Comparative DEGs table. Evaluated differentially expressed genes considering all pairwaise comparsons between *D. buzzatii* and *D. koepferae* (q<0.01) using *D. mojavensis* reference and the respective *D. melanogaster* homologs

D. mojavensis	D. mojavensis		D. melanogaster	Description
gene symbol SLC	gene id	gene symbol	gene id	<u>'</u>
Dmoj\GI17747	FBgn0140488	Nha1	FBgn0031865	Na[+]/H[+] hydrogen antiporter 1
Dmoj\GI17747	FBgn0136938	dmGlut	FBgn0010497	Dietary and metabolic glutamate transporter
Dmoj\GI19091	FBgn0141830	CG8468	FBgn0033913	Dietary and metabolic glutamate transporter
Dmoj\GI21795	FBgn0144525	Mct1	FBgn0023549	Monocarboxylate transporter 1
Dmoj\Gl22971	FBgn0145698	CG7333	FBgn0038715	monocarboxylate danoportor 1
Dmoj\GI12888	FBgn0135645	blot	FBgn0027660	bloated tubules
Dmoj\GI19618	FBgn0142355	Oatp58Da	FBgn0050277	Organic anion transporting polypeptide 58Da
Dmoj\GI13032	FBgn0135789	CG8177	FBgn0036043	
Dmoj\GI24359	FBgn0147082	CG6125	FBgn0038337	
Dmoj\GI10539	FBgn0133303	MFS9	FBgn0038799	Major Facilitator Superfamily Transporter 9
Dmoj\GI22204	FBgn0144933	CG1607	FBgn0039844	
Dmoj\GI24622	FBgn0147344	CG6231	FBgn0038720	
Dmoj\GI18111	FBgn0140851	CG3036	FBgn0031645	
Dmoj\GI13648	FBgn0136403	CG10960	FBgn0036316	
Dmoj\GI10224	FBgn0132989	CG11665	FBgn0033028	
Dmoj\Gl22338	FBgn0145066	Zip99C	FBgn0039714	Zinc/iron regulated transporter-related protein 99C
Dmoj\GI17795	FBgn0140536	Ndae1	FBgn0259111	Na[+]-driven anion exchanger 1
Dmoj\GI13105	FBgn0135862	CG7255	FBgn0036493	
Dmoj\Gl21812	FBgn0144542	Ctr1A	FBgn0062413	Copper transporter 1A
Dmoj\GI24495	FBgn0147217	NKCC	FBgn0051547	sodium potassium chloride cotransporter
Dmoj\GI19615	FBgn0142352	Oatp58Dc	FBgn0034716	Organic anion transporting polypeptide 58Dc
Dmoj\GI20333	FBgn0143069	CG9826	FBgn0034784	
Dmoj\GI11052	FBgn0133815	NAAT1	FBgn0029762	Nutrient Amino Acid Transporter 1
Dmoj\GI23108	FBgn0145835	CG7342	FBgn0038716	
Dmoj\GI20348	FBgn0143084	List	FBgn0034381	Lithium-inducible SLC6 transporter
Dmoj\GI13064	FBgn0135821	Catsup	FBgn0002022	Catecholamines up
Dmoj\GI17357	FBgn0140101	CG13384	FBgn0032036	
Dmoj\GI10643	FBgn0133407	CG9717	FBgn0039789	
Dmoj\GI10092	FBgn0132858	CG6126	FBgn0038407	
Dmoj\GI19603	FBgn0142340	ZnT41F	FBgn0025693	Zinc transporter 41F
Dmoj\GI20339	FBgn0143075	MFS14	FBgn0010651	Major Facilitator Superfamily Transporter 14
Dmoj\GI11098	FBgn0133860	sesB	FBgn0003360	stress-sensitive B
Dmoj\Gl21320	FBgn0144050	CG3394	FBgn0034999	
Dmoj\GI13371	FBgn0136128	CG9706	FBgn0036662	
Dmoj\GI19350	FBgn0142088	sut1	FBgn0028563	sugar transporter 1
Dmoj\GI19756	FBgn0142493	CG8249	FBgn0034045	
Dmoj\Gl24592	FBgn0147314	sll	FBgn0038524	slalom
Dmoj\GI17852	FBgn0140593	Ent1	FBgn0031250	Equilibrative nucleoside transporter 1
Dmoj\GI14799	FBgn0137550	CG4991	FBgn0030817	
Dmoj\GI16685	FBgn0139433	CG42269	FBgn0259164	La una ala in
Dmoj\Gl22265	FBgn0144993	kar	FBgn0001296	karmoisin
Dmoj\GI19616 Dmoj\GI15324	FBgn0142353 FBgn0138073	Oatp58Da Ucp4A	FBgn0050277 FBgn0030872	Organic anion transporting polypeptide 58Da
Dmoj\GI16224	FBgn0138973	Nhe1	FBgn0030872 FBgn0026787	Na[+1/H[+1 hydrogen eychanger 1
Dmoj\GI16224 Dmoj\GI16741	FBgn0138973 FBgn0139489	CG33282	FBgn0026787 FBgn0053282	Na[+]/H[+] hydrogen exchanger 1
Dmoj\G123107	FBgn0139489 FBgn0145834	CG33282 CG7333	FBgn0038715	
Dmoj\Gl23107	FBgn0143983	CG12866	FBgn0033955	
Dmoj\Gl20336	FBgn0143072	CG12806 CG15096	FBgn0034394	
Dmoj\Gl24704	FBgn0147426	Glut3	FBgn0015230	Glucose transporter type 3
Dmoj\GI18804	FBgn0141543	CG14744	FBgn0033324	
Dmoj\GI12576	FBgn0135333	CG7514	FBgn0035567	
Dmoj\GI10986	FBgn0133749	CG8028	FBgn0031010	
Dmoj\GI17535	FBgn0140279	Ucp4B	FBgn0031758	Uncoupling protein 4B
Dmoj\GI14188	FBgn0136941	CG18788	FBgn0042126	
Dmoj\Gl21883	FBgn0144612	Ntl	FBgn0267326	Neurotransmitter transporter-like
Dmoj\GI19787	FBgn0142524	MFS12	FBgn0033234	Major Facilitator Superfamily Transporter 12
-	-		_	
Dmoj\GI17977	FBgn0140717	MFS3	FBgn0031307	Major Facilitator Superfamily Transporter 3
Dmoj\Gl21513	FBgn0144243	CG5254	FBgn0040383	Mitochondrial phocphate carrier protein 4
Dmoj\GI20792	FBgn0143527	Mpcp1	FBgn0034497	Mitochondrial phosphate carrier protein 1
Dmoj\GI12085	FBgn0134842	path	FBgn0036007	pathetic Fathy solid (long chain) transport protein
Dmoj\GI17838	FBgn0140579	Fatp	FBgn0267828	Fatty acid (long chain) transport protein

FBgn0140628	ine	FBgn0011603	inebriated
FBgn0136713	Indy	FBgn0036816	I'm not dead yet
FBgn0145487	aralar1	FBgn0028646	
FBgn0136440	NaPi-III	FBgn0260795	Na[+]-dependent inorganic phosphate cotransporter type III
FBgn0133234	CG7084	FBgn0038938	
FBgn0136632	CG13907	FBgn0035173	
FBgn0142416	Tpc2	FBgn0035078	Thiamine pyrophosphate carrier protein 2
FBgn0138137	CG3168	FBgn0029896	
FBqn0134250	Mpcp2	FBqn0026409	Mitochondrial phosphate carrier protein 2
-			Trehalose transporter 1-2
-		-	Solute carrier family 45 member 1
_		-	
-			fear-of-intimacy
-		-	bedraggled
-	-	-	bedraggied
-		-	Organic anion transporting polypeptide 33Eb
-		-	
-		-	Organic anion transporting polypeptide 33Ea
-		-	
-		-	
-		-	
-			
⊢Bgn0144060	vmat	⊦Bgn0260964	Vesicular monoamine transporter
FBgn0138045	mvs	FBqn0004657	myospheroid
	-	-	alpha-Esterase-7
-			a.pa 2001000 /
-		-	
-		-	alaba Fatarana 2
-		_	alpha-Esterase-2
-		-	Esterase 6
-		-	Esterase 6
-		_	Esterase 6
FBgn0145202	Est-6	FBgn0000592	Esterase 6
FBgn0142889	Jheh2	FBgn0034405	Juvenile hormone epoxide hydrolase 2
FBgn0138902	CG6414	FBgn0029690	
FBgn0146202	alpha-Est5	FBgn0261393	alpha-Esterase-5
FBgn0134304	CG7365	FBgn0036939	
FBgn0132887	CG4757	FBgn0027584	
FBgn0139957	CG6523	FBgn0032509	
FBgn0144274	spidey	FBgn0029975	
FBgn0139593	ND-13B	FBgn0047038	NADH dehydrogenase (ubiquinone) 13 kDa B subunit
FBgn0147046	CG9743	FBgn0039756	
FBgn0147047	CG15531	FBgn0039755	
-		FBgn0033101	
_		-	
5		FBgn0003965	vermilion
FBqn0139005	V		
FBgn0139005 FBgn0064059	v	-	
FBgn0064059	ry	FBgn0003308	rosy
FBgn0064059 FBgn0139801	ry CG17928	FBgn0003308 FBgn0032603	rosy
FBgn0064059 FBgn0139801 FBgn0141126	ry CG17928 Fmo-2	FBgn0003308 FBgn0032603 FBgn0033079	
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669	ry CG17928 Fmo-2 CG11796	FBgn0003308 FBgn0032603 FBgn0033079 FBgn0036992	rosy
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669 FBgn0145167	ry CG17928 Fmo-2 CG11796 CG4836	FBgn0003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925	rosy
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669 FBgn0145167 FBgn0146078	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1	FBgn003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095	rosy Flavin-containing monooxygenase 2
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0138008	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A	FBgn003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095 N/A	rosy
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0138008 FBgn0146077	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1	FBgn003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095 N/A FBgn0038095	rosy Flavin-containing monooxygenase 2 N/A
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0138008 FBgn0146077 FBgn0141433	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2	FBgn0003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095 N/A FBgn0038095 FBgn0014469	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0138008 FBgn0146077 FBgn0141433 FBgn0142964	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2	FBgn0003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn0000473	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2 Cytochrome P450-6a2
FBgn0064059 FBgn0139801 FBgn01341126 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0138008 FBgn0146077 FBgn0141433 FBgn0142964 FBgn0142966	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2 Cyp6a2	FBgn0003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn000473 FBgn0000473	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2
FBgn0064059 FBgn0139801 FBgn0141126 FBgn0145167 FBgn0145167 FBgn0146078 FBgn0138008 FBgn0146077 FBgn0141433 FBgn0142964 FBgn0142966 FBgn0135704	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2 Cyp6a2 CG6034	FBgn0003308 FBgn0032603 FBgn0033079 FBgn0036992 FBgn0270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn000473 FBgn000473 FBgn0036750	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2 Cytochrome P450-6a2
FBgn0064059 FBgn0139801 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0146077 FBgn0144033 FBgn0142964 FBgn0135704 FBgn0139736	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2 Cyp6a2 Cyp6a2 CG6034 Cyp309a1	FBgn0003308 FBgn0032603 FBgn0032699 FBgn00270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn000473 FBgn0000473 FBgn0036750 FBgn0031432	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2 Cytochrome P450-6a2 Cytochrome P450-6a2
FBgn0064059 FBgn0139801 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0146077 FBgn0144077 FBgn0141433 FBgn0142964 FBgn0135704 FBgn0139736 FBgn0146945	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2 Cyp6a2 CG6034 Cyp309a1 Uro	FBgn0003308 FBgn0032603 FBgn00326992 FBgn00270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn0000473 FBgn0000473 FBgn0036750 FBgn0031432 FBgn0003961	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2 Cytochrome P450-6a2 Cytochrome P450-6a2 Urate oxidase
FBgn0064059 FBgn0139801 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0138008 FBgn0146077 FBgn0141433 FBgn0142964 FBgn0135704 FBgn0139736 FBgn0146945 FBgn0134730	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2 Cyp6a2 CG6034 Cyp309a1 Uro GstO3	FBgn0003308 FBgn0032603 FBgn0032603 FBgn0036992 FBgn0270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn000473 FBgn000473 FBgn0036750 FBgn0031432 FBgn0003961 FBgn0035904	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2 Cytochrome P450-6a2 Cytochrome P450-6a2 Urate oxidase Glutathione S transferase O3
FBgn0064059 FBgn0139801 FBgn0134669 FBgn0145167 FBgn0146078 FBgn0146077 FBgn0144077 FBgn0141433 FBgn0142964 FBgn0135704 FBgn0139736 FBgn0146945	ry CG17928 Fmo-2 CG11796 CG4836 Cyp304a1 N/A Cyp304a1 Cyp4e2 Cyp6a2 Cyp6a2 CG6034 Cyp309a1 Uro	FBgn0003308 FBgn0032603 FBgn00326992 FBgn00270925 FBgn0038095 N/A FBgn0038095 FBgn0014469 FBgn0000473 FBgn0000473 FBgn0036750 FBgn0031432 FBgn0003961	rosy Flavin-containing monooxygenase 2 N/A Cytochrome P450-4e2 Cytochrome P450-6a2 Cytochrome P450-6a2 Urate oxidase
	FBgn0136713 FBgn0145487 FBgn0136440 FBgn0133234 FBgn0136632 FBgn0142416 FBgn0138137 FBgn0134250 FBgn0143111 FBgn0135643 FBgn0135663 FBgn0144030 FBgn0135663 FBgn0144030 FBgn0135663 FBgn0144030 FBgn0133203 FBgn0143795 FBgn0133203 FBgn0143795 FBgn0138045 FBgn0138045 FBgn0148204 FBgn0138045 FBgn014900 FBgn014900 FBgn014900 FBgn014900 FBgn0145201 FBgn0145202 FBgn0145204 FBgn0145204 FBgn0145204 FBgn0145205 FBgn0145204 FBgn0145204 FBgn0145205 FBgn0145207 FBgn0144274 FBgn0144274 FBgn0144296 FBgn0144418	FBgn0136713 Indy FBgn0145487 aralar1 FBgn0136440 NaPi-III FBgn0133234 CG7084 FBgn0136632 CG13907 FBgn0142416 Tpc2 FBgn0138137 CG3168 FBgn0134250 Mpcp2 FBgn0135443 Slc45-1 FBgn0135663 foi FBgn0135663 foi FBgn0144030 bdg FBgn0134204 CG7888 FBgn0137202 Oatp33Eb FBgn0139865 Oatp33Ea FBgn0143795 CG3790 FBgn0133517 CG5646 FBgn0133203 CG15553 FBgn0144060 Vmat FBgn0138045 mys FBgn0138045 mys FBgn0138045 mys FBgn0146204 alpha-Est7 FBgn0138902 CG6414 FBgn0145204 Est-6 FBgn0145204 Est-6 FBgn0145205 Est-6 FBgn0145205 Est-6 FBgn0145206 Est-6 FBgn0145207 CG6414 FBgn0145208 Est-6 FBgn0145209 Est-6 FBgn0145201 Est-6 FBgn0145202 Est-6 FBgn0145203 CG6414 FBgn0145204 Est-6 FBgn0145205 Est-6 FBgn0145205 Est-6 FBgn0145206 Est-6 FBgn0145207 CG6523 FBgn01389957 CG6523 FBgn0139957 CG6523 FBgn0139593 ND-13B FBgn0147046 CG9743 FBgn0142968 CG9436 FBgn01444418 CG33178	FBgn0136713 Indy FBgn0036816 FBgn0145487 aralar1 FBgn0028646 FBgn0136440 NaPi-III FBgn0260795 FBgn0133234 CG7084 FBgn0038938 FBgn0136632 CG13907 FBgn0035173 FBgn0138137 CG3168 FBgn0026409 FBgn0134250 Mpcp2 FBgn0026409 FBgn0134431 Tret1-2 FBgn0033644 FBgn0135443 Slc45-1 FBgn0035968 FBgn0143070 CG15096 FBgn0034994 FBgn0135663 foi FBgn0034494 FBgn0134204 CG7888 FBgn0034049 FBgn0137202 Oatp33Eb FBgn0032433 FBgn0139865 Oatp33Ea FBgn0032433 FBgn0146223 CG14606 FBgn0033778 FBgn0133517 CG5646 FBgn0039525 FBgn0133203 CG15553 FBgn0039817 FBgn014600 Vmat FBgn004657 FBgn0138902 CG6414 FBgn000592 FBgn0146847 alpha-Est7 FBgn000592 <t< td=""></t<>

Dmci\C 1110F2	FD==0424644	CCEDAG	FD~~0036344	
Dmoj\GI11853	FBgn0134611	CG5946	FBgn0036211	
Dmoj\GI16636	FBgn0139384	Jafrac2	FBgn0040308	Thioredoxin peroxidase 2
Dmoj\Gl22127	FBgn0144856	Cyp4c3	FBgn0015032	Cytochrome P450-4c3
Dmoj\Sod1	FBgn0134384	Sod1	FBgn0003462	Superoxide dismutase 1
Dmoj\GI19114	FBgn0141853	CG5009	FBgn0027572	
Dmoj\GI19074	FBgn0141813	Acox57D-p	FBgn0034628	acyl-Coenzyme A oxidase at 57D proximal
Dmoj\Gl20052	FBgn0142788	Cyp9b1	FBgn0015038	Cytochrome P450-9b1
Dmoj\GI11492	FBgn0134252	shd	FBgn0003388	shade
Dmoj\Gl22465	FBgn0145193	CG4860	FBgn0037999	
Dmoj\GI13547	FBgn0136302	CG9629	FBgn0036857	
Dmoj\GI18497	FBgn0141236	CG8303	FBqn0034143	
Dmoj\GI17568	FBgn0140312	Prx6005	FBqn0031479	Peroxiredoxin 6005
Dmoj\GI15383	FBgn0138132	CG9519	FBgn0030589	1 GIOAN COOK
-	-		_	Home evigences
Dmoj\Gl24074	FBgn0146797	Ho	FBgn0037933	Heme oxygenase
Dmoj\GI14867	FBgn0137618	CG5321	FBgn0030575	
Dmoj\Gl21510	FBgn0144240	CG7322	FBgn0030968	
Dmoj\GI16989	FBgn0139735	Cyp309a2	FBgn0041337	
Dmoj\GI10473	FBgn0133237	rdhB	FBgn0038946	retinol dehydrogenase B
Dmoj\Gl20619	FBgn0143354	hgo	FBgn0040211	homogentisate 1,2-dioxygenase
Dmoj\GI15481	FBgn0138230	GstT3	FBgn0031117	Glutathione S transferase T3
Dmoj\GI23178	FBgn0145905	CG31559	FBgn0051559	
Dmoj\Desat2	FBgn0133248	Desat2	FBgn0043043	Desaturase 2
Dmoj\GI17664	FBgn0140405	Mco1	FBgn0032116	Multicopper oxidase-1
Dmoj\GI20008	FBgn0142744	CG17807	FBgn0034748	
Dmoj\GI17645	FBgn0140387	Adhr	FBgn0000056	Adh-related
Dmoj\GI12408	FBgn0135165	CG9314	FBgn0032061	,
-	-		-	Cutashrama D450 4d4
Dmoj\GI16117	FBgn0138866	Cyp4d1	FBgn0005670	Cytochrome P450-4d1
Dmoj\GI16116	FBgn0138865	Cyp4d1	FBgn0005670	Cytochrome P450-4d1
Dmoj\GI15376	FBgn0138125	CG9509	FBgn0030594	
Dmoj\GI11003	FBgn0133766	CG2254	FBgn0029994	
Dmoj\GI19441	FBgn0142179	CG10814	FBgn0033830	
Dmoj\GI17558	FBgn0140302	Cyp28a5	FBgn0028940	
Dmoj\GI11498	FBgn0134258	CG13473	FBgn0036442	
Dmoj\GI12897	FBgn0135654	ND-24L	FBgn0036706	NADH dehydrogenase (ubiquinone) 24 kDa subunit-like
Dmoj\Gl20357	FBgn0143093	COX5BL	FBgn0031831	Cytochrome c oxidase subunit 5B-like
Dmoj\Gl26293	FBgn0281030	ND-51L1	FBgn0034251	NADH dehydrogenase (ubiquinone) 51 kDa subunit-like 1
Dmoj\GI15072	FBgn0137822	COX4L	FBgn0033020	Cytochrome c oxidase subunit 4-like
Dmoj\GI14832	FBgn0137583	firl	FBgn0032405	firelighter
Dmoj\Gl24212	FBgn0146935	CG7950	FBgn0260468	
Dmoj\Gl21924	FBgn0144653	Cyp4ac1	FBgn0031693	
Dmoj\GI21924 Dmoj\GI17672	FBgn0140413	CG7311	FBgn0028848	
-			_	
Dmoj\GI20316	FBgn0143052	CG30275	FBgn0050275	
Dmoj\Gl24862	FBgn0147582	CG9698	FBgn0039784	
Dmoj\GI17187	FBgn0139931	CG3597	FBgn0031417	
Dmoj\GI14956	FBgn0137707	CG40486	FBgn0263830	
Dmoj\GI19073	FBgn0141812	Acox57D-p	FBgn0034628	acyl-Coenzyme A oxidase at 57D proximal
Dmoj\GI18704	FBgn0141443	Cyp6a17	FBgn0015714	Cytochrome P450-6a17
Dmoj\GI24047	FBgn0146770	Cyp12e1	FBgn0037817	
Dmoj\GI12291	FBgn0135048	CG13902	FBgn0035166	
Dmoj\GI19238	FBgn0141977	Cyp6d2	FBgn0034756	
Dmoj\Gl24131	FBgn0146854	sgll	FBgn0051472	sugarlethal
Dmoj\GI11604	FBgn0134364	CG7560	FBgn0036157	
Dmoj\Gl21555	FBgn0144285	Cyp4s3	FBgn0030615	
Dmoj\GI21333	FBgn0135705	CG6034	_	
-	-		FBgn0036750	conia
Dmoj\GI11974	FBgn0134731	Se Cureat	FBgn0086348	sepia
Dmoj\GI20590	FBgn0143325	Cyp6g1	FBgn0025454	
Dmoj\GI15186	FBgn0137936	CG3842	FBgn0029866	
Dmoj\GI10146	FBgn0132912	COX5A	FBgn0019624	Cytochrome c oxidase subunit 5A
Dmoj\GI17627	FBgn0140371	cype	FBgn0015031	cyclope
Dmoil@122740	FBgn0145476	COX7A	FBgn0040529	Cytochrome c oxidase subunit 7A
Dmoj\Gl22749		T 0	FBgn0040070	thioredoxin-2
Dmoj\Gl22749 Dmoj\Trx-2	FBgn0133846	Trx-2	. Egillet let le	
-	FBgn0133846 FBgn0135007	CG12262	FBgn0035811	
Dmoj\Trx-2	-		_	clot
Dmoj\Trx-2 Dmoj\GI12250	FBgn0135007	CG12262	FBgn0035811	clot
Dmoj\Trx-2 Dmoj\GI12250 Dmoj\GI17989	FBgn0135007 FBgn0140729	CG12262	FBgn0035811 FBgn0000318	clot

Dmoj\GI11589	FBgn0134349	CG6084	FBgn0086254	
Dmoj\GI18148	FBgn0140887	CG6287	FBgn0032350	
Dmoj\Gl24641	FBgn0147363	glob1	FBgn0027657	globin 1
Dmoj\GI18385	FBgn0141124	wal	FBgn0010516	walrus
Dmoj\GI15562	FBgn0138311	CG10962	FBgn0030073	Wallas
Dmoj\GI20198	FBgn0142934	Prx2540-2	FBgn0033518	Peroxiredoxin 2540-2
Dmoj\Gl23694	FBgn0146420	pug	FBgn0020385	pugilist
Dmoj\Gl26579	FBgn0281316	CG30022	FBgn0050022	pagiliot
Dmoj\Gl23165	FBgn0145892	CG9331	FBgn0032889	
Dmoj\GI17168	FBgn0139912	L2HGDH	FBgn0032729	L-2-hydroxyglutarate dehydrogenase
Dmoj\GI23880	FBgn0146605	Prx5	FBgn0038570	Peroxiredoxin 5
Dmoj\GI11069	FBgn0133832	Pgd	FBgn0004654	Phosphogluconate dehydrogenase
Dmoj\GI22545	FBgn0145273	Hph	FBgn0264785	HIF prolyl hydroxylase
Dmoj\GI11068	FBgn0133831	D2hgdh	FBgn0023507	D-2-hydroxyglutaric acid dehydrogenase
Dmoj\Gl22727	FBgn0145454	CG5346	FBgn0038981	D E Hydroxyglatano dola denyarogonado
Dmoj\GI19146	FBgn0141885	CG8399	FBgn0034067	
Dmoj\GI19860	FBgn0142597	Nox	FBgn0085428	NADPH oxidase
Dmoj\GI24725	FBgn0147447	Cyp9f3	FBgn0038034	
Dmoj\GI14608	FBgn0137359	GstT4	FBgn0030484	Glutathione S transferase T4
Dmoj\Gl23851	FBgn0146576	Pxd	FBgn0004577	Peroxidase
Dmoj\Gl23244	FBgn0145971	GILT1	FBgn0038149	Gamma-interferon-inducible lysosomal thiol reductase 1
Dmoj\GI13115	FBgn0135872	Eip71CD	FBgn0000565	Ecdysone-induced protein 28/29kD
Dmoj\Gl24757	FBgn0147479	Mdh2	FBgn0262559	Malate dehydrogenase 2
Dmoj\GI11930	FBgn0134688	CG9674	FBgn0036663	
Dmoj\GI16236	FBgn0138985	CG17896	FBgn0023537	
Dmoj\GI20341	FBgn0143077	CG15093	FBgn0034390	
Dmoj\GI11829	FBgn0134588	CG3902	FBgn0036824	
Dmoj\Desat1	FBgn0133249	Desat1	FBgn0086687	Desaturase 1
Dmoj\Gl22276	FBgn0145004	Gdh	FBgn0001098	Glutamate dehydrogenase
Dmoj\GI14919	FBgn0137670	Cyt-b5-r	FBgn0000406	Cytochrome b5-related
Dmoj\Adh2	FBgn0012567	Adh	FBgn0000055	Alcohol dehydrogenase
Dmoj\GI12682	FBgn0135439	ple	FBgn0005626	pale
BillojiGri2002	1 Egilo 100400		1 Egilococce	·
Dmoj\GI13499	FBgn0136255	ND-51	FBgn0031771	NADH dehydrogenase (ubiquinone) 51 kDa subunit
Dmoj\GI11696	FBgn0134455	CG7430	FBgn0036762	
Dmoj\Gl22037	FBgn0144766	ldh3b	FBgn0038922	Isocitrate dehydrogenase 3b
Dmoj\GI16551	FBgn0139299	UQCR-C2	FBgn0250814	Ubiquinol-cytochrome c reductase core protein 2
Dmoj\Gl23793	FBgn0146518	ND-MLRQ	FBgn0052230	NADH dehydrogenase (ubiquinone) MLRQ subunit
Dmoj\GI17532	FBgn0140276	Gpdh	FBgn0001128	Glycerol 3 phosphate dehydrogenase
Dmoj\GI13905	FBgn0136659	Tet	FBgn0263392	Ten-Eleven Translocation (TET) family protein
Dmoj\GI12153	FBgn0134910	CG1275	FBgn0035321	
Dmoj\GI16833	FBgn0139580	Dhpr	FBgn0035964	Dihydropteridine reductase
Dmoj\Gl24244	FBgn0146967	Men	FBgn0002719	Malic enzyme
Dmoj\GI12765	FBgn0135522	CG10863	FBgn0027552	
Dmoj\GI12904	FBgn0135661	ND-SGDH	FBgn0011455	NADH dehydrogenase (ubiquinone) SGDH subunit
Dmoj\GI11212	FBgn0133973	UQCR-14L	FBgn0039576	Ubiquinol-cytochrome c reductase 14 kDa subunit-like
Dmoj\GI19303	FBgn0142042	SdhB	FBgn0014028	Succinate dehydrogenase, subunit B (iron-sulfur)
Dmoj\Gl22813	FBgn0145540	Prx3	FBgn0038519	Peroxiredoxin 3
Dmoj\GI14857	FBgn0137608	CG12116	FBgn0030041	
Dmoj\GI19578	FBgn0142315	CG5493	FBgn0034364	
Dmoj\GI19152	FBgn0141891	Asph	FBgn0034075	Aspartyl beta-hydroxylase
Dmoj\GI16427	FBgn0139175	COX6B	FBgn0031066	Cytochrome c oxidase subunit 6B
Dmoj\CtBP	FBgn0146084	CtBP	FBgn0020496	C-terminal Binding Protein
Dmoj\Gl23201	FBgn0145928	Men-b	FBgn0029155	Malic enzyme b
Dmoj\GI13568	FBgn0136323	CG7470	FBgn0037146	
Dmoj\Gl23575	FBgn0146301	CG3999	FBgn0037801	
Dmoj\GI11531	FBgn0134291	Cpr	FBgn0015623	Cytochrome P450 reductase
Dmoj\Adh1	FBgn0012566	Adh	FBgn0000055	Alcohol dehydrogenase
Dmoj\GI12492	FBgn0135249	ldh	FBgn0001248	Isocitrate dehydrogenase
Dmoj\GI13269	FBgn0136026	Nc73EF	FBgn0010352	Neural conserved at 73EF
Dmoj\GI13115	FBgn0135872	Eip71CD	FBgn0000565	Ecdysone-induced protein 28/29kD
Dmoj\GI17663	FBgn0140404	Aldh	FBgn0012036	Aldehyde dehydrogenase
Dmoj\Adh2	FBgn0012567	Adh	FBgn0000055	Alcohol dehydrogenase
Dmoj\GI17665	FBgn0140406	Fbp2	FBgn0000640	Fat body protein 2
Dmoj\Gl22321	FBgn0145049	CG1544	FBgn0039827	
Dmoj\GI16197	FBgn0138946	CG6762	FBgn0030876	
Dmoj\GI12839	FBgn0135596	Pxn	FBgn0011828	Peroxidasin

D "01440=0				
Dmoj\GI14070	FBgn0136824	ND-B22	FBgn0032511	NADH dehydrogenase (ubiquinone) B22 subunit
Dmoj\GI17845	FBgn0140586	CG9265	FBgn0032910	
Dmoj\GI22556	FBgn0145284	Kdm2	FBgn0037659	Lysine (K)-specific demethylase 2
Dmoj\GI18700	FBgn0141439	N/A	N/A	N/A
Dmoj\GI15385	FBgn0138134	CG12539	FBgn0030586	
Dmoj\GI19427	FBgn0142165	PPO2	FBgn0033367	Prophenoloxidase 2
Dmoj\Gl22625	FBgn0145353	SdhC	FBgn0037873	Succinate dehydrogenase, subunit C
Dmoj\GI10356	FBgn0133120	Ssadh	FBgn0039349	Succinic semialdehyde dehydrogenase
Dmoj\GI24093	FBgn0146816	ND-23	FBgn0017567	NADH dehydrogenase (ubiquinone) 23 kDa subunit
Dmoj\GI20381	FBgn0143117	CG2065	FBgn0033204	
Dmoj\GI17642	FBgn0140386	N/A	N/A	N/A
Dmoj\GI24740	FBgn0147462	Cyp6d5	FBgn0038194	
Dmoj\GI15204	FBgn0137953	Eo	FBgn0030597	Ecdysone oxidase
Dmoj\GI14858	FBgn0137609	Sptr	FBgn0014032	Sepiapterin reductase
-	_	CG9522	_	Sepiapteriii reductase
Dmoj\GI15384	FBgn0138133		FBgn0030587	
Dmoj\GI17433	FBgn0140177	Cyp6t1	FBgn0031182	
Dmoj\GI15906	FBgn0138655	CG4586	FBgn0029924	
Dmoj\GI19287	FBgn0142026	PPO1	FBgn0283437	Prophenoloxidase 1
Dmoj\GI13193	FBgn0135950	lyd	FBgn0036125	lodotyrosine deiodinase
Dmoj\GI21687	FBgn0144417	CG33177	FBgn0053177	
Dmoj\GI18650	FBgn0141389	Cyp4p1	FBgn0015037	Cytochrome P450-4p1
Dmoj\GI23955	FBgn0146679	Pxt	FBgn0261987	Peroxinectin-like
Dmoj\GI14225	FBgn0136978	AlkB	FBgn0065035	
Dmoj\Gl24027	FBgn0146750	CG42331	FBgn0259233	
Dmoj\GI23191	FBgn0145918	GILT2	FBgn0039099	Gamma-interferon-inducible lysosomal thiol reductase 2
Dmoj\GI10563	FBgn0133327	lox	FBgn0039848	lysyl oxidase-like
Dmoj\GI15373	FBgn0138122	CG12398	FBgn0030596	
Dmoj\GI17600	FBgn0140344	CG10495	FBgn0032750	
Dmoj\Gl22554	FBgn0145282	CG14882	FBgn0038429	
Dmoj\GI21954	FBgn0144683	N/A	N/A	N/A
Dmoj\GI15210	FBgn0137959	Cyp28c1	FBgn0030339	N/A
-	-	CG13833	-	
Dmoj\GI10766	FBgn0133529		FBgn0039040	One hit had been deep and a second of
Dmoj\GI22938	FBgn0145665	Sodh-2	FBgn0022359	Sorbitol dehydrogenase-2
Dmoj\GI14338	FBgn0137091	CG12206	FBgn0029662	
Dmoj\GI18701	FBgn0141440	Cyp6a22	FBgn0013773	
GST	I			
Dmoj\GI14608	FBgn0137359	GstT4	FBgn0030484	Glutathione S transferase T4
Dmoj\GI20122	FBgn0142858	GstE8	FBgn0063492	Glutathione S transferase E8
Dmoj\GI11973	FBgn0134730	GstO3	FBgn0035904	Glutathione S transferase O3
Dmoj\GI23596	FBgn0146322	GstZ2	FBgn0037697	Glutathione S transferase Z2
Dmoj\GI24379	FBgn0147102	GstD1	FBgn0001149	Glutathione S transferase D1
Dmoj\GI22354	FBgn0145082	GstD10	FBgn0042206	Glutathione S transferase D10
Dmoj\GI15481	FBgn0138230	GstT3	FBgn0031117	Glutathione S transferase T3
Dmoj\GI20123	FBgn0142859	GstE8	FBgn0063492	Glutathione S transferase E8
Dmoj\GI20133	FBgn0142869	GstE11	FBgn0034354	Glutathione S transferase E11
Dmoj\Gl20124	FBgn0142860	GstE9	FBgn0063491	Glutathione S transferase E9
Dmoj\GI24095	FBgn0146818	GstE1	FBgn0034335	Glutathione S transferase E1
Dmoj\GI11974	FBgn0134731	se	FBgn0086348	sepia
Dmoj\GI20121	FBgn0142857	GstE4	FBgn0063496	Glutathione S transferase E4
	g 12001		290000400	
GT				
	EPap012F040	ngant4	EDan0051056	nelypontide CalNAs transferace 4
Dmoj\GI13183	FBgn0135940	pgant4	FBgn0051956	polypeptide GalNAc transferase 4
Dmoj\GI19414	FBgn0142152	GalNAc-T1	FBgn0034025	Taskalasa O akasakata suutkasa 4
Dmoj\Gl22062	FBgn0144791	Tps1	FBgn0027560	Trehalose-6-phosphate synthase 1
	FBgn0132970	kkv	FBgn0001311	krotzkopf verkehrt
Dmoj\GI11811	FBgn0134570	CG3797	FBgn0036842	
Dmoj\GI15586	FBgn0138335	GalNAc-T2	FBgn0030930	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2
Dmoj\GI17059	FBgn0139804	Ugt36Bc	FBgn0040260	
Dmoj\GI17058	FBgn0139803	Ugt37c1	FBgn0026754	UDP-glycosyltransferase 37c1
.,	FBgn0145358	N/A	N/A	N/A
Dmoj\Gl22630	3			
-	FBgn0140190	CG8708	FBgn0033271	
Dmoj\GI22630	-	CG8708 CG17324	FBgn0033271 FBgn0027074	
Dmoj\Gl22630 Dmoj\Gl17446	FBgn0140190 FBgn0143677		FBgn0027074	
Dmoj\Gl22630 Dmoj\Gl17446 Dmoj\Gl20943	FBgn0140190	CG17324	-	

-	<u> </u>	_	
FBgn0143818	alpha4GT2	FBgn0039378	
FBan0132968	CG10166	FBan0032799	
-		-	
_		-	
-		_	
_	-	-	Glycogen synthase
-	-	-	alpha1,6-fucosyltransferase
_		_	aipha 1,0-rucosyruansierase
-		-	Dheanhatidulineaital glucan anghar hisayathasia alaga M
-		-	Phosphatidylinositol glycan anchor biosynthesis class M
-		-	
-		-	h.dam.
-		-	twiggy
FBgn0137392	GlyP	FBgn0004507	Glycogen phosphorylase
			arms
-		-	Multidrug resistance protein 4
-		-	
-	CG10505	FBgn0034612	
FBgn0140692	CG9664	FBgn0031515	
FBgn0280797	Mdr50	FBgn0010241	Multi drug resistance 50
FBgn0133059	Mrp4	FBgn0263316	Multidrug resistance protein 4
FBgn0140015	CG17646	FBgn0264494	
FBgn0138962	CG4822	FBgn0031220	
FBgn0144967	CG9990	FBgn0039594	
FBgn0143840	CG8908	FBgn0034493	
FBgn0138087	Pmp70	FBgn0031069	Peroxisomal Membrane Protein 70 kDa
FBgn0142308	Mdr49	FBgn0004512	Multi drug resistance 49
-	MRP	FBgn0032456	Multidrug-Resistance like Protein 1
-	E23	-	Early gene at 23
-		_	
_	000040	_	
-		-	NA 101
_		-	Multidrug resistance protein 4
_		_	
_		-	
-		-	
-		-	
-		-	Early gene at 23
		-	
FBgn0138549	CG1703	FBgn0030321	
ED0440004	00-0	ED0000470	O. d. aharara
-		-	Cytochrome P450-6a2
		-	Cytochrome P450-6a2
-		-	la mi
-		-	Multidrug resistance protein 4
-		-	Urate oxidase
FBgn0135704		FBgn0036750	
		-	
FBgn0147462	Cyp6d5	-	
FBgn0253215	N/A	N/A	N/A
FBgn0253216	N/A	N/A	N/A
FBgn0253199	N/A	N/A	N/A
FBgn0253266	N/A	N/A	N/A
FBgn0253226	N/A	N/A	N/A
	CG3699	FBgn0040349	
FBgn0137748	00000		
FBgn0137748 FBgn0280296	CG34330	FBgn0085359	
-		FBgn0085359 FBgn0025583	Immune induced molecule 2
FBgn0280296	CG34330	-	Immune induced molecule 2
FBgn0280296 FBgn0142851	CG34330 IM2	FBgn0025583	Immune induced molecule 2 N/A
FBgn0280296 FBgn0142851 FBgn0145193	CG34330 IM2 CG4860	FBgn0025583 FBgn0037999	
FBgn0280296 FBgn0142851 FBgn0145193 FBgn0280701	CG34330 IM2 CG4860 N/A	FBgn0025583 FBgn0037999 N/A	
FBgn0280296 FBgn0142851 FBgn0145193 FBgn0280701 FBgn0142304	CG34330 IM2 CG4860 N/A CG13323	FBgn0025583 FBgn0037999 N/A FBgn0033788	
FBgn0280296 FBgn0142851 FBgn0145193 FBgn0280701 FBgn0142304 FBgn0141777	CG34330 IM2 CG4860 N/A CG13323 CG3292	FBgn0025583 FBgn0037999 N/A FBgn0033788 FBgn0034710	N/A
	FBgn0132968 FBgn0280907 FBgn0134261 FBgn0132886 FBgn0139068 FBgn0139068 FBgn0139716 FBgn0136709 FBgn0134051 FBgn0134051 FBgn0137392 FBgn0133058 FBgn0139902 FBgn0142482 FBgn0140692 FBgn013059 FBgn013059 FBgn013059 FBgn013059 FBgn0140015 FBgn0138867 FBgn0138962 FBgn014967 FBgn0138087 FBgn013809 FBgn013059 FBgn013059 FBgn0145400 FBgn013059 FBgn014500 FBgn013059 FBgn014500	FBgn0143818 alpha4GT2 FBgn0132968 CG10166 FBgn0280907 CG31915 FBgn0132861 CG9384 FBgn0132886 Ugt86Da FBgn0139068 FucT6 FBgn0139068 FucT6 FBgn0139716 PIG-M FBgn0136709 CG11306 FBgn0134051 CG9109 FBgn0133052 GlyP FBgn0137392 GlyP FBgn0133058 Mrp4 FBgn0139902 CG7806 FBgn0140692 CG9664 FBgn0139902 CG9664 FBgn013059 Mrp4 FBgn0133059 Mrp4 FBgn0133059 Mrp4 FBgn0138962 CG4822 FBgn0138962 CG4822 FBgn0138087 Pmp70 FBgn0142308 Mdr49 FBgn0138667 E23 FBgn0138667 E23 FBgn0138940 CG3164 FBgn0138940 CG3164 FBgn013867 E23 <t< td=""><td>FBgn0143818 alpha4GT2 FBgn0039378 FBgn0132968 CG10166 FBgn0032799 FBgn0280907 CG31915 FBgn0051915 FBgn0134261 CG9384 FBgn0036446 FBgn0132886 Ugt86Da FBgn0036446 FBgn0139068 FucT6 FBgn0030327 FBgn0139068 FucT6 FBgn0039273 FBgn0139716 PIG-M FBgn0034649 FBgn0137709 CG11306 FBgn0037108 FBgn0134051 CG9109 FBgn0037108 FBgn0133058 Mrp4 FBgn0032018 FBgn0133058 Mrp4 FBgn0032018 FBgn0133058 Mrp4 FBgn0032018 FBgn0142482 CG10505 FBgn0032018 FBgn0142482 CG10505 FBgn0034612 FBgn0133059 Mrp4 FBgn0263316 FBgn0133059 Mrp4 FBgn0263316 FBgn0133059 Mrp4 FBgn0263316 FBgn0142462 CG4822 FBgn0032202 FBgn01438962 CG4822 FBgn0033493 FBgn0143</td></t<>	FBgn0143818 alpha4GT2 FBgn0039378 FBgn0132968 CG10166 FBgn0032799 FBgn0280907 CG31915 FBgn0051915 FBgn0134261 CG9384 FBgn0036446 FBgn0132886 Ugt86Da FBgn0036446 FBgn0139068 FucT6 FBgn0030327 FBgn0139068 FucT6 FBgn0039273 FBgn0139716 PIG-M FBgn0034649 FBgn0137709 CG11306 FBgn0037108 FBgn0134051 CG9109 FBgn0037108 FBgn0133058 Mrp4 FBgn0032018 FBgn0133058 Mrp4 FBgn0032018 FBgn0133058 Mrp4 FBgn0032018 FBgn0142482 CG10505 FBgn0032018 FBgn0142482 CG10505 FBgn0034612 FBgn0133059 Mrp4 FBgn0263316 FBgn0133059 Mrp4 FBgn0263316 FBgn0133059 Mrp4 FBgn0263316 FBgn0142462 CG4822 FBgn0032202 FBgn01438962 CG4822 FBgn0033493 FBgn0143

Dmoi\GI10427	EPan01/2165	PPO2	EPan0033367	Prophopolovidaco 2
Dmoj\GI19427	FBgn0142165		FBgn0033367	Prophenoloxidase 2
Dmoj\GI13492	FBgn0136248	CG6839	FBgn0036831	WDEL (
Dmoj\GI20444	FBgn0143180	KdelR	FBgn0267330	KDEL receptor
Dmoj\GI13547	FBgn0136302	CG9629	FBgn0036857	
Dmoj\GI24288	FBgn0147011	CG31233	FBgn0051233	
Dmoj\GI13889	FBgn0136643	CG14989	FBgn0035495	
Dmoj\GI18891	FBgn0141630	Cht4	FBgn0022700	Chitinase 4
Dmoj\GI11911	FBgn0134669	CG11796	FBgn0036992	
Dmoj\GI23284	FBgn0146011	CG6277	FBgn0039475	
Dmoj\GI23290	FBgn0146017	CG6283	FBgn0039474	
Dmoj\GI24854	FBgn0147574	CG15533	FBgn0039768	
Dmoj\GI23289	FBgn0146016	N/A	N/A	N/A
Dmoj\Gl24076	FBgn0146799	CG14718	FBgn0037939	TWA
-	-		-	
Dmoj\Gl20529	FBgn0143265	loopin-1	FBgn0259795	
Dmoj\Gl22439	FBgn0145167	CG4836	FBgn0270925	
Dmoj\GI23599	FBgn0146325	N/A	N/A	N/A
Dmoj\GI12914	FBgn0135671	S-Lap1	FBgn0035915	Sperm-Leucylaminopeptidase 1
Dmoj\GI18241	FBgn0140980	b	FBgn0000153	black
Dmoj\GI10839	FBgn0133602	е	FBgn0000527	ebony
Dmoj\GI11700	FBgn0134459	CG7906	FBgn0036417	
Dmoj\GI11699	FBgn0134458	CG7924	FBgn0036416	
Dmoj\GI24176	FBgn0146899	CG13606	FBgn0039161	
Dmoj\GI20472	FBgn0143208	CG9822	FBgn0034623	
Dmoj\GI13134	FBgn0135891	Eip78C	FBgn0004865	Ecdysone-induced protein 78C
Dmoj\GI20285	FBgn0143021	Achl	FBgn0033936	Achilles
-	-			Actinics
Dmoj\GI17558	FBgn0140302	Cyp28a5	FBgn0028940	
Dmoj\GI16995	FBgn0139741	CG34447	FBgn0085476	
Dmoj\GI11772	FBgn0134531	N/A	N/A	N/A
Dmoj\GI13604	FBgn0136359	CG34244	FBgn0085273	
Dmoj\Gl22046	FBgn0144775	CG4408	FBgn0039073	
Dmoj\GI20266	FBgn0143002	kermit	FBgn0010504	
Dmoj\GI10664	FBgn0133428	N/A	N/A	N/A
Dmoj\GI10838	FBgn0133601	е	FBgn0000527	ebony
Dmoj\GI12534	FBgn0135291	CG16758	FBgn0035348	
Dmoj\GI23038	FBgn0145765	Npc2b	FBgn0038198	Niemann-Pick type C-2b
Dmoj\GI11471	FBgn0134232	S-Lap3	FBgn0045770	Sperm-Leucylaminopeptidase 3
Dmoj\GI21075	FBgn0143807	CG9253	FBgn0032919	CPOINT 2000).ummopopuudoo 0
Dmoj\GI17762	FBgn0140503	CG3513	FBgn0031559	
-				NI/A
Dmoj\GI11244	FBgn0134005	N/A	N/A	N/A
Dmoj\GI12225	FBgn0134982	mex1	FBgn0004228	midgut expression 1
Dmoj\GI23278	FBgn0146005	Cpr97Eb	FBgn0039481	Cuticular protein 97Eb
Dmoj\GI16741	FBgn0139489	CG33282	FBgn0053282	
Dmoj\Gl25812	FBgn0280549	N/A	N/A	N/A
Dmoj\Gl26094	FBgn0280831	CecC	FBgn0000279	Cecropin C
Dmoj\Gl26684	FBgn0281421	N/A	N/A	N/A
Dmoj\GI24064	FBgn0146787	CG12224	FBgn0037974	
Dmoj\GI20159	FBgn0142895	CG15098	FBgn0034398	
Dmoj\GI10658	FBgn0133422	N/A	N/A	N/A
Dmoj\GI20083	FBgn0142819	CG13748	FBgn0033355	
Dmoj\Gl22425	FBgn0145153	N/A	N/A	N/A
				1973
Dmoj\Gl23250	FBgn0145977	poly	FBgn0086371	DNA askersana II alamatina fastar
Dmoj\GI17330	FBgn0140074	TfIIS	FBgn0010422	RNA polymerase II elongation factor
Dmoj\GI19218	FBgn0141957	CG33998	FBgn0053998	
Dmoj\GI13751	FBgn0136506	Cpr78E	FBgn0037114	Cuticular protein 78E
Dmoj\GI10662	FBgn0133426	N/A	N/A	N/A
Dmoj\GI14221	FBgn0136974	Gbeta13F	FBgn0001105	G protein beta-subunit 13F
Dmoj\Gl23249	FBgn0145976	Lip3	FBgn0023495	
Dmoj\GI12604	FBgn0135361	CG7953	FBgn0028533	
Dmoj\GI17402	FBgn0140146	CG8997	FBgn0028920	
Dmoj\GI23291	FBgn0146018	CG6295	FBgn0039471	
Dmoj\GI12593	FBgn0135350	CG7916	FBgn0028534	
Dmoj\G172393 Dmoj\G127063	FBgn0281800	N/A	N/A	N/A
-				
Dmoj\Gl26336	FBgn0281073	N/A	N/A	N/A
Dmoj\Gl20122	FBgn0142858	GstE8	FBgn0063492	Glutathione S transferase E8
Dmoj\GI12734	FBgn0135491	CG11350	FBgn0035552	
Dmoj\GI12733	FBgn0135490	CG7465	FBgn0035551	

Dmoj\GI15268	FBgn0138017	Shmt	FBgn0029823	Serine hydroxymethyl transferase
Dmoj\GI21533	FBgn0144263	Sec61gamma	FBgn0031049	Sec61 gamma subunit
Dmoj\Gl26057	FBgn0280794	N/A	N/A	N/A
Dmoj\GI22470	FBgn0145198	CG43673	FBgn0263748	
Dmoj\GI19511	FBgn0142249	Lcp9	FBgn0025578	Larval cuticle protein 9
Dmoj\GI20905	FBgn0143639	Lcp1	FBgn0002531	Larval cuticle protein 1
Dmoj\Gl25804	FBgn0280541	Lcp1	FBgn0002531	Larval cuticle protein 1
Dmoj\GI20899	FBgn0143633	Lcp1	FBgn0002531	Larval cuticle protein 1
Dmoj\GI16761	FBgn0139509	LysB	FBgn0004425	Lysozyme B
Dmoj\Gl20904	FBgn0143638	Lcp1	FBgn0002531	Larval cuticle protein 1
Dmoj\GI12682	FBgn0135439	ple	FBgn0005626	pale