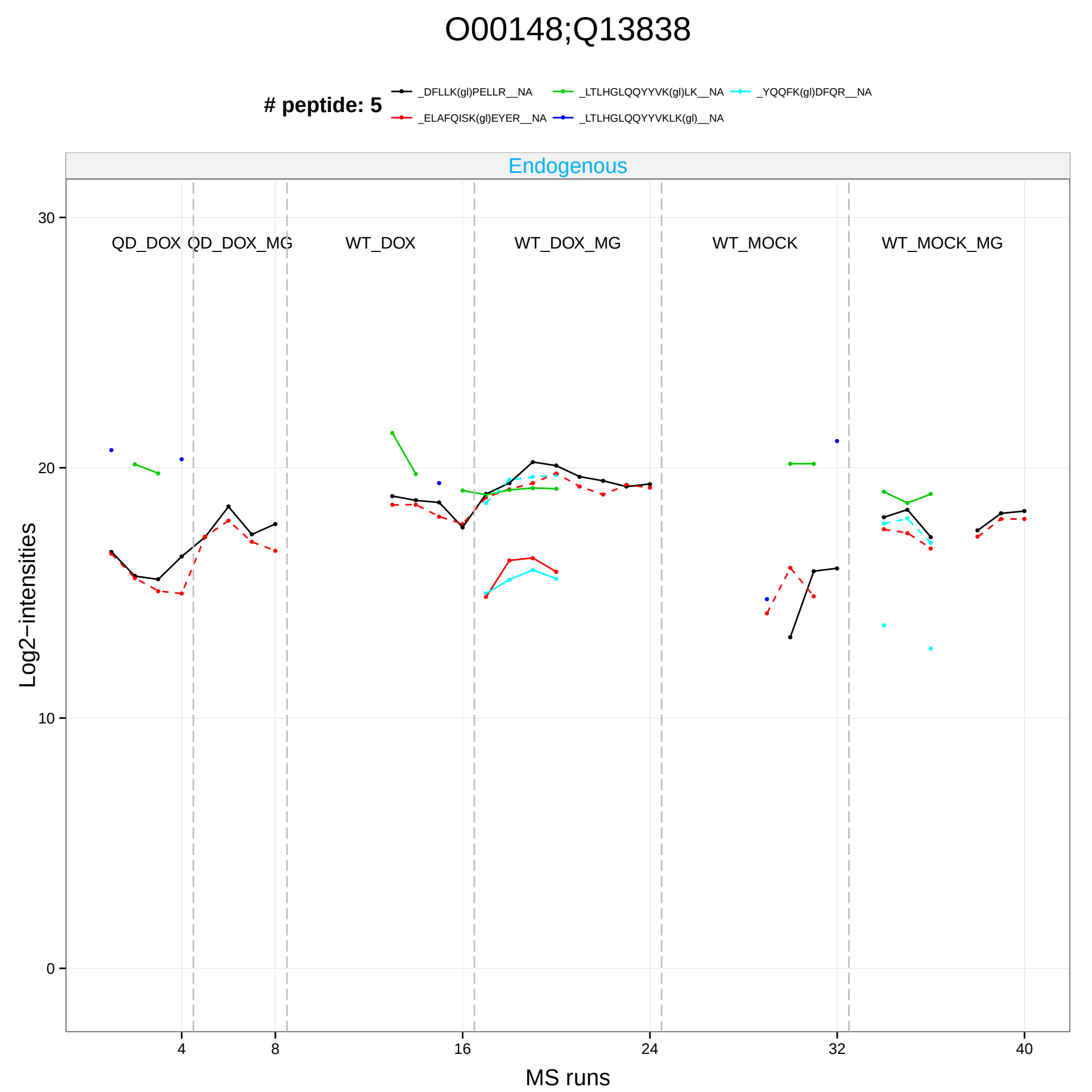
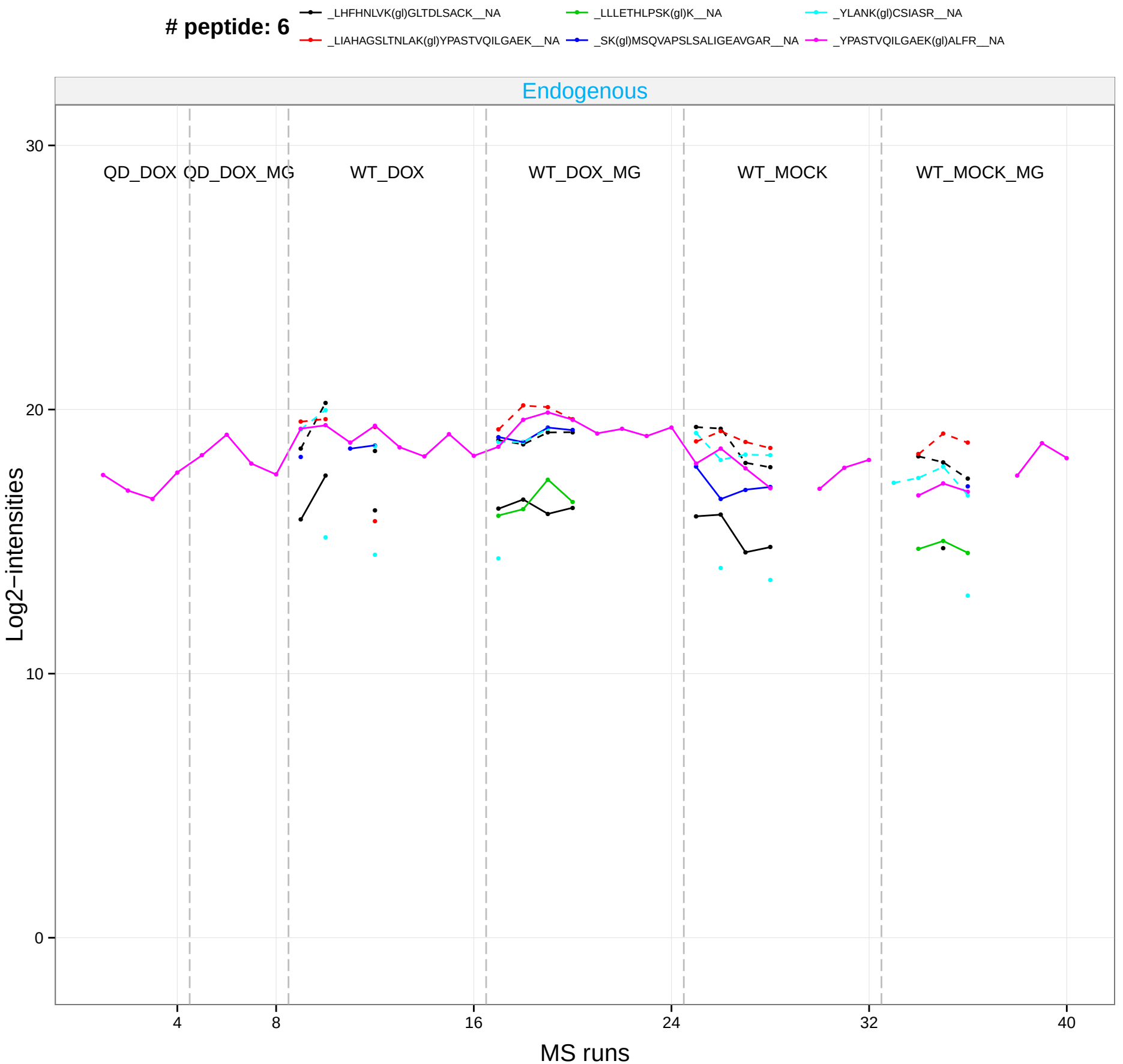
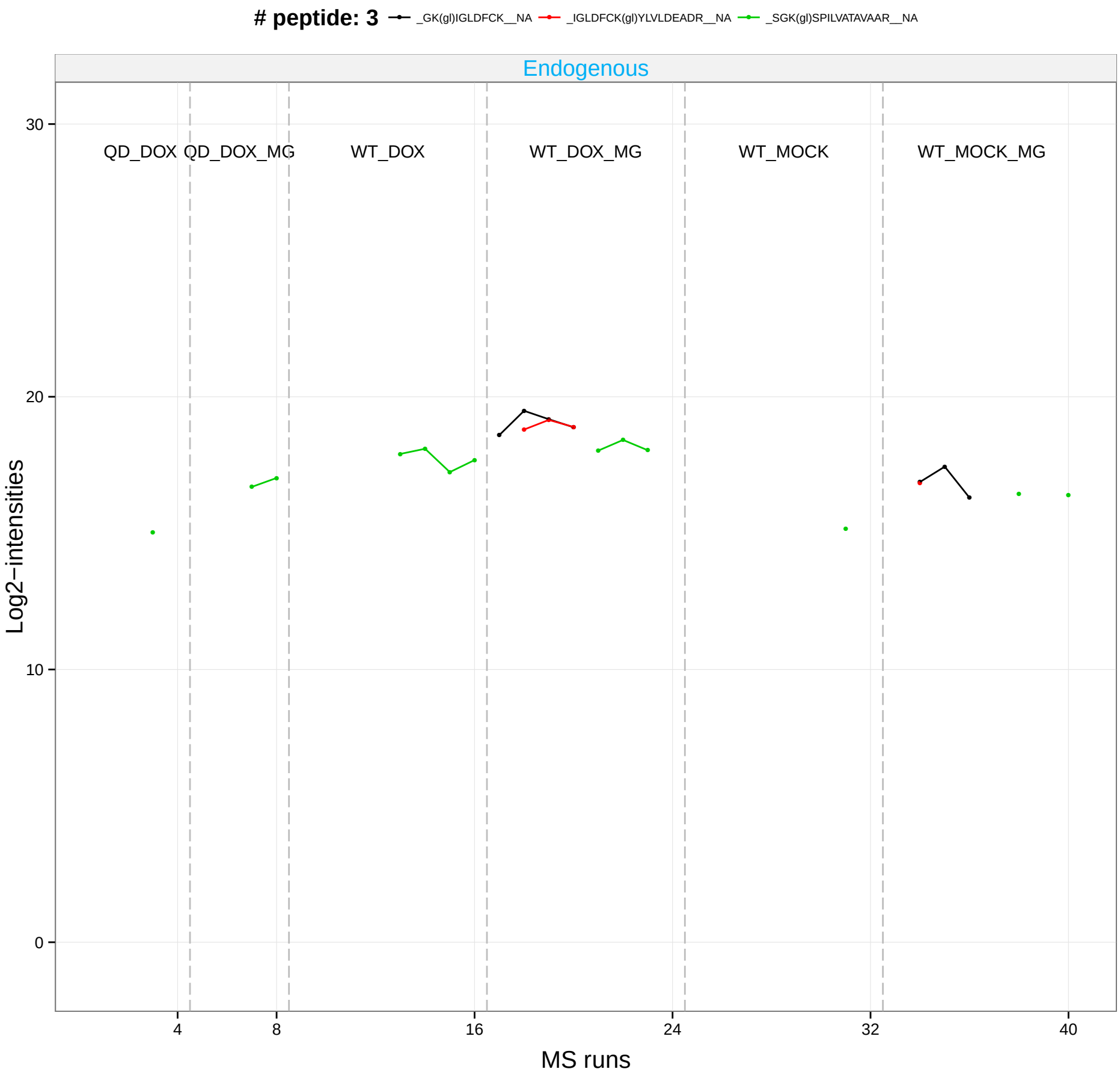

O00148;Q13838
peptide: 5
_DFLLK(gl)PELLR__NA
_LTLHGLQQYYVK(gl)LK__NA
_YQQFK(gl)DFQR__NA
_ELAFQISK(gl)EYER__NA
_LTLHGLQQYYVKLK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

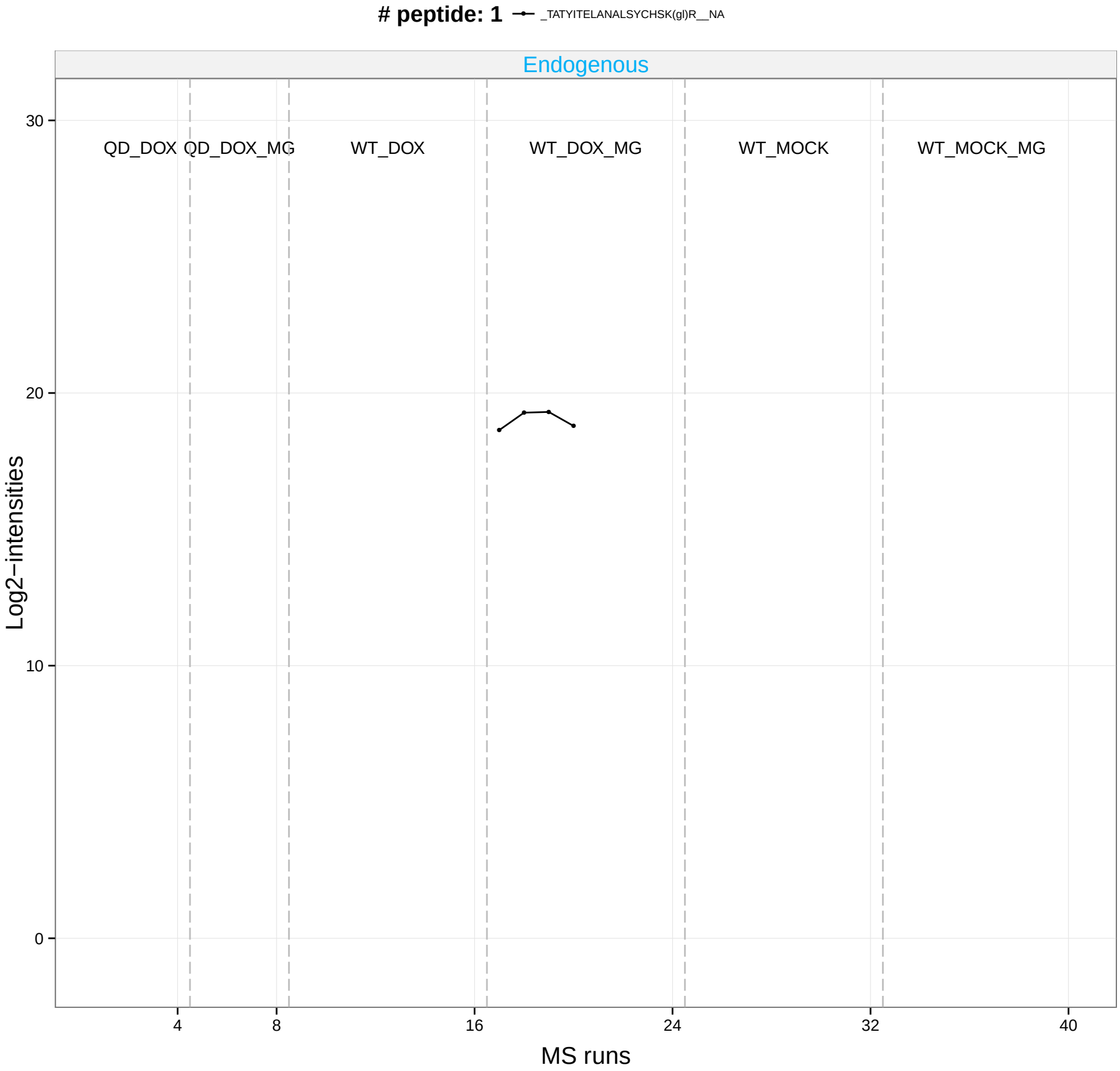
O00567


peptide: 6
_LHFHNLVK(gl)GLTDLSACK__NA
_LLLETHLPSK(gl)K__NA
_YLANK(gl)CSIASR__NA
_LIAHAGSLTNLAK(gl)YPASTVQILGAEK__NA
_SK(gl)MSQVAPSLSALIGEAVGAR__NA
_YPASTVQILGAEK(gl)ALFR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

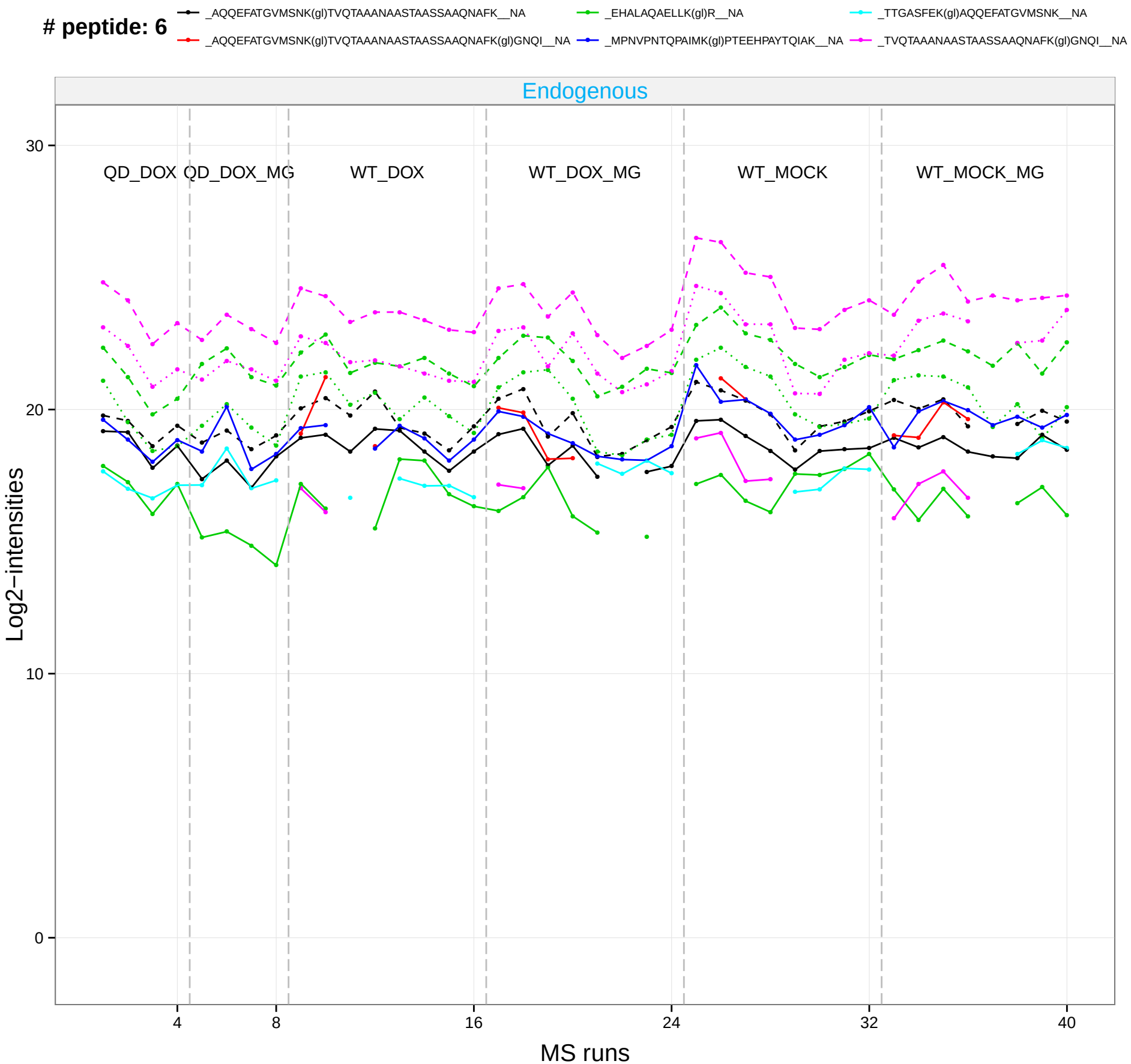
O00571;O15523



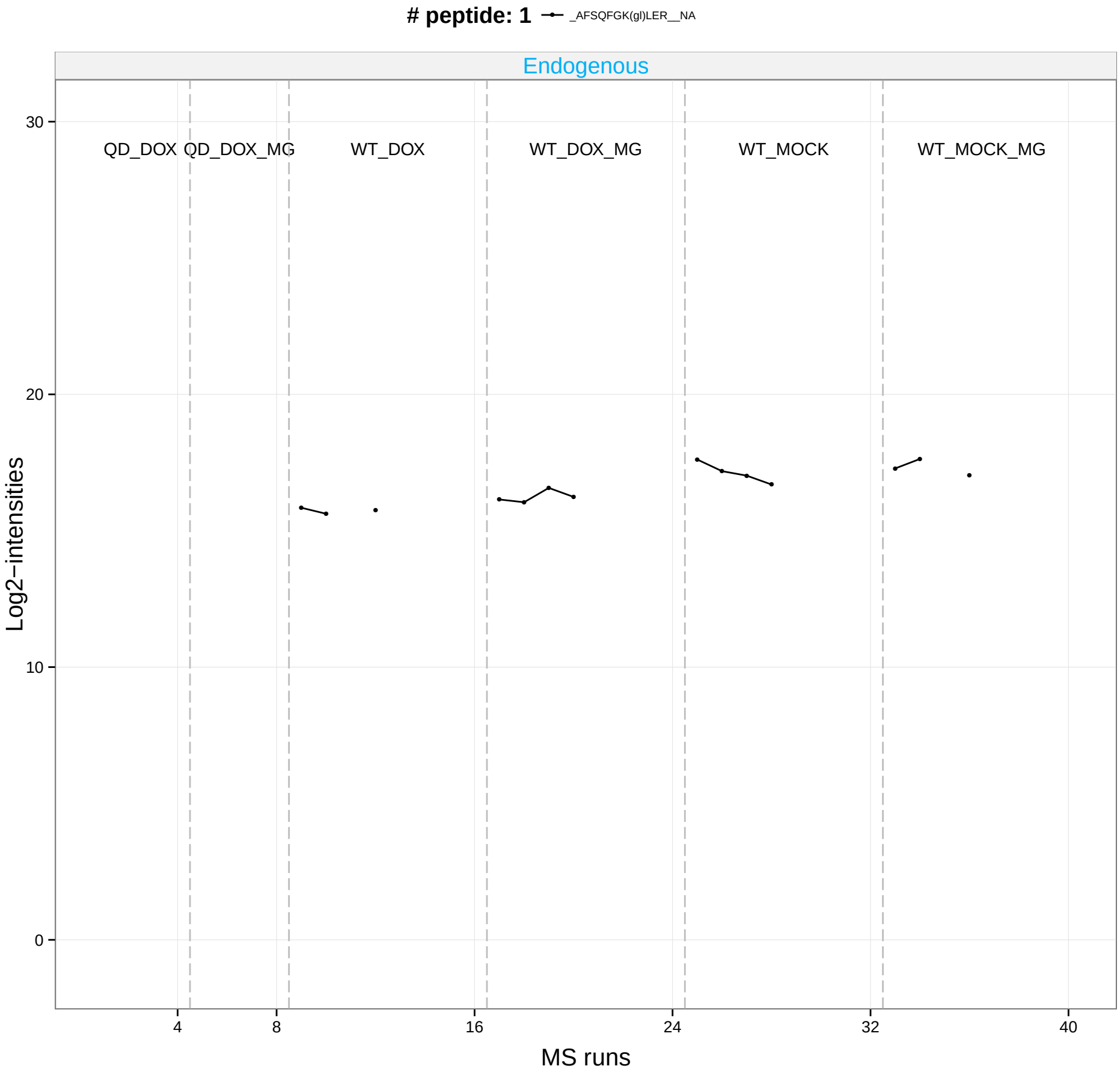
O14965


peptide: 1
_TATYITELANALSYCHSK(gl)R__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

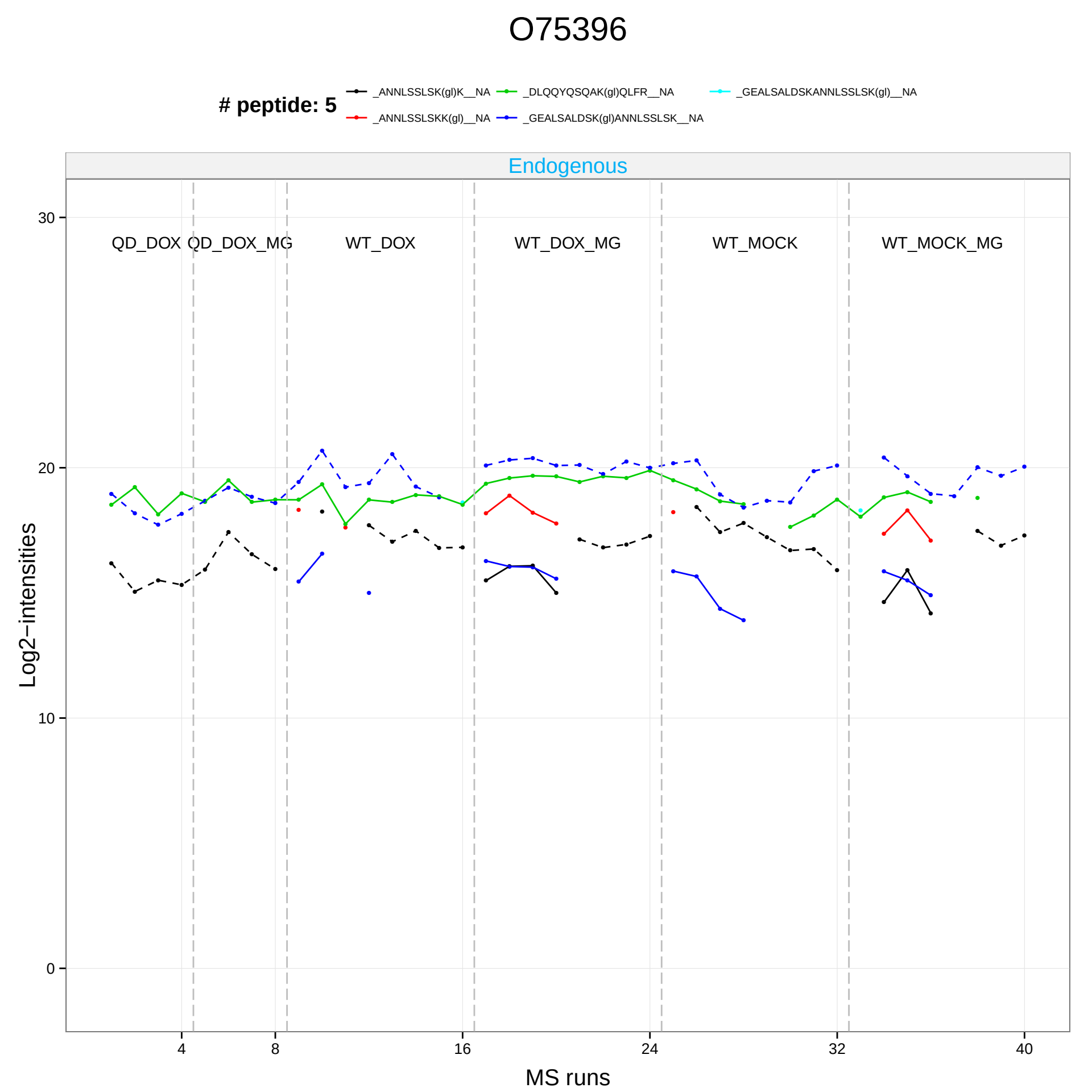
O15126


peptide: 6
_AQQEFATGVMSNK(gl)TVQTAAANAASTAASSAAQNAFK__NA
_EHALAQAELLK(gl)R__NA
_TTGASFEK(gl)AQQEFATGVMSNK__NA
_AQQEFATGVMSNK(gl)TVQTAAANAASTAASSAAQNAFK(gl)GNQI__NA
_MPNVPNTQPAIMK(gl)PTEEHPAYTQIAK__NA
_TVQTAAANAASTAASSAAQNAFK(gl)GNQI__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

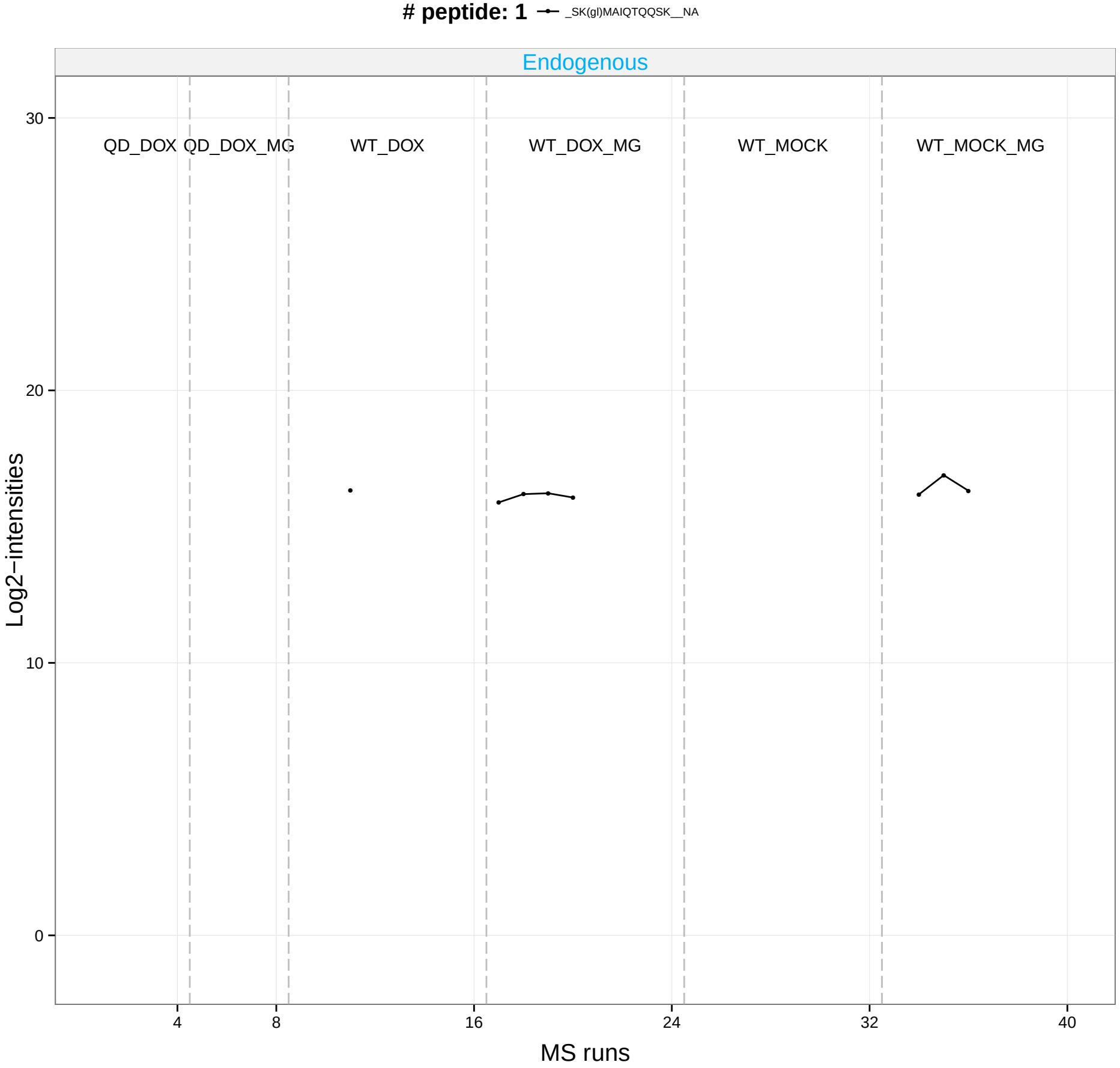
O60506


peptide: 1
_AFSQFGK(gl)LER__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

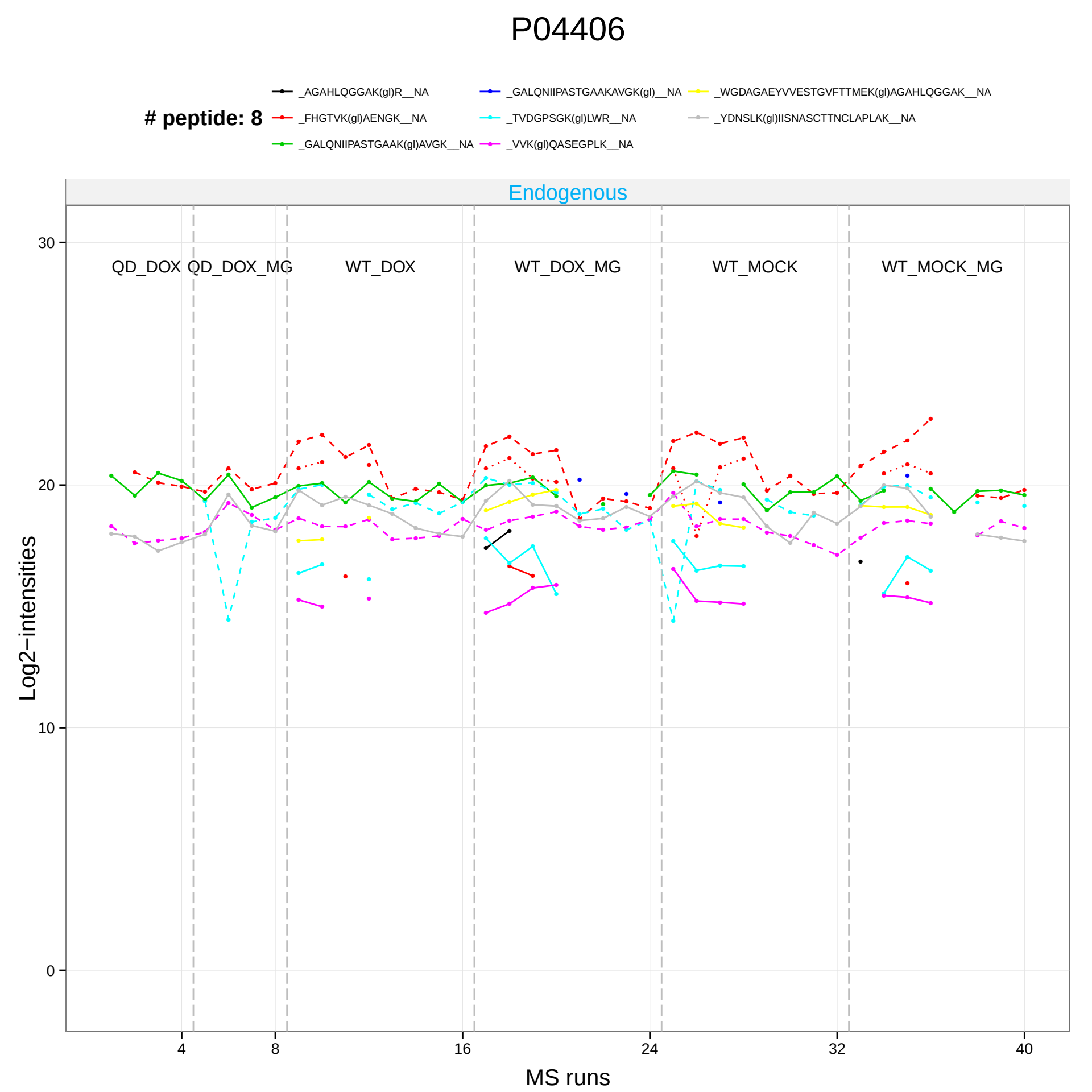
O75396


peptide: 5
_ANNLSSLSK(gl)K__NA
_DLQQYQSQAK(gl)QLFR__NA
_GEALSALDSKANNLSSLSK(gl)__NA
_ANNLSSLSKK(gl)__NA
_GEALSALDSK(gl)ANNLSSLSK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

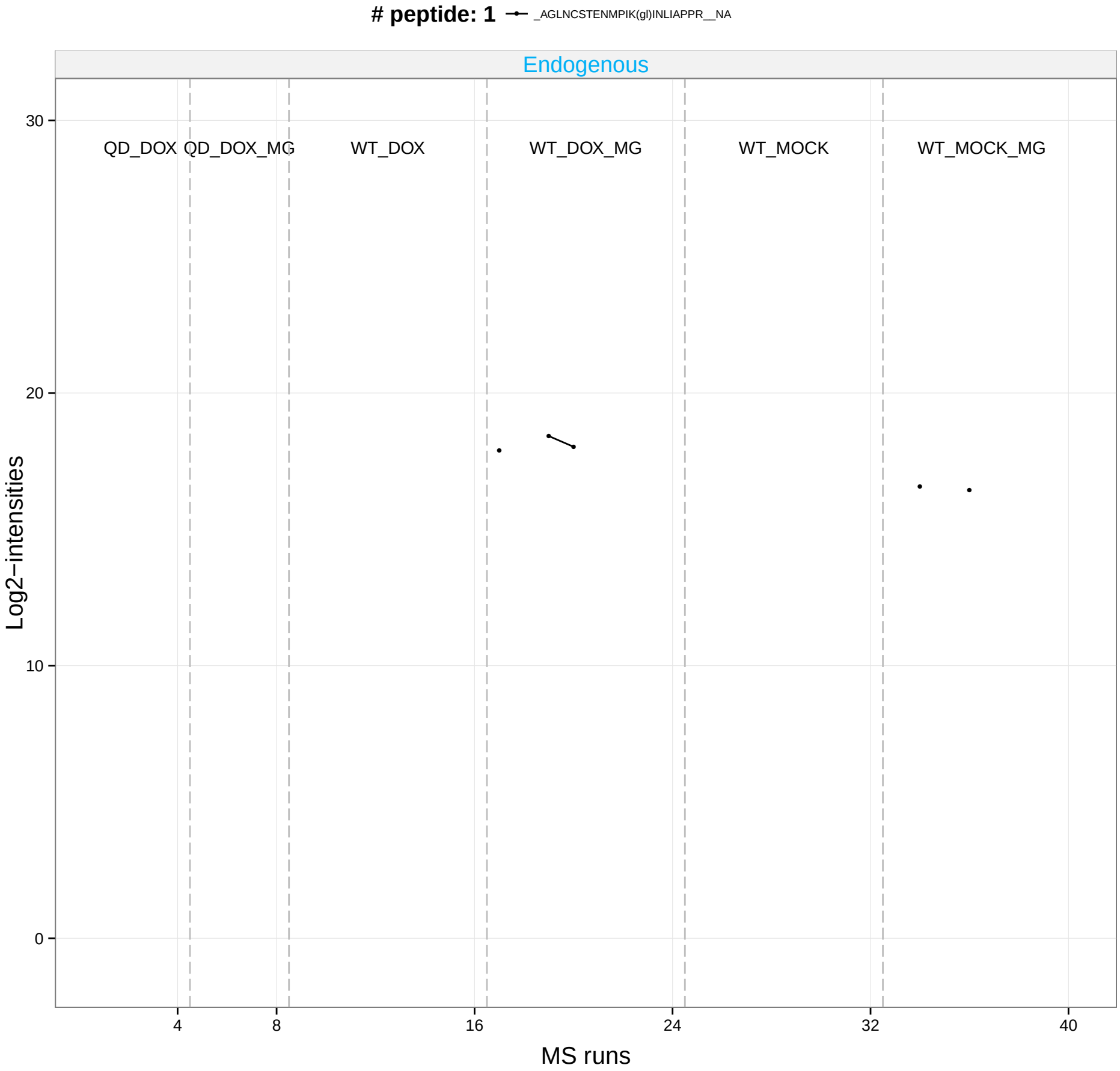
O96008

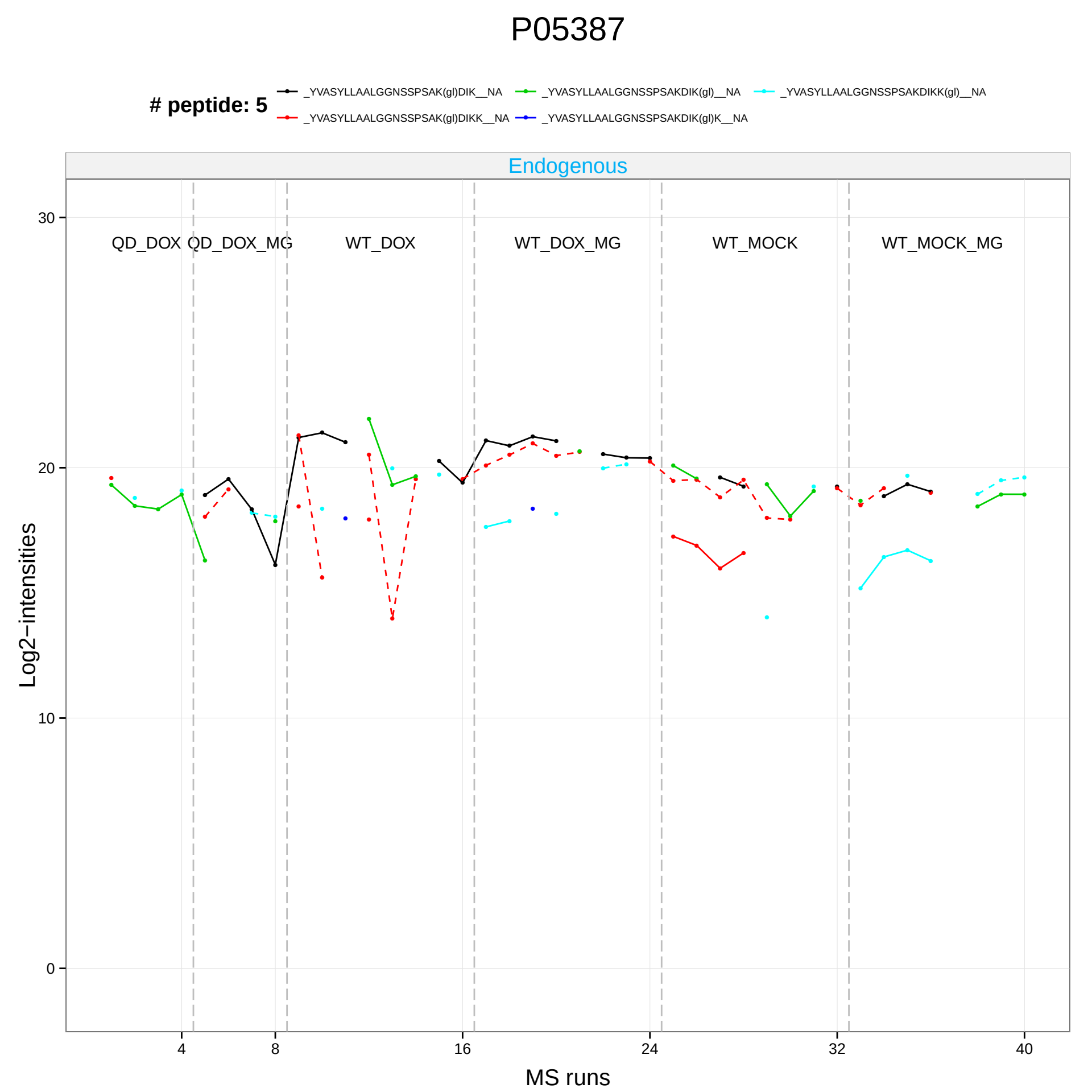

peptide: 1
_SK(gl)MAIQTQQSK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

P04406


peptide: 8
_AGAHLQGGAK(gl)R__NA
_FHGTVK(gl)AENGK__NA
_GALQNIIPASTGAAK(gl)AVGK__NA
_GALQNIIPASTGAAKAVGK(gl)__NA
_TVDGPSGK(gl)LWR__NA
_VVK(gl)QASEGPLK__NA
_WGDAGAEYVVESTGVFTTMEK(gl)AGAHLQGGAK__NA
_YDNSLK(gl)IISNASCTTNCLAPLAK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

P05198

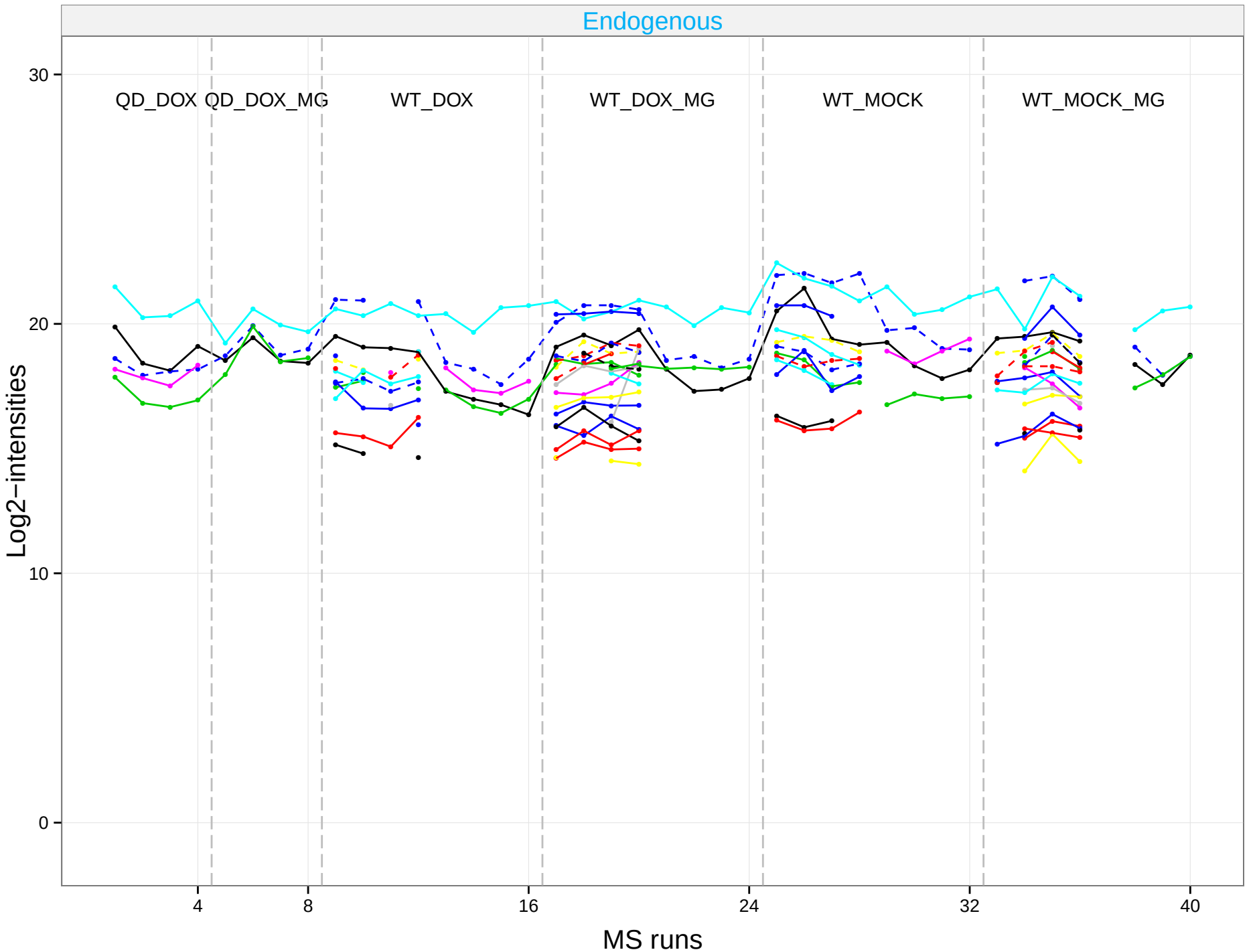

peptide: 1
_AGLNCSTENMPIK(gl)INLIAPPR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs


P05387
peptide: 5
_YVASYLLAALGGNSSPSAK(gl)DIK__NA
_YVASYLLAALGGNSSPSAKDIK(gl)__NA
_YVASYLLAALGGNSSPSAKDIKK(gl)__NA
_YVASYLLAALGGNSSPSAK(gl)DIKK__NA
_YVASYLLAALGGNSSPSAKDIK(gl)K__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

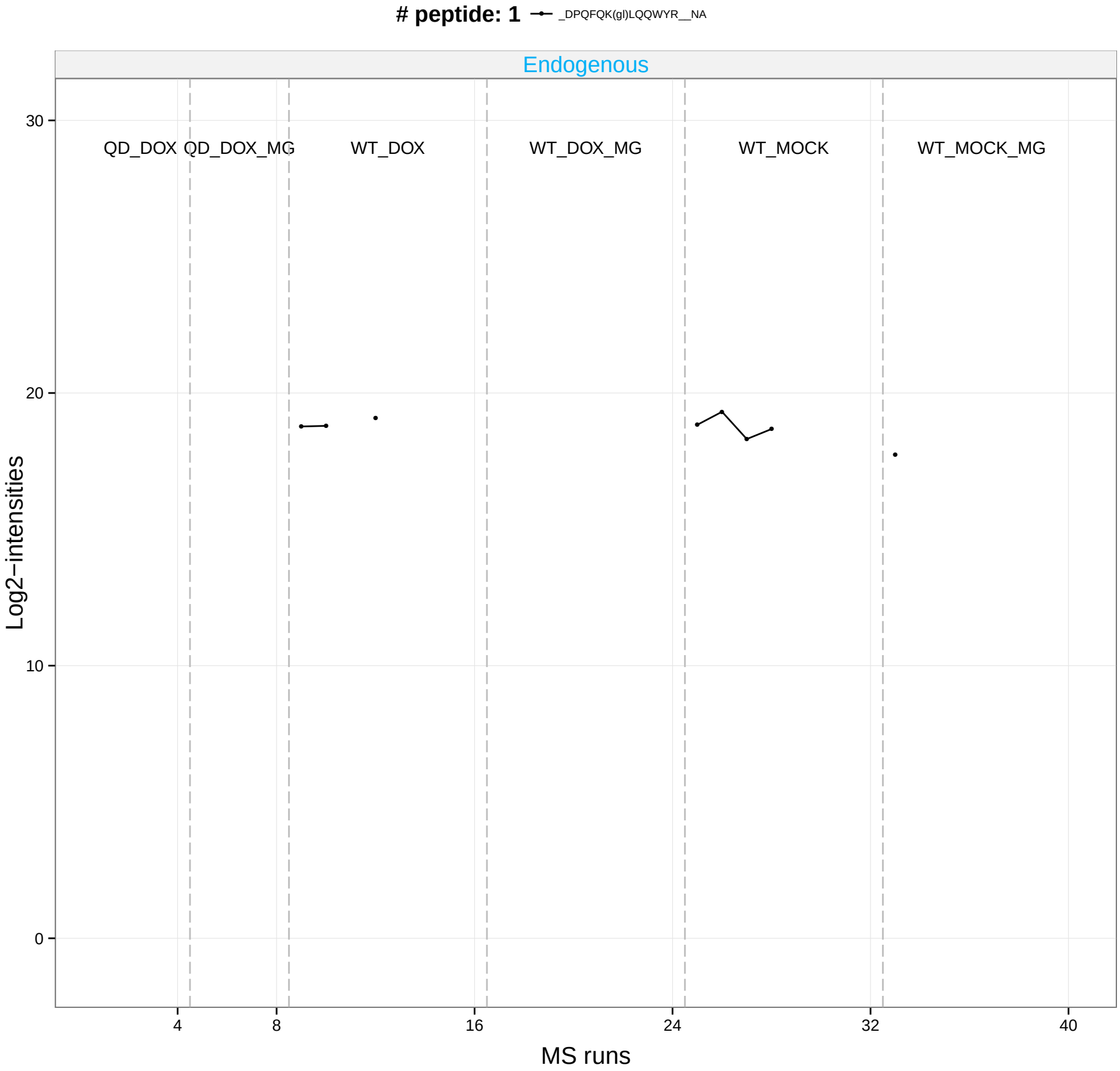
P06733

 # peptide: 21

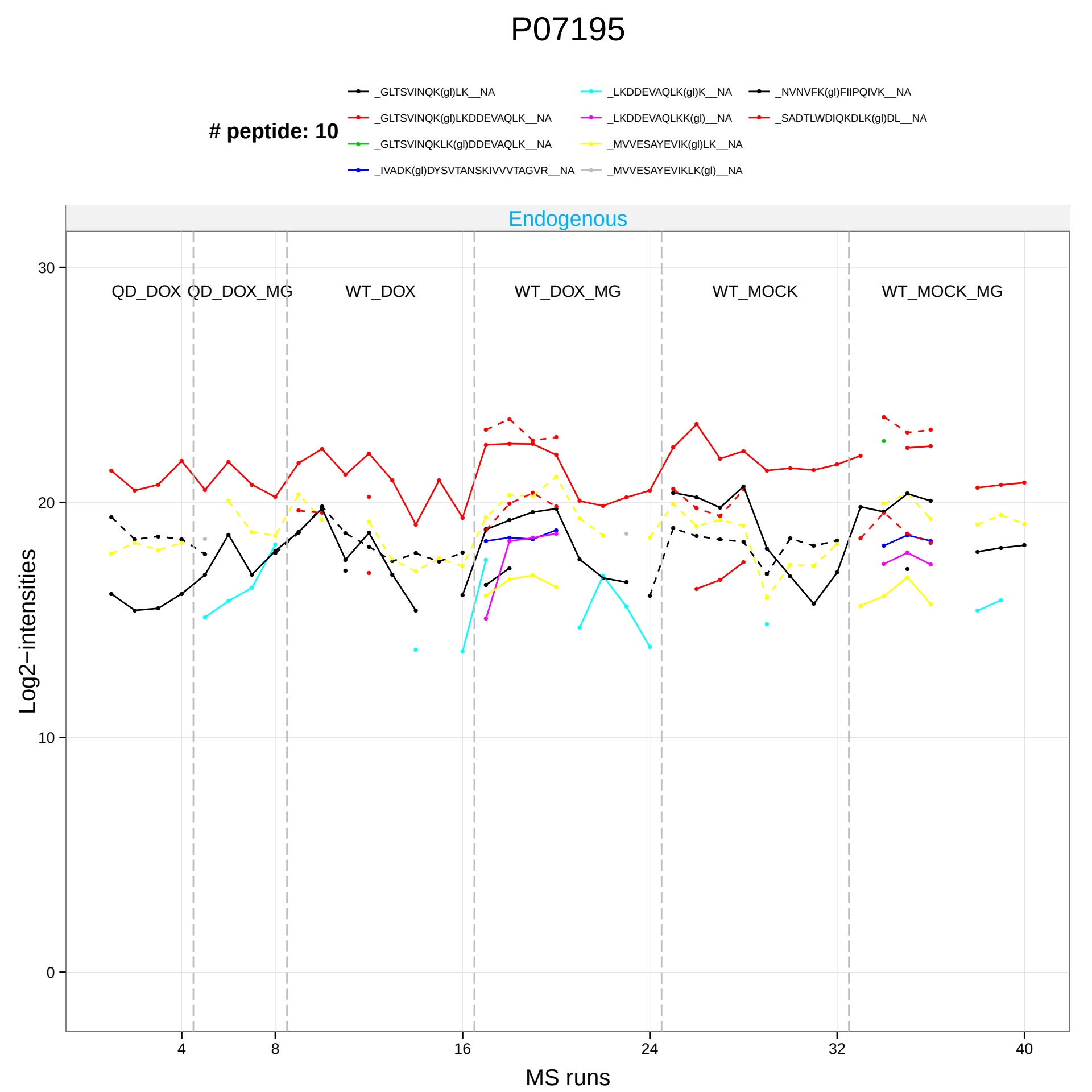
- _(ac)SILK(gl)IHAR__NA
- _AGAVEK(gl)GVPLYR__NA
- _AVEHINK(gl)TIAPALVSK__NA
- _AVNEK(gl)SCNCLLLK__NA
- _FGANAILGVSLAVCK(gl)AGAVEK__NA
- _FGANAILGVSLAVCKAGAVEK(gl)__NA
- _GVSK(gl)AVEHINK__NA
- _IEEELGSK(gl)AK__NA
- _IEEELGSKAK(gl)__NA
- _IGAEVYHNLK(gl)NVIK__NA
- _IGAEVYHNLKNVIK(gl)__NA
- _LAK(gl)YNQLLR__NA
- _SCNCLLLK(gl)VNQIGSVTESLQACK__NA
- _SFIK(gl)DYPVVSIEDPFDQDDWGAWQK__NA
- _SGK(gl)YDLDFK__NA
- _SGKYDLDFK(gl)SPDDPSR__NA
- _TAIGK(gl)AGYTDK__NA
- _TIAPALVSK(gl)K__NA
- _TIAPALVSKK(gl)__NA
- _VNQIGSVTESLQACK(gl)LAQANGWGVMVSHR__NA
- _YMGK(gl)GVSK__NA



P06744


peptide: 1
_DPQFQK(gl)LQQWYR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

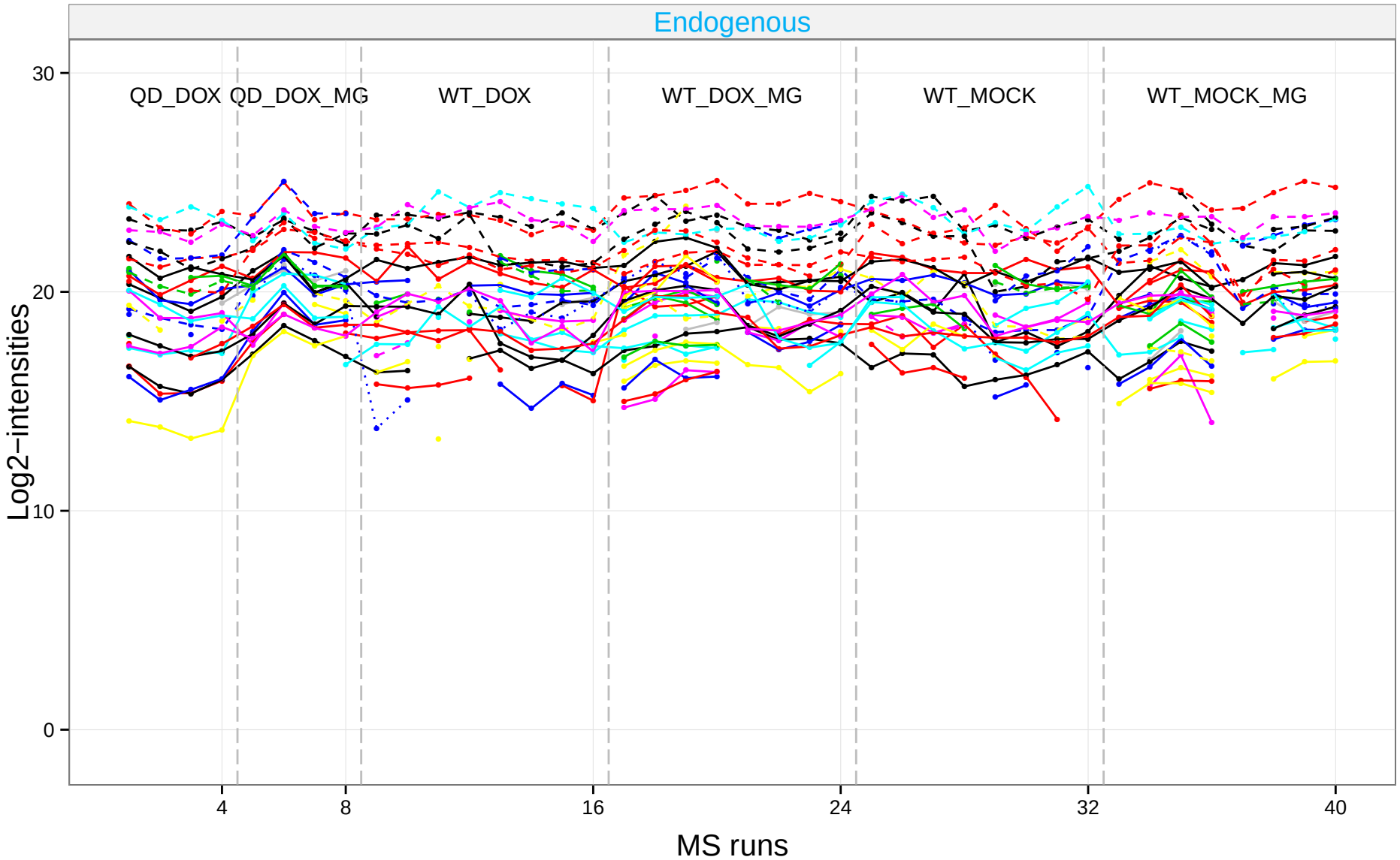
P07195

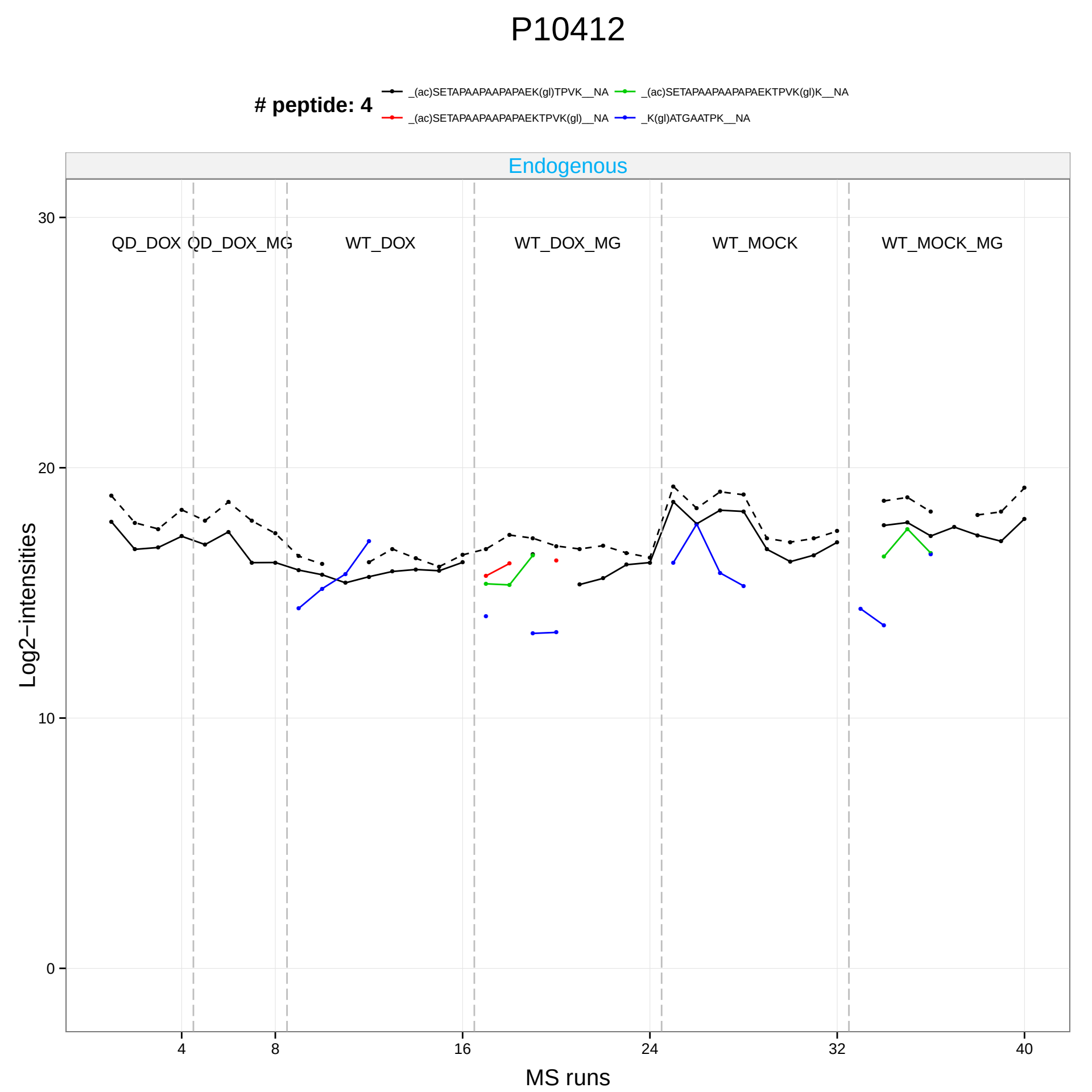

_GLTSVINQK(gl)LK__NA
_GLTSVINQK(gl)LKDDEVAQLK__NA
_GLTSVINQKLK(gl)DDEVAQLK__NA
_IVADK(gl)DYSVTANSKIVVVTAGVR__NA
_LKDDEVAQLK(gl)K__NA
_LKDDEVAQLKK(gl)__NA
_MVVESAYEVIK(gl)LK__NA
_MVVESAYEVIKLK(gl)__NA
_NVNVFK(gl)FIIPQIVK__NA
_SADTLWDIQKDLK(gl)DL__NA
peptide: 10
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
MS runs
30
20
10
0
4
8
16
24
32
40

P08107

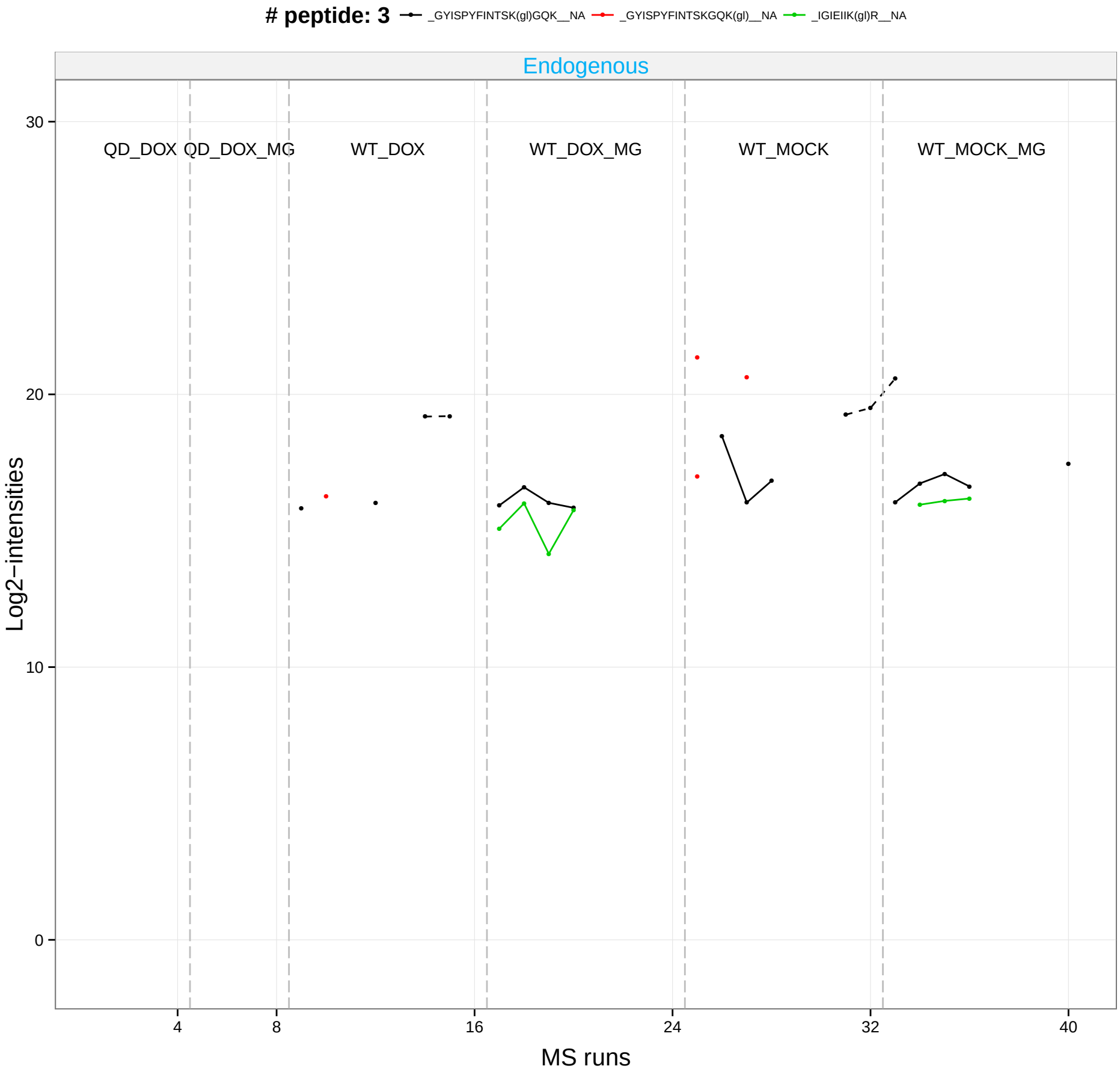
 # peptide: 39

- _(ac)AK(gl)AAAIGIDLGTTYSCVGVFQHGK__NA
- _AFYPEEISSMVLTK(gl)MK__NA
- _AFYPEEISSMVLTKM(ox)K(gl)__NA
- _CQEVISWLDANTLAEK(gl)DEFEHKR__NA
- _CQEVISWLDANTLAEKDEFEHK(gl)R__NA
- _DAK(gl)LDKAQIHDLVLVGGSTR__NA
- _DAKLDK(gl)AQIHDLVLVGGSTR__NA
- _DISQNK(gl)R__NA
- _ELEQVCNPIISGLYQGAGGPGPGGFGAQGPK(gl)GGSGSGPTIEEVD__NA
- _FGDPVVQSDMK(gl)HWPFQVINDGDKPK__NA
- _FGDPVVQSDMKHWPFQVINDGDK(gl)PK__NA
- _FGDPVVQSDMKHWPFQVINDGDKPK(gl)__NA
- _GETK(gl)AFYPEEISSMVLTK__NA
- _HWPFQVINDGDK(gl)PKVQVSYK__NA
- _HWPFQVINDGDKPK(gl)VQVSYK__NA
- _HWPFQVINDGDKPKVQVSYK(gl)__NA
- _LDK(gl)AQIHDLVLVGGSTR__NA
- _LIGDAAK(gl)NQVALNPQNTVFDAK__NA
- _LIGDAAK(gl)NQVALNPQNTVFDAK(gl)R__NA
- _LVNHFVEEFK(gl)R__NA
- _LVNHFVEEFKRK(gl)__NA
- _MVQEAEK(gl)YK(gl)AEDEVQR__NA
- _MVQEAEK(gl)YKAEDEVQR__NA
- _MVQEAEKYK(gl)AEDEVQR__NA
- _NALESYAFNMK(gl)SAVEDEGLK__NA
- _NQVALNPQNTVFDAK(gl)R__NA
- _NSTIPTK(gl)QTQIFTTYSDNQPGVLIQVYEGER__NA
- _QATK(gl)DAGVIAGLNVLR__NA
- _RK(gl)ELEQVCNPIISGLYQGAGGPGPGGFGAQGPK
- _RNSTIPTK(gl)QTQIFTTYSDNQPGVLIQVYEGER__N
- _SAVEDEGLK(gl)GK__NA
- _SENVQDLLLLDVAPLSLGLETAGGVMTALIK(gl)R__N
- _VQK(gl)LLQDFFNGR__NA
- _VQVSYK(gl)GETK__NA
- _VQVSYK(gl)GETK(gl)AFYPEEISSMVLTK__NA
- _VQVSYK(gl)GETKAFYPEEISSMVLTK__NA
- _VSAK(gl)NALESYAFNM(ox)K__NA
- _VSAK(gl)NALESYAFNMK__NA
- _VSAK(gl)NALESYAFNMK(gl)SAVEDEGLK__NA

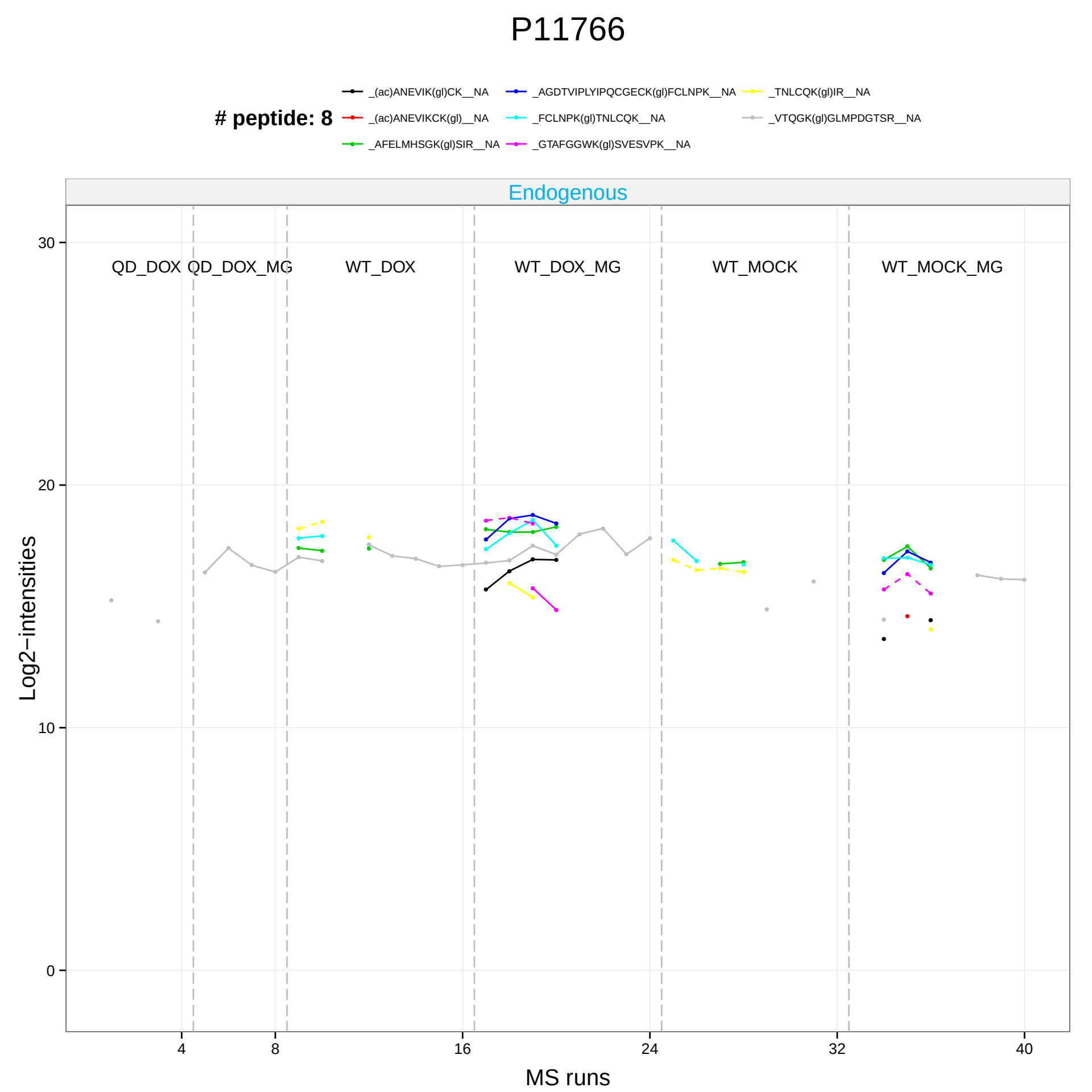



P10412
peptide: 4
_(ac)SETAPAAPAAPAPAEK(gl)TPVK__NA
_(ac)SETAPAAPAAPAPAEKTPVK(gl)K__NA
_(ac)SETAPAAPAAPAPAEKTPVK(gl)__NA
_K(gl)ATGAATPK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

P10809


peptide: 3
_GYISPYFINTSK(gl)GQK__NA
_GYISPYFINTSKGQK(gl)__NA
_IGIEIIK(gl)R__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

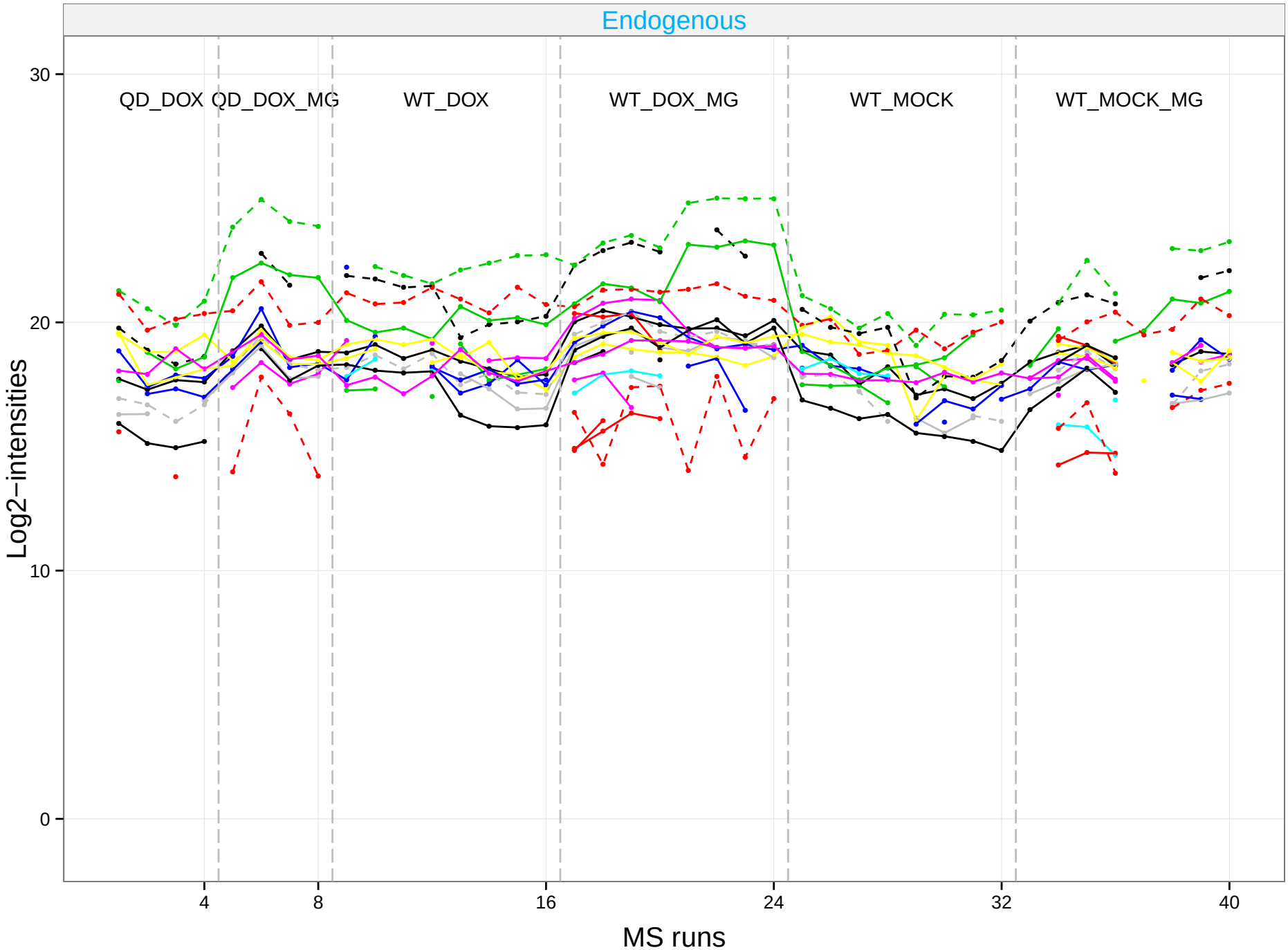
P11766

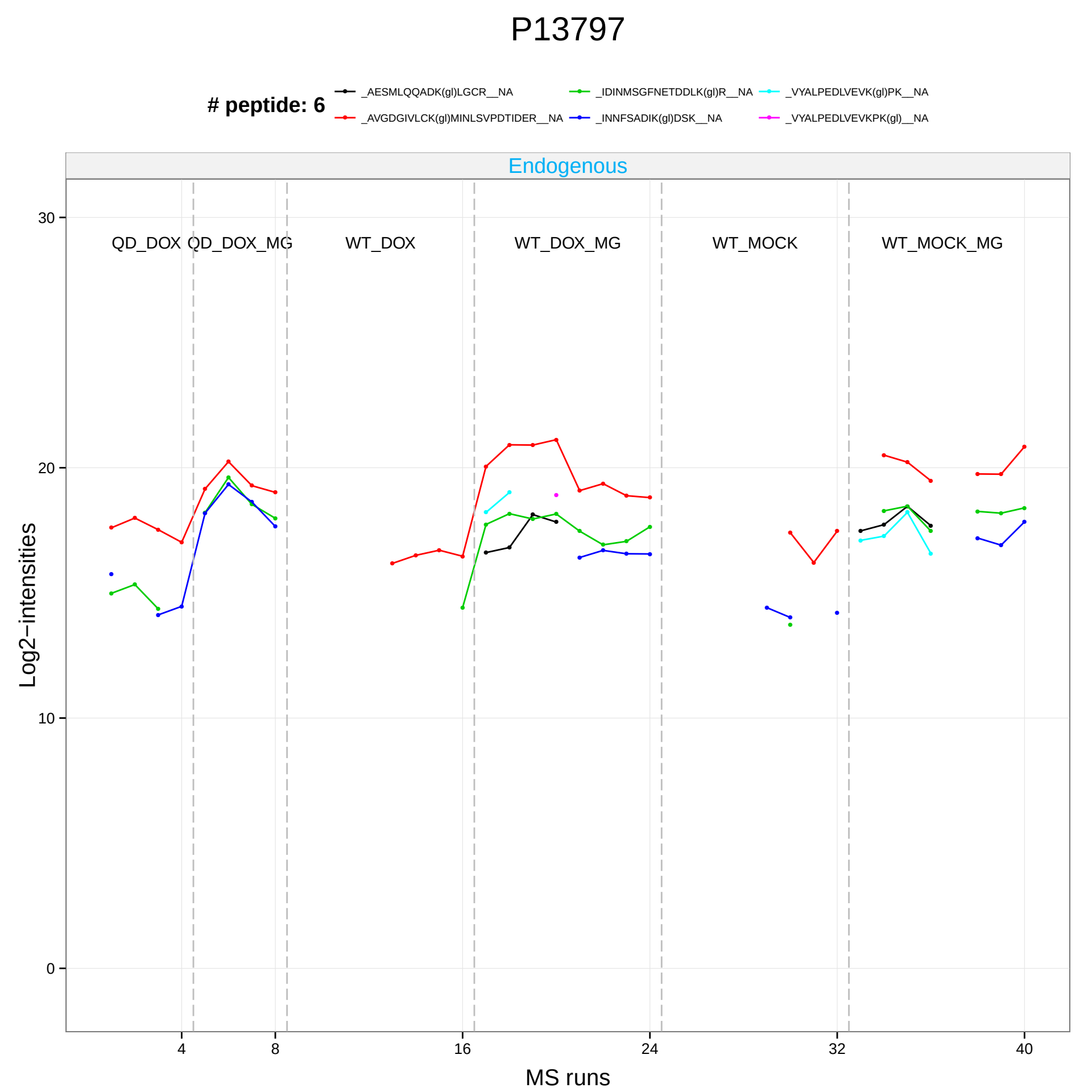

peptide: 8
_(ac)ANEVIK(gl)CK__NA
_(ac)ANEVIKCK(gl)__NA
_AFELMHSGK(gl)SIR__NA
_AGDTVIPLYIPQCGECK(gl)FCLNPK__NA
_FCLNPK(gl)TNLCQK__NA
_GTAFGGWK(gl)SVESVPK__NA
_TNLCQK(gl)IR__NA
_VTQGK(gl)GLMPDGTSR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

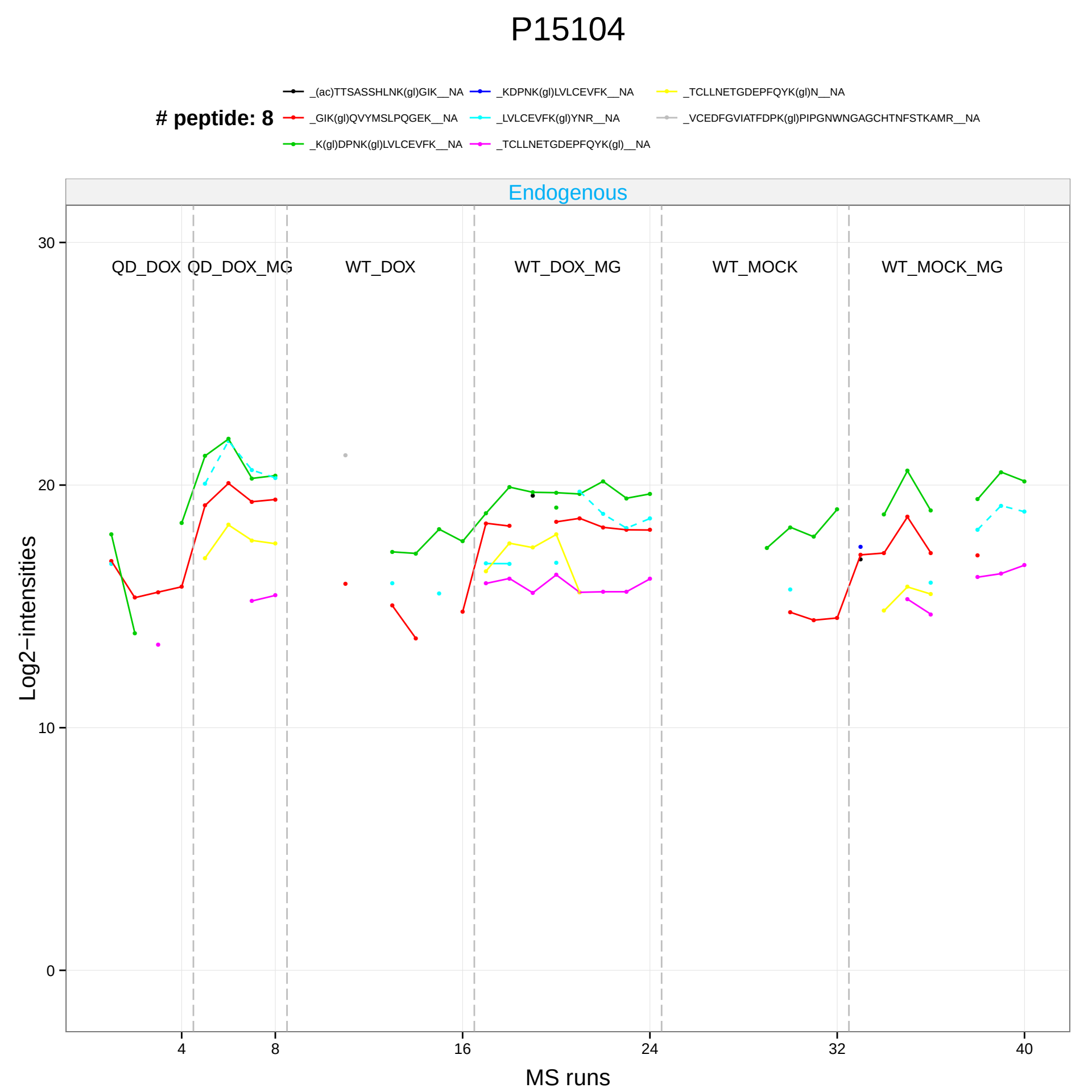
P13010

 # peptide: 24

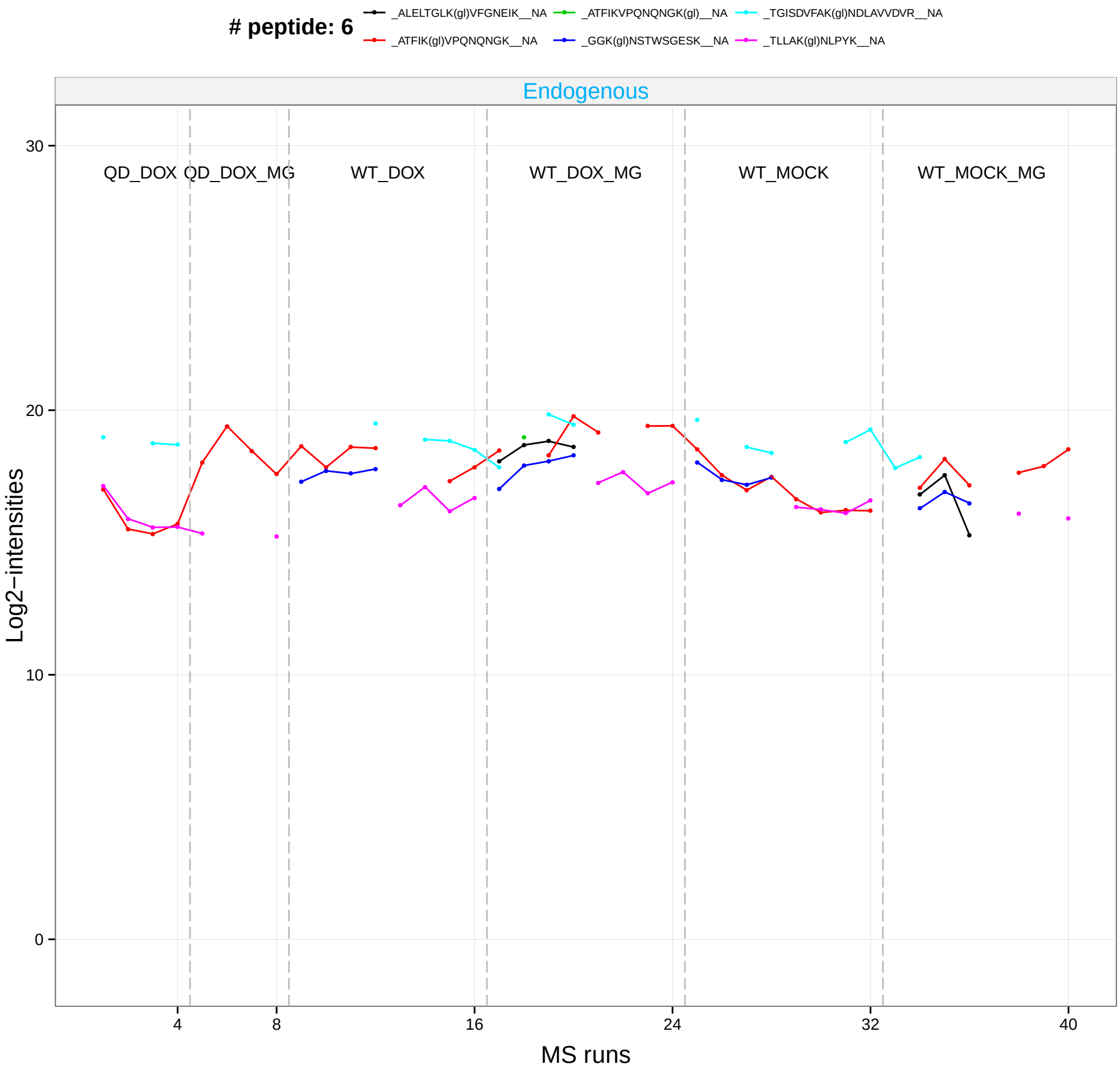
_CFSVLGFCK(gl)SSQVQR__NA
_EPLPPIQQHIWNMLNPPAEVTTK(gl)SQIPLSK__NA
_EPLPPIQQHIWNMLNPPAEVTTKSQIPLSK(gl)__NA
_ETVYCLNDDDETEVLK(gl)EDIIQGFR__NA
_FFM(ox)GNQVLK(gl)VFAAR__NA
_FFMGNQVLK(gl)VFAAR__NA
_FNNFLK(gl)ALQEK__NA
_FSK(gl)SQLDIIIHSLK__NA
_IAAYK(gl)SILQER__NA
_K(gl)VITMFVQR__NA
_LGGHGPSFPLK(gl)GITEQQK__NA
_LGGHGPSFPLKGITEQQK(gl)__NA
_QYMFSSLK(gl)NSK__NA
_QYMFSSLKNSK(gl)__NA
_SQIPLSK(gl)IK__NA
_SQLDIIIHSLK(gl)K__NA
_SQLDIIIHSLKK(gl)__NA
_TDTLEDLFPTTK(gl)IPNPR__NA
_TLFPLIEAKK(gl)__NA
_TWTVVDAK(gl)TLK__NA
_TWTVVDAK(gl)TLKK__NA
_YGSDIVPFSK(gl)VDEEQMK__NA
_YGSDIVPFSKVDEEQMK(gl)__NA
_YKSEGK(gl)CFSVLGFCK__NA

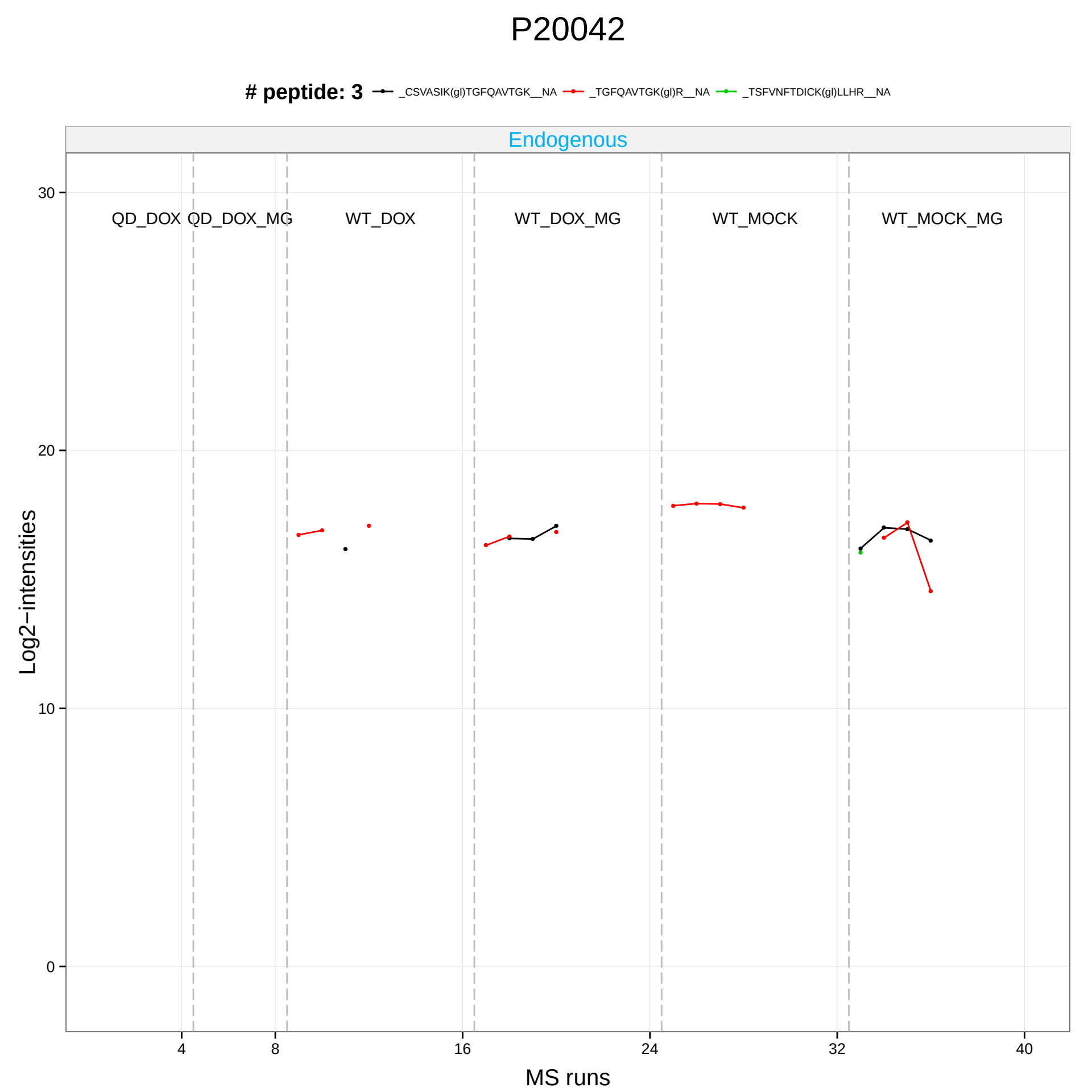



P13797
peptide: 6
_AESMLQQADK(gl)LGCR__NA
_IDINMSGFNETDDLK(gl)R__NA
_VYALPEDLVEVK(gl)PK__NA
_AVGDGIVLCK(gl)MINLSVPDTIDER__NA
_INNFSADIK(gl)DSK__NA
_VYALPEDLVEVKPK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs


P15104
peptide: 8
_(ac)TTSASSHLNK(gl)GIK__NA
_KDPNK(gl)LVLCEVFK__NA
_TCLLNETGDEPFQYK(gl)N__NA
_GIK(gl)QVYMSLPQGEK__NA
_LVLCEVFK(gl)YNR__NA
_VCEDFGVIATFDPK(gl)PIPGNWNGAGCHTNFSTKAMR__NA
_K(gl)DPNK(gl)LVLCEVFK__NA
_TCLLNETGDEPFQYK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

P19338

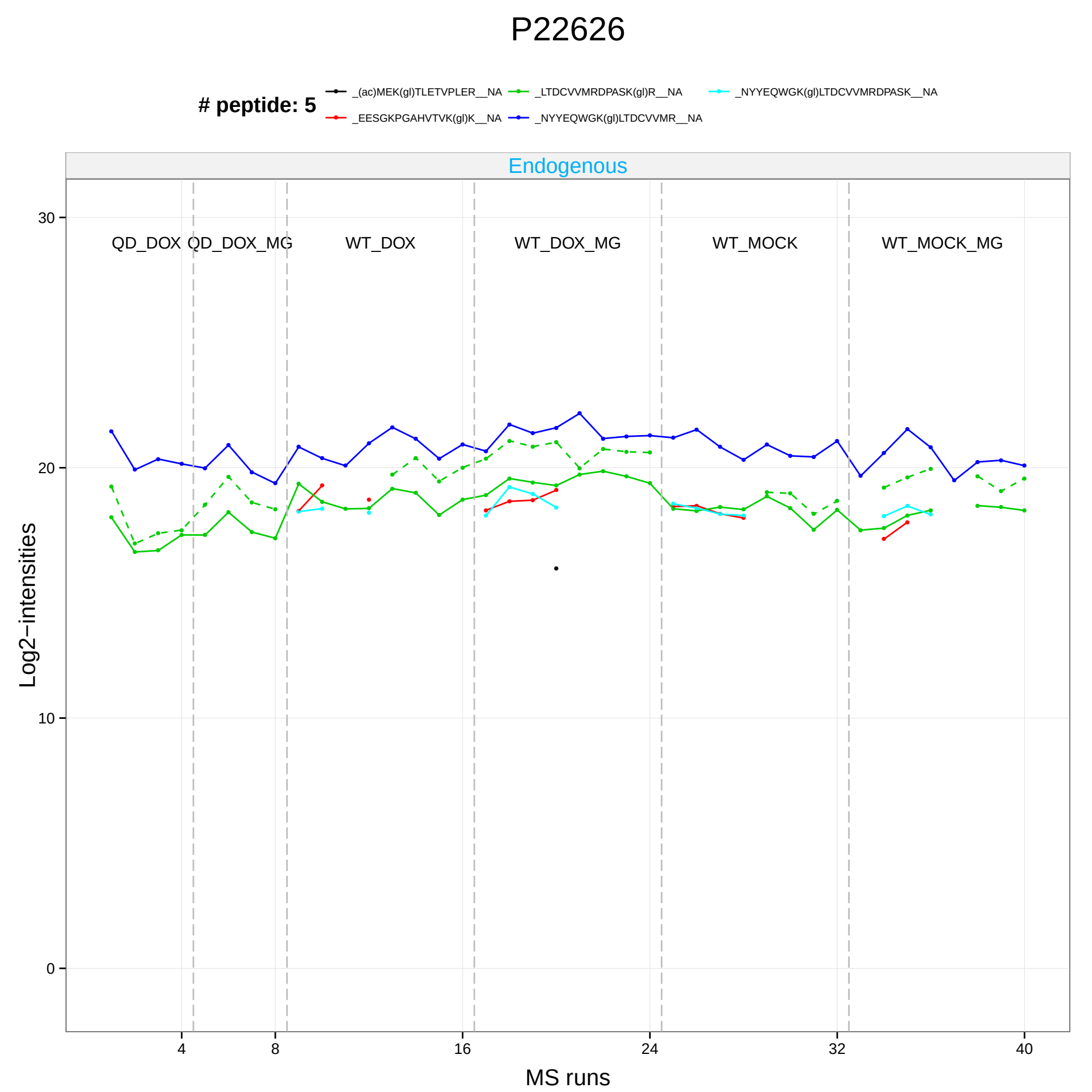

peptide: 6
_ALELTGLK(gl)VFGNEIK__NA
_ATFIKVPQNQNGK(gl)__NA
_TGISDVFAK(gl)NDLAVVDVR__NA
_ATFIK(gl)VPQNQNGK__NA
_GGK(gl)NSTWSGESK__NA
_TLLAK(gl)NLPYK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs


P20042
peptide: 3
_CSVASIK(gl)TGFQAVTGK__NA
_TGFQAVTGK(gl)R__NA
_TSFVNFTDICK(gl)LLHR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

P22102

 # peptide: 20

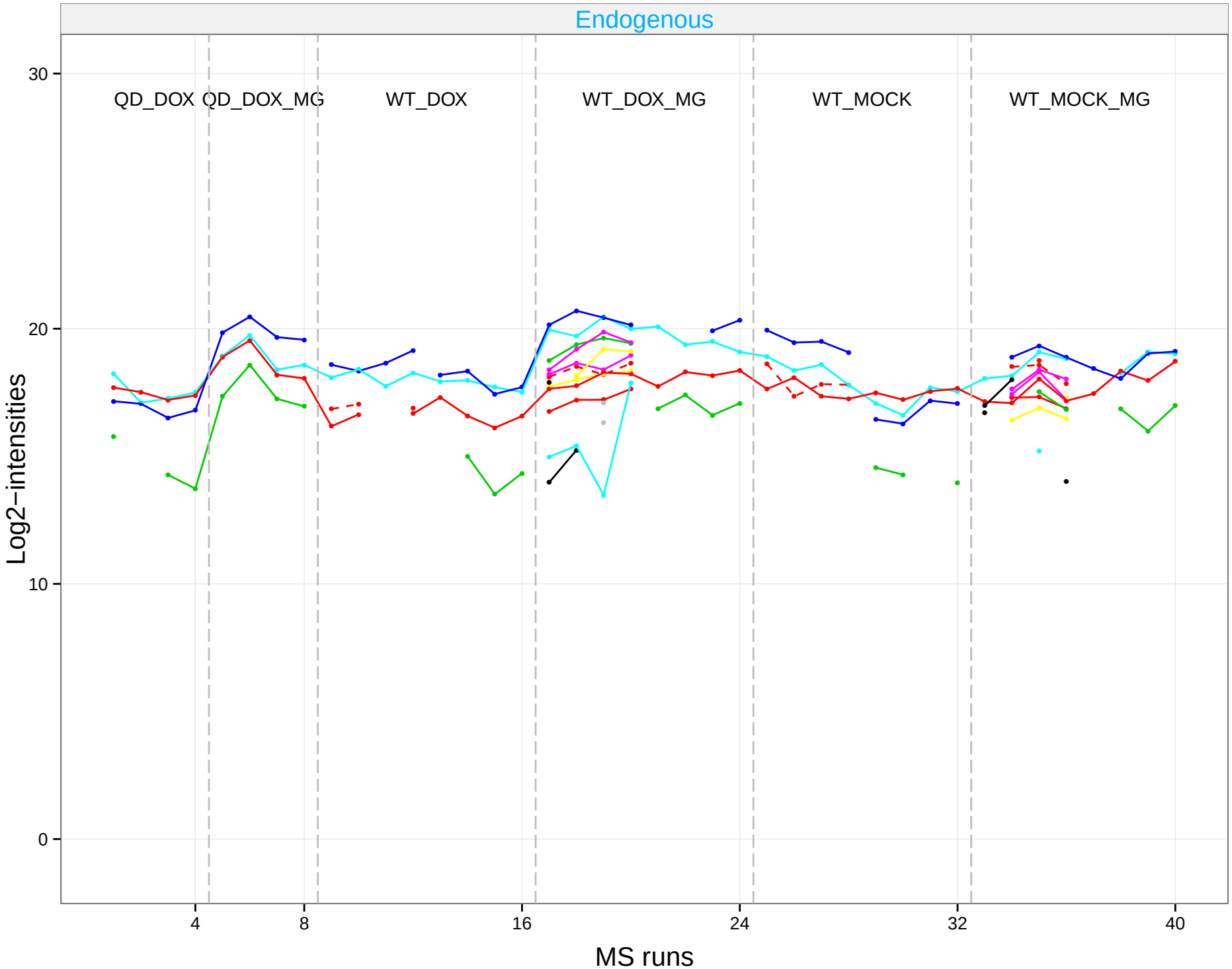
- _AFTKPEEACSFILSADFPALVVK(gl)ASGLAAGK__NA
- _ASGLAAGK(gl)GVIVAK__NA
- _ENLISALEEAK(gl)K__NA
- _ENLISALEEAKK(gl)__NA
- _GLAAIK(gl)FEGAIYR__NA
- _GVIVAK(gl)SK__NA
- _GVIVAK(gl)SKEEACK__NA
- _HGIPTAQWK(gl)AFTKPEEACSFILSADFPALVVK__NA
- _IK(gl)DTVLQR__NA
- _ISNTAISISDHTALAQFCK(gl)EK__NA
- _ISNTAISISDHTALAQFCKEK(gl)__NA
- _LAQSHHVK(gl)QVLVAPGNAGTACSEK__NA
- _LLEGDGGPNTGGMGAYCPAPQVSNDLLLK(gl)IK__NA
- _NGPK(gl)VLEFNCR__NA
- _QVLVAPGNAGTACSEK(gl)ISNTAISISDHTALAQFCK__NA
- _SAGVQCFGPTAEAAQLESSK(gl)R__NA
- _SLTYK(gl)ESGVDIAAGNMLVK__NA
- _TFNCGVGAVLVVSK(gl)EQTEQILR__NA
- _TVDGMQQEGTPYTGILYAGIMLTK(gl)NGPK__NA
- _TVDGMQQEGTPYTGILYAGIMLTKNGPK(gl)__NA


P22626
peptide: 5
_(ac)MEK(gl)TLETVPLER__NA
_LTDCVVMRDPASK(gl)R__NA
_NYYEQWGK(gl)LTDCVVMRDPASK__NA
_EESGKPGAHVTVK(gl)K__NA
_NYYEQWGK(gl)LTDCVVMR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

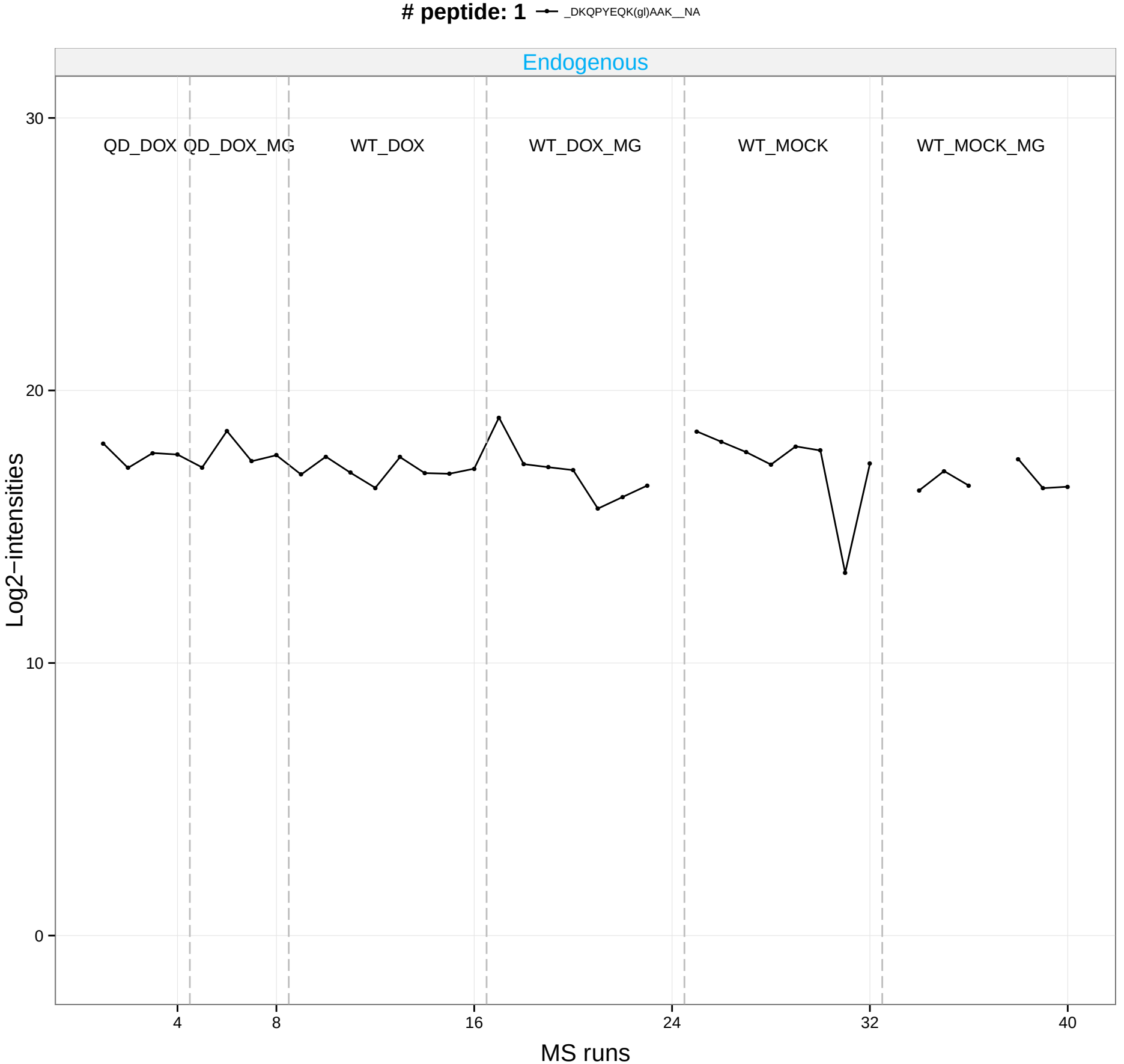
P24928

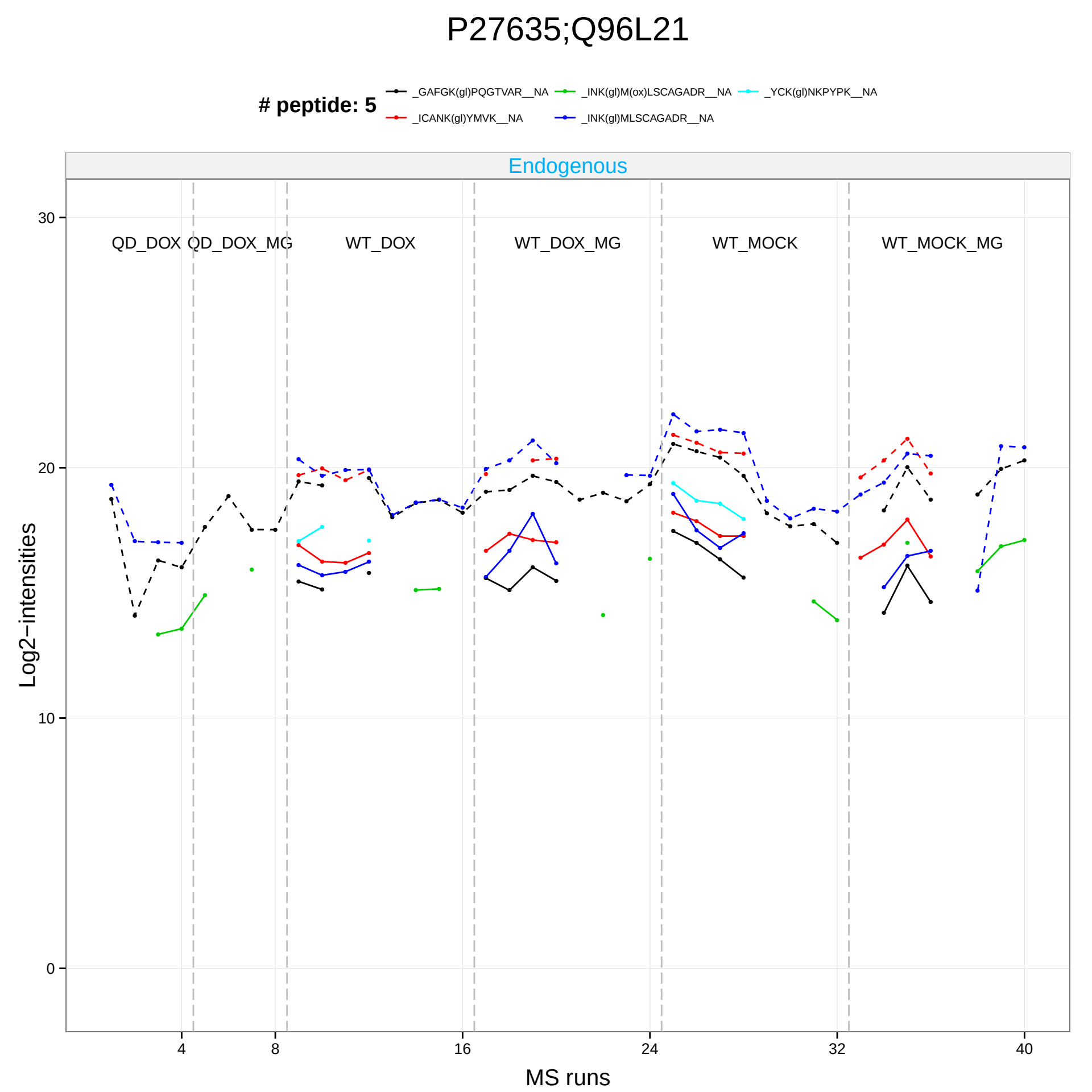
 # peptide: 18

- _DK(gl)TGSSAQKSLSEYNNFK__NA
- _DKTGSSAQK(gl)SLSEYNNFK__NA
- _EGLIDTAVK(gl)TAETGYIQR__NA
- _IMNSDENK(gl)MQEEEEVVDKMDDDVFLR__NA
- _INISQVIAVVGQQNVEGK(gl)R__NA
- _LIK(gl)SMESVMVK__NA
- _LK(gl)ELINISK__NA
- _SMVVSGAK(gl)GSK__NA
- _SMVVSGAKGSK(gl)__NA
- _TGSSAQK(gl)SLSEYNNFK__NA
- _TYQDIQNTIK(gl)K__NA
- _VIFPTGDSK(gl)VVLPCNLLR__NA
- _VLSEK(gl)DVDPVR__NA
- _VVVENGELIMGILCK(gl)K__NA
- _YGEDGLAGESVEFQNLATLK(gl)PSNKAFEK__NA
- _YGEDGLAGESVEFQNLATLKPSNK(gl)__NA
- _YGEDGLAGESVEFQNLATLKPSNK(gl)AFEK__NA
- _YGEDGLAGESVEFQNLATLKPSNKAFEK(gl)__NA



P26583

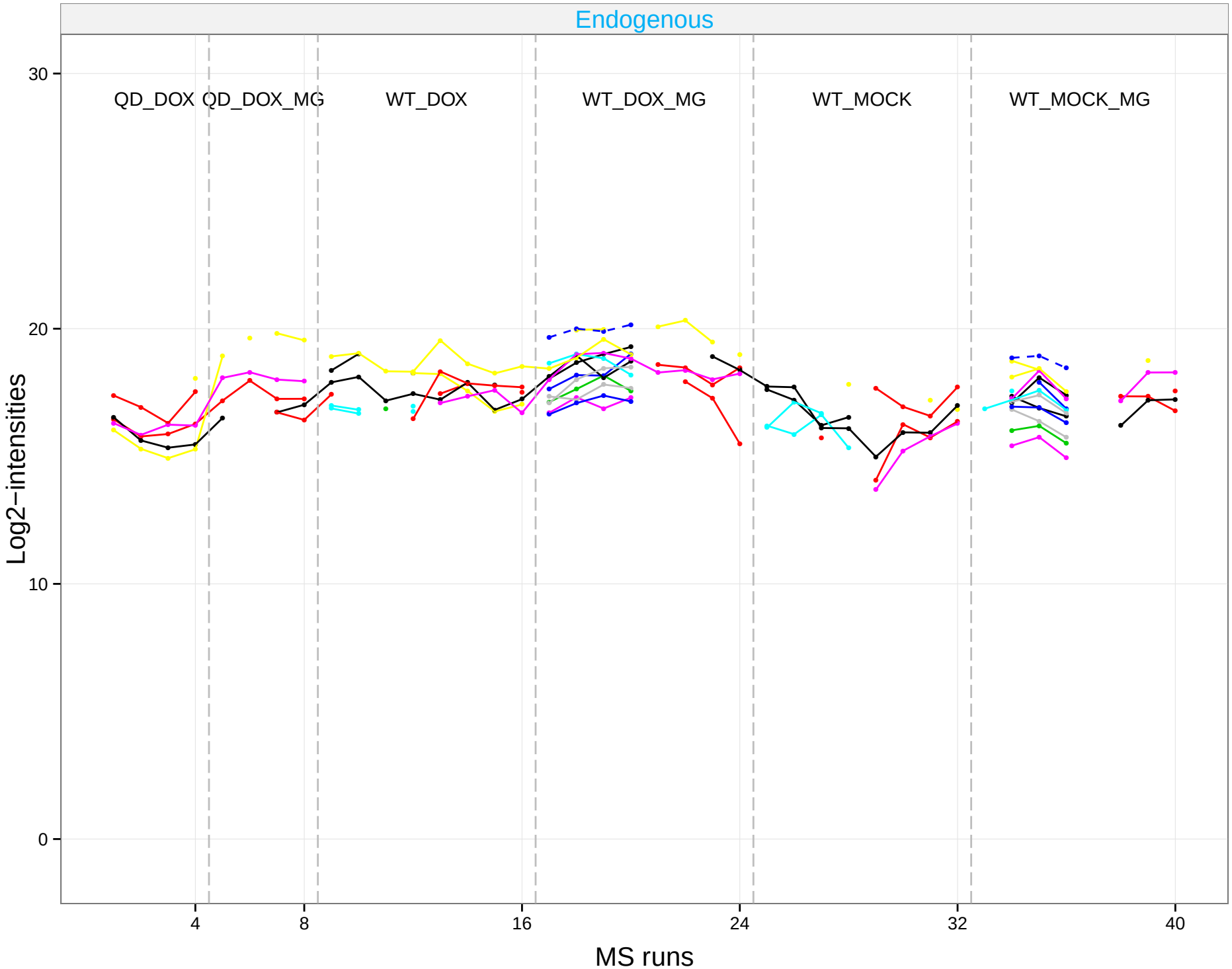

peptide: 1
_DKQPYEQK(gl)AAK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

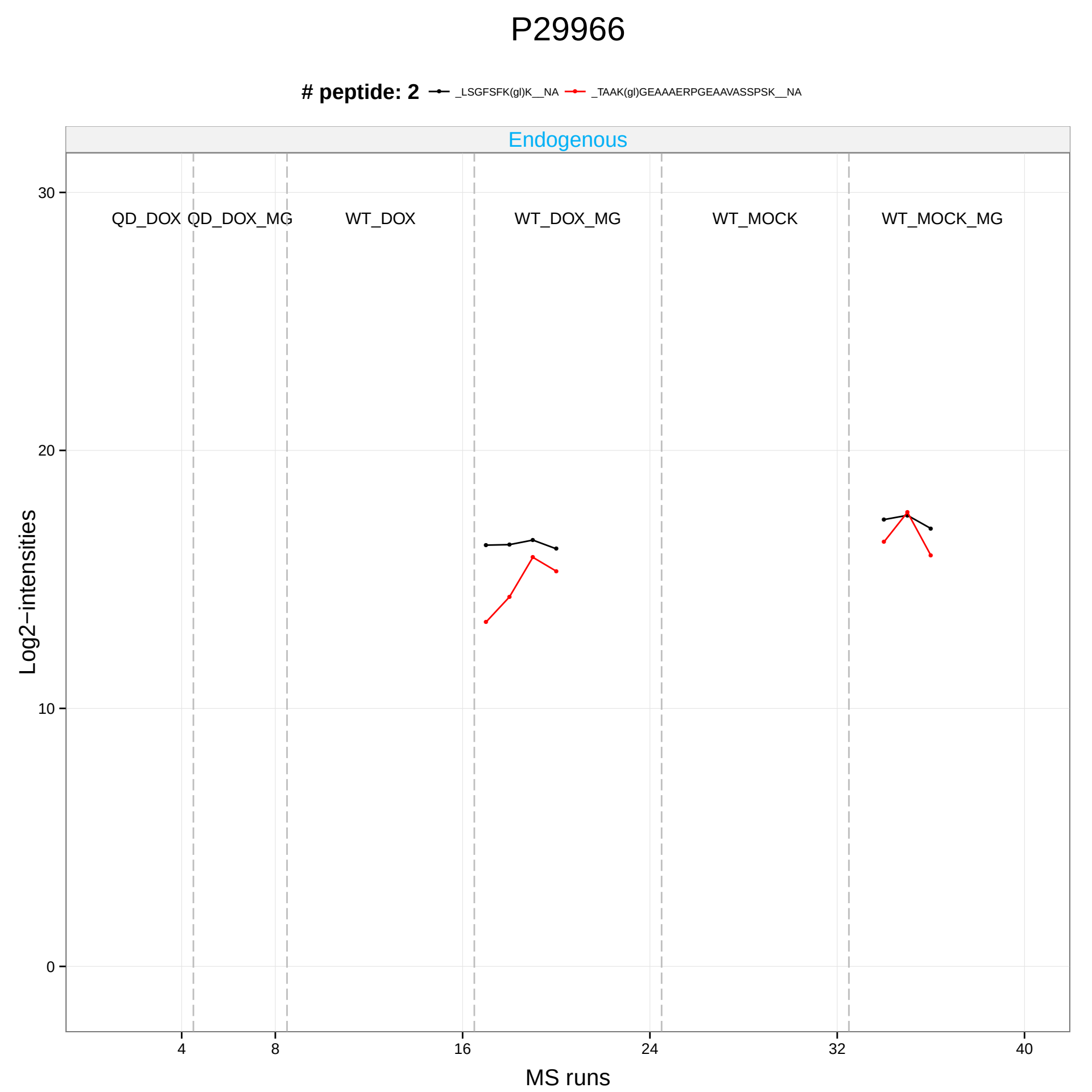

P27635;Q96L21
peptide: 5
_GAFGK(gl)PQGTVAR__NA
_INK(gl)M(ox)LSCAGADR__NA
_YCK(gl)NKPYPK__NA
_ICANK(gl)YMVK__NA
_INK(gl)MLSCAGADR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

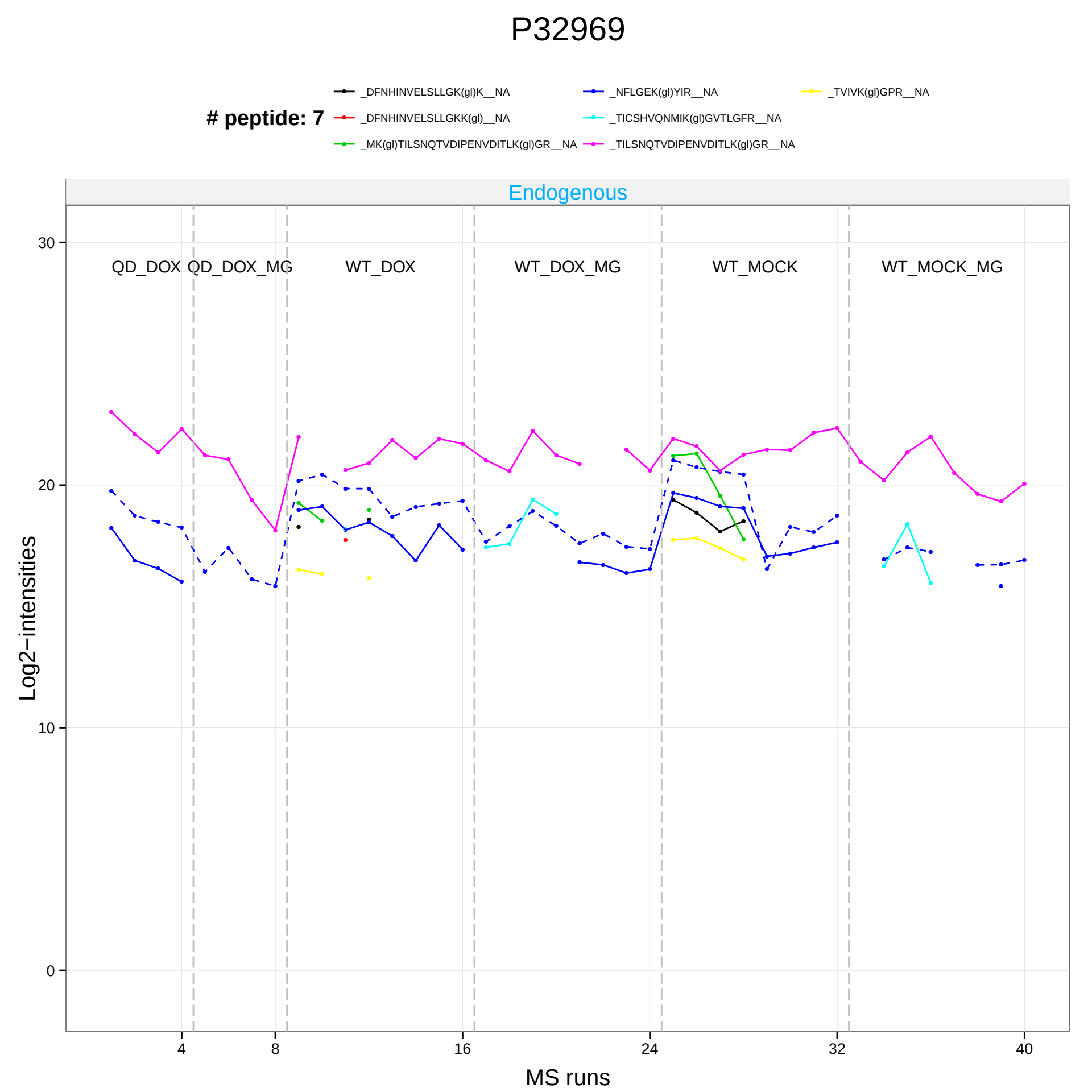
P27708

 # peptide: 16

- _AMLSTGFK(gl)IPK__NA
- _AMLSTGFKIPK(gl)__NA
- _CEAYLK(gl)AMLSTGFK__NA
- _FLSSAAAVSK(gl)EHPVVISK__NA
- _FLSSAAAVSKEHPVVISK(gl)__NA
- _HPQPGAVELAAK(gl)HCR__NA
- _ILALDCGLK(gl)YNQIR__NA
- _K(gl)NILLTIGSYK__NA
- _KNILLTIGSYK(gl)NK__NA
- _LAADFSVPLIIDIK(gl)CTK__NA
- _LAADFSVPLIIDIKCTK(gl)__NA
- _LSSFVTK(gl)GYR__NA
- _NSVTGGTAAFEPSVDYCVVK(gl)IPR__NA
- _SFEEAFQK(gl)ALR__NA
- _SFPFVSK(gl)TLGVDLVALATR__NA
- _WDLSK(gl)FLR__NA



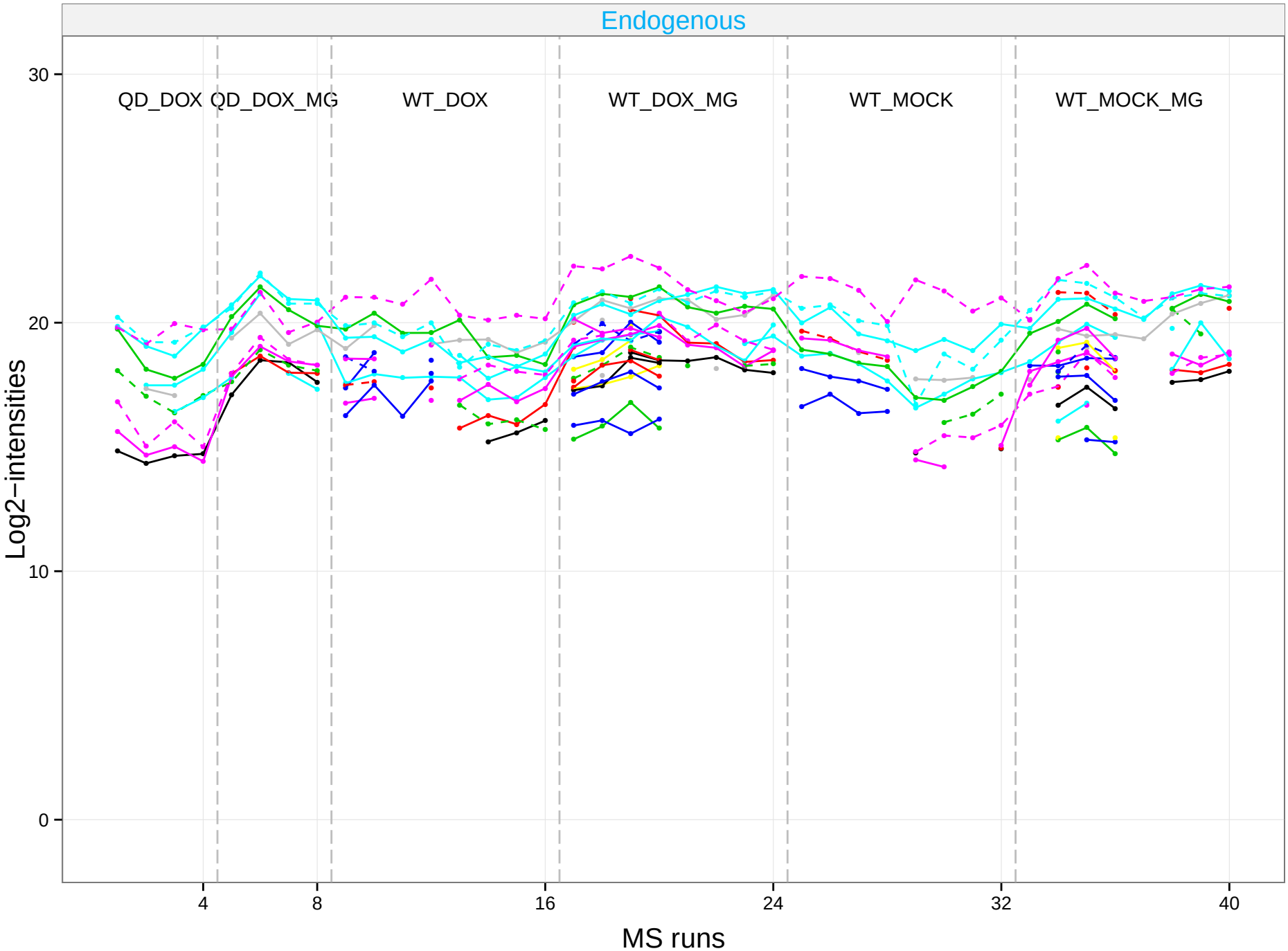

P29966
peptide: 2
_LSGFSFK(gl)K__NA
_TAAK(gl)GEAAAERPGEAAVASSPSK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs


P32969
peptide: 7
_DFNHINVELSLLGK(gl)K__NA
_DFNHINVELSLLGKK(gl)__NA
_MK(gl)TILSNQTVDIPENVDITLK(gl)GR__NA
_NFLGEK(gl)YIR__NA
_TICSHVQNMIK(gl)GVTLGFR__NA
_TILSNQTVDIPENVDITLK(gl)GR__NA
_TVIVK(gl)GPR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

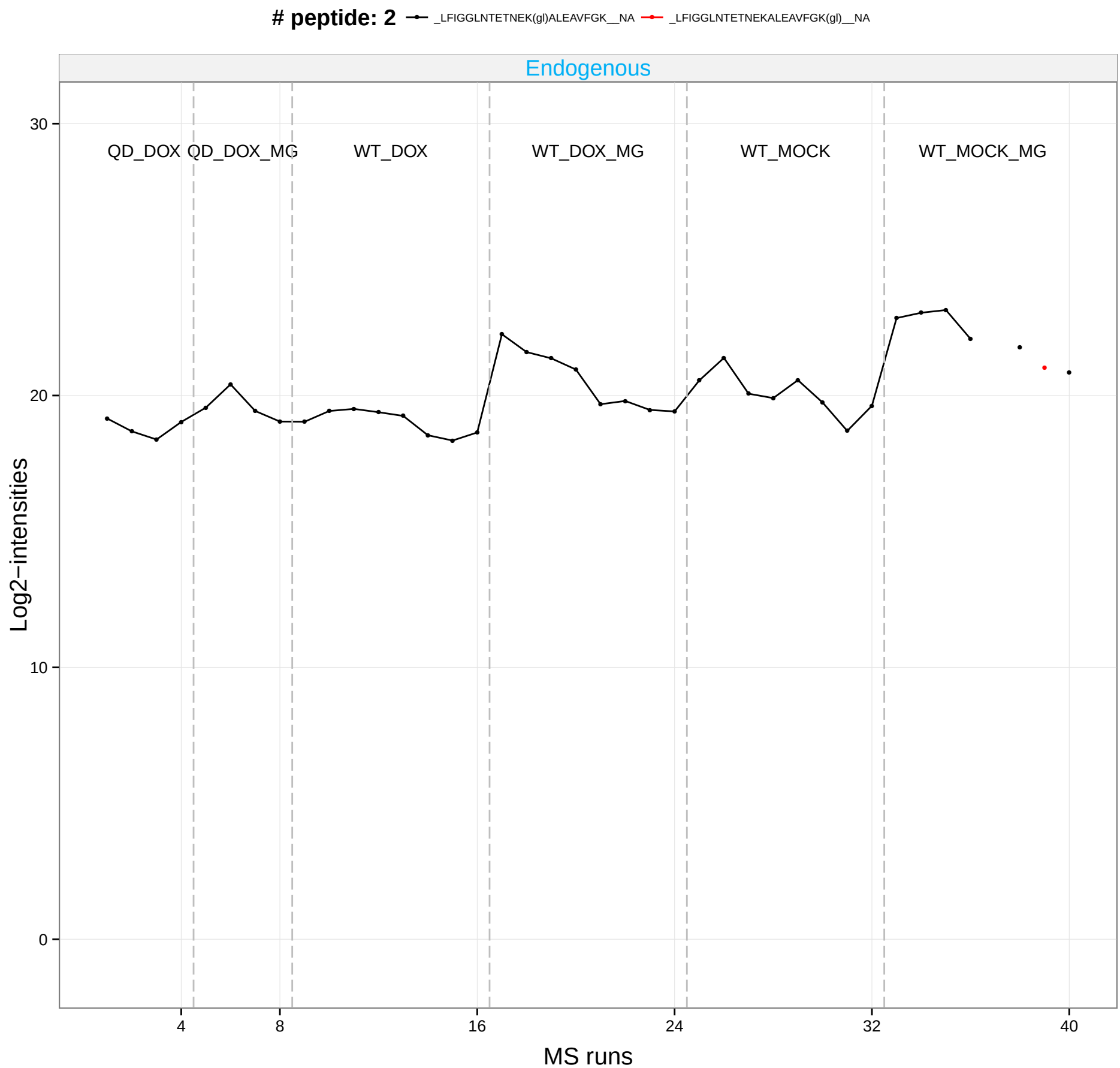
P33993

 # peptide: 22

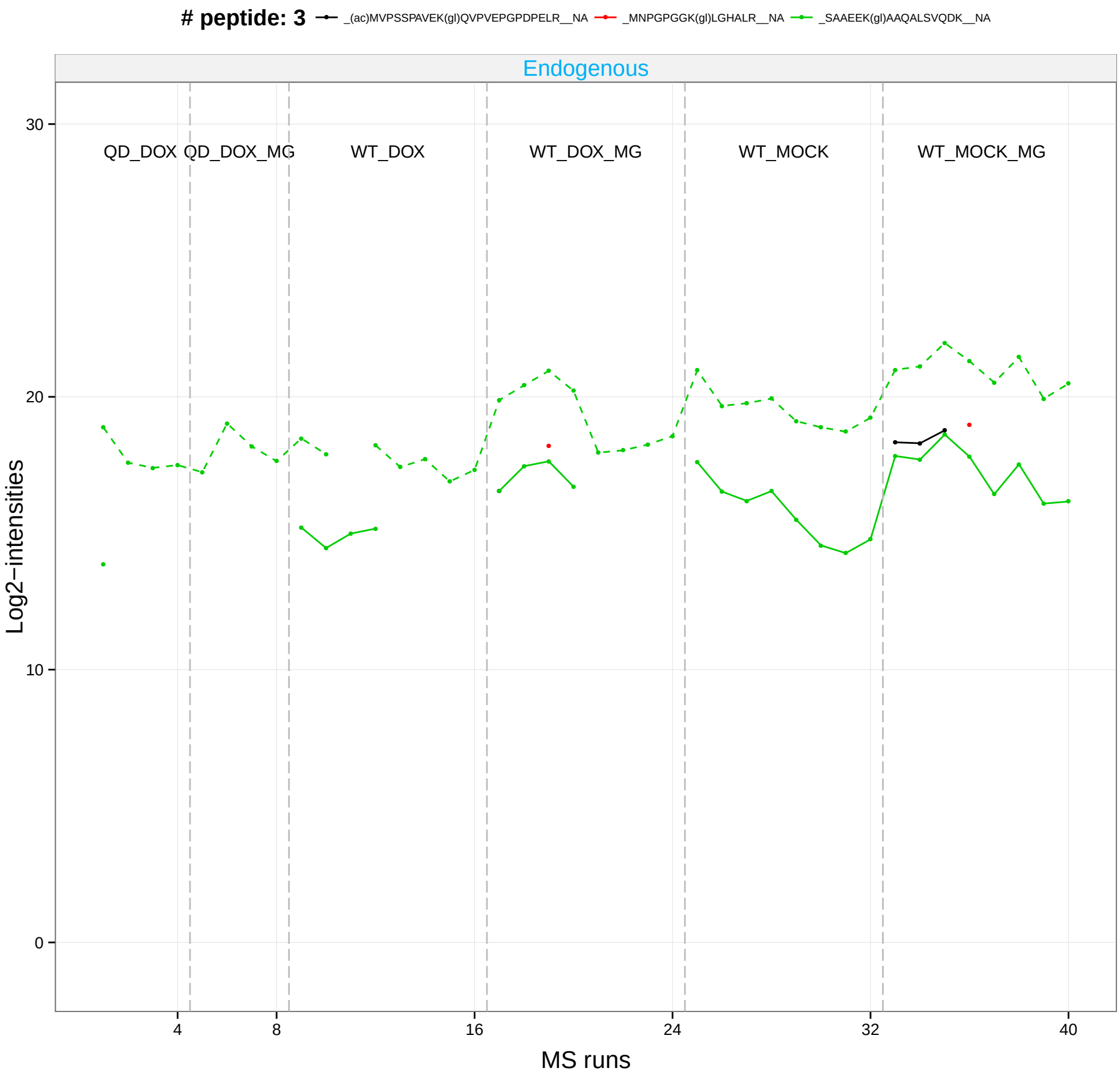
- _(ac)ALK(gl)DYALEK__NA
- _(ac)ALKDYALEK(gl)EK__NA
- _ADSVGK(gl)LVTVR__NA
- _EAWASK(gl)DATYTSAR__NA
- _FELYFQGPSSNK(gl)PR__NA
- _FIK(gl)FQEMK__NA
- _FLQEFYQDDELGK(gl)K__NA
- _FLQEFYQDDELGK(gl)KQFK__NA
- _FLQEFYQDDELGKK(gl)__NA
- _FLQEFYQDDELGKK(gl)QFK__NA
- _FLQEFYQDDELGKKQFK(gl)__NA
- _FQEMK(gl)MQEHSDQVPVGNIPR__NA
- _GNINICLMGDPGVAK(gl)SQLLSYIDR__NA
- _LAASIAPEIYGHEDVK(gl)K__NA
- _LAASIAPEIYGHEDVKK(gl)__NA
- _LFADAVQELLPQYK(gl)ER__NA
- _LMEMSK(gl)DSLLGDK__NA
- _LMEMSK(gl)DSLLGDKGQTAR__NA
- _QPPSQFEPLDMK(gl)LMR__NA
- _REAWASK(gl)DATYTSAR__NA
- _RFELYFQGPSSNK(gl)PR__NA
- _TAIHEVMEQQTISIAK(gl)AGILTTLNAR__NA



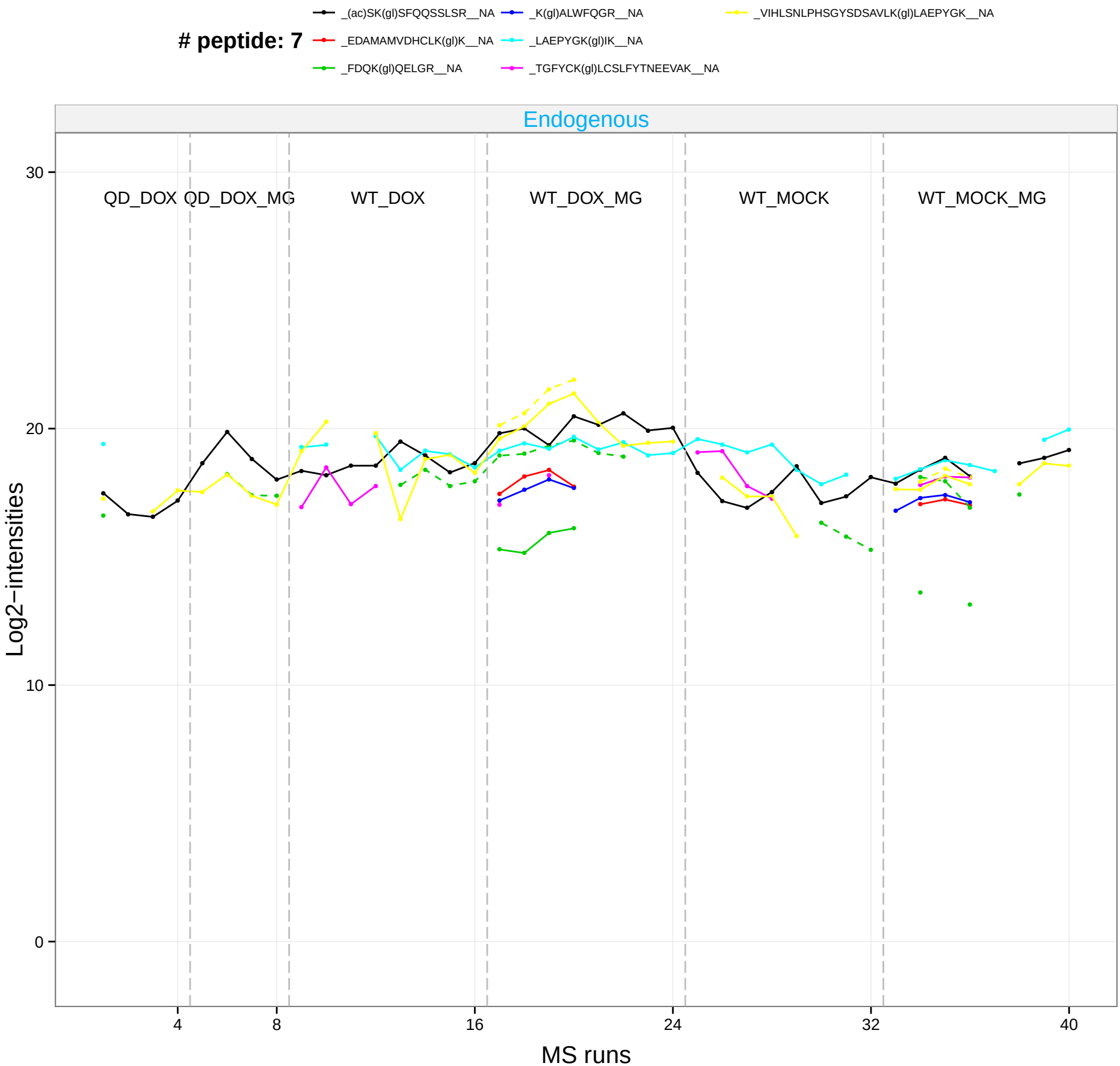
P38159

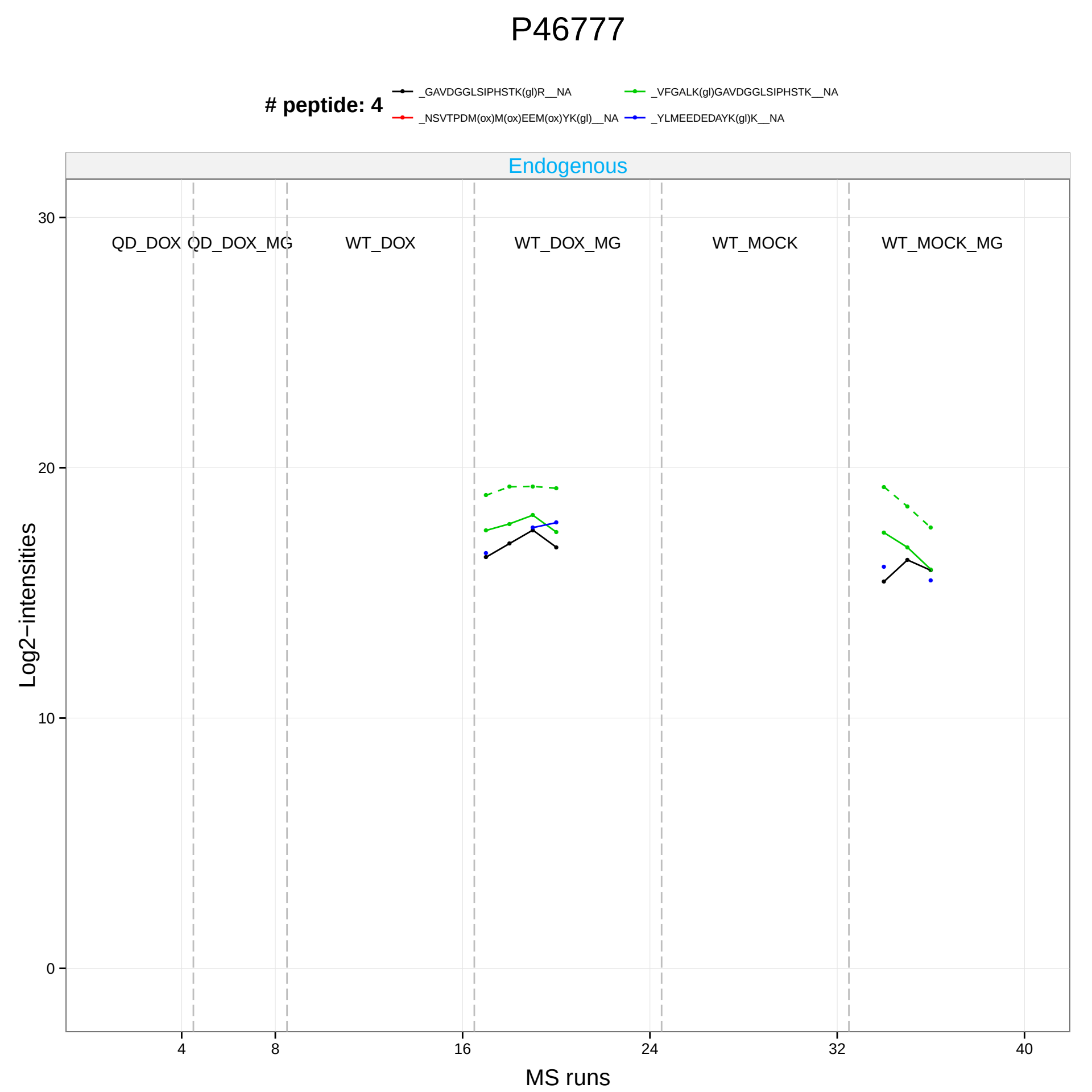

peptide: 2
_LFIGGLNTETNEK(gl)ALEAVFGK__NA
_LFIGGLNTETNEKALEAVFGK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

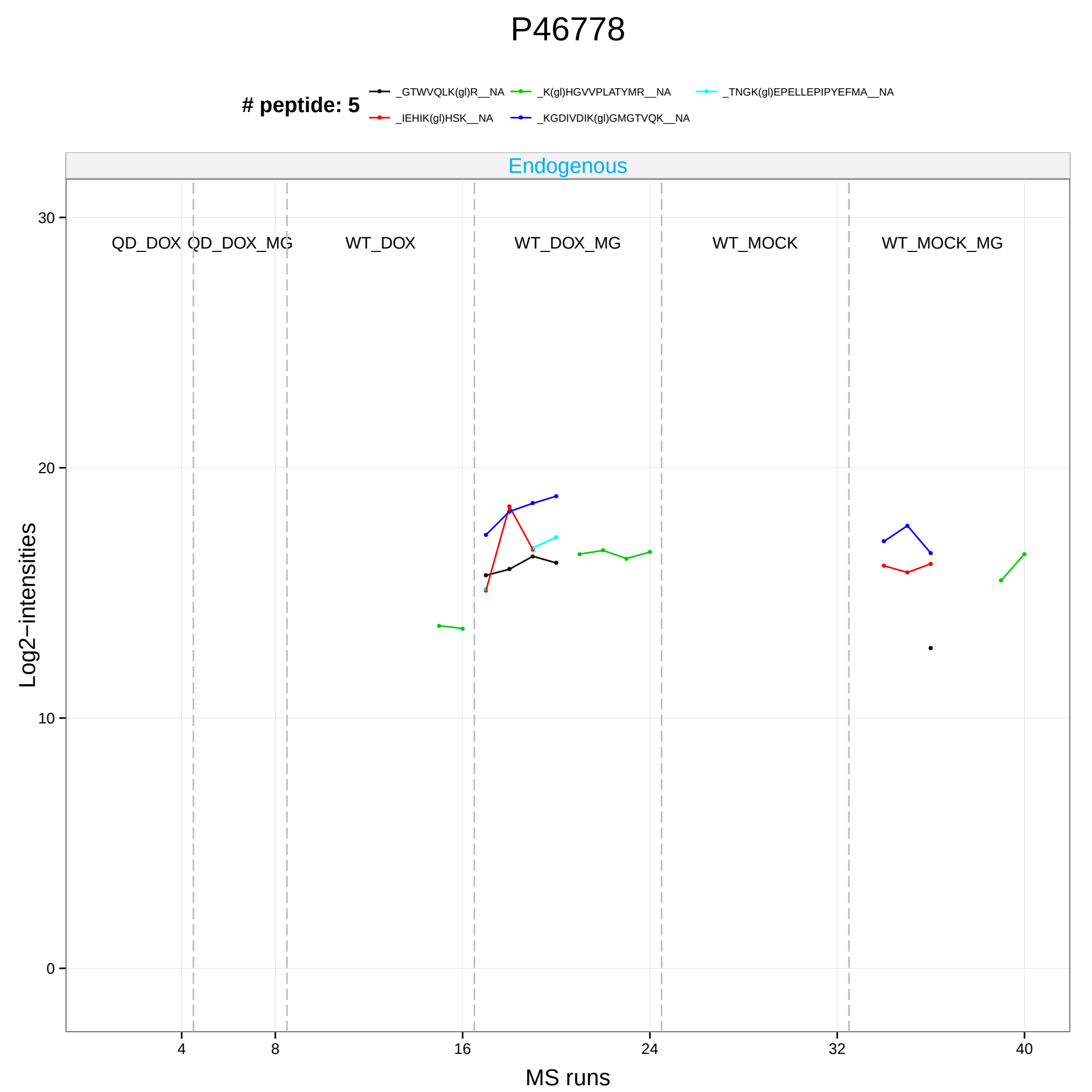
P41440


peptide: 3
_(ac)MVPSSPAVEK(gl)QVPVEPGPDPELR__NA
_MNPGPGGK(gl)LGHALR__NA
_SAAEEK(gl)AAQALSVQDK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

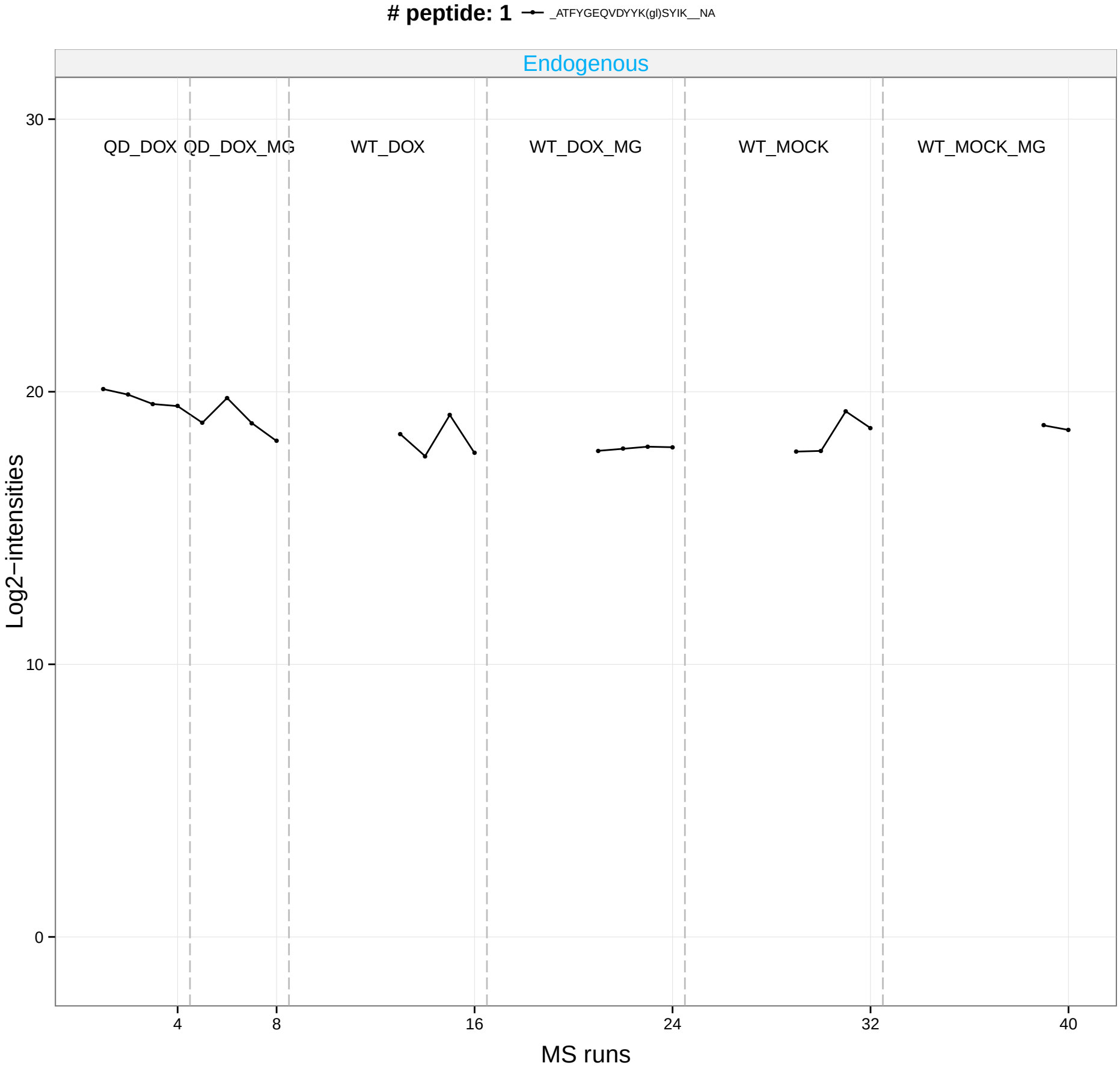
P43243

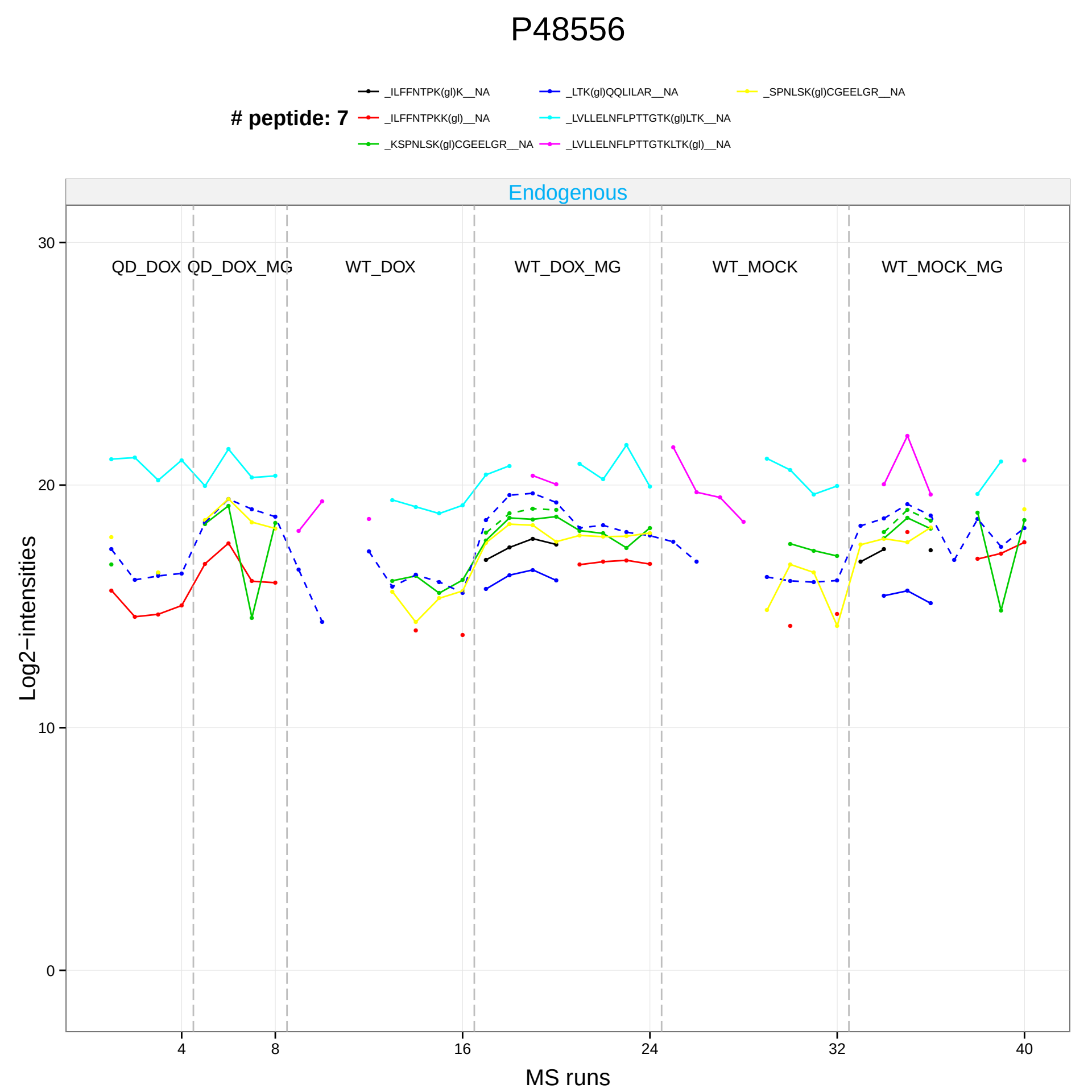

peptide: 7
_(ac)SK(gl)SFQQSSLSR__NA
_K(gl)ALWFQGR__NA
_VIHLSNLPHSGYSDSAVLK(gl)LAEPYGK__NA
_EDAMAMVDHCLK(gl)K__NA
_LAEPYGK(gl)IK__NA
_FDQK(gl)QELGR__NA
_TGFYCK(gl)LCSLFYTNEEVAK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs


P46777
peptide: 4
_GAVDGGLSIPHSTK(gl)R__NA
_VFGALK(gl)GAVDGGLSIPHSTK__NA
_NSVTPDM(ox)M(ox)EEM(ox)YK(gl)__NA
_YLMEEDEDAYK(gl)K__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

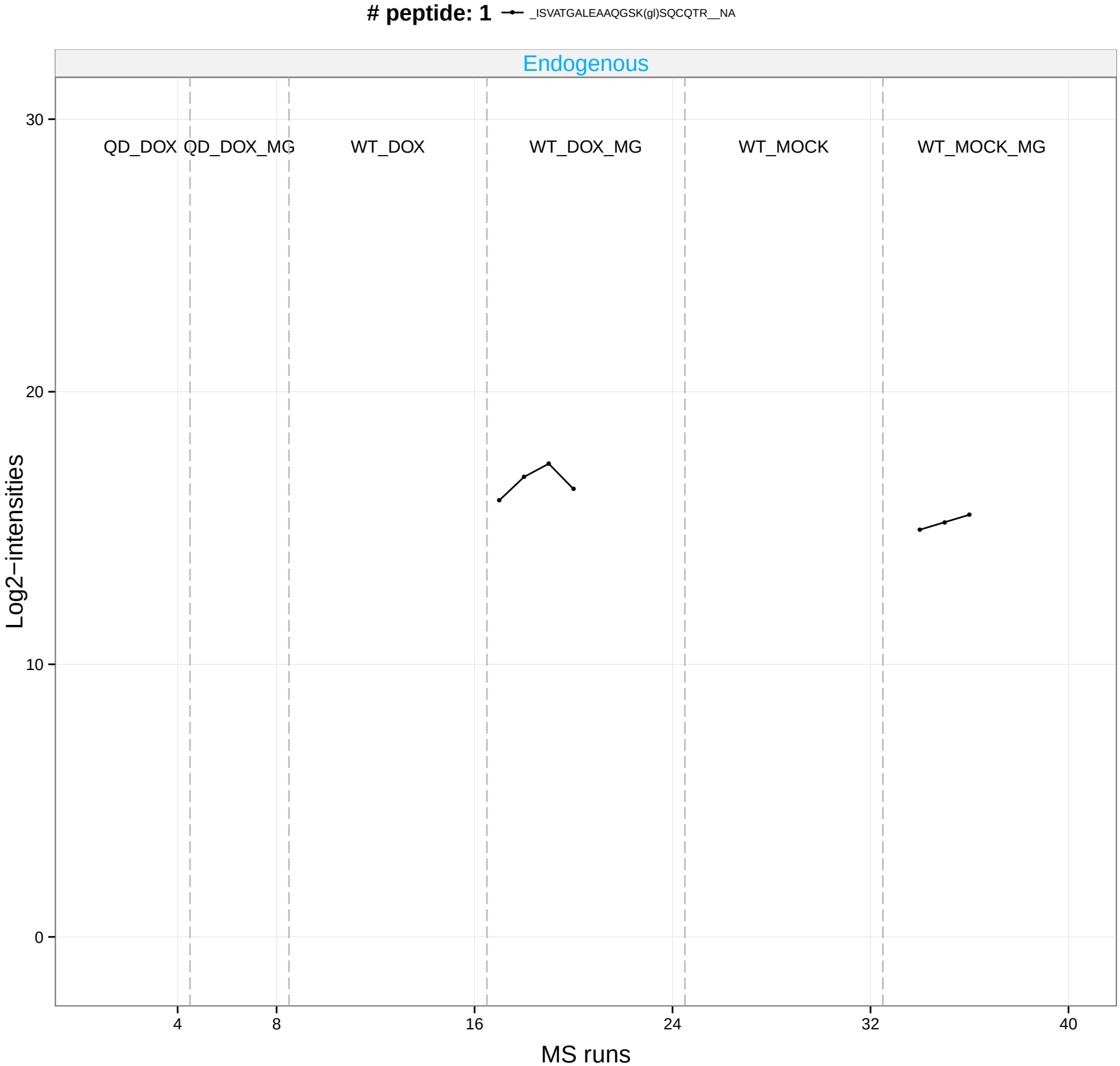

P46778
peptide: 5
_GTWVQLK(gl)R__NA
_K(gl)HGVVPLATYMR__NA
_TNGK(gl)EPELLEPIPYEFMA__NA
_IEHIK(gl)HSK__NA
_KGDIVDIK(gl)GMGTVQK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

P46940


peptide: 1
_ATFYGEQVDYYK(gl)SYIK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs


P48556
peptide: 7
_ILFFNTPK(gl)K__NA
_ILFFNTPKK(gl)__NA
_KSPNLSK(gl)CGEELGR__NA
_LTK(gl)QQLILAR__NA
_LVLLELNFLPTTGTK(gl)LTK__NA
_LVLLELNFLPTTGTKLTK(gl)__NA
_SPNLSK(gl)CGEELGR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

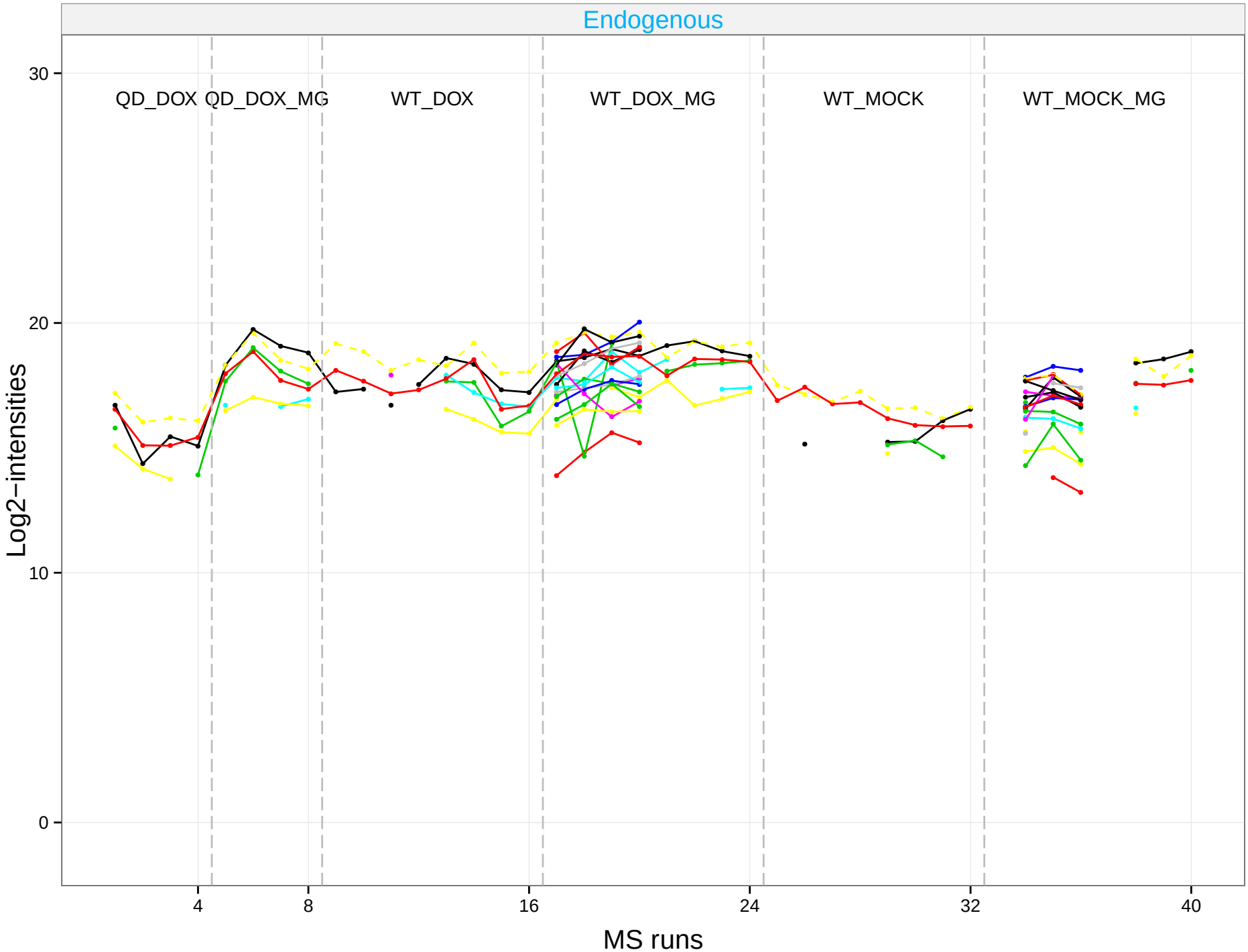
P51610


peptide: 1
_ISVATGALEAAQGSK(gl)SQCQTR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

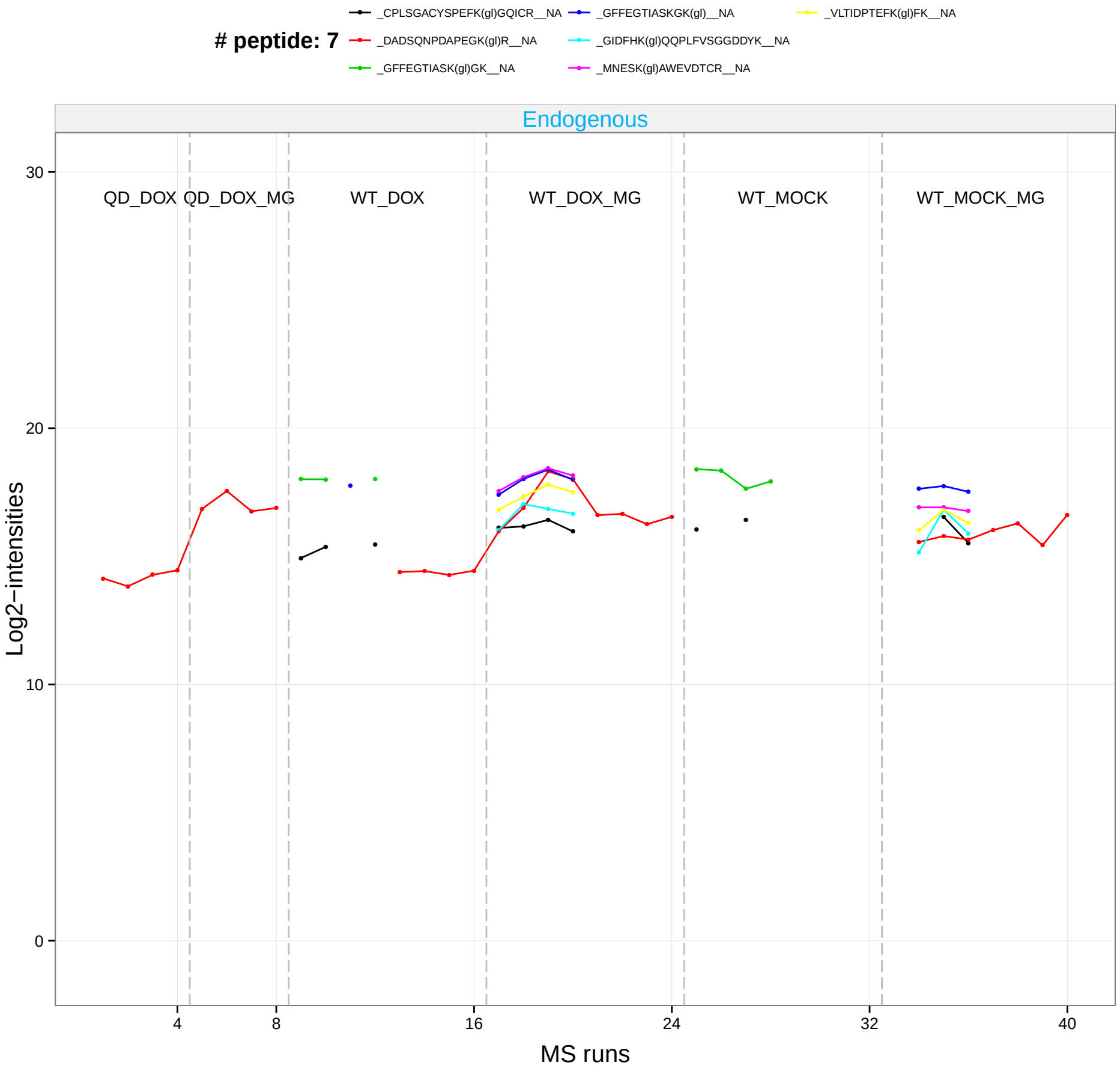
P52701

 # peptide: 19

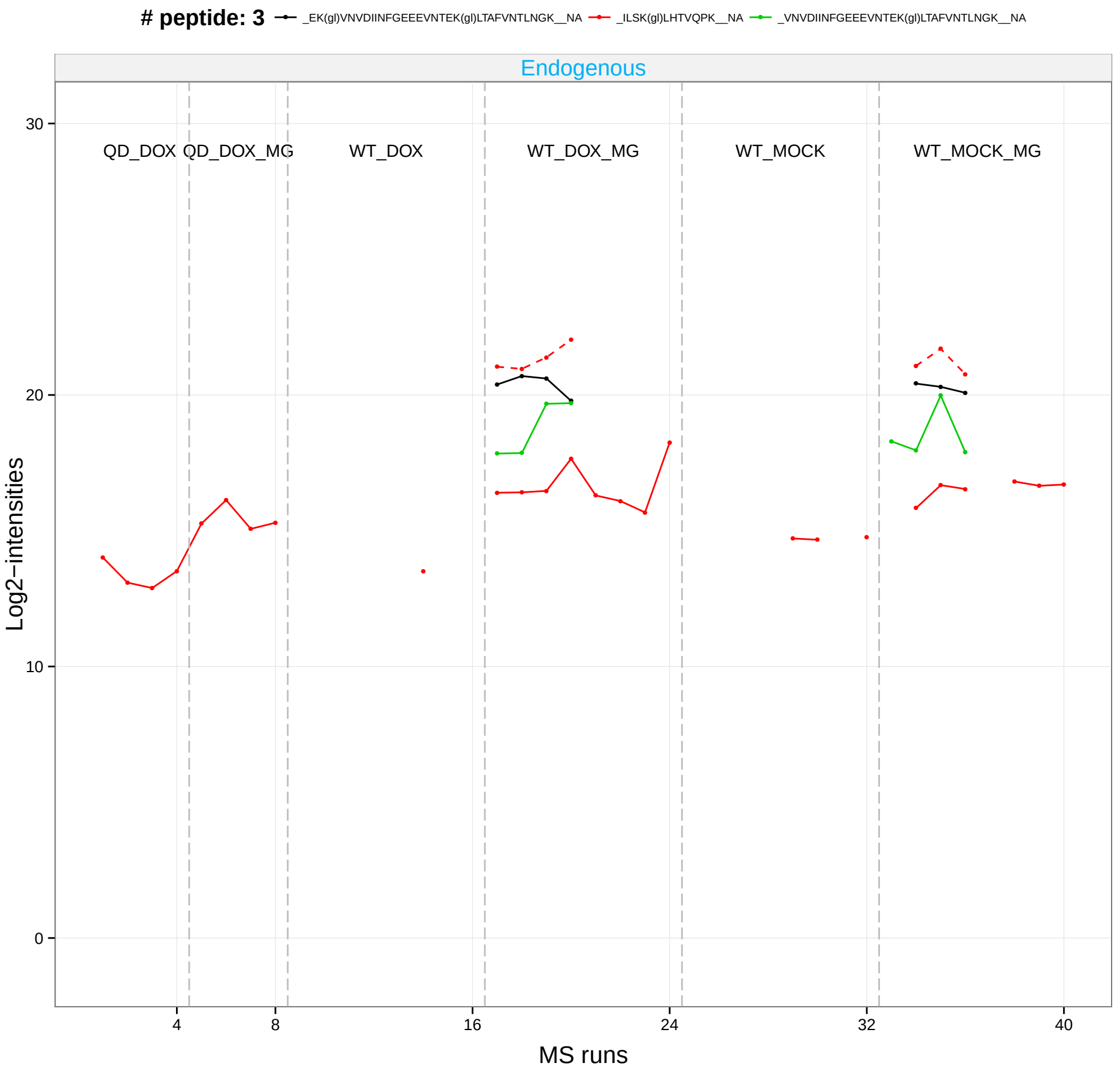
- _AYCVLVTGPNMGGK(gl)STLMR__NA
- _DVSLK(gl)DCMR__NA
- _GACPK(gl)SYGFNAAR__NA
- _GGDGPMCRPVILLPEDTPPFLELK(gl)GSR__NA
- _GTQTYSVLEGDPSENYSK(gl)YLLSLK__NA
- _IIGIMEEVADGFKSK(gl)__NA
- _IITK(gl)GTQTYSVLEGDPSENYSK__NA
- _ILK(gl)QVISLQTK__NA
- _LANLPEEVIQK(gl)GHR__NA
- _LLK(gl)QWLCAPLCNHYAINDR__NA
- _LLSK(gl)IHNVGSPLK__NA
- _QATSISSETK(gl)NTLR__NA
- _QVISLQTK(gl)NPEGR__NA
- _SELALSALGGCVFYLK(gl)K__NA
- _SGAIFTK(gl)AYQR__NA
- _SSLSCSLQEGLIPGSQFWDASK(gl)TLR__NA
- _TILK(gl)SSLSCSLQEGLIPGSQFWDASK__NA
- _YSDSLVQK(gl)GYK__NA
- _YWTK(gl)TIEK__NA



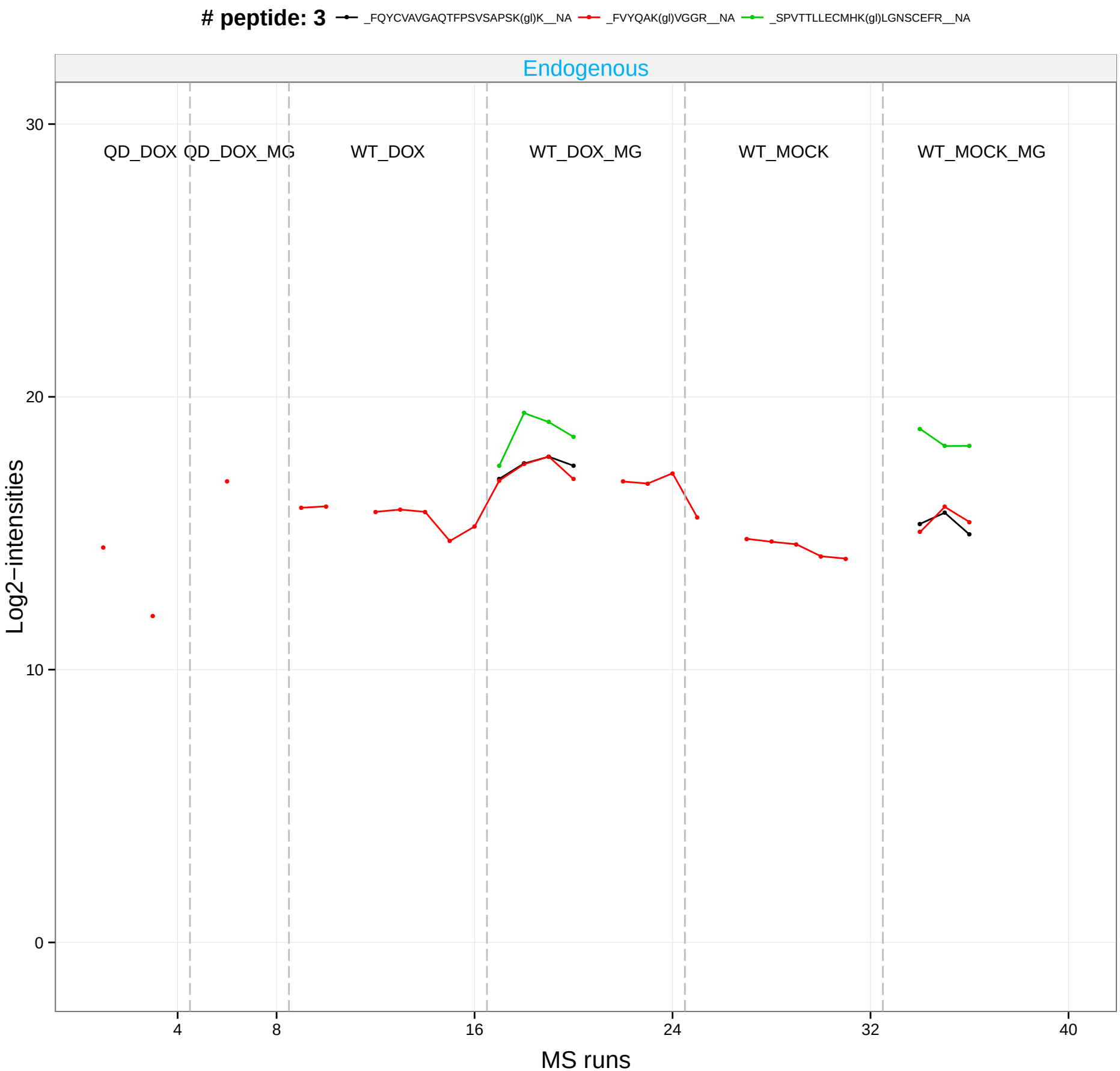
P53621


peptide: 7
_CPLSGACYSPEFK(gl)GQICR__NA
_GFFEGTIASKGK(gl)__NA
_VLTIDPTEFK(gl)FK__NA
_DADSQNPDAPEGK(gl)R__NA
_GIDFHK(gl)QQPLFVSGGDDYK__NA
_GFFEGTIASK(gl)GK__NA
_MNESK(gl)AWEVDTCR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

P55036;A2A3N6



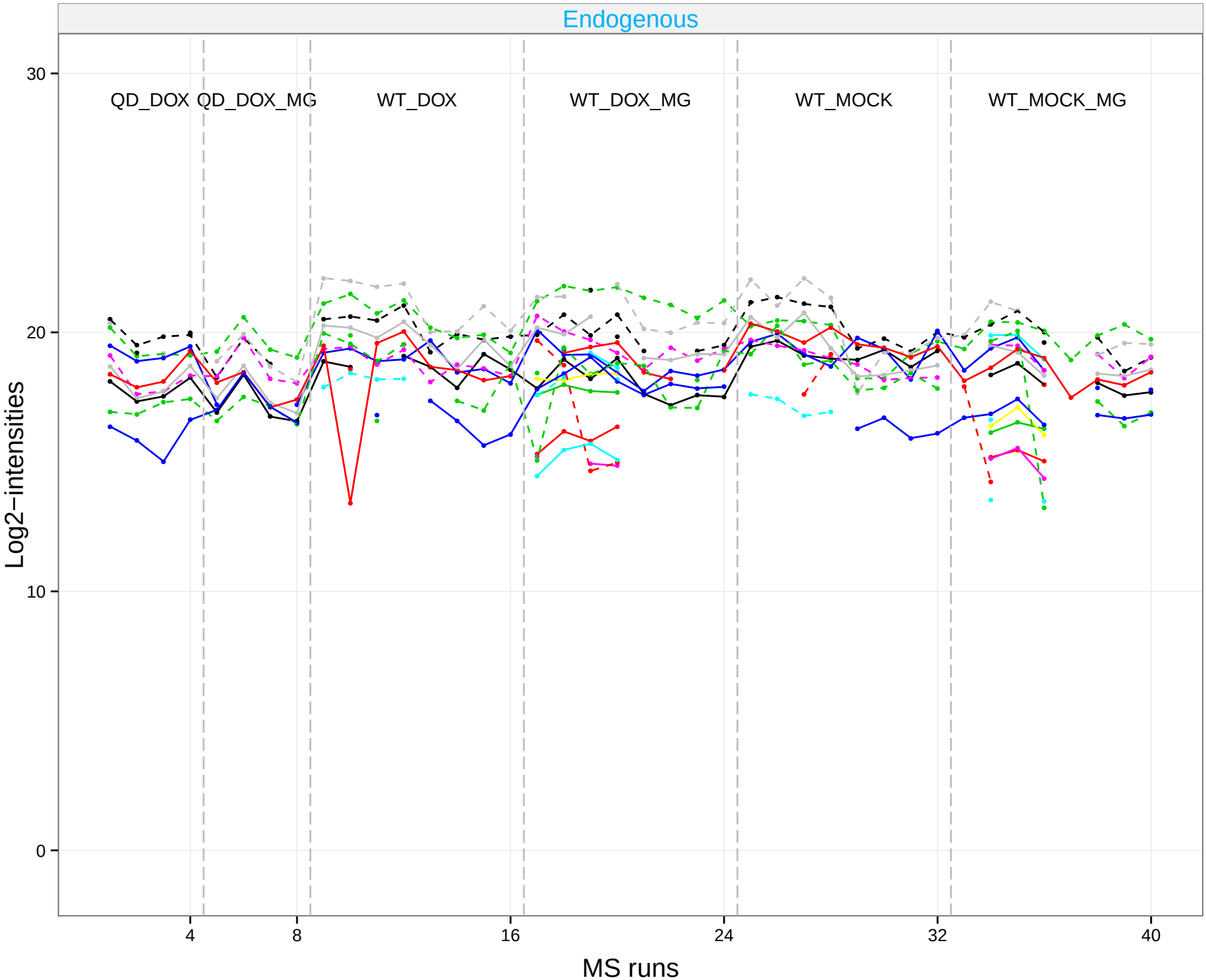
P55265


peptide: 3
_FQYCVAVGAQTFPSVSAPSK(gl)K__NA
_FVYQAK(gl)VGGR__NA
_SPVTTLLECMHK(gl)LGNSCEFR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

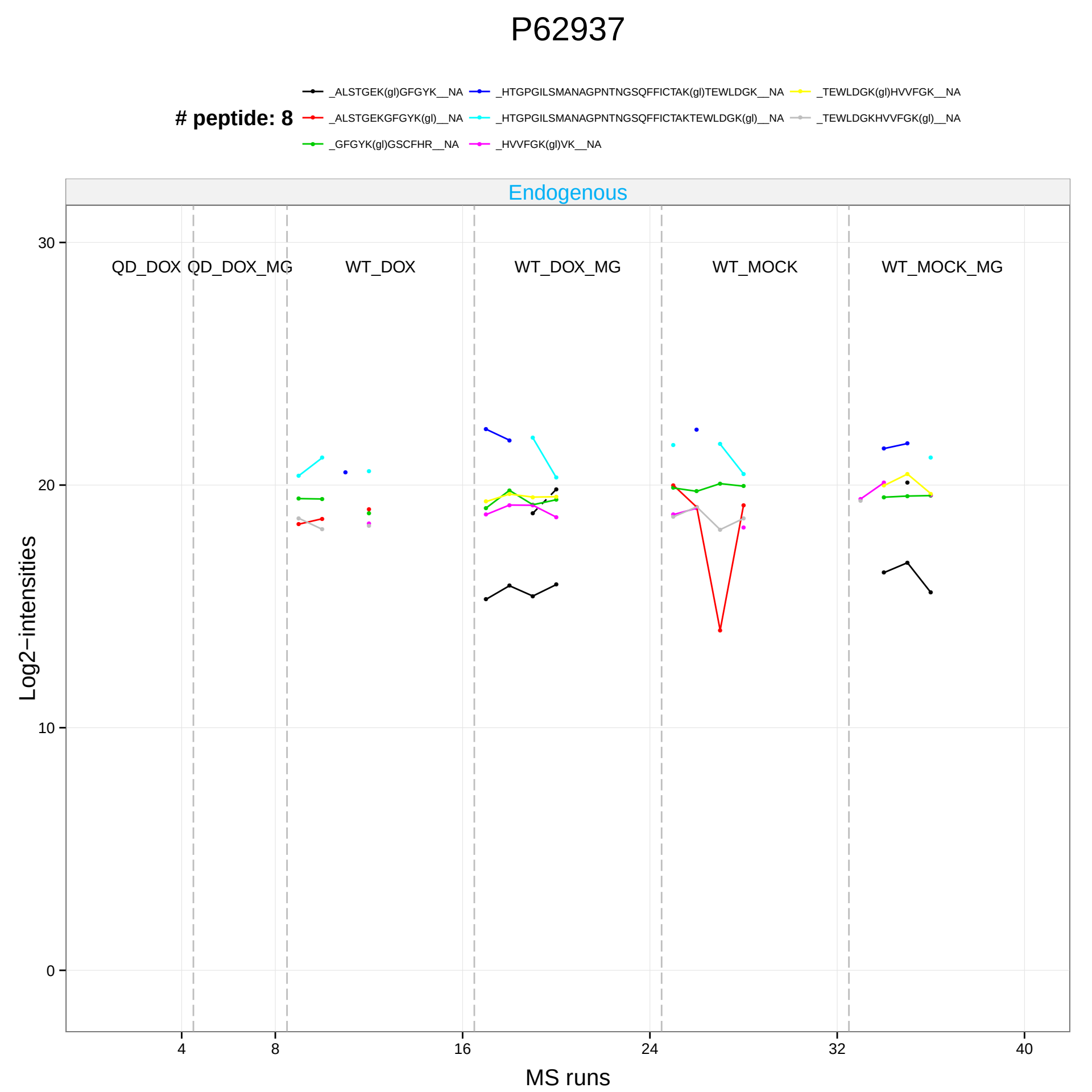
P61978

 # peptide: 13

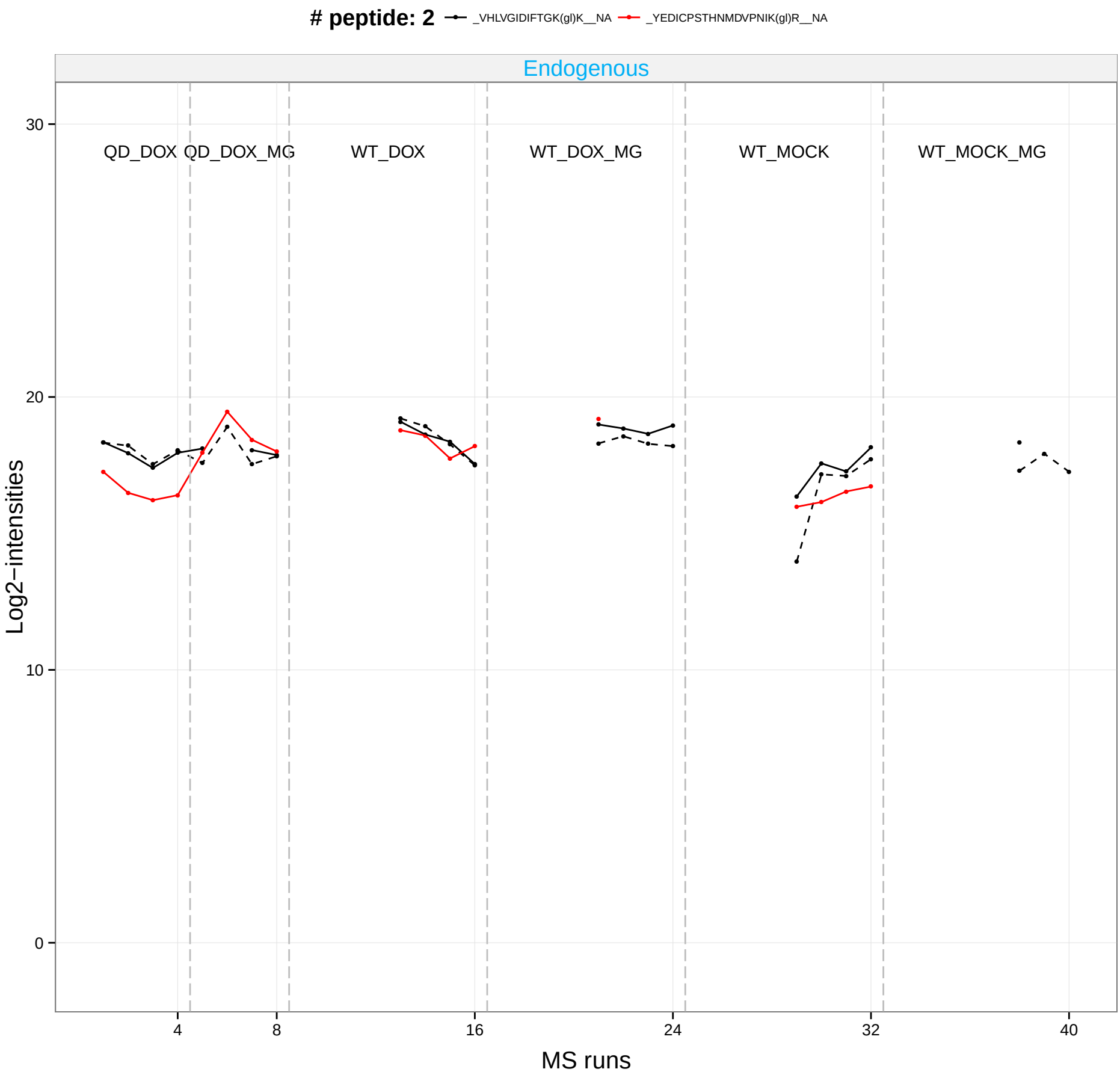
_DLAGSIIGK(gl)GGQR__NA
_ENTQTTIK(gl)LFQECCPHSTDR__NA
_IILDLISESPIK(gl)GR__NA
_IITITGTQDQIQNAQYLLQNSVK(gl)QYSGK__NA
_IITITGTQDQIQNAQYLLQNSVKQYSGK(gl)__NA
_ILLQSK(gl)NAGAVIGK__NA
_ILSISADIETIGEILK(gl)K__NA
_LLIHQSLAGGIIGVK(gl)GAK__NA
_LLIHQSLAGGIIGVKGAK(gl)__NA
_NAGAVIGK(gl)GGK__NA
_NAGAVIGKGGK(gl)__NA
_RPAEDMEEEQAFK(gl)R__NA
_VVLIGGK(gl)PDR__NA



P62937


peptide: 8
_ALSTGEK(gl)GFGYK__NA
_ALSTGEKGFGYK(gl)__NA
_GFGYK(gl)GSCFHR__NA
_HTGPGILSMANAGPNTNGSQFFICTAK(gl)TEWLDGK__NA
_HTGPGILSMANAGPNTNGSQFFICTAKTEWLDGK(gl)__NA
_HVVFGK(gl)VK__NA
_TEWLDGK(gl)HVVFGK__NA
_TEWLDGKHVVFGK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

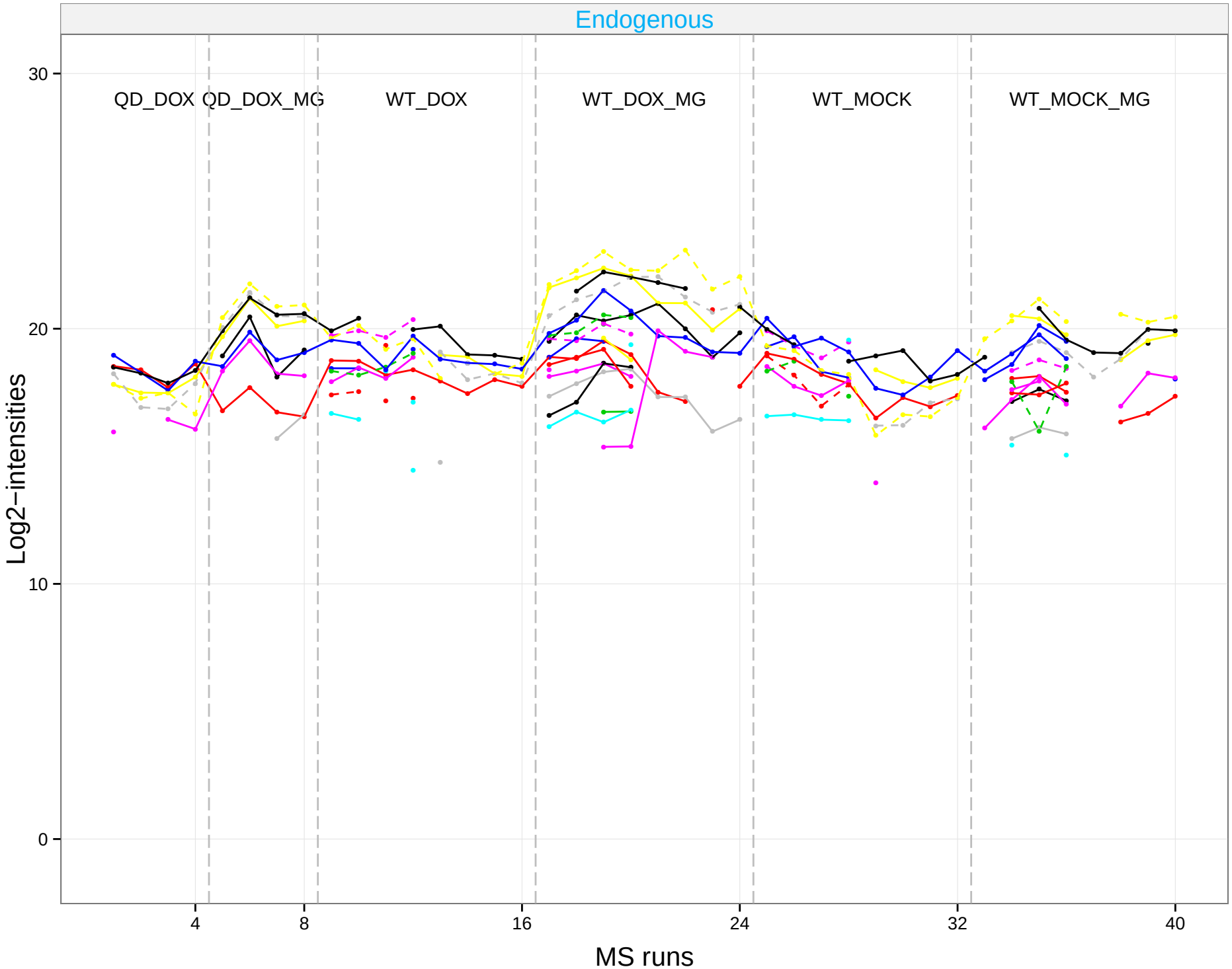
P63241;Q6IS14;Q9GZV4



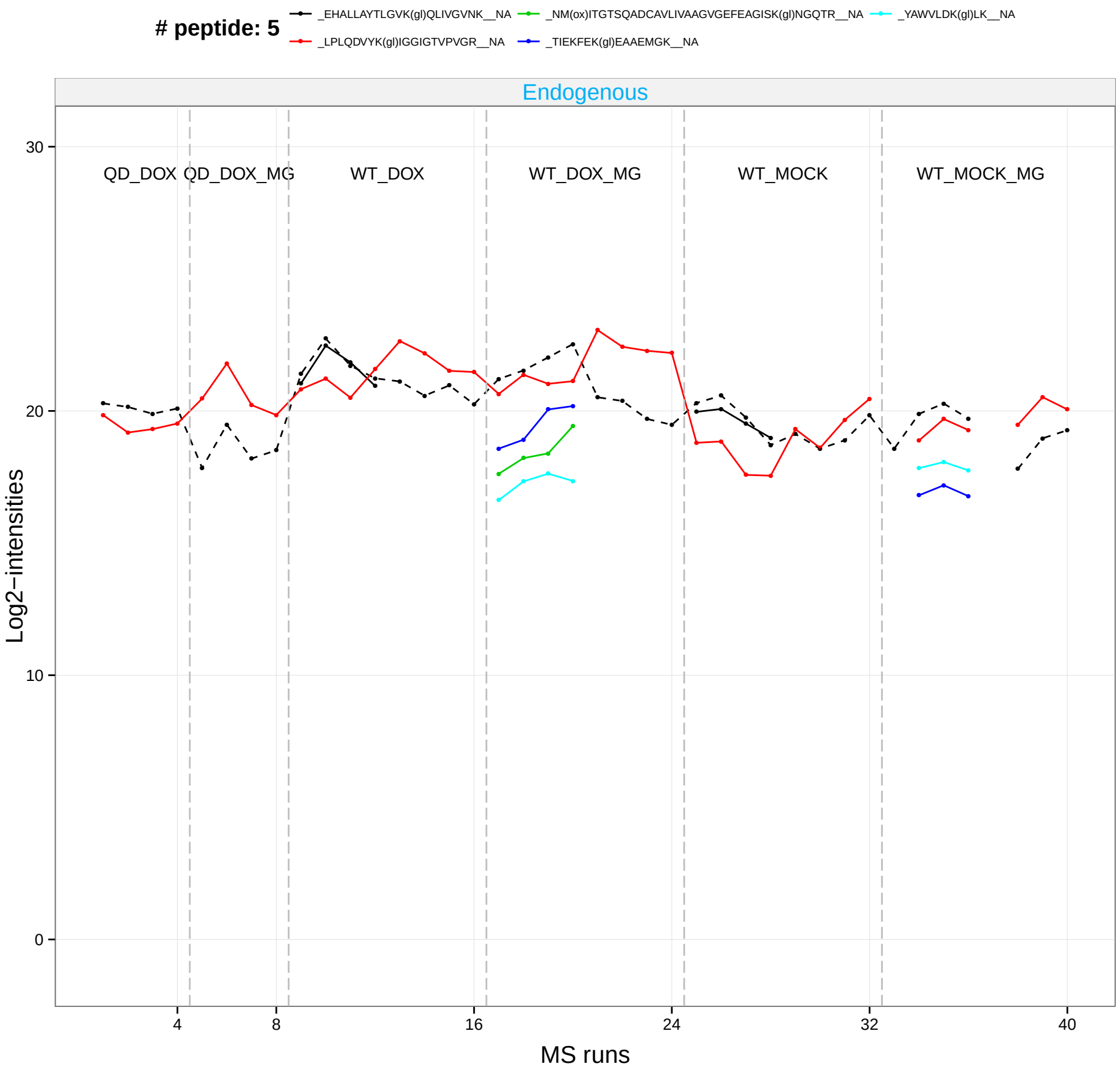
P63244

 # peptide: 18

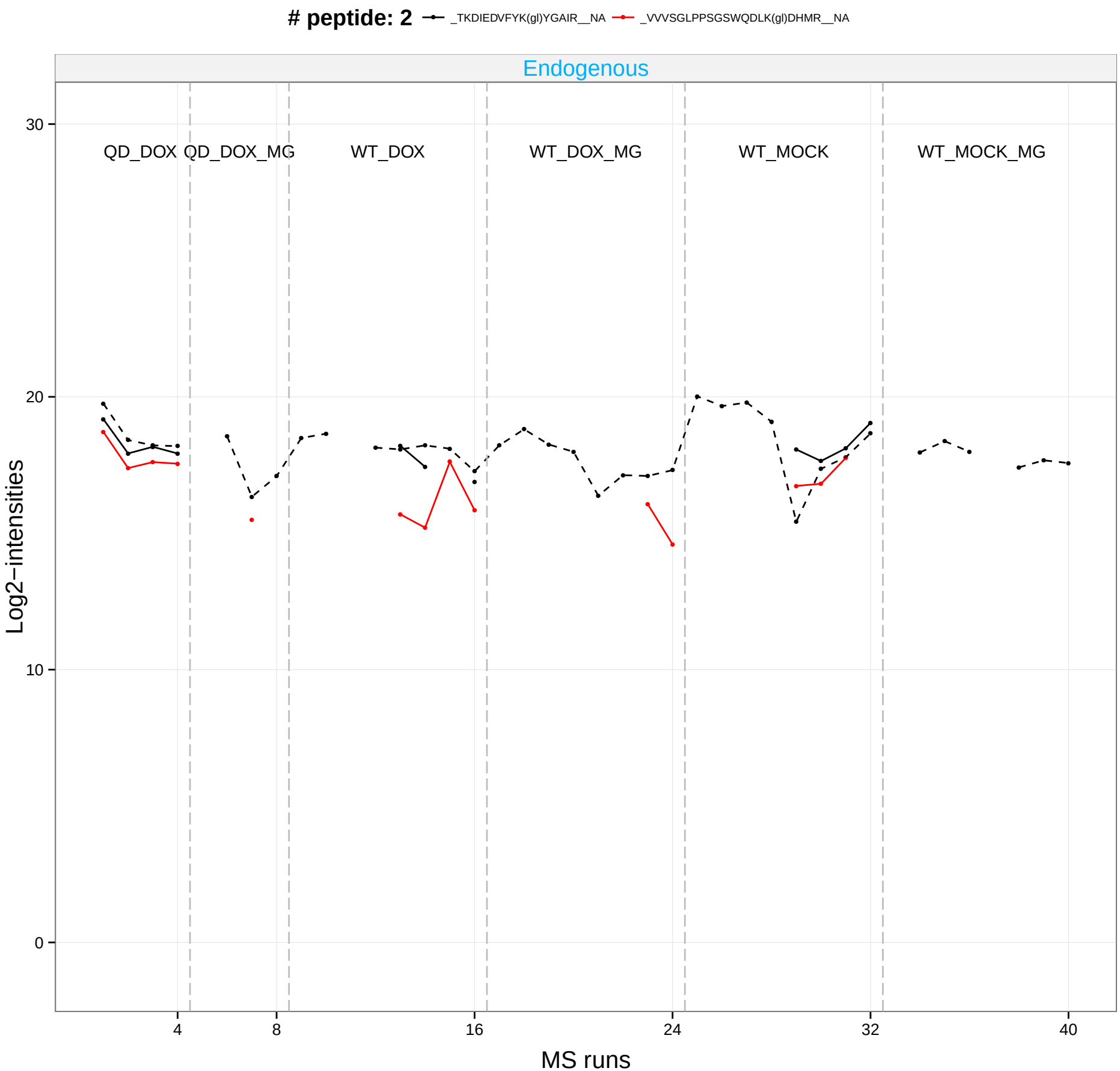
_DGQAMLWDLNEGK(gl)HLYTLDGGDIINALCFSPNR__NA
_DKTIIMWK(gl)LTR__NA
_DKTIK(gl)LWNTLGVCK__NA
_FSPNSSNPIIVSCGWDK(gl)LVK__NA
_FSPNSSNPIIVSCGWDKLVK(gl)__NA
_FVGHTK(gl)DVLSVAFSSDNR__NA
_GTLK(gl)GHNGWVTQIATTPQFPDMILSASR__NA
_IIVDELK(gl)QEVISTSSK__NA
_IWDLEGK(gl)IIVDELKQEVISTSSK__NA
_LK(gl)TNHIGHTGYLNTVTVSPDGSLCASGGK__NA
_LVK(gl)VWNLANCK__NA
_LWNTLGVCK(gl)YTVQDESHSEWVSCVR__NA
_RFVGHTK(gl)DVLSVAFSSDNR__NA
_TIK(gl)LWNTLGVCK__NA
_VWNLANCK(gl)LK__NA
_VWNLANCKLK(gl)__NA
_YWLCAATGPSIK(gl)IWDLEGK__NA
_YWLCAATGPSIKIWDLEGK(gl)__NA



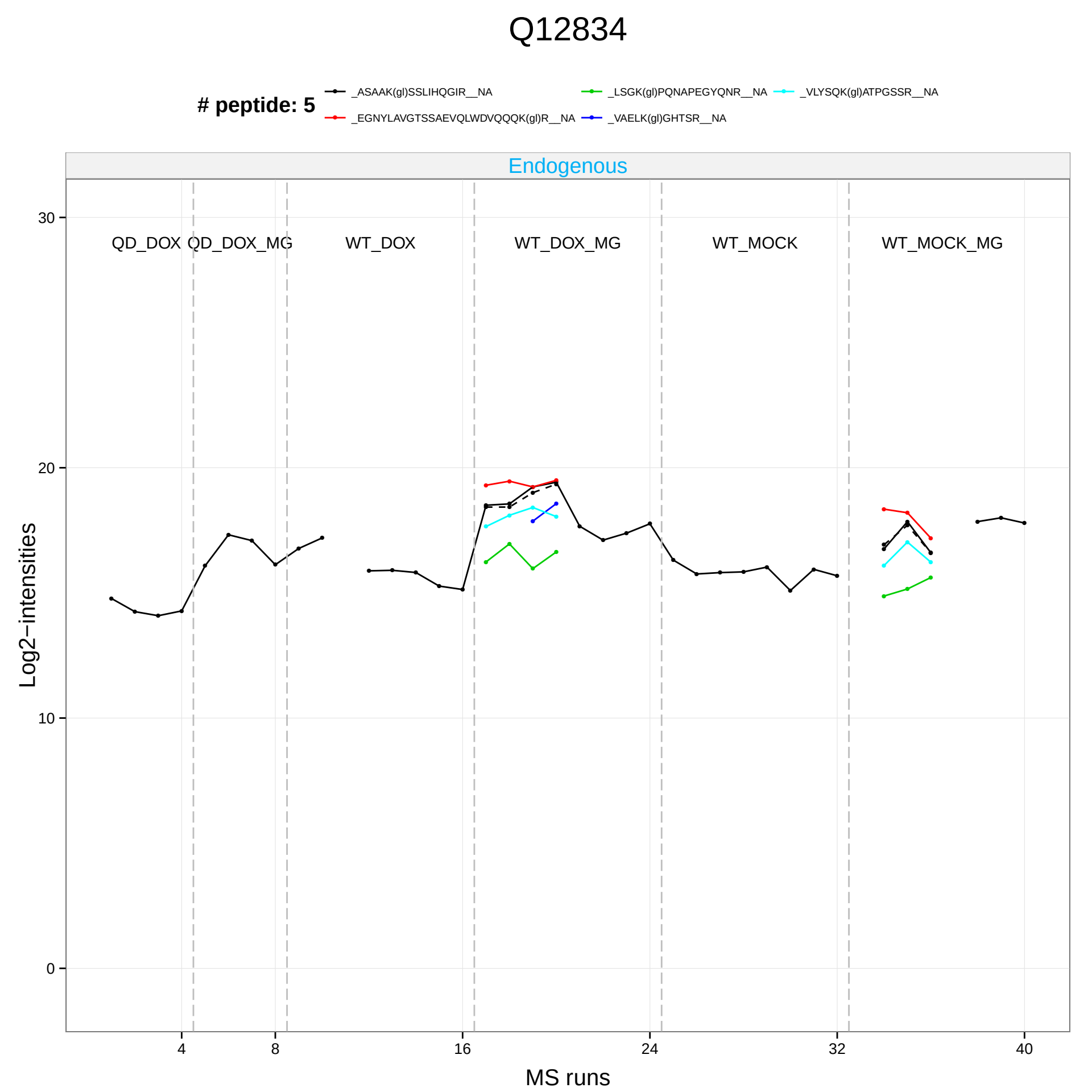
P68104;Q5VTE0;Q05639



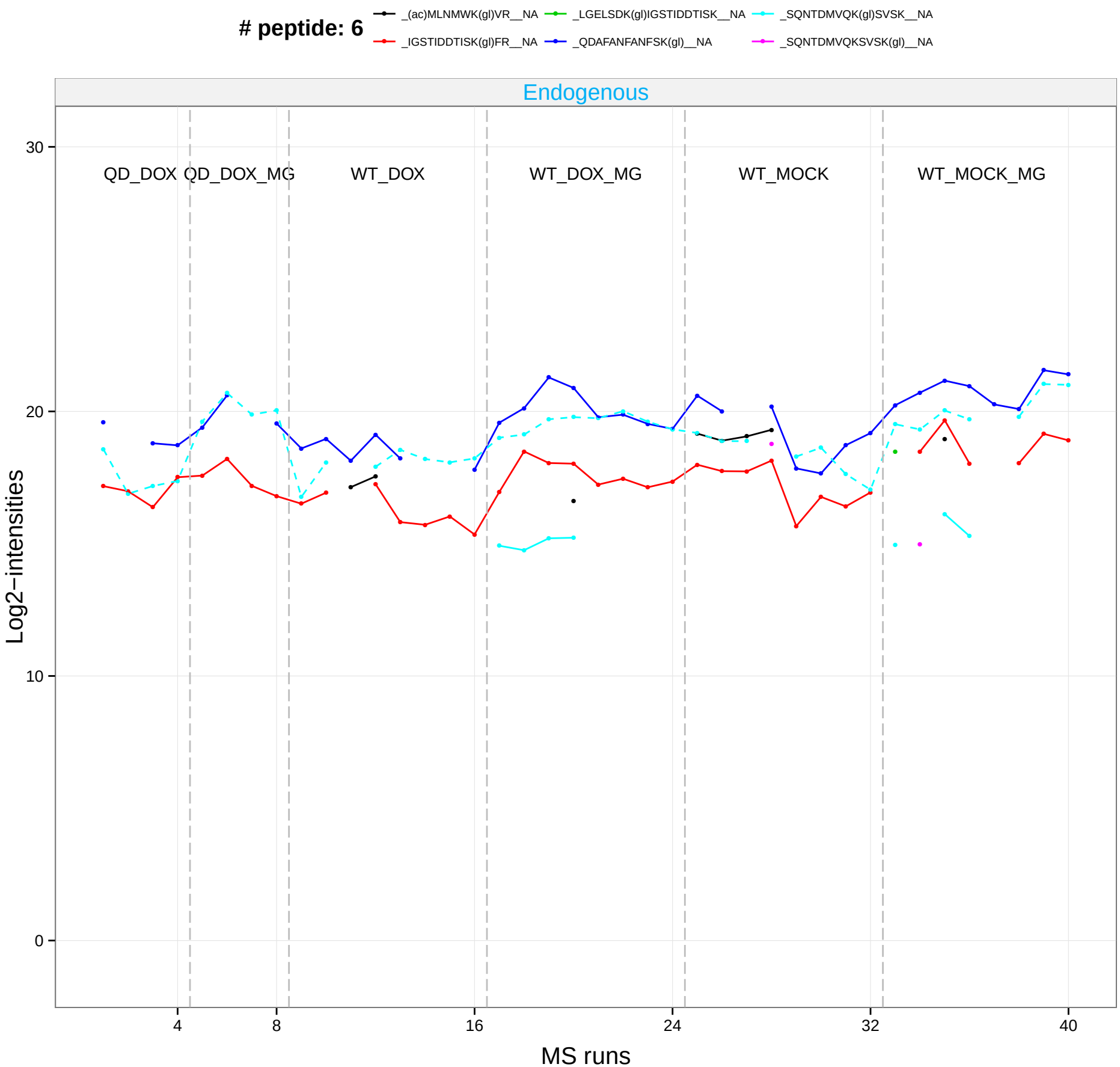
Q07955


peptide: 2
_TKDIEDVFYK(gl)YGAIR__NA
_VVVSGLPPSGSWQDLK(gl)DHMR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

Q12834


peptide: 5
_ASAAK(gl)SSLIHQGIR__NA
_LSGK(gl)PQNAPEGYQNR__NA
_VLYSQK(gl)ATPGSSR__NA
_EGNYLAVGTSSAEVQLWDVQQQK(gl)R__NA
_VAELK(gl)GHTSR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

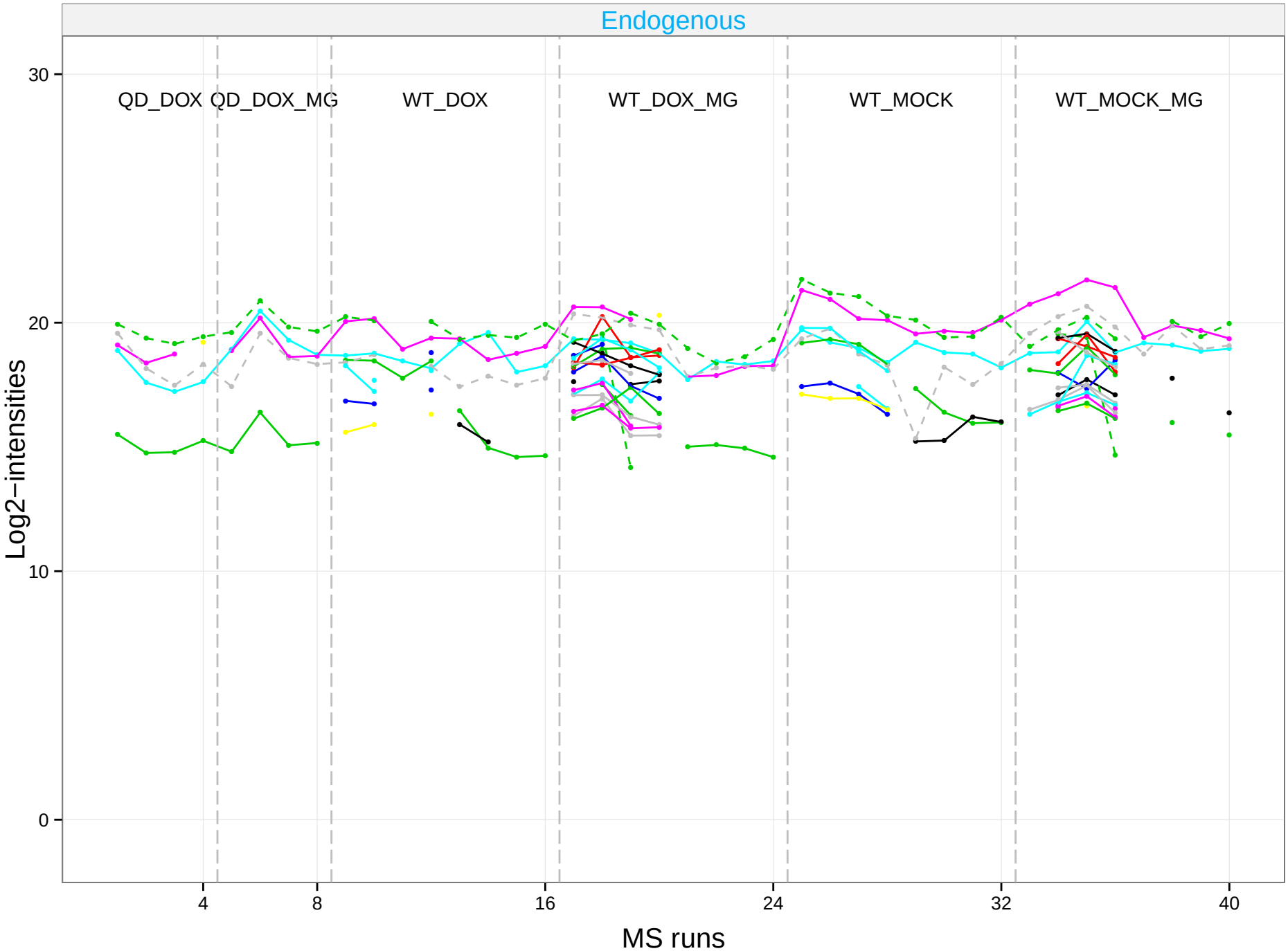
Q14677


peptide: 6
_(ac)MLNMWK(gl)VR__NA
_LGELSDK(gl)IGSTIDDTISK__NA
_SQNTDMVQK(gl)SVSK__NA
_IGSTIDDTISK(gl)FR__NA
_QDAFANFANFSK(gl)__NA
_SQNTDMVQKSVSK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

Q15154

 # peptide: 24

_DPQQEPMEEIENLK(gl)K__NA
_DPQQEPMEEIENLKK(gl)__NA
_DTGVNEK(gl)AR__NA
_EKFYEAK(gl)LQQQQR__NA
_FYEAK(gl)LQQQQR__NA
_IQALQTACPDLQLSAASVGNCPTK(gl)K__NA
_IQALQTACPDLQLSAASVGNCPTKK(gl)__NA
_K(gl)PFNFLPMQINTNK__NA
_KPFNFLPMQINTNK(gl)__NA
_LIDIQEK(gl)IQALQTACPDLQLSAASVGNCPTKK__NA
_LK(gl)QMLNELMR__NA
_LMAAK(gl)QK__NA
_LMQDLDNNSITVK(gl)QR__NA
_LNNMDWGAQQK(gl)K__NA
_LNNMDWGAQQKK(gl)__NA
_LPDEK(gl)VELFSK__NA
_LPDEKVELFSK(gl)MR__NA
_LQK(gl)LNEVR__NA
_NNCPFSADENYRPLAK(gl)TR__NA
_QLGSILQDSLAK(gl)FAGR__NA
_TPWLYEQEGEVEKPFIK(gl)TGFSVSVEK__NA
_VELFSK(gl)MR__NA
_WK(gl)NNCPFSADENYRPLAK__NA
_YMPAVTSTPTVNQHETSTSK(gl)SVFEPEDSSIVDNELWSEMR__NA



Q16186

 # peptide: 11

_AMQNNAK(gl)PEQK__NA
_GASNK(gl)YLVEFR__NA
_GLVYIQQTDDSLIHFCWK(gl)DR__NA
_GTTVTPDK(gl)R__NA
_KGLVYIQQTDDSLIHFCWK(gl)DR__NA
_M(ox)SLK(gl)GTTVTPDK__NA
_M(ox)SLK(gl)GTTVTPDKR__NA
_MSLK(gl)GTTVTPDK__NA
_MSLK(gl)GTTVTPDK(gl)R__NA
_MSLK(gl)GTTVTPDKR__NA
_TSGNVEDDLIIFPDDCEFK(gl)R__NA

Endogenous

QD_DOX QD_DOX_MG WT_DOX WT_DOX_MG WT_MOCK WT_MOCK_MG

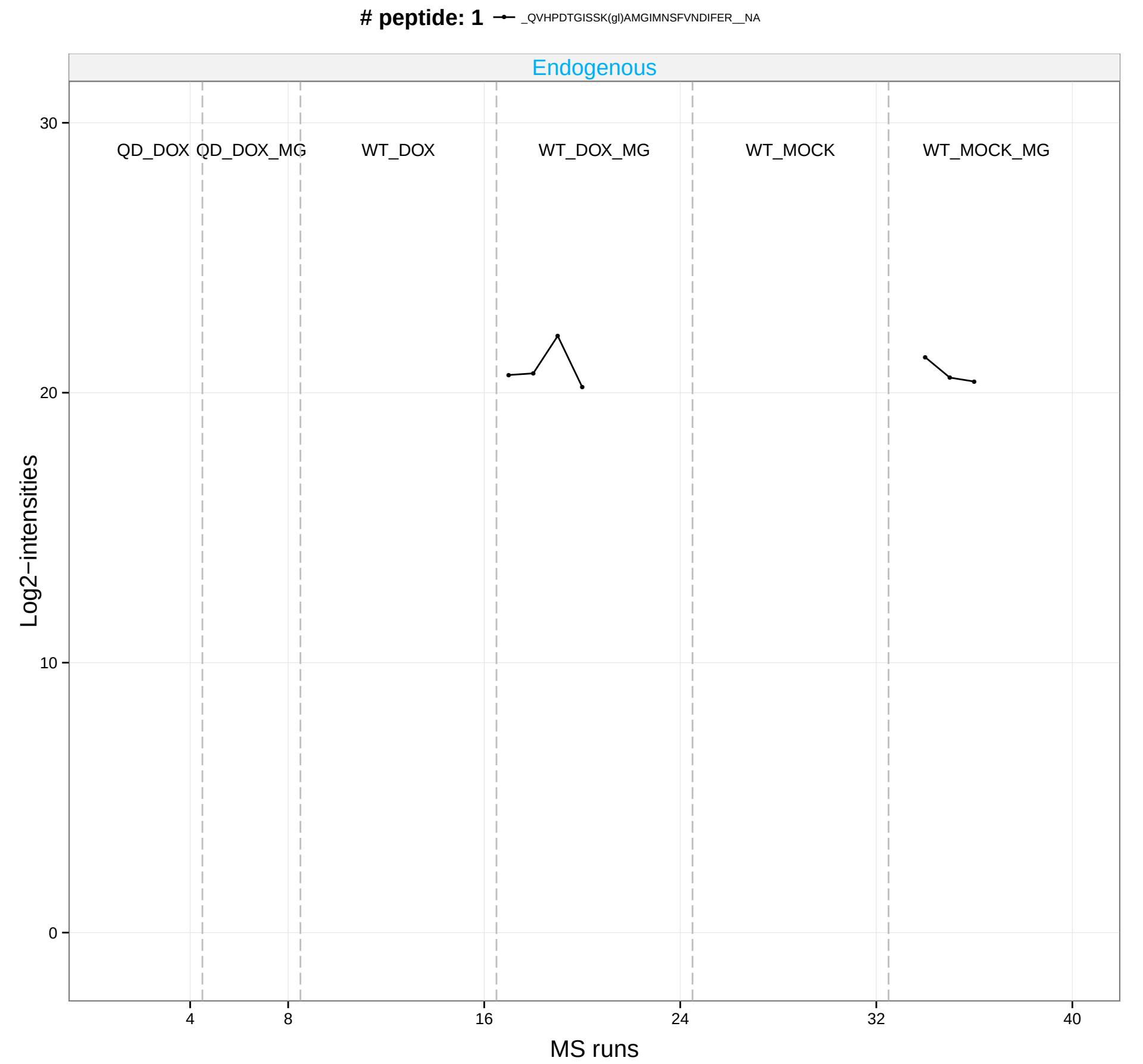
Log2−intensities

30
20
10
0

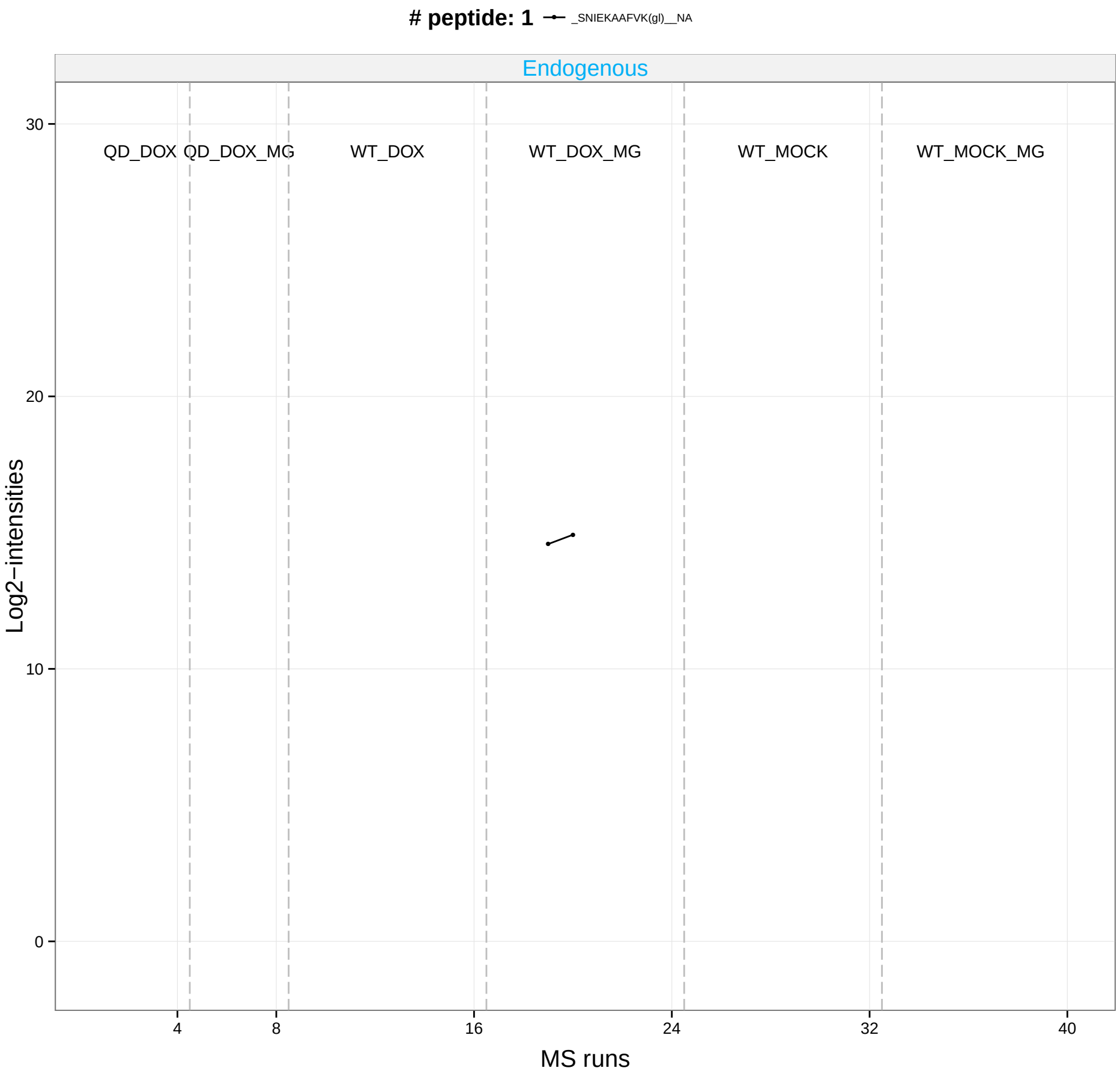
4 8 16 24 32 40

MS runs

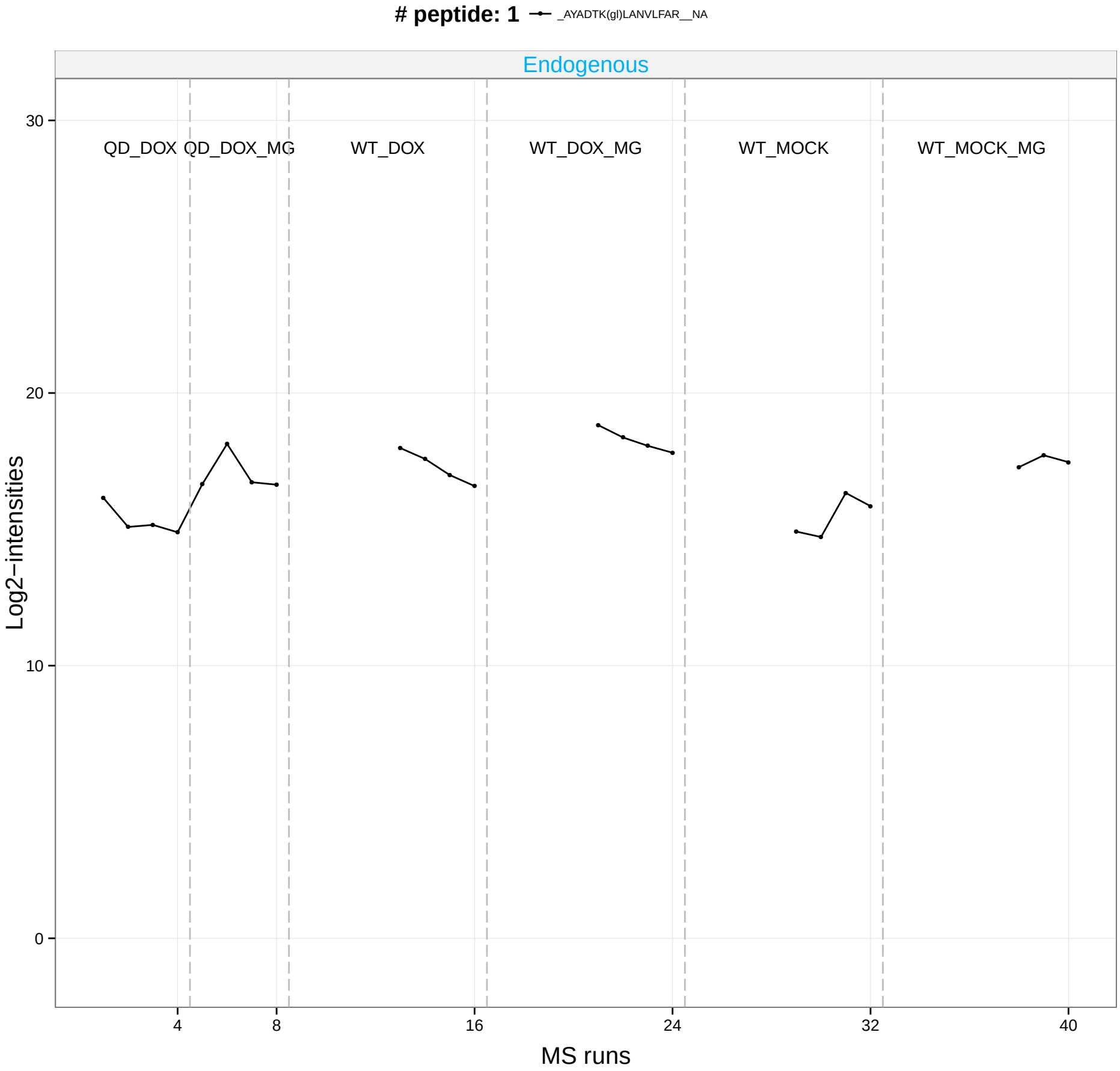
3079;Q99877;Q99879;Q99880;P23527;P33778;Q8N257;P58876;O60814



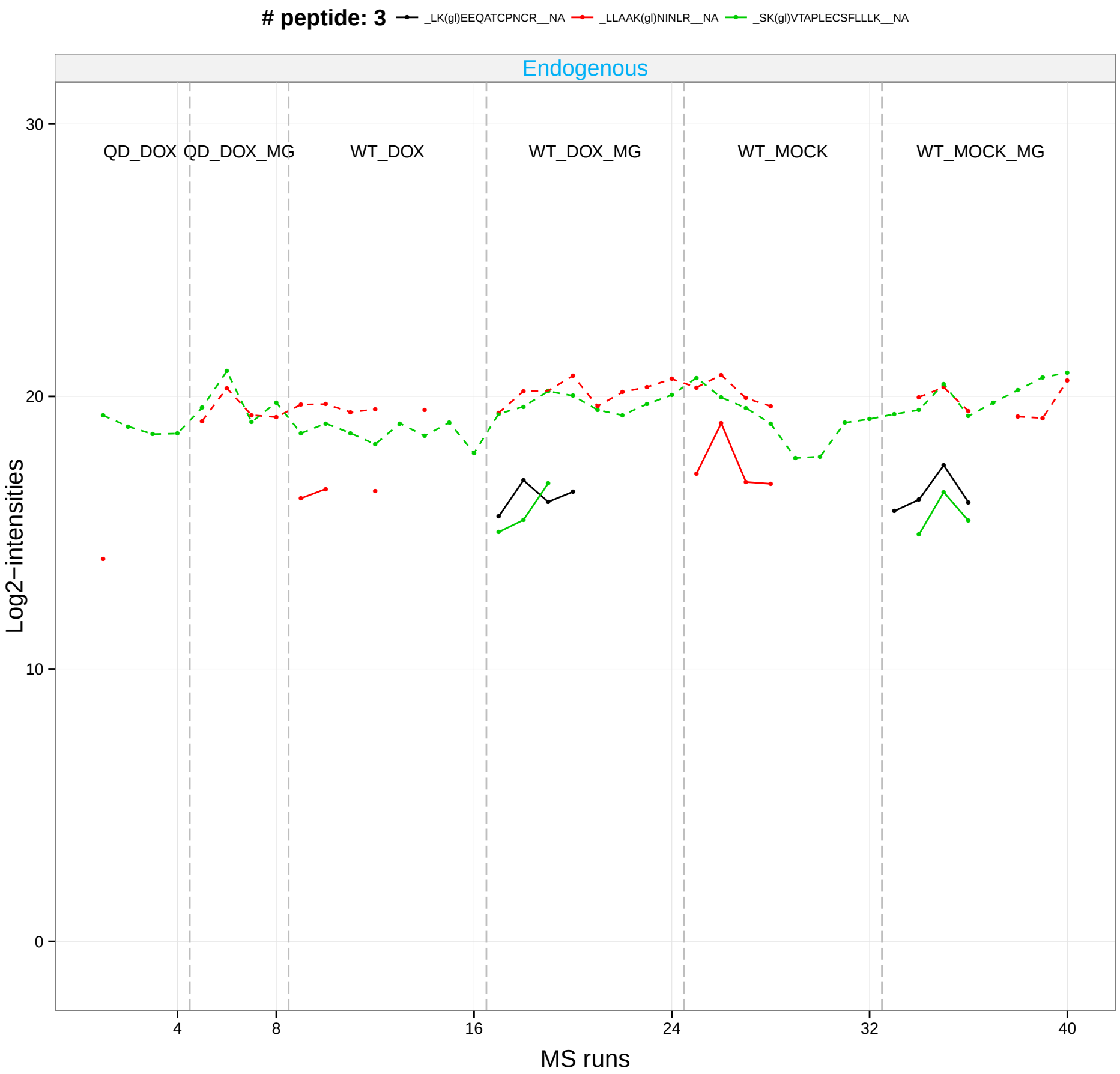
Q5VW36


peptide: 1
_SNIEKAAFVK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

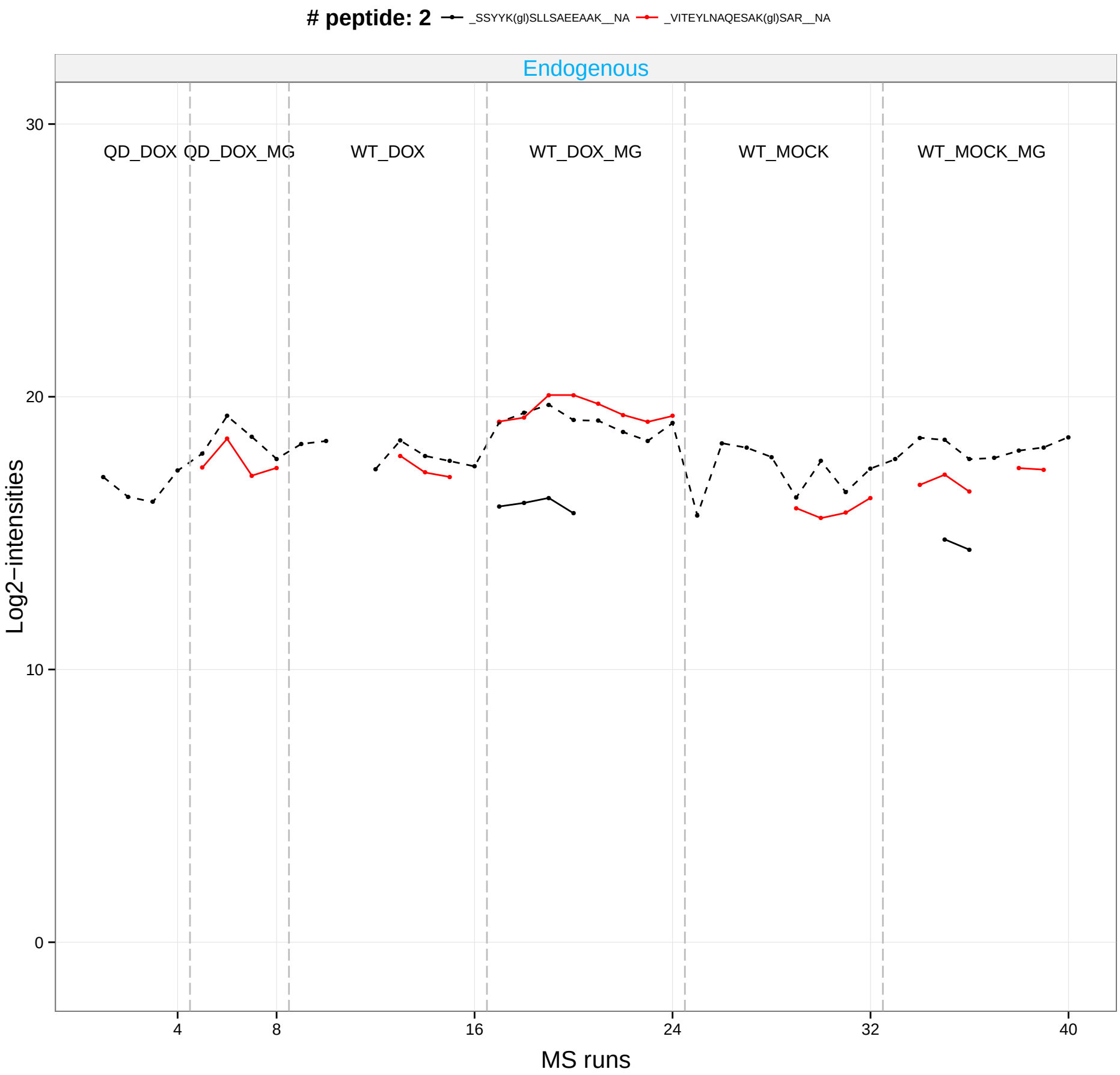
Q6UX07


peptide: 1
_AYADTK(gl)LANVLFAR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

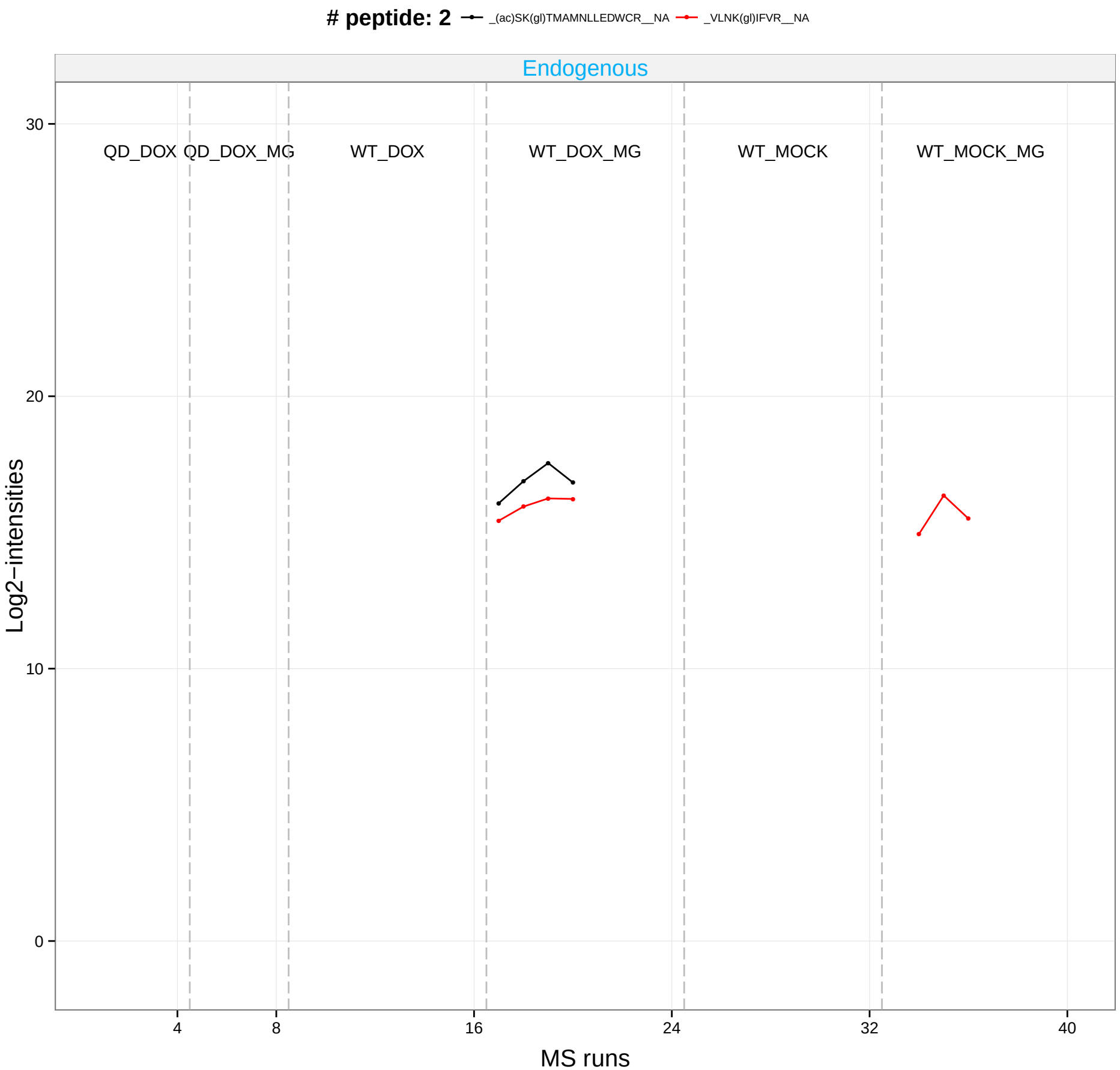
Q6ZMK1


peptide: 3
_LK(gl)EEQATCPNCR__NA
_LLAAK(gl)NINLR__NA
_SK(gl)VTAPLECSFLLLK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

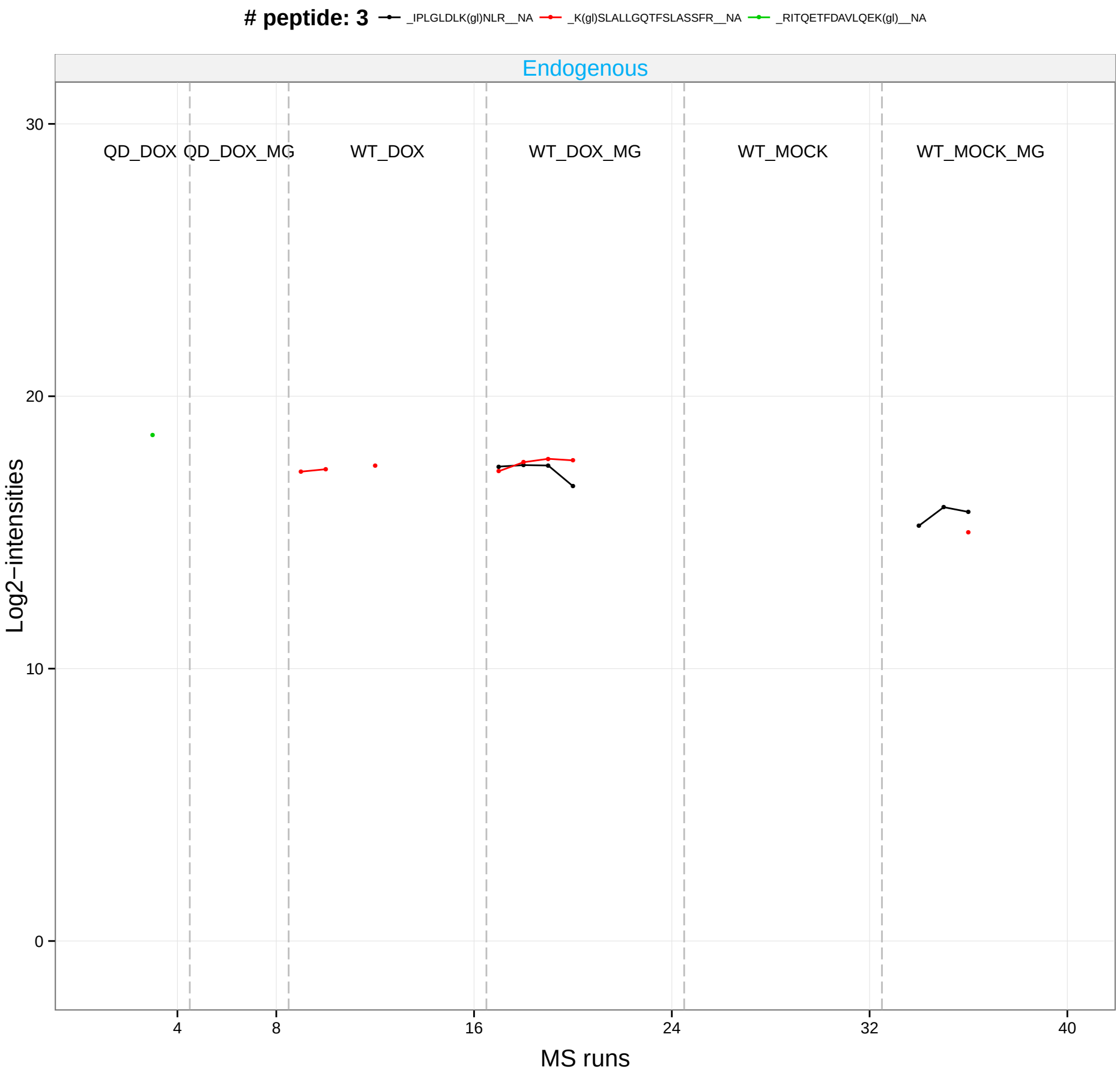
Q7KZF4


peptide: 2
_SSYYK(gl)SLLSAEEAAK__NA
_VITEYLNAQESAK(gl)SAR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

Q86V59


peptide: 2
_(ac)SK(gl)TMAMNLLEDWCR__NA
_VLNK(gl)IFVR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

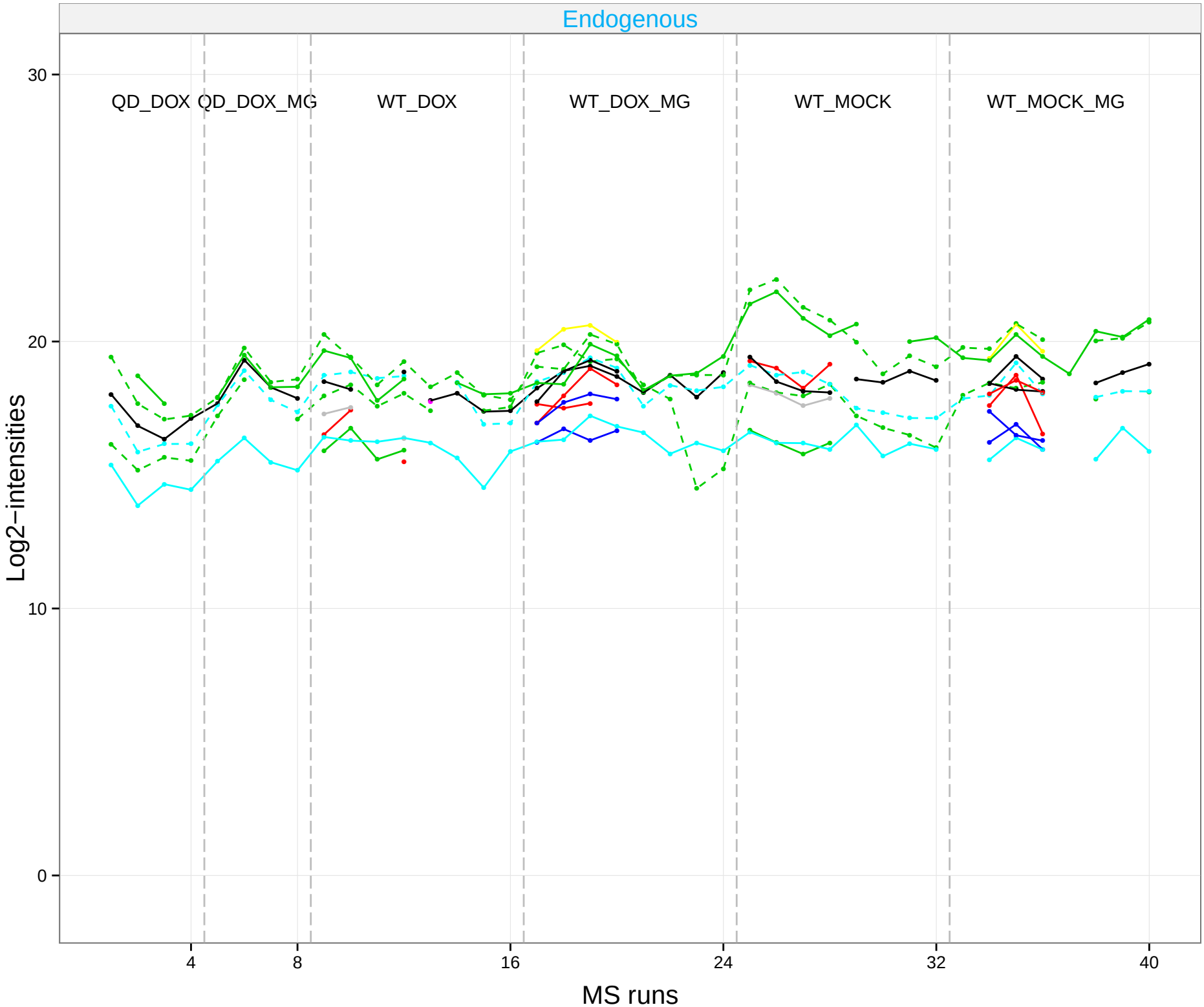
Q8IX01


peptide: 3
_IPLGLDLK(gl)NLR__NA
_K(gl)SLALLGQTFSLASSFR__NA
_RITQETFDAVLQEK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

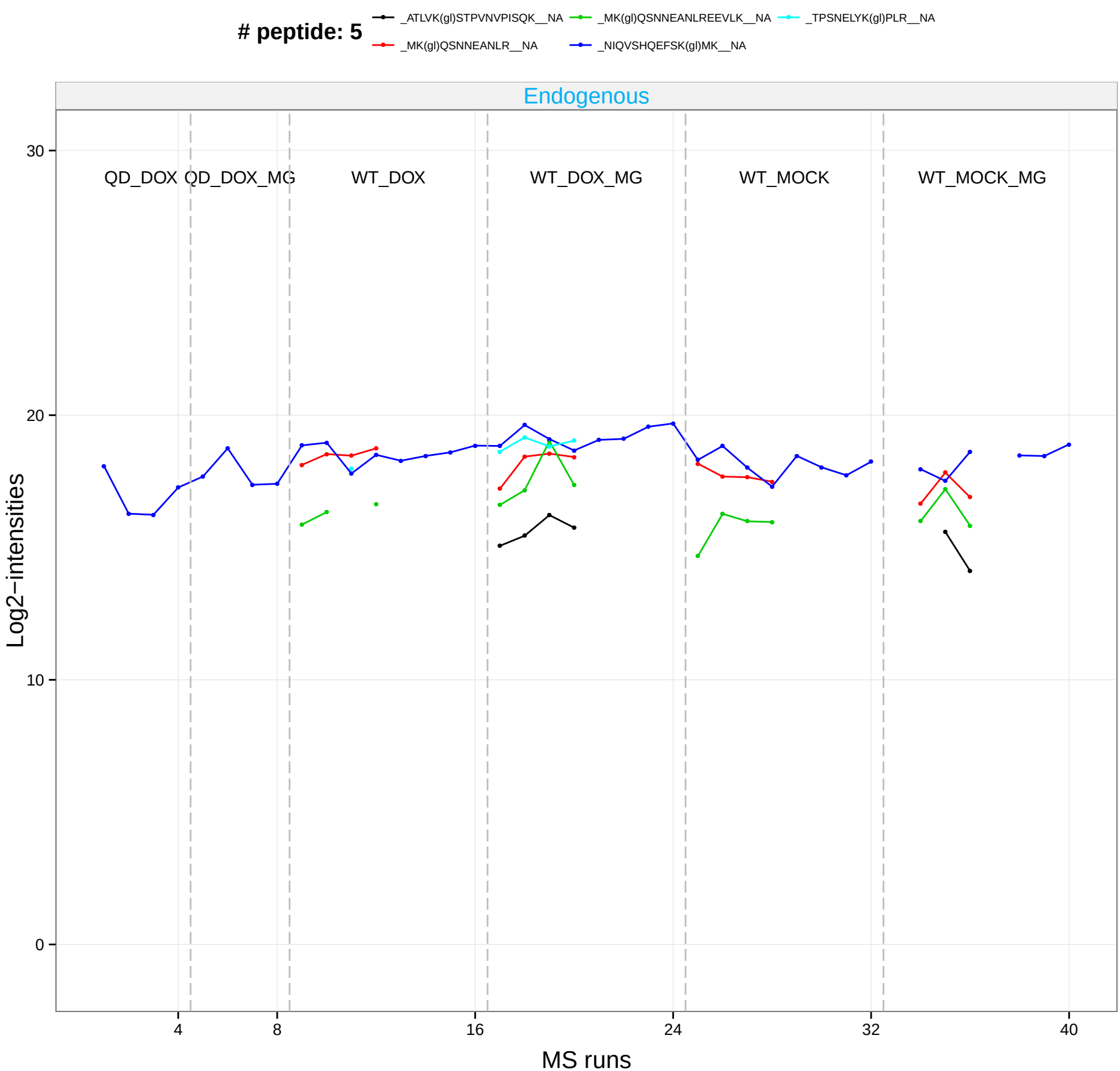
Q8N0X7

 # peptide: 12

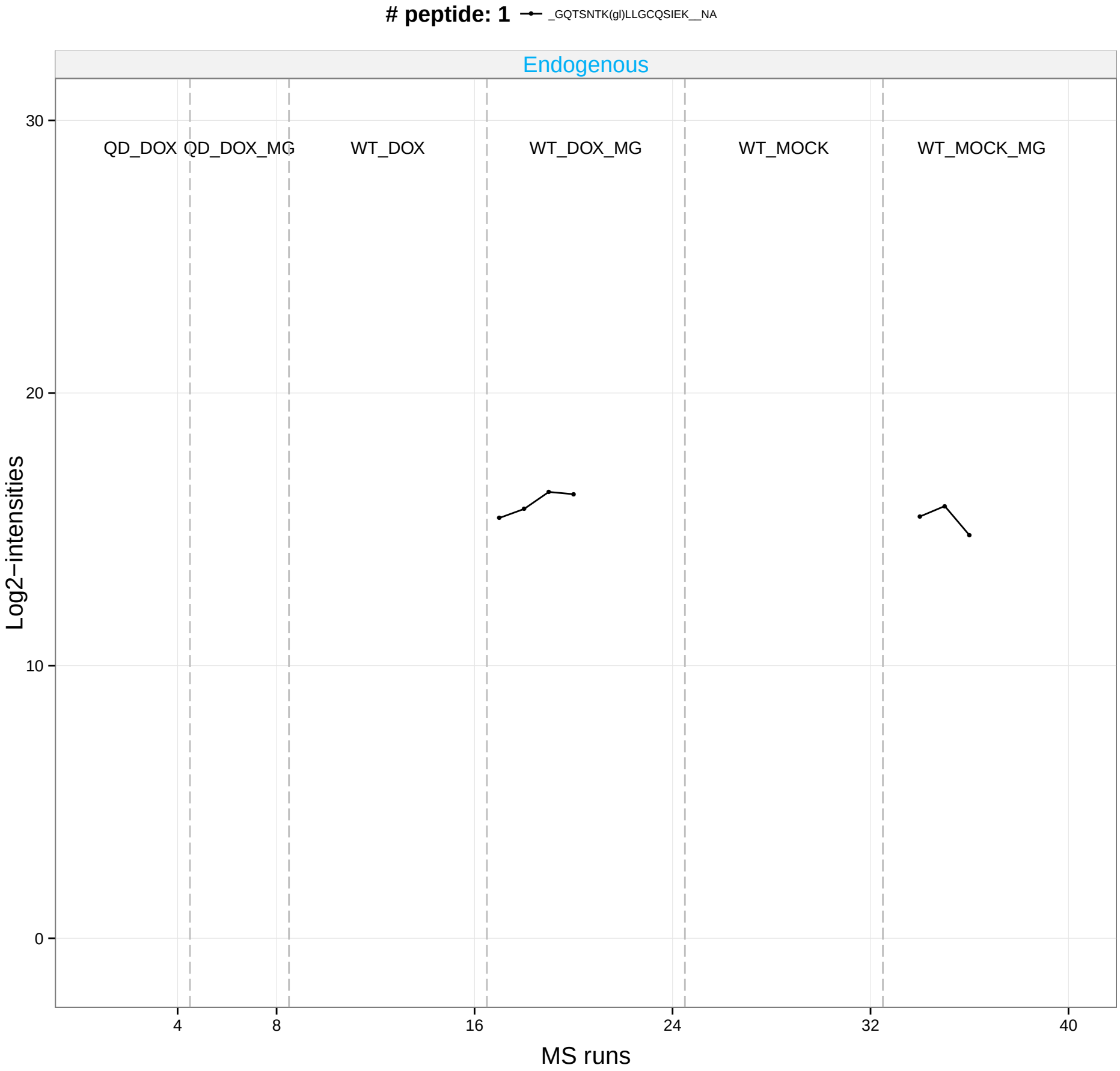
_EASGTDVK(gl)QLDQGNK__NA
_EASGTDVK(gl)QLDQGNKDVR__NA
_GAEITGK(gl)AIQK__NA
_GISISSK(gl)ESEHTGPGWESAR__NA
_GLYIAK(gl)QATGGAAK__NA
_GLYIAKQATGGAAK(gl)__NA
_IQPEEKPVEVSPAVTK(gl)GLYIAK__NA
_NYYK(gl)QGIGHLLR__NA
_QATGGAAK(gl)VSQFLVDGVCTVANCVGK__NA
_QLDQGNK(gl)DVR__NA
_TRPSSDQLK(gl)EASGTDVK__NA
_VAHNILSGASWVSWGLVK(gl)GAEITGK__NA



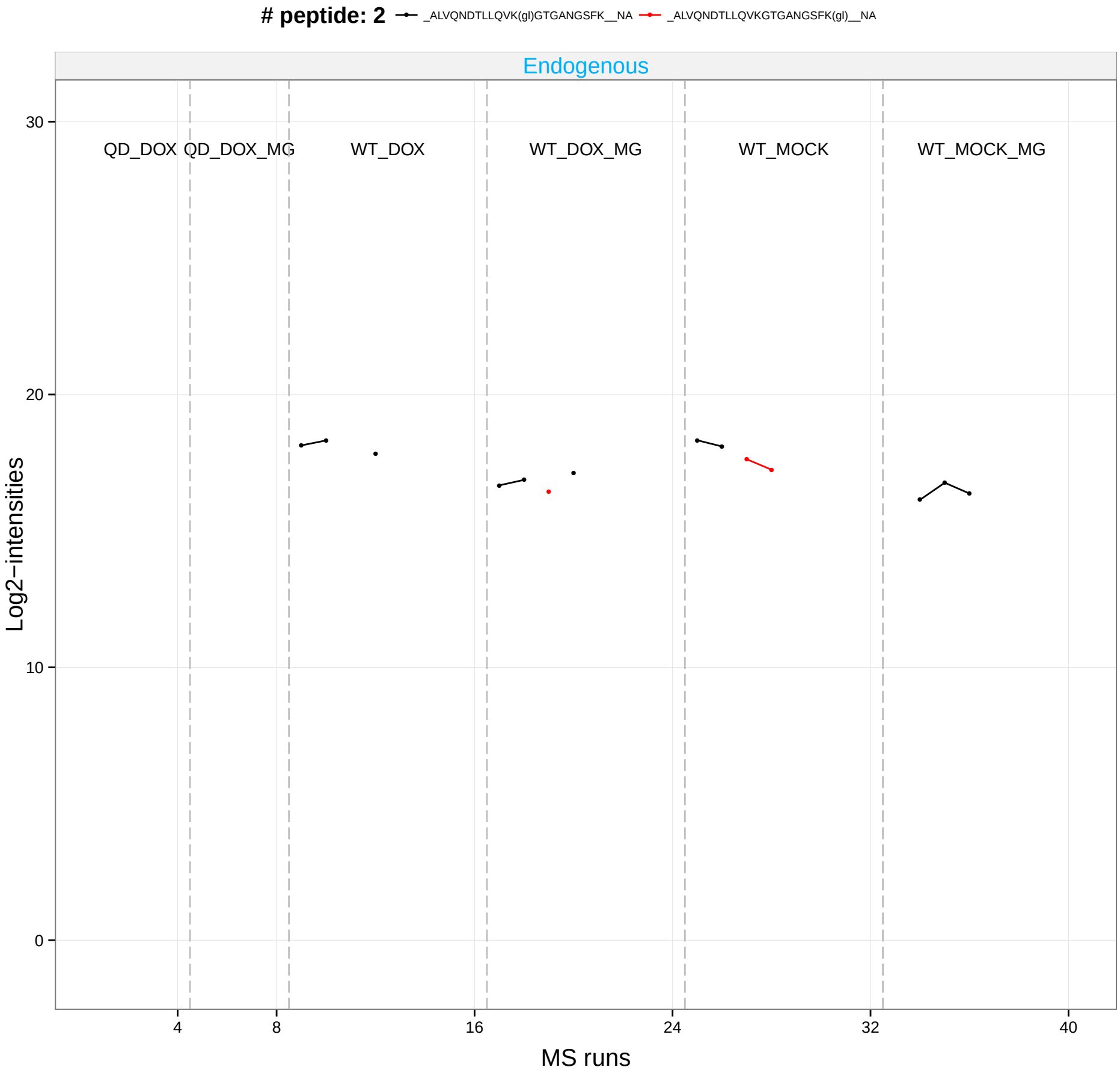
Q8WUM4


peptide: 5
_ATLVK(gl)STPVNVPISQK__NA
_MK(gl)QSNNEANLREEVLK__NA
_TPSNELYK(gl)PLR__NA
_MK(gl)QSNNEANLR__NA
_NIQVSHQEFSK(gl)MK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

Q8WYP5


peptide: 1
_GQTSNTK(gl)LLGCQSIEK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

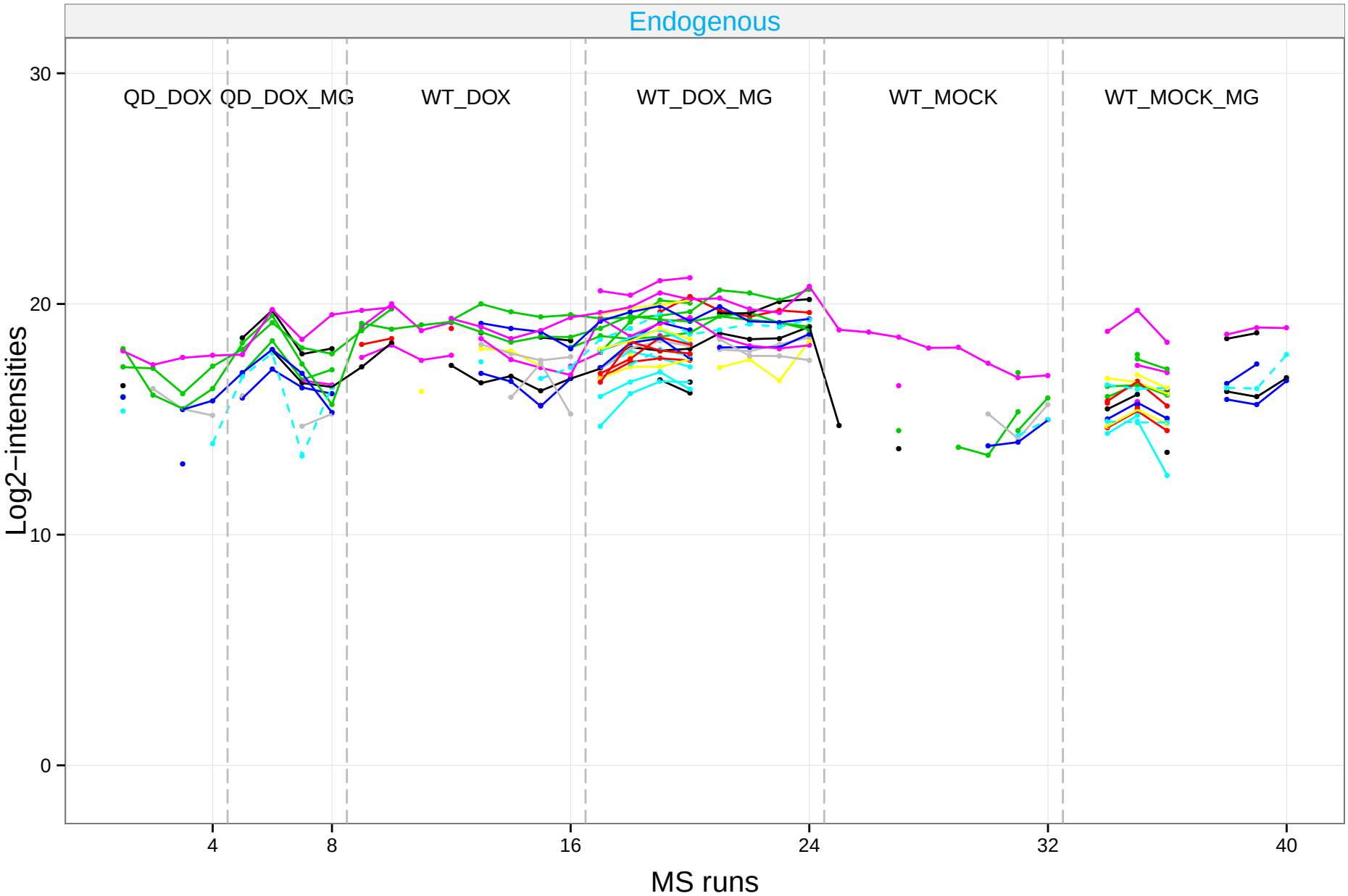
Q92522


peptide: 2
_ALVQNDTLLQVK(gl)GTGANGSFK__NA
_ALVQNDTLLQVKGTGANGSFK(gl)__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

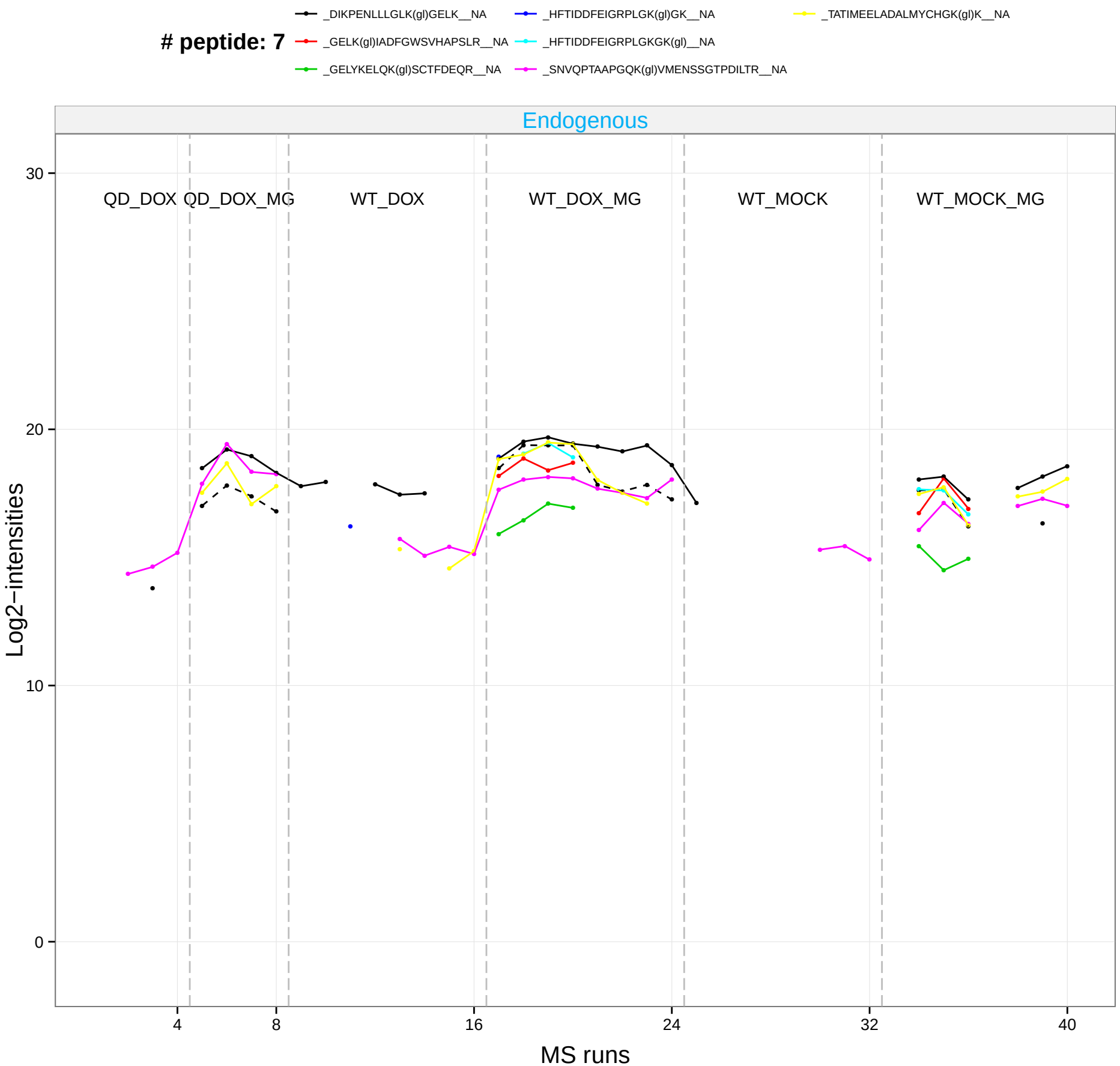
Q92616

 # peptide: 31

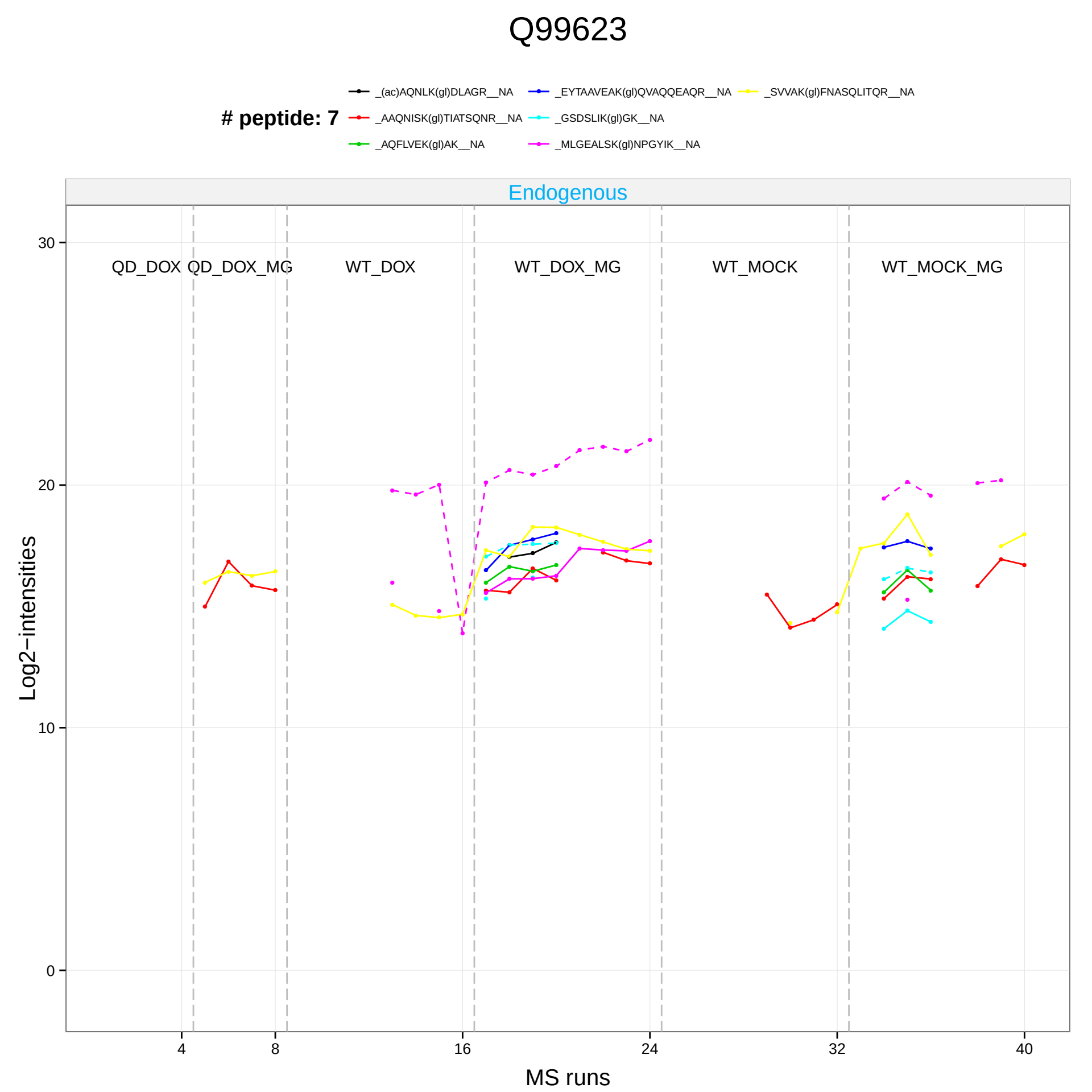
- _(ac)AADTQVSETLK(gl)R__NA
- _AYSDQAIVNLLK(gl)MR__NA
- _DLILPTIQK(gl)SLLR__NA
- _GAAYGLAGLVK(gl)GLGILSLK__NA
- _GLAGHLK(gl)SNSPR__NA
- _HLGVILPAVMLALK(gl)EK__NA
- _HLGVILPAVMLALKEK(gl)__NA
- _IK(gl)NPFLSLAACVMPSR__NA
- _K(gl)GAAYGLAGLVK__NA
- _K(gl)TQKCLQTLLDTK__NA
- _KLLSSLGGFK(gl)LAHGLLEELK__NA
- _KTQK(gl)CLQTLLDTK__NA
- _LAHGLLEELK(gl)TVLSSHK__NA
- _LFNDSSPVVLEESWDALNAITK(gl)K__NA
- _LK(gl)AADALGK__NA
- _LK(gl)ALGTLVSHVTLR__NA
- _LMEIYQEK(gl)LYRPPPVLDALGR__NA
- _LSSFWQLIVDEK(gl)K__NA
- _LSSLFVK(gl)CLQNPSSDIR__NA
- _MK(gl)IDPEAFITR__NA
- _NLLHSLQSSGIGSK(gl)AGVPSK__NA
- _QGVCIGLSEIMK(gl)STSR__NA
- _QLDDEEVSEFALDGLK(gl)QVMAIK__NA
- _QSVVVLMGSLAK(gl)HLDK__NA
- _QVMAIK(gl)SR__NA
- _SK(gl)ADYTSHLR__NA
- _SWCQEELSVAVK(gl)R__NA
- _TK(gl)AGSVELLGAMAYCAPK__NA
- _TVSAK(gl)ALGAMVK__NA
- _VGK(gl)GEPGAAPLSAPAFSLVFPFLK__NA
- _VQK(gl)AGQQALR__NA



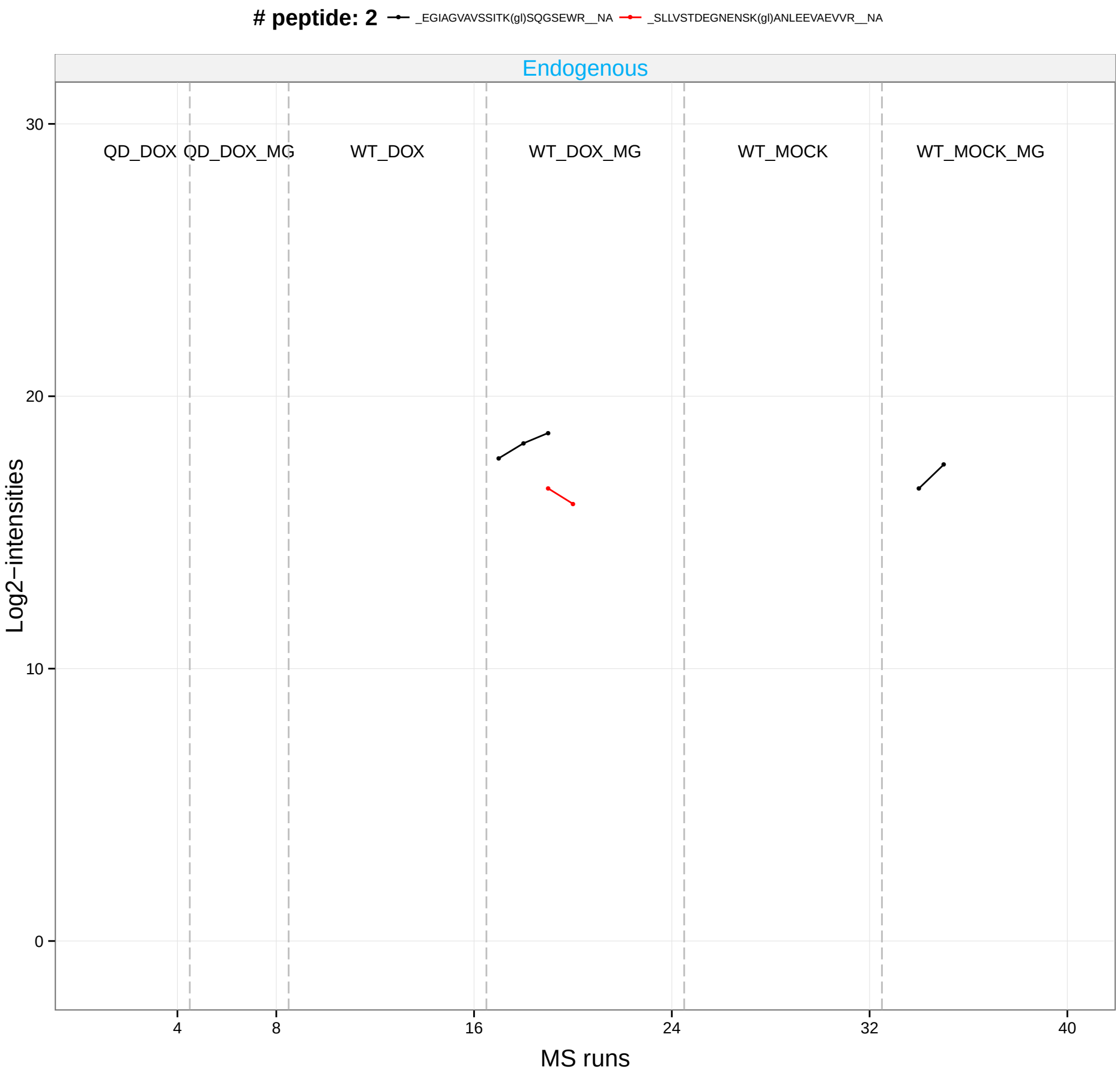
Q96GD4


peptide: 7
_DIKPENLLLGLK(gl)GELK__NA
_GELK(gl)IADFGWSVHAPSLR__NA
_GELYKELQK(gl)SCTFDEQR__NA
_HFTIDDFEIGRPLGK(gl)GK__NA
_HFTIDDFEIGRPLGKGK(gl)__NA
_SNVQPTAAPGQK(gl)VMENSSGTPDILTR__NA
_TATIMEELADALMYCHGK(gl)K__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

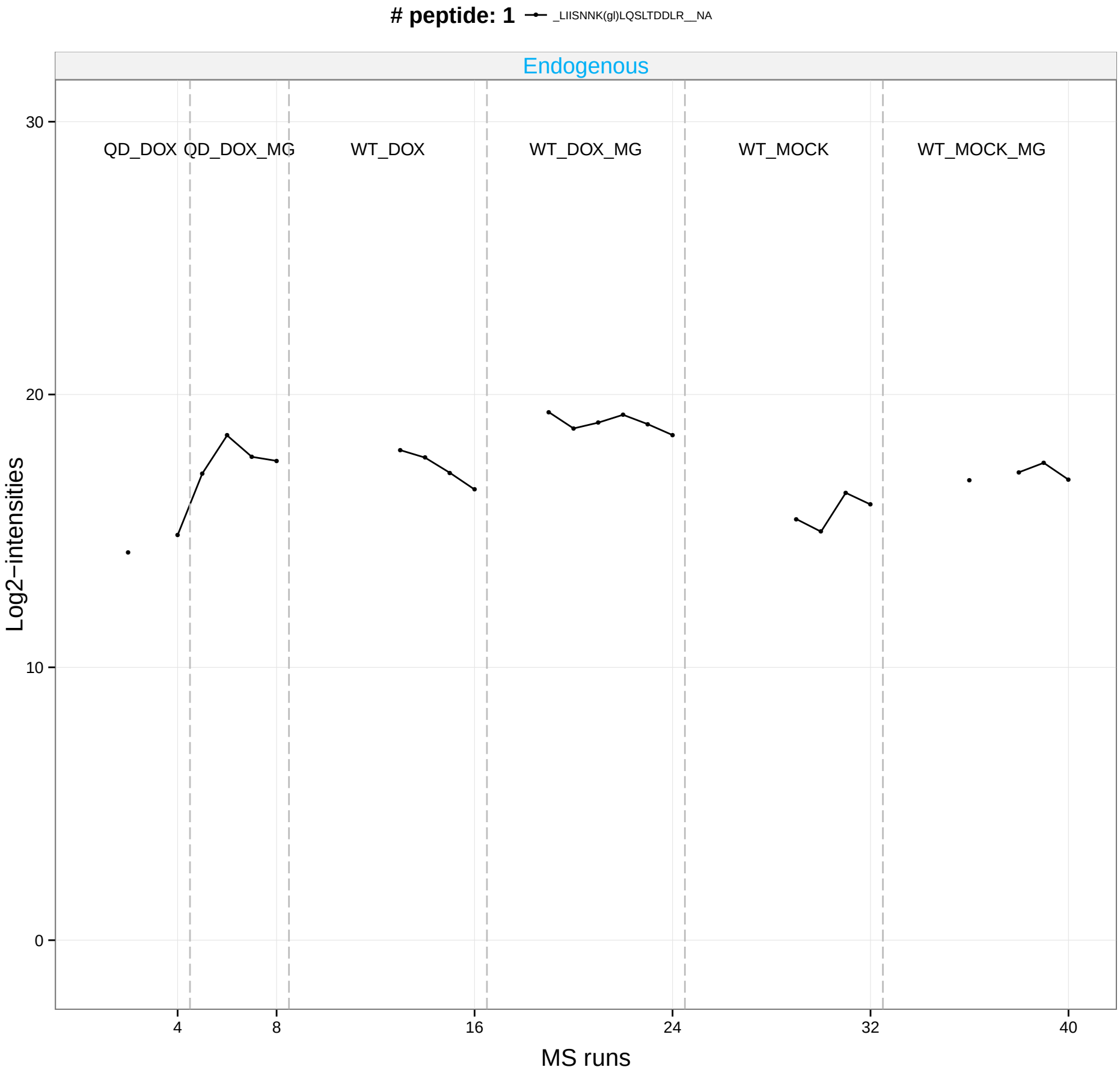
Q99623


peptide: 7
_(ac)AQNLK(gl)DLAGR__NA
_EYTAAVEAK(gl)QVAQQEAQR__NA
_SVVAK(gl)FNASQLITQR__NA
_AAQNISK(gl)TIATSQNR__NA
_GSDSLIK(gl)GK__NA
_AQFLVEK(gl)AK__NA
_MLGEALSK(gl)NPGYIK__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

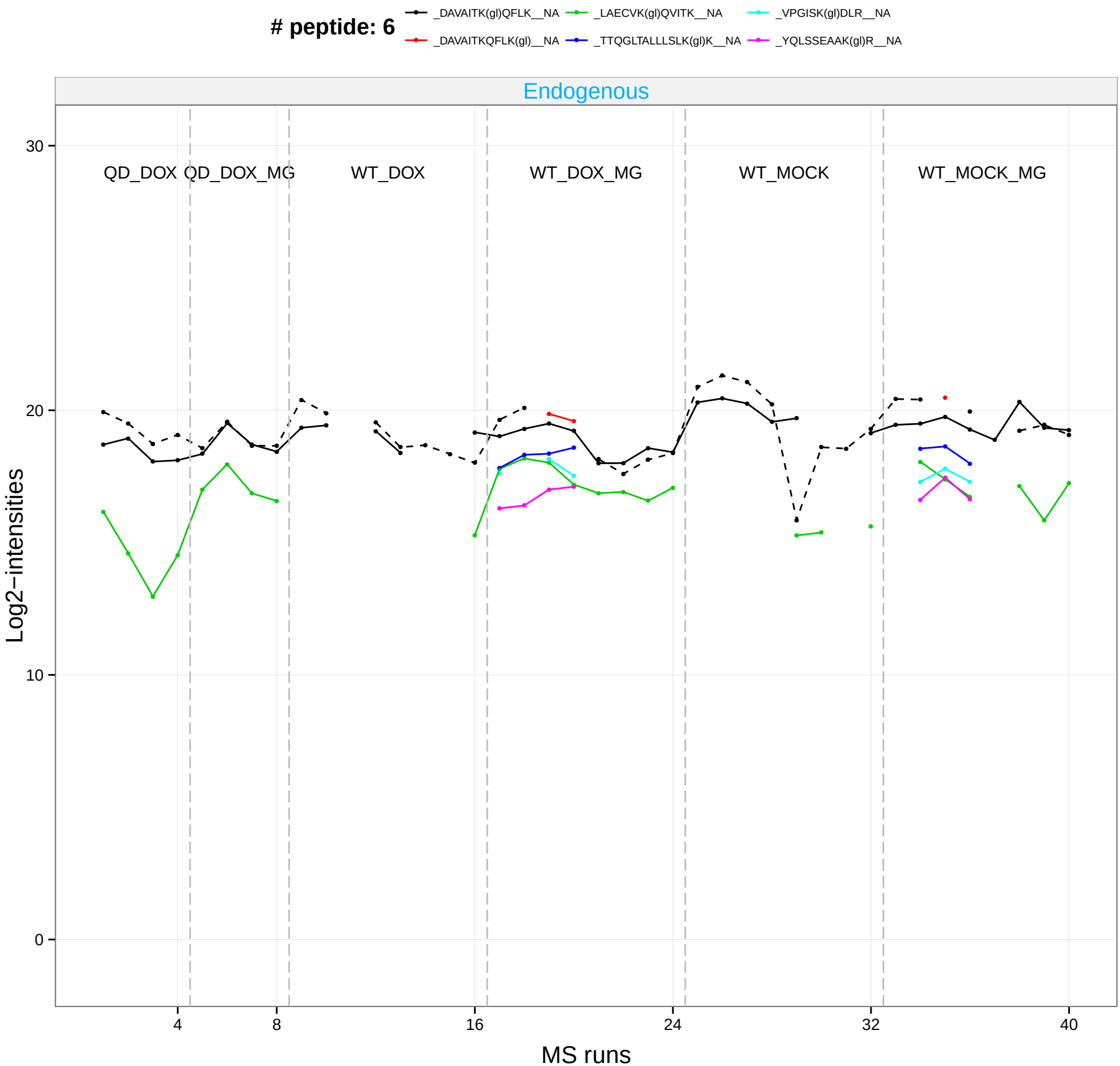
Q9BYK8


peptide: 2
_EGIAGVAVSSITK(gl)SQGSEWR__NA
_SLLVSTDEGNENSK(gl)ANLEEVAEVVR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

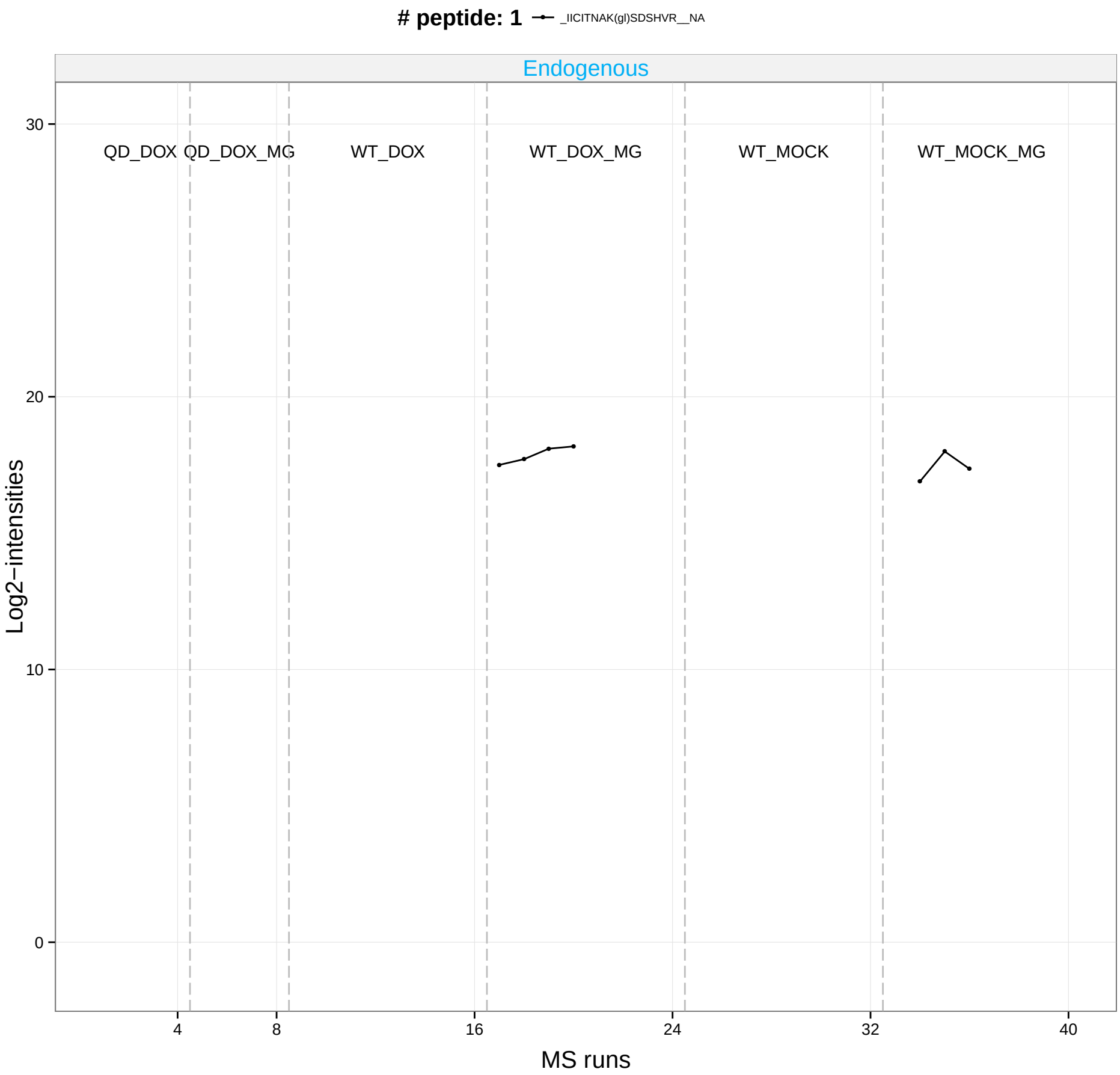
Q9H9A6


peptide: 1
_LIISNNK(gl)LQSLTDDLR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

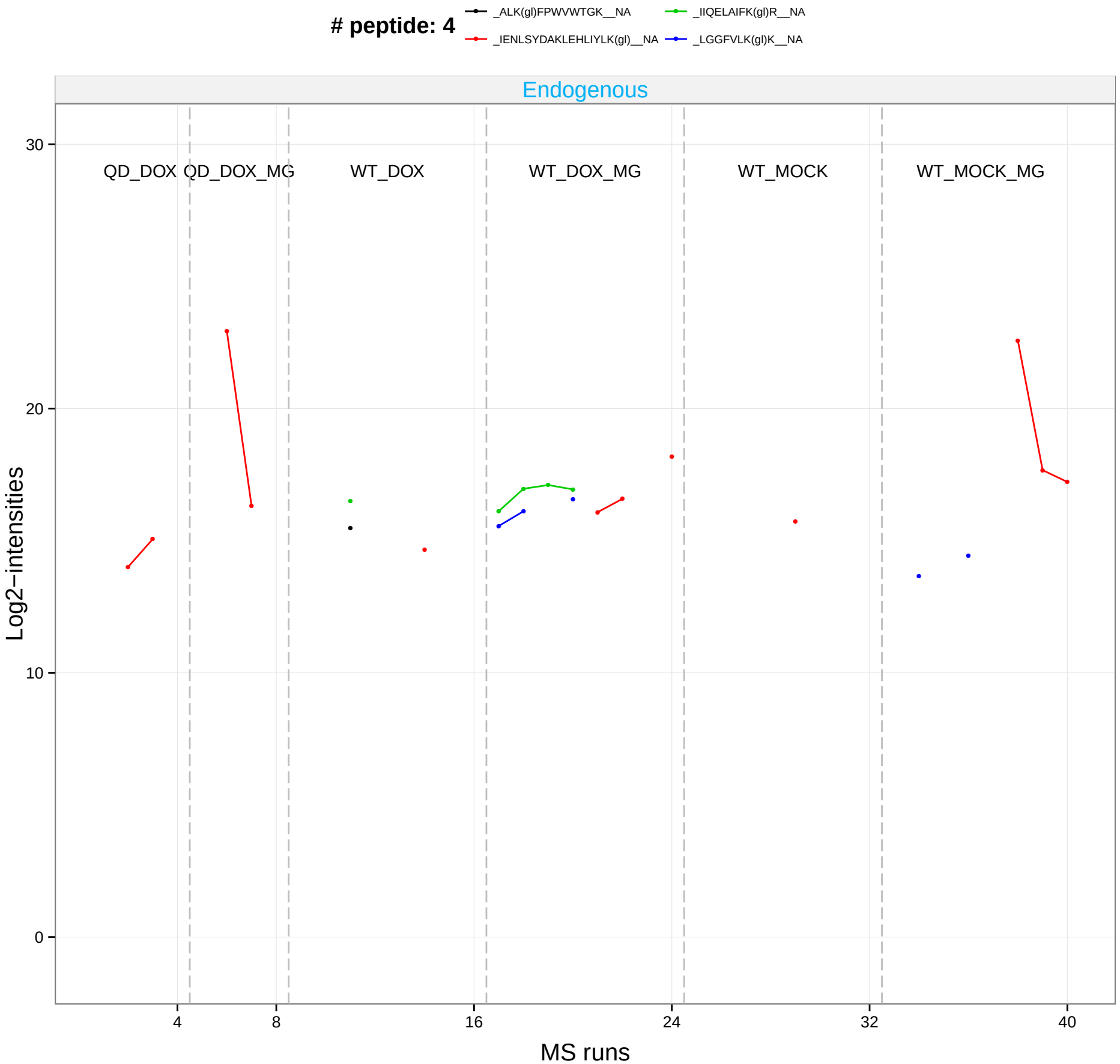
Q9NRW7


peptide: 6
_DAVAITK(gl)QFLK__NA
_LAECVK(gl)QVITK__NA
_VPGISK(gl)DLR__NA
_DAVAITKQFLK(gl)__NA
_TTQGLTALLLSLK(gl)K__NA
_YQLSSEAAK(gl)R__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

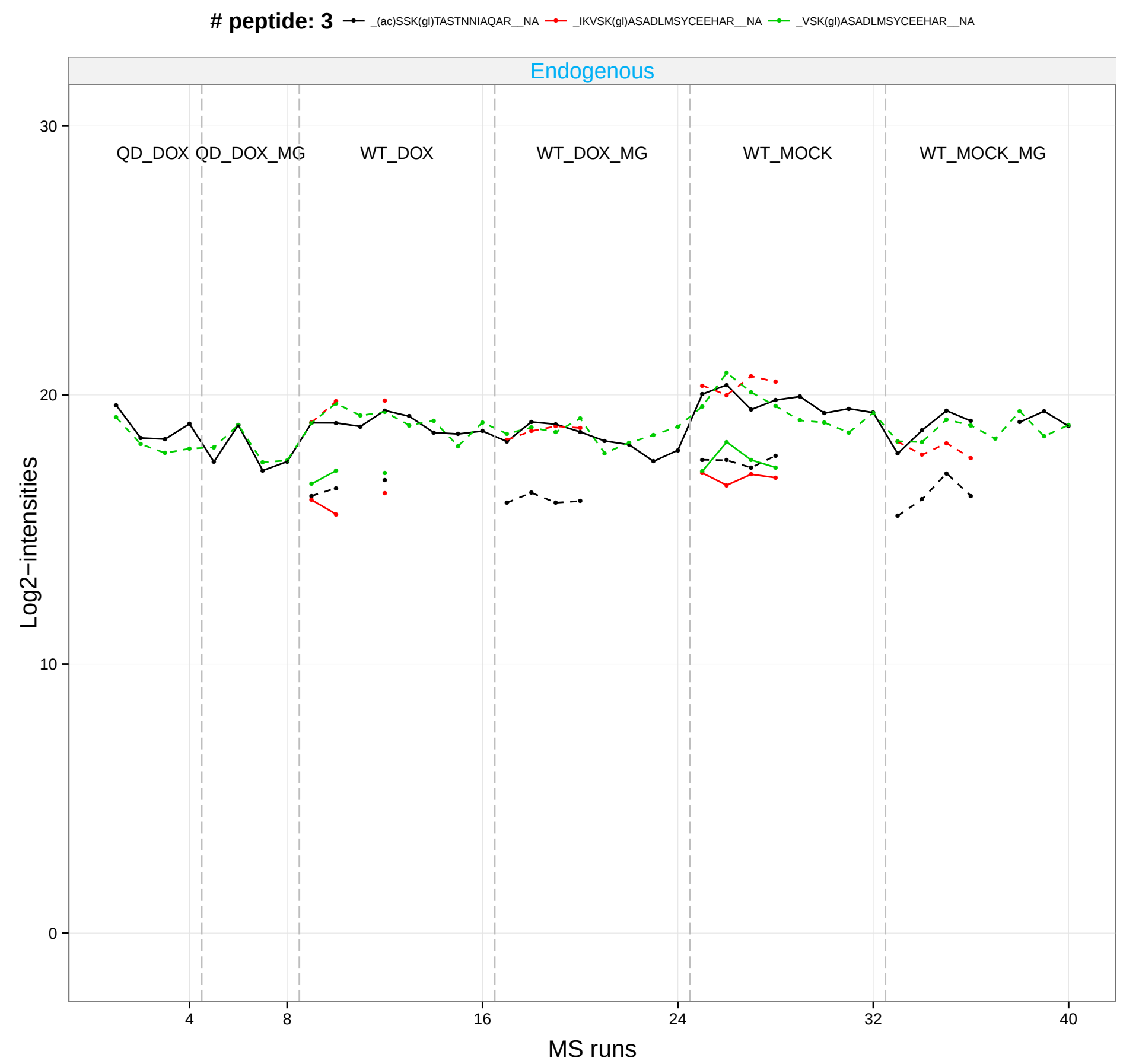
Q9NVM9


peptide: 1
_IICITNAK(gl)SDSHVR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs

Q9NZJ4


peptide: 4
_ALK(gl)FPWVWTGK__NA
_IIQELAIFK(gl)R__NA
_IENLSYDAKLEHLIYLK(gl)__NA
_LGGFVLK(gl)K__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
Log2−intensities
30
20
10
0
4
8
16
24
32
40
MS runs

Q9UBI6


peptide: 3
_(ac)SSK(gl)TASTNNIAQAR__NA
_IKVSK(gl)ASADLMSYCEEHAR__NA
_VSK(gl)ASADLMSYCEEHAR__NA
Endogenous
QD_DOX
QD_DOX_MG
WT_DOX
WT_DOX_MG
WT_MOCK
WT_MOCK_MG
30
20
10
0
Log2−intensities
4
8
16
24
32
40
MS runs