00	19.15	
CG32313	FBgn0052313	3.00		
spir	FBgn0003475	3.00	2.15	

<i>tyn</i>	FBgn0029128	3.00		
<i>hdm</i>	FBgn0029977	2.98		
<i>CG42822</i>	FBgn0262004	2.98		
<i>CG32944</i>	FBgn0052944	2.97		
<i>CG33346</i>	FBgn0053346	2.97		
<i>alpha-Est3</i>	FBgn0015571	2.97		
<i>CG5191</i>	FBgn0038803	2.97	0.403	
<i>CG4297</i>	FBgn0031258	2.97	0.429	
<i>dys</i>	FBgn0039411	2.96		
<i>Ef1alpha100E</i>	FBgn0000557	2.95		
<i>QC</i>	FBgn0052412	2.95		
<i>stv</i>	FBgn0086708	2.94		2.53
<i>ZnT77C</i>	FBgn0037000	2.92		
<i>CG8738</i>	FBgn0033321	2.91		
<i>CG3251</i>	FBgn0031622	2.90		
<i>d</i>	FBgn0262029	2.90		
<i>ldgf2</i>	FBgn0020415	2.90	2.56	
<i>CG33494</i>	FBgn0053494	2.89		
<i>CG16749</i>	FBgn0037678	2.88		
<i>HDC20377</i>		2.86		
<i>CG33144</i>	FBgn0053144	2.86		
<i>Mdr49</i>	FBgn0004512	2.86	3.38	
<i>Cyp6a8</i>	FBgn0013772	2.83		
<i>Cyp6a2</i>	FBgn0000473	2.80		30.5
<i>RapGAP1</i>	FBgn0264895	2.80	0.339	
<i>CG6495</i>	FBgn0027550	2.80		
<i>CG32982</i>	FBgn0052982	2.79	2.33	
<i>Cyp4p1</i>	FBgn0015037	2.79		
<i>GstD5</i>	FBgn0010041	2.78	20.38	
<i>Het-A</i>		2.75		
<i>Kaz1-ORFB</i>	FBgn0063923	2.75	4.27	
<i>CG30456</i>	FBgn0050456	2.74		
<i>wisp</i>	FBgn0260780	2.74		
<i>CG8349</i>	FBgn0032003	2.73		
<i>CHKov2</i>	FBgn0039328	2.72		
<i>CG8046</i>	FBgn0033388	2.71		
<i>R1</i>		2.71		
<i>Rbf</i>	FBgn0015799	2.71		
<i>CG4741</i>	FBgn0035040	2.70		
<i>CG2217</i>	FBgn0027544	2.69	0.431	
<i>Ugt86Da</i>	FBgn0040259	2.68	8.97	2.28
<i>btn</i>	FBgn0014949	2.68		
<i>Jhl-26</i>	FBgn0028424	2.68	7.00	3.84
<i>AOX1</i>	FBgn0267408	2.67	14.75	2.60
<i>stai</i>	FBgn0266521	2.67		
<i>lr41a</i>	FBgn0040849	2.66		
<i>Ugt86Dd</i>	FBgn0040256	2.65	4.15	
<i>PKD</i>	FBgn0038603	2.65		2.04
<i>Thor</i>	FBgn0261560	2.63		
<i>CG4398</i>	FBgn0034126	2.62	0.052	
<i>CG14692</i>	FBgn0037836	2.61		
<i>CG16712</i>	FBgn0031561	2.60	7.20	2.48

<i>Arc1</i>	FBgn0033926	2.59	5.93	
<i>Rhp</i>	FBgn0026374	2.57	0.328	
<i>COX7AL</i>	FBgn0037579	2.57		
<i>CG8066</i>	FBgn0038243	2.56	5.94	
<i>Muc14A</i>	FBgn0052580	2.56		
<i>CG15611</i>	FBgn0034194	2.56		
<i>Gs2</i>	FBgn0001145	2.55	3.09	
<i>CG33784</i>	FBgn0053784	2.55		
<i>KFase</i>	FBgn0031821	2.53		
<i>Gnmt</i>	FBgn0038074	2.53	8.13	
<i>CG17549</i>	FBgn0032774	2.52		
<i>pirk</i>	FBgn0034647	2.51		
<i>Doc2</i>	FBgn0035956	2.50		
<i>Hsp67Bc</i>	FBgn0001229	2.49	2.74	0.200
<i>egr</i>	FBgn0033483	2.49		
<i>CG7191</i>	FBgn0031945	2.48		
<i>CG3397</i>	FBgn0037975	2.48		0.162
<i>GstE7</i>	FBgn0063493	2.48		8.94
<i>GstE10</i>	FBgn0063499	2.48		
<i>CG11897</i>	FBgn0039644	2.48	5.52	2.14
<i>ome</i>	FBgn0259175	2.47		
<i>CG30022</i>	FBgn0050022	2.47		
<i>CG8008</i>	FBgn0033387	2.46		
<i>CG4822</i>	FBgn0031220	2.45	0.410	
<i>CG5783</i>	FBgn0032670	2.45	0.388	
<i>spri</i>	FBgn0085443	2.45		
<i>Traf4</i>	FBgn0026319	2.44		
<i>CG4829</i>	FBgn0030796	2.43		
<i>be</i>	FBgn0052594	2.42		
<i>CG8870</i>	FBgn0038144	2.41		
<i>Ptth</i>	FBgn0013323	2.39		
<i>KP78b</i>	FBgn0026063	2.38	6.25	
<i>CG14692</i>	FBgn0037836	2.38		
<i>Ror</i>	FBgn0010407	2.38		
<i>CG14642</i>	FBgn0037222	2.37	3.23	
<i>pdm3</i>	FBgn0261588	2.37		
<i>CG17029</i>	FBgn0036551	2.37		
<i>CG1358</i>	FBgn0033196	2.36		
<i>CG13813</i>	FBgn0036956	2.36		
<i>egr</i>	FBgn0033483	2.36		
<i>Inx3</i>	FBgn0265274	2.35		
<i>HDC20468</i>		2.35	8.59	
<i>CG6287</i>	FBgn0032350	2.35		0.382
<i>CG43064</i>	FBgn0262366	2.34		
<i>Pask</i>	FBgn0034950	2.32		
<i>DptB</i>	FBgn0034407	2.32		
<i>CG32521</i>	FBgn0052521	2.32		
<i>mtt</i>	FBgn0050361	2.32		
<i>Mef2</i>	FBgn0011656	2.32		
<i>CG30022</i>	FBgn0050022	2.31		
<i>CG18547</i>	FBgn0037973	2.30	4.73	
<i>Mmp1</i>	FBgn0035049	2.30	15.55	0.264

<i>GstD2</i>	FBgn0010038	2.30	9.61	0.235
<i>Nha2</i>	FBgn0263390	2.29	2.28	6.11
<i>IP3K2</i>	FBgn0266375	2.29		
<i>CG13116</i>	FBgn0032139	2.29		0.454
<i>Sip1</i>	FBgn0010620	2.29	0.396	
<i>CG4502</i>	FBgn0031896	2.28		
<i>Ect4</i>	FBgn0262579	2.28		2.06
<i>yellow-f</i>	FBgn0041710	2.27	0.330	0.358
<i>CG31743</i>	FBgn0032618	2.27		2.60
<i>tilB</i>	FBgn0014395	2.27		
<i>CG13893</i>	FBgn0035146	2.27		
<i>l(2)03659</i>	FBgn0010549	2.26	3.15	
<i>CR32207</i>	FBgn0052207	2.26	12.38	
<i>GstE2</i>	FBgn0063498	2.25		
<i>CG7255</i>	FBgn0036493	2.24	0.448	
<i>CG31253</i>	FBgn0051253	2.24		
<i>eater</i>	FBgn0243514	2.24	8.00	
<i>HDC20224</i>		2.23	5.63	
<i>CG7203</i>	FBgn0031942	2.23		
<i>CG13912</i>	FBgn0035186	2.23		2.27
<i>CG42240</i>	FBgn0250869	2.23	0.403	
<i>CG13698</i>	FBgn0036773	2.23		
<i>CG11686</i>	FBgn0040551	2.22		
<i>CG44325</i>	FBgn0265413	2.21		
<i>rst</i>	FBgn0003285	2.21	13.17	
<i>GstT4</i>	FBgn0030484	2.21	2.14	2.11
<i>CG7627</i>	FBgn0032026	2.21		
<i>IP3K2</i>	FBgn0266375	2.21		
<i>CG30485</i>	FBgn0050485	2.20		
<i>CG31038</i>	FBgn0051038	2.20		
<i>regucalcin</i>	FBgn0030362	2.20		
<i>CG14857</i>	FBgn0038262	2.20		
<i>GlyP</i>	FBgn0004507	2.19		
<i>Lsd-1</i>	FBgn0039114	2.19		
<i>CG3168</i>	FBgn0029896	2.19		
<i>Dyrk2</i>	FBgn0016930	2.18		0.448
<i>Tsp42Ea</i>	FBgn0029508	2.17		
<i>CG18063</i>	FBgn0028856	2.17		
<i>dnr1</i>	FBgn0260866	2.17		
<i>Rab23</i>	FBgn0037364	2.16	3.31	
<i>CG4404</i>	FBgn0030432	2.16	2.28	
<i>CG14291</i>	FBgn0038660	2.16		
<i>bbg</i>	FBgn0087007	2.15	0.392	
<i>Hsp23</i>	FBgn0001224	2.15		
<i>clos</i>	FBgn0261016	2.15		
<i>CG32368</i>	FBgn0052368	2.13		
<i>CG43164</i>	FBgn0262720	2.12		
<i>CG13654</i>	FBgn0039290	2.12	2.49	
<i>CG12428</i>	FBgn0039543	2.12		
<i>sick</i>	FBgn0263873	2.11		
<i>CG31436</i>	FBgn0051436	2.11		
<i>CG44251</i>	FBgn0265186	2.11	2.21	

<i>Inx2</i>	FBgn0027108	2.11		
<i>CG2065</i>	FBgn0033204	2.11	9.00	0.163
<i>CDase</i>	FBgn0039774	2.10	2.07	
<i>PGRP-SB1</i>	FBgn0043578	2.10		0.108
<i>CG12825</i>	FBgn0033221	2.10	2.64	
<i>brp</i>	FBgn0259246	2.09		
<i>CG13650</i>	FBgn0039277	2.08		
<i>MESK2</i>	FBgn0043070	2.08		
<i>CG31274</i>	FBgn0051274	2.08	0.326	0.418
<i>CG10283</i>	FBgn0032681	2.08		
<i>ebd2</i>	FBgn0037076	2.08		
<i>Ela</i>	FBgn0013949	2.08		
<i>pyd</i>	FBgn0262614	2.06		
<i>CG6231</i>	FBgn0038720	2.06		
<i>CecB</i>	FBgn0000278	2.06		
<i>CG14636</i>	FBgn0037217	2.05	0.355	
<i>veil</i>	FBgn0034225	2.05		
<i>CG43340</i>	FBgn0263077	2.05		2.22
<i>CG30460</i>	FBgn0050460	2.05		
<i>Cyp6d5</i>	FBgn0038194	2.05		0.490
<i>CG31705</i>	FBgn0028490	2.05		
<i>PGRP-LA</i>	FBgn0035975	2.05		
<i>CG6701</i>	FBgn0033889	2.05		
<i>GstD9</i>	FBgn0038020	2.04		
<i>pdgy</i>	FBgn0027601	2.03		
<i>spri</i>	FBgn0085443	2.03		
<i>mlt</i>	FBgn0265512	2.03	4.24	
<i>GstE6</i>	FBgn0063494	2.03	43.50	4.17
<i>CG32436</i>	FBgn0052436	2.03		
<i>Cyp4e2</i>	FBgn0014469	2.03		
<i>Gel</i>	FBgn0010225	2.02		
<i>Sans</i>	FBgn0033785	2.02		
<i>Tep4</i>	FBgn0041180	2.02	3.86	
<i>fat-spondin</i>	FBgn0026721	2.01		
<i>HDC03722</i>		2.00		
<i>HP1D3csd</i>	FBgn0030994	2.00		
<i>su(r)</i>	FBgn0086450	2.00	3.20	2.17
<i>lh</i>	FBgn0263397	2.00	2.34	
HP1 depletion effect: DOWN				
<i>CG32850</i>	FBgn0052850	0.498	4.43	
<i>CG41128</i>	FBgn0069923	0.490	0.338	
<i>CG3262</i>	FBgn0032986	0.486		
<i>Dbp80</i>	FBgn0024804	0.480		
<i>CG32350</i>	FBgn0052350	0.480		
<i>Acf1</i>	FBgn0027620	0.480		
<i>Fas2</i>	FBgn0000635	0.479		2.11
<i>ND-AGGG</i>	FBgn0058002	0.477		
<i>PlexB</i>	FBgn0025740	0.474		
<i>COX4L</i>	FBgn0033020	0.464		
<i>PlexB</i>	FBgn0025740	0.455		
<i>CG34330</i>	FBgn0085359	0.452	4.33	
<i>CG6329</i>	FBgn0033872	0.435		

<i>Nipped-A</i>	FBgn0053554	0.430		
<i>CG2709</i>	FBgn0024977	0.425		
<i>I(2)41Ab</i>	FBgn0262123	0.417	4.24	
<i>dob</i>	FBgn0030607	0.402	3.62	
<i>TpnC41C</i>	FBgn0013348	0.398		
<i>CG3829</i>	FBgn0035091	0.394		
<i>HDC20114</i>		0.370		
<i>Ser</i>	FBgn0004197	0.366	0.385	
<i>Transpac</i>	FBte0001377	0.363		
<i>CG41099</i>	FBgn0039955	0.362		
<i>CG3788</i>	FBgn0034800	0.353		
<i>CG17698</i>	FBgn0040056	0.347		
<i>br</i>	FBgn0000210	0.327		
<i>slgA</i>	FBgn0003423	0.316	0.345	
<i>mos</i>	FBgn0033773	0.309		
<i>Caps</i>	FBgn0053653	0.284		
<i>CR43242</i>	FBgn0262887	0.252		0.097
<i>CG40160</i>	FBgn0058160	0.239	0.441	
<i>CR43242</i>	FBgn0262887	0.239		0.097
<i>CG40006</i>	FBgn0058006	0.230	28.55	
<i>CG40294</i>	FBgn0058294	0.157		0.398
<i>CG30440</i>	FBgn0050440	0.132		
<i>Maf1</i>	FBgn0267861	0.129		
<i>Caps</i>	FBgn0053653	0.116		
<i>CR45923</i>	FBgn0267585	0.093	6.95	
<i>CG17698</i>	FBgn0040056	0.093		
<i>CG32017</i>	FBgn0052017	0.058	5.13	
<i>Cht3</i>	FBgn0250907	0.029		
<i>Su(var)205</i>	FBgn0003607	0.025		

Transcripts that are up- or down-regulated by RNAi-mediated depletion of HP1 in Kc cells and H1 in L3 salivary glands or a homozygous null mutation of *Chd1* in L3 larvae. Fold change is calculated from Affymetrix microarray data relative to wild-type control. Transposons, repetitive and presumed heterochromatic sequences (Chr Xh, Chr 2h, Chr 3h and Chr4) are shown in red type.