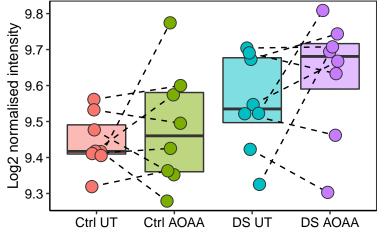
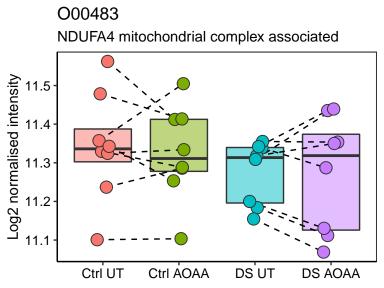
O00217

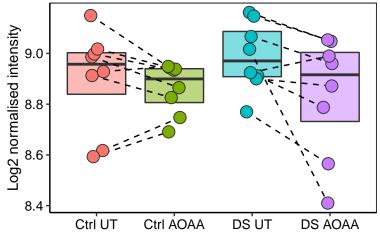
NADH:ubiquinone oxidoreductase core subunit S8



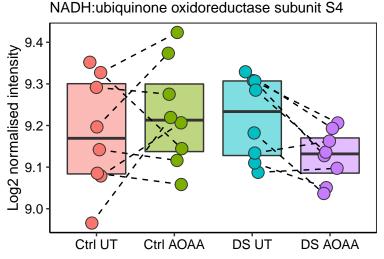


O14561

NADH:ubiquinone oxidoreductase subunit AB1



O43181

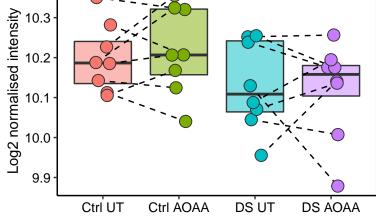


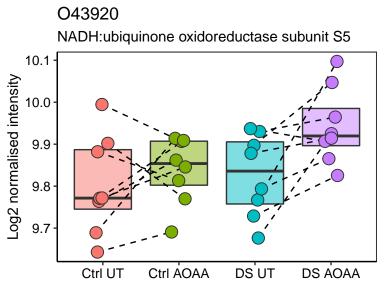
O43674 NADH: ubiquinone oxidoreductase subunit B5 og2 normalised intensity 10.1 10.0 9.9 9.8 Ctrl UT Ctrl AOAA DS AOAA DS UT

O43676 NADH: ubiquinone oxidoreductase subunit B3 10.4 Log2 normalised intensity 10.2 -10.0 9.8 Ctrl UT Ctrl AOAA DS UT DS AOAA

O43677 NADH: ubiquinone oxidoreductase subunit C1 Log2 normalised intensity 2.2.4 6.75 Ctrl AOAA DS UT Ctrl UT DS AOAA

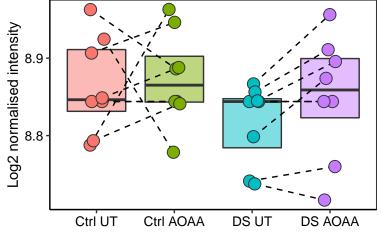
O43678 NADH: ubiquinone oxidoreductase subunit A2



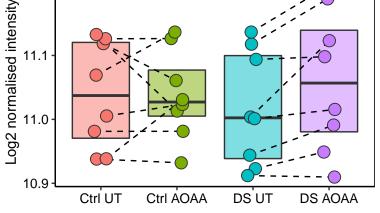


O75251

NADH:ubiquinone oxidoreductase core subunit S7



O75306 NADH:ubiquinone oxidoreductase core subunit S2 11.2 11.1



O75380 NADH: ubiquinone oxidoreductase subunit S6 10.6 og2 normalised intensity 10.5 10.4 -10.3-

Ctrl AOAA

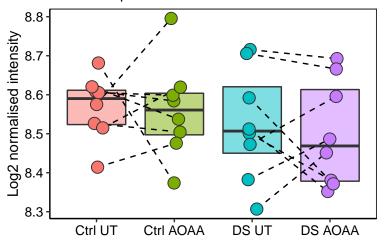
DS UT

DS AOAA

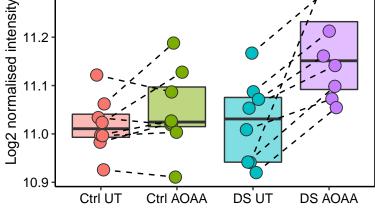
Ctrl UT

O75438

NADH:ubiquinone oxidoreductase subunit B1

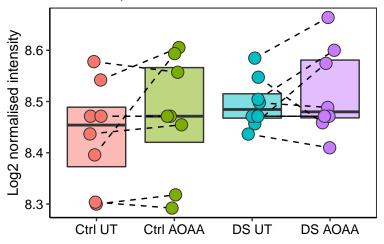


O75489 NADH:ubiquinone oxidoreductase core subunit S3 11.3 11.2

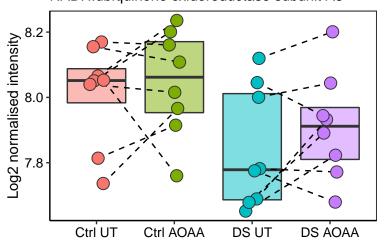


O95139

NADH:ubiquinone oxidoreductase subunit B6

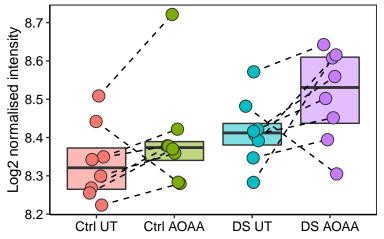


O95167
NADH:ubiquinone oxidoreductase subunit A3



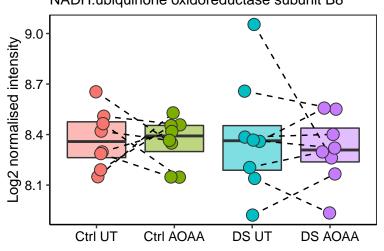
O95168

NADH:ubiquinone oxidoreductase subunit B4

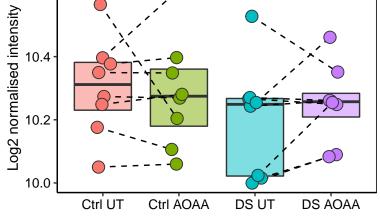


O95169

NADH:ubiquinone oxidoreductase subunit B8

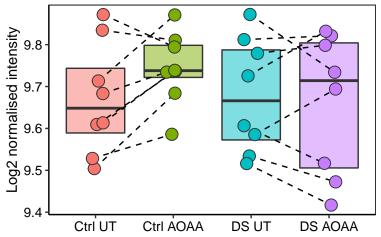


O95182
NADH:ubiquinone oxidoreductase subunit A7



O95298

NADH:ubiquinone oxidoreductase subunit C2



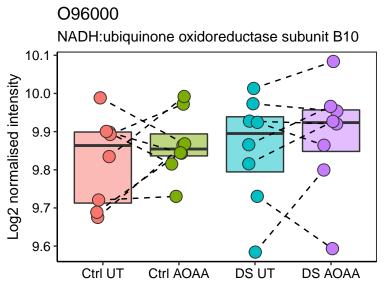
O95299 NADH: ubiquinone oxidoreductase subunit A10 11.2 -og2 normalised intensity 11.0 -10.9 -

Ctrl AOAA

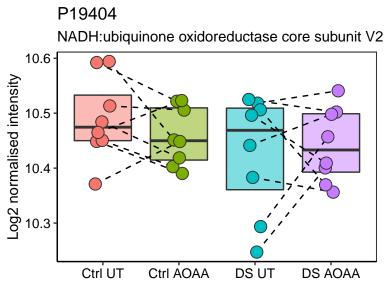
Ctrl UT

DS UT

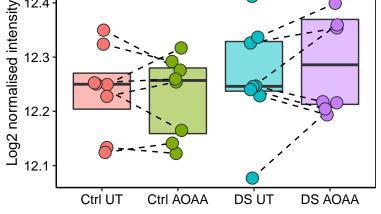
DS AOAA



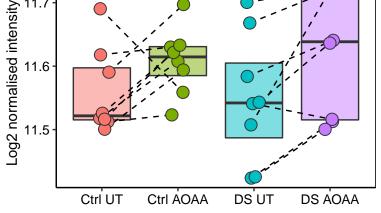
P17568 NADH: ubiquinone oxidoreductase subunit B7 10.5 Log2 normalised intensity 10.3 10.1 9.9 9.7 Ctrl AOAA DS UT DS AOAA Ctrl UT



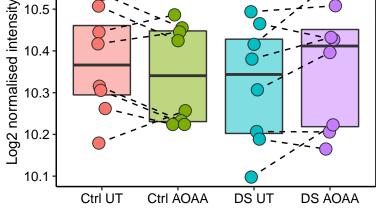
P28331
NADH:ubiquinone oxidoreductase core subunit S1

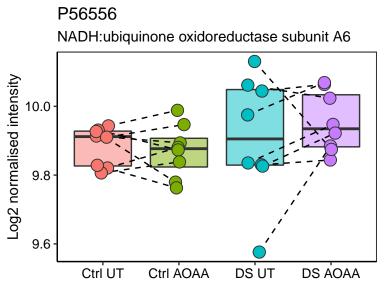


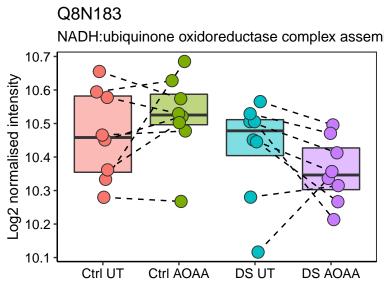
P49821
NADH:ubiquinone oxidoreductase core subunit V1



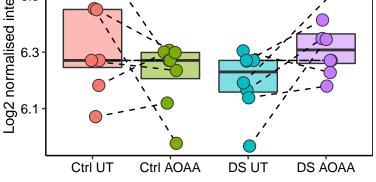
P51970 NADH:ubiquinone oxidoreductase subunit A8 10.5 10.4 -10.3



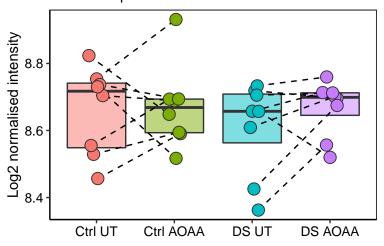




Q9BU61 NADH:ubiquinone oxidoreductase complex assemb -og2 normalised intensity 19 9 59 19 69 6.5 6.3

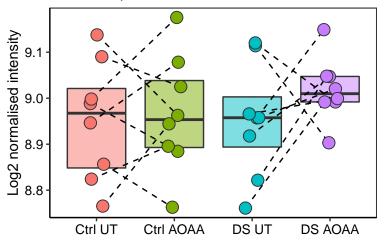


Q9NX14
NADH:ubiquinone oxidoreductase subunit B11

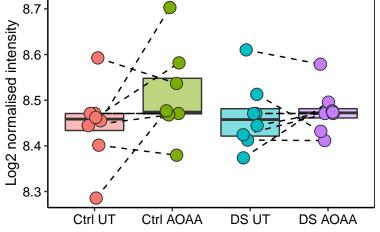


Q9P0J0

NADH:ubiquinone oxidoreductase subunit A13

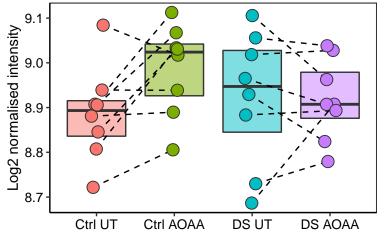


Q9P032
NADH:ubiquinone oxidoreductase complex assemb



Q9UI09

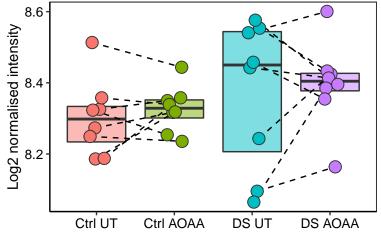
NADH: ubiquinone oxidoreductase subunit A12



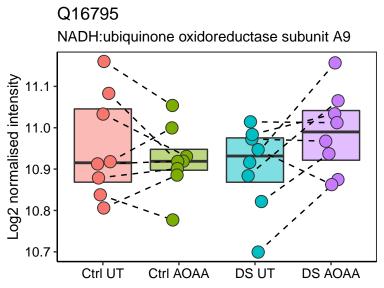
Q9Y6M9 NADH:ubiquinone oxidoreductase subunit B9 9.5 Log2 normalised intensity 9.4 9.3 9.2 Ctrl UT Ctrl AOAA DS AOAA DS UT

Q86Y39

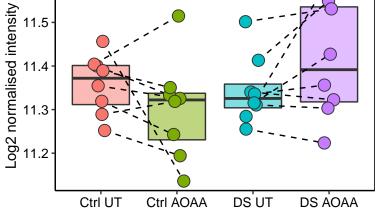
NADH:ubiquinone oxidoreductase subunit A11

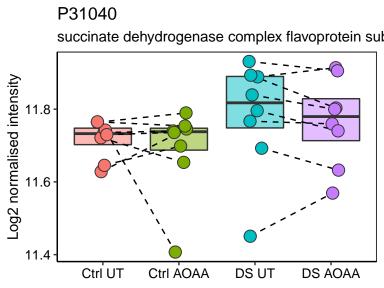


Q16718 NADH: ubiquinone oxidoreductase subunit A5 10.5 -Log2 normalised intensity
0.01
7.01
7.02 10.4 10.2 Ctrl AOAA Ctrl UT DS UT DS AOAA

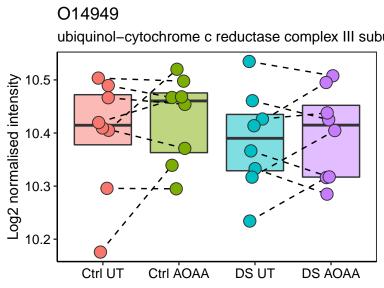


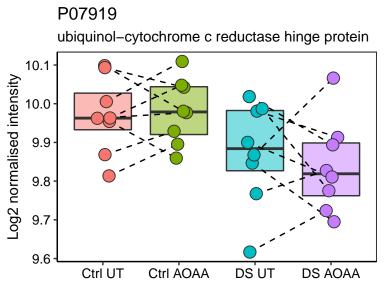
P21912 succinate dehydrogenase complex iron sulfur subu

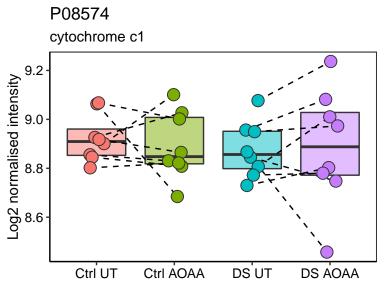


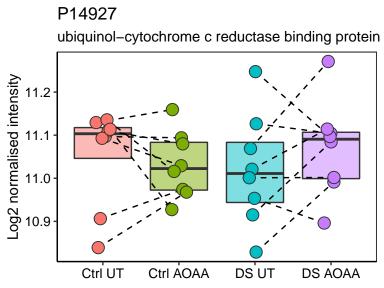


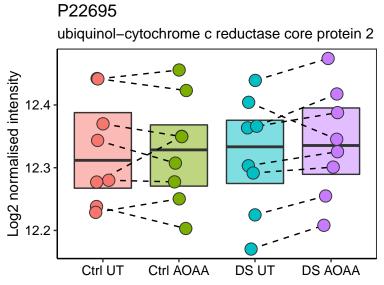
Q99643 succinate dehydrogenase complex subunit C 7.4 -0g2 normalised intensity 2.2 2.1 2.0 7.2 6.9 Ctrl UT Ctrl AOAA DS UT DS AOAA

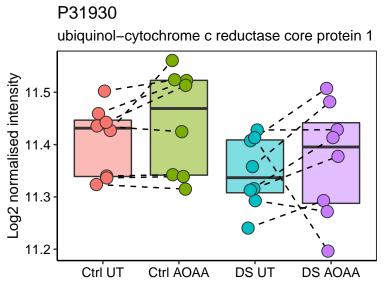


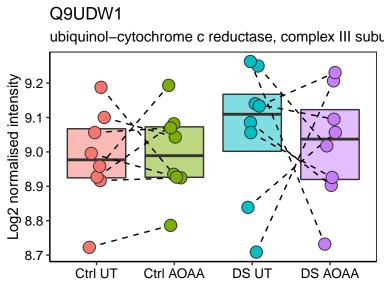


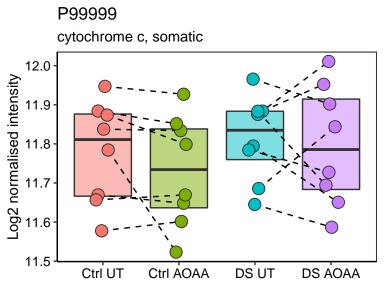




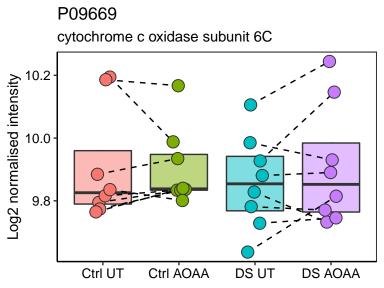


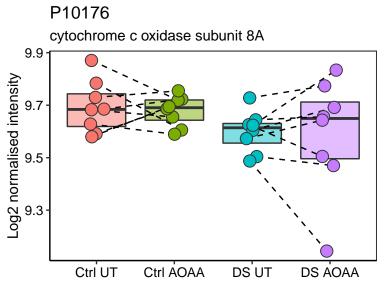


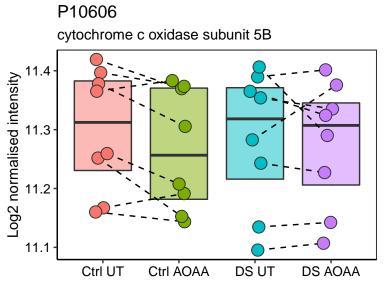




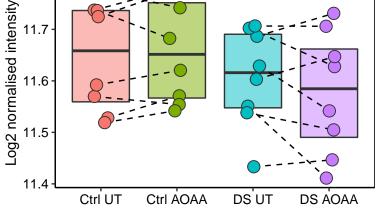
O14548 cytochrome c oxidase subunit 7A2 like 10.3 -og2 normalised intensity 10.2 10.1 10.0 -9.9 Ctrl UT Ctrl AOAA DS AOAA DS UT

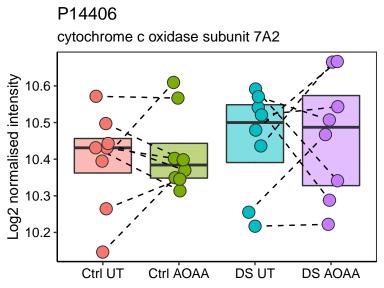


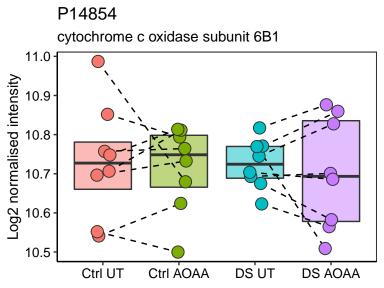




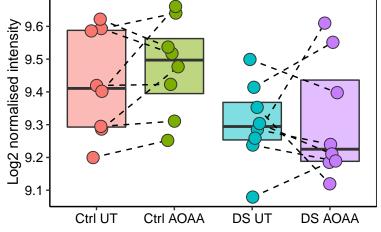
P13073 cytochrome c oxidase subunit 411



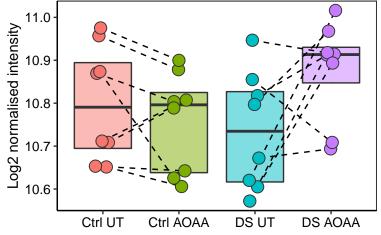


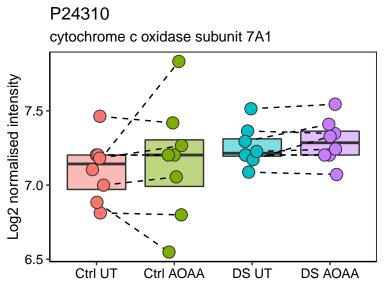


P15954 cytochrome c oxidase subunit 7C

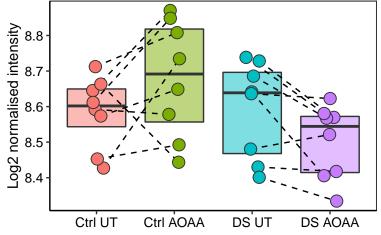


P20674 cytochrome c oxidase subunit 5A





P24311 cytochrome c oxidase subunit 7B



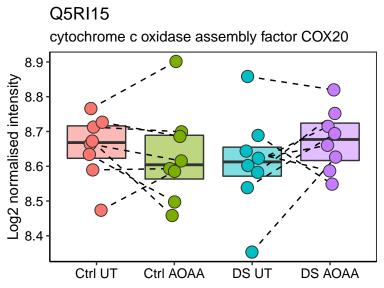
Q5JTJ3 cytochrome c oxidase assembly factor 6

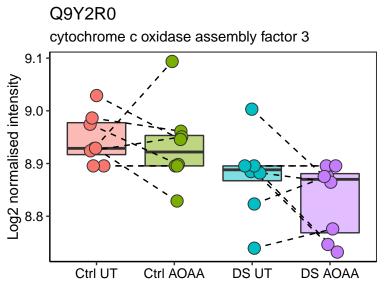
Ctrl AOAA

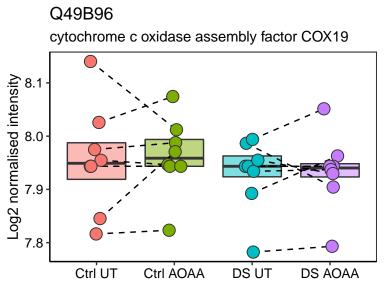
DS UT

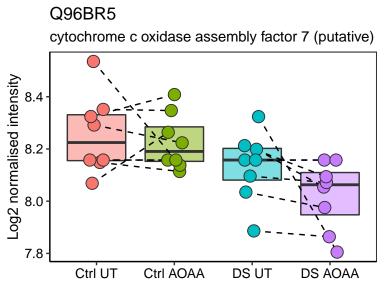
DS AOAA

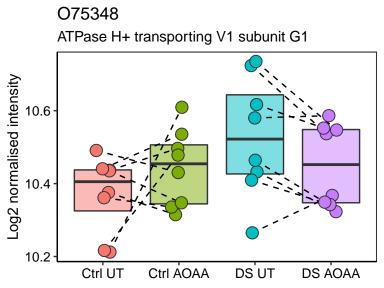
Ctrl UT



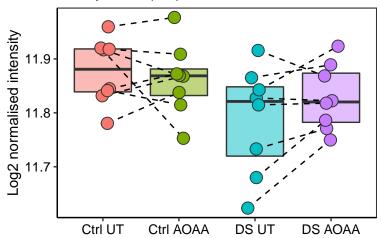


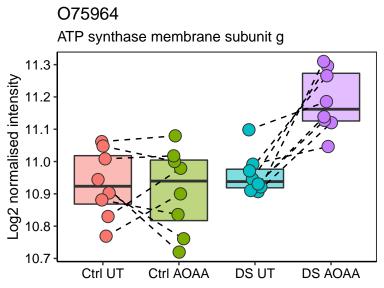






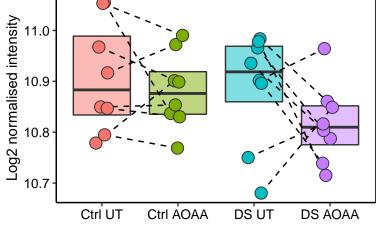
O75947 ATP synthase peripheral stalk subunit d



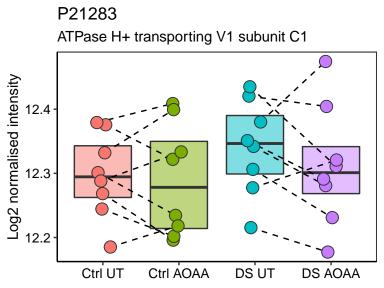


P06576 ATP synthase F1 subunit beta Log2 normalised intensity
13.2. Ctrl UT Ctrl AOAA DS UT DS AOAA

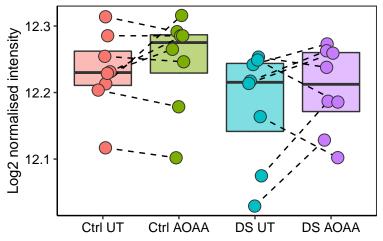
P18859 chr21 ATP synthase peripheral stalk subunit F6

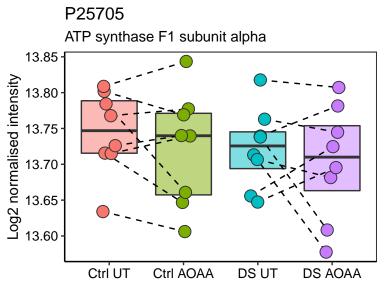


P21281 ATPase H+ transporting V1 subunit B2 12.4 -Log2 normalised intensity
15
7
7
7
7 Ctrl UT DS UT DS AOAA Ctrl AOAA

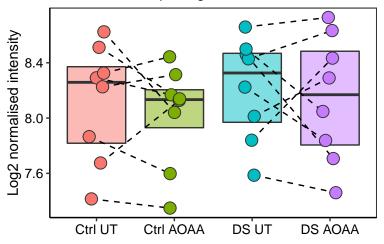


P24539
ATP synthase peripheral stalk–membrane subunit

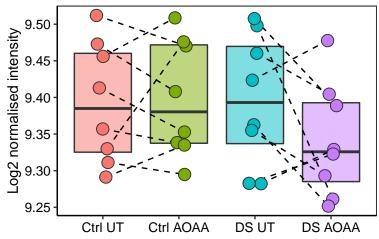




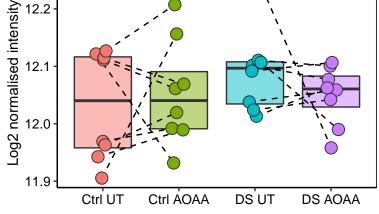
P27449
ATPase H+ transporting V0 subunit c

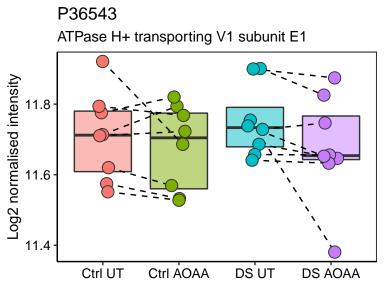


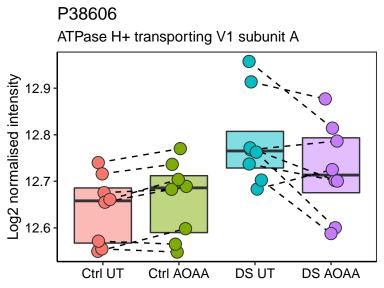
P30049
ATP synthase F1 subunit delta



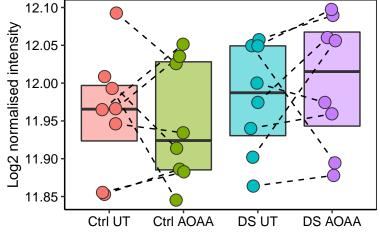
P36542 ATP synthase F1 subunit gamma



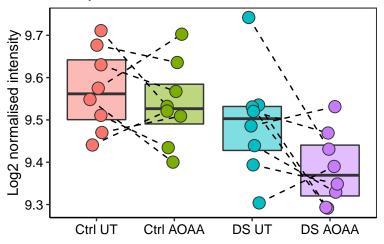




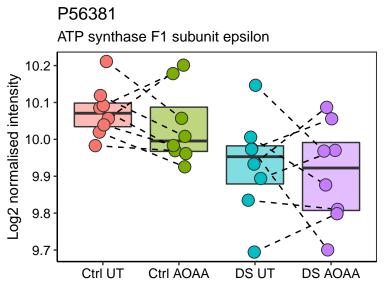
P48047 chr21
ATP synthase peripheral stalk subunit OSCP



P48201
ATP synthase membrane subunit c locus 3



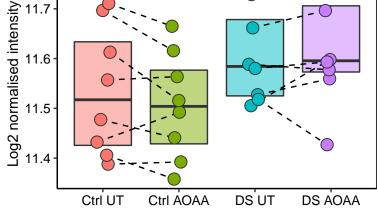
P56134 ATP synthase membrane subunit f Log2 normalised intensity 9.4 9.2 9.0 Ctrl UT Ctrl AOAA DS UT DS AOAA

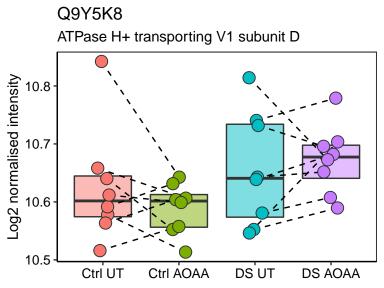


P56385 ATP synthase membrane subunit e 10.8 Log2 normalised intensity
0
7
9
9 10.5 Ctrl UT Ctrl AOAA DS UT DS AOAA

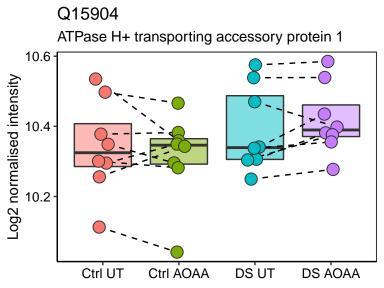
P61421 ATPase H+ transporting V0 subunit d1 Log2 normalised intensity 12.0 11.9 11.8 11.7 Ctrl AOAA DS UT DS AOAA Ctrl UT

Q9UI12 ATPase H+ transporting V1 subunit H 11.6-11.5 -

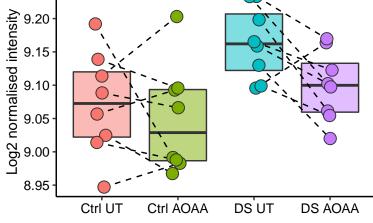




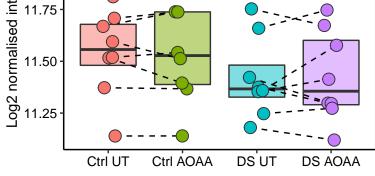
Q9Y487 ATPase H+ transporting V0 subunit a2 10.6 -Log2 normalised intensity 10.2 Ctrl AOAA DS UT DS AOAA Ctrl UT



Q16864 ATPase H+ transporting V1 subunit F



Q93050 ATPase H+ transporting V0 subunit a1 12.00 Log2 normalised intensity 11.75 - 11.50 - 11.25 - 11.2



Q99766 distal membrane arm assembly complex 2 like

