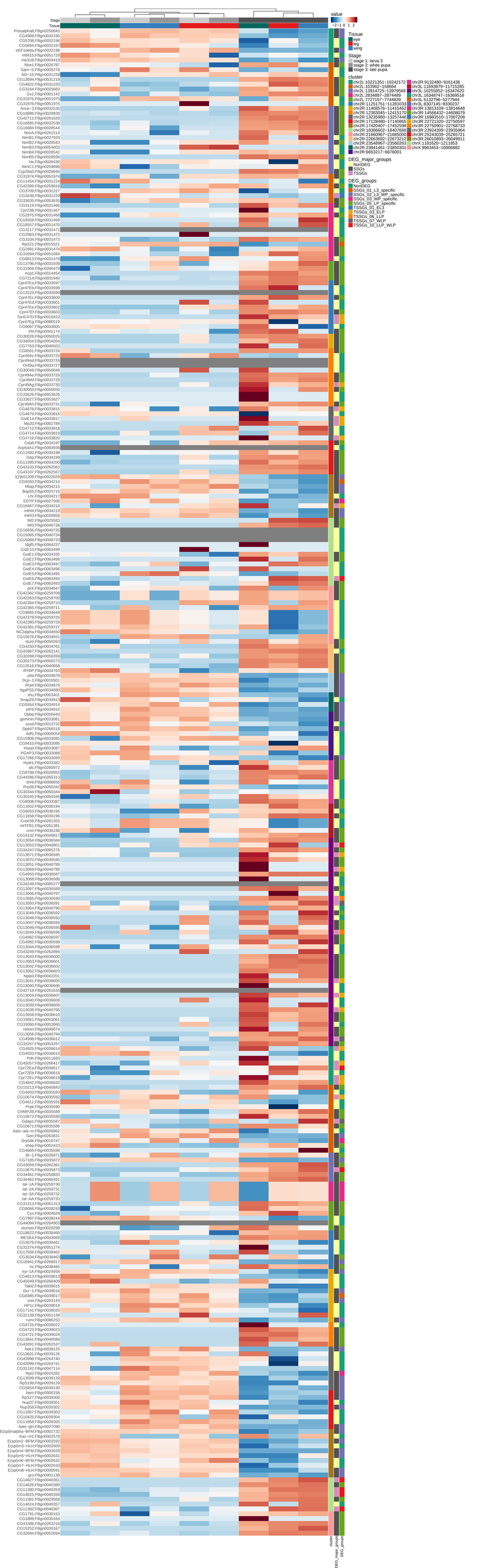
Stage
Tissue
Prosalphaδ;FBgn0250843
CG4908;FBgn0032195
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eEF1delta;FBgn0032198
mfrl15;FBgn0051720
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Nhe1;FBgn0026787
Sam−S;FBgn0005278
ND−15;FBgn0031228
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Gs1;FBgn0001142
CG3197;FBgn0051975
CG3197E;FBgn0051976
Arce−3;FBgn0032536
CG16886;FBgn0028938
CG42711;FBgn0261628
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NimA;FBgn0261514
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He;FBgn0028430
NimC1;FBgn0259896
Cyp28a5;FBgn0028940
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CG42399;FBgn0259818
CG3709;FBgn0031227
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CG3119;FBgn0031466
Cpr23B;FBgn0031467
CG2975;FBgn0031468
CG18558;FBgn0031469
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CG3117;FBgn0031471
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CG31904;FBgn0260479
Acp1;FBgn0014454
CG7214;FBgn0031940
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Cpr47Ec;FBgn0033600
Cpr47Ed;FBgn0033601
Cpr47Ee;FBgn0033602
Cpr47EF;FBgn0033603
TpnC47D;FBgn0010423
Cpr47Eg;FBgn0088519
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Vhl;FBgn0041174
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GstE14;FBgn0033817
Mp20;FBgn0002789
CG4712;FBgn0033818
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Cda9;FBgn0033197
Acp54A1;FBgn0083936
CG11400;FBgn0034198
Gbp;FBgn0034199
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stum;FBgn0050263
CG4250;FBgn0034761
CG42867;FBgn0262141
CG30269;FBgn0050269
CG30273;FBgn0050273
CG13516;FBgn0040658
RnBP;FBgn0034763
pita;FBgn0034878
Dcp−1;FBgn0010501
Rrp4;FBgn0034879
IgaPS5;FBgn0044883
shu;FBgn0003401
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eIF6;FBgn0034915
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geminin;FBgn0033081
sced;FBgn0013732
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Hsap;FBgn0033087
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Hydr1;FBgn0033382
alc;FBgn0260972
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CG44286;FBgn0265313
shrb;FBgn0086656
Pxp38;FBgn0050342
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CG8008;FBgn0033387
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Ccdc56;FBgn0261353
mtFER1;FBgn0261381
crim;FBgn0036198
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Br−17;FBgn0035871
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Taiz2;FBgn0039015
Dcr−1;FBgn0039016
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bam;FBgn0000158
Rps27;FBgn0039300
Nup37;FBgn0039301
Nup358;FBgn0039302
CG11857;FBgn0039303
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Aats−gln;FBgn0027090
E(sp)malpha−BFM;FBgn0002732
Kao−m17;FBgn0002578
E(sp)m2−BFM;FBgn0002592
E(sp)m3−HLH;FBgn0002609
E(sp)m4−BFM;FBgn0002629
E(sp)m5−HLH;FBgn0002631
E(sp)m6−BFM;FBgn0002632
E(sp)m7−HLH;FBgn0002633
E(sp)m8−HLH;FBgn0000591
gro;FBgn0001139
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CG11380;FBgn0040359
CG14625;FBgn0040358
CG11381;FBgn0029568
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CG11382;FBgn0040367
CG1791;FBgn0030163
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CG43386;FBgn0263216
CG15252;FBgn0030167
CG32694;FBgn0052694
value
−2 −1 0 1 2
Tissue
eye
leg
wing
Stage
stage 1: larva 3
stage 2: white pupa
stage 3: late pupa
cluster
chr2L:10221351−10242172
chr2L:1039962−158664
chr2L:13914725−13979569
chr2L:2834887−2874489
chr2L:7727157−7744839
chr2R:11251751−11281033
chr2R:12385576−11415462
chr2R:12383045−12415170
chr2R:13235980−13257446
chr2R:17128480−17140865
chr2R:17420407−17452598
chr2R:18386602−18407688
chr2R:21660967−21685000
chr2R:22663692−22673212
chr2R:23548987−23560263
chr2R:23841461−23850301
chr2R:6652217−6676001
chr2R:9132480−9161436
chr3L:11693879−11715285
chr3L:16255852−16347620
chr3L:16348747−16369518
chr3L:5132796−5277944
chr3L:8307145−8330237
chr3R:13813109−13814648
chr3R:14566432−14608079
chr3R:16983510−17007244
chr3R:22721320−22750597
chr3R:22750901−22768733
chr3R:23924399−23935964
chr3R:25243039−25265721
chr3R:26010893−26049911
chrX:1181620−1211853
chrX:9963453−10006882
DEG_major_groups
NonDEG
SSGs
TSSGs
DEG_groups
NonDEG
SSGs_01_L3_specific
SSGs_02_L3_WP_specific
SSGs_03_WP_specific
SSGs_05_LP_specific
TSSGs_01_EL3
TSSGs_03_ELP
TSSGs_06_LLP
TSSGs_07_WLP
TSSGs_10_LLP_WLP
cluster
DEG_major_groups
DEG_groups