

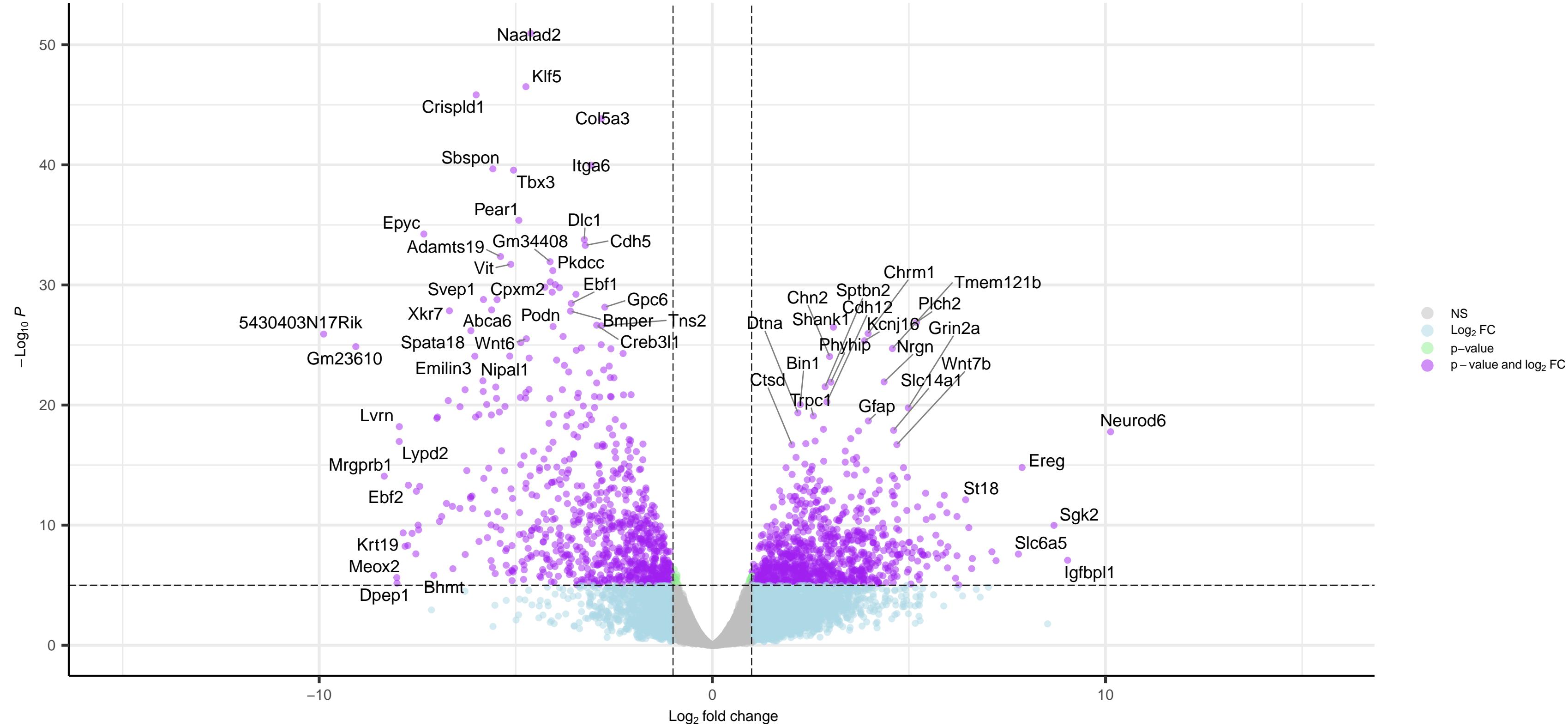
# Nerve sheath tumor, low grade vs. Spindle and epithelioid tumor, high grade

Of genotype: 4ko



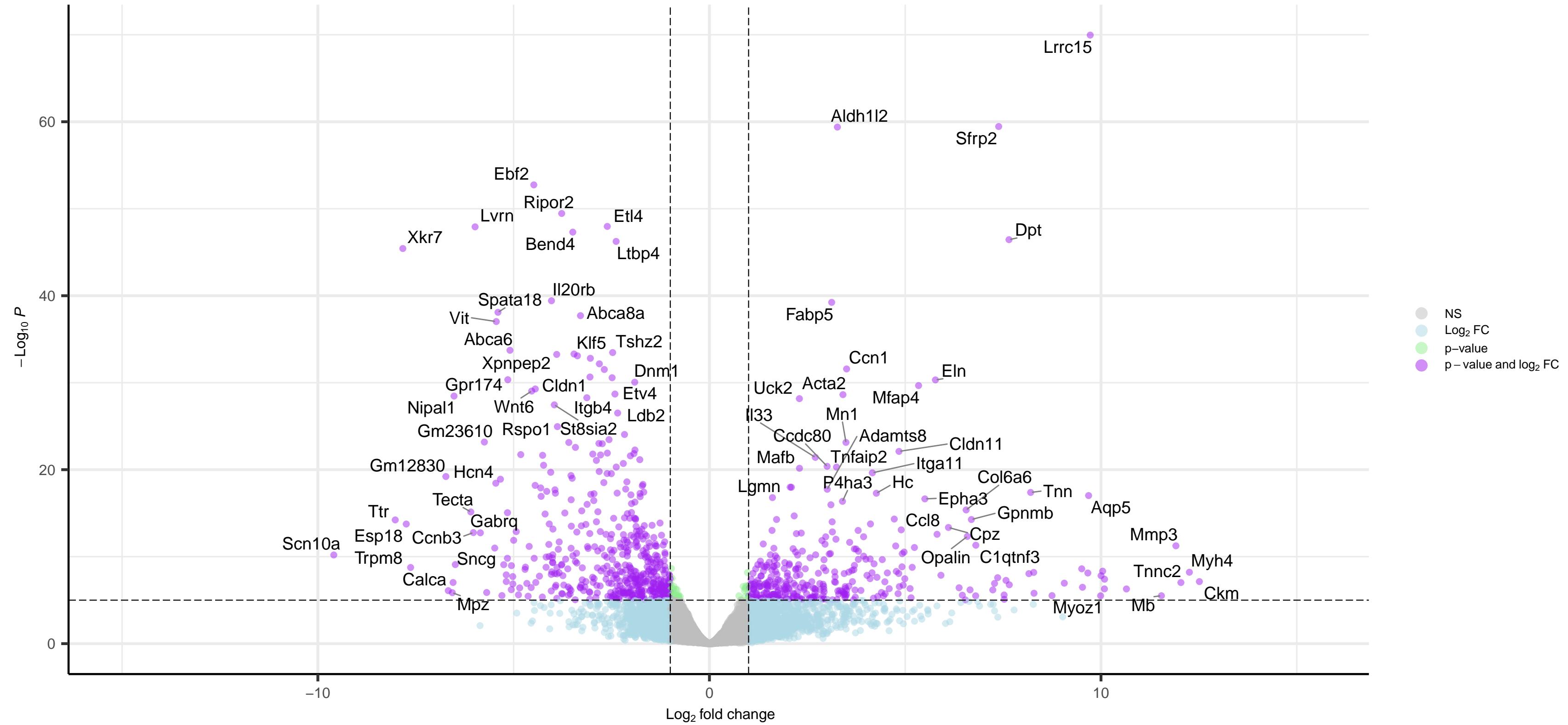
# Nerve sheath tumors vs. Gliomas

Of genotype: 4ko



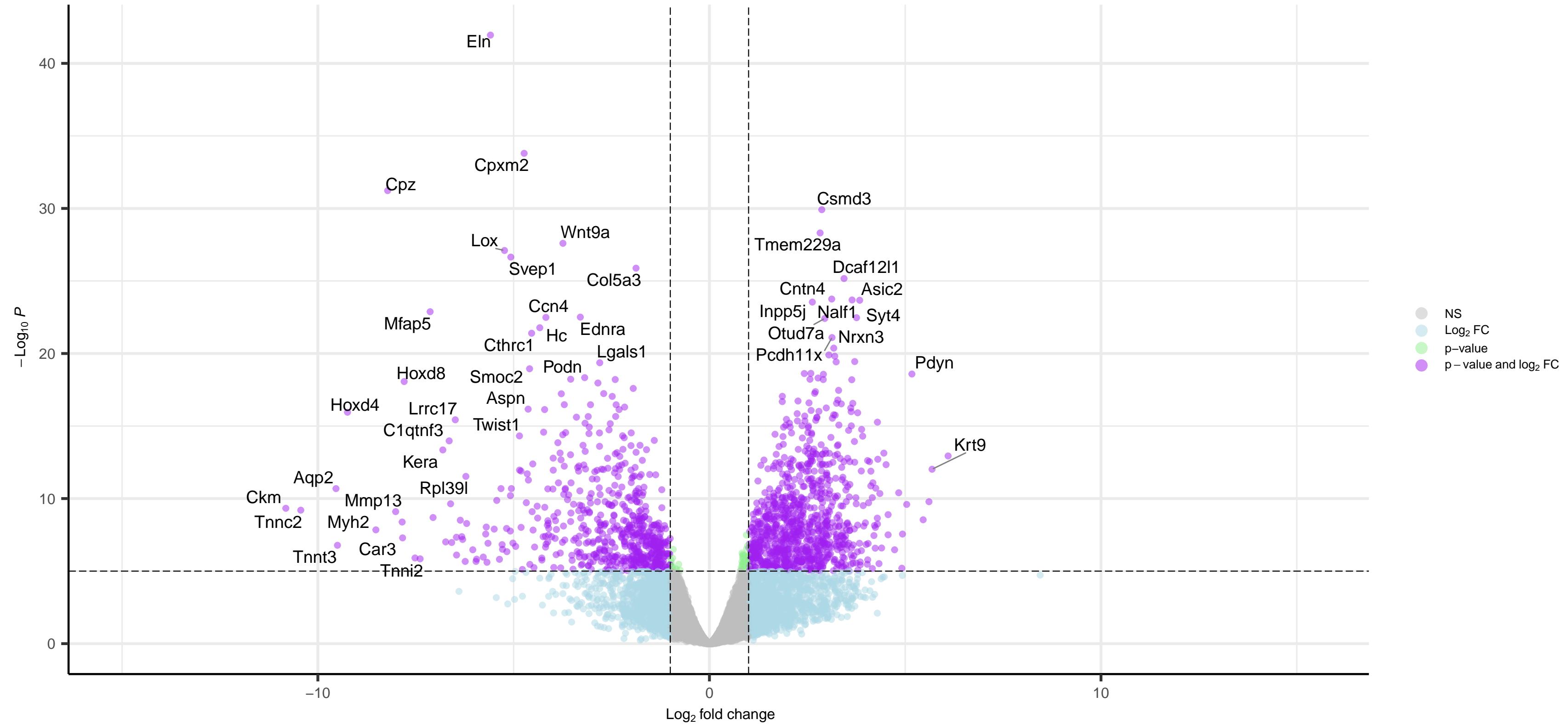
# Nerve sheath tumors vs. Spindle and epithelioid tumors

Of genotype: 4ko



## **Spindle and epithelioid tumors vs. Glioma**

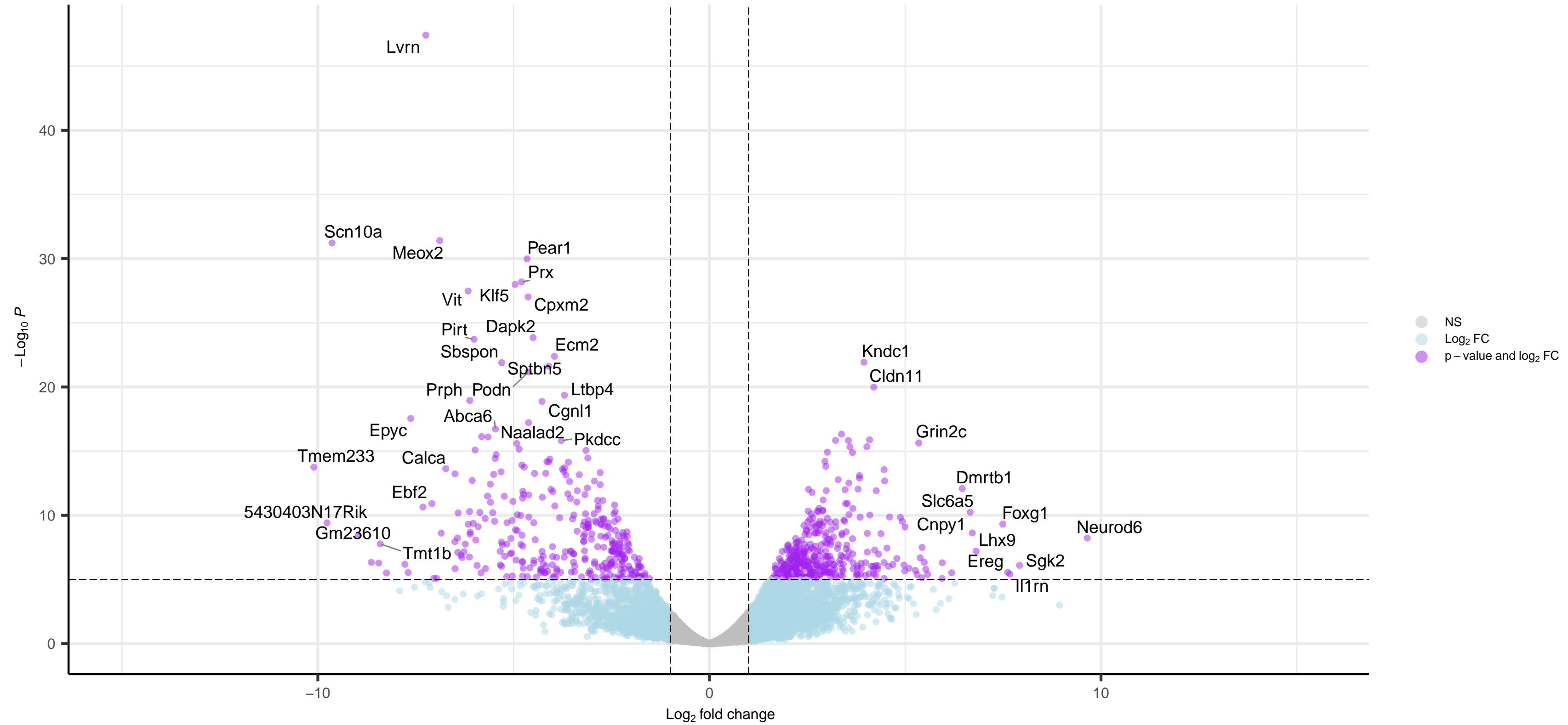
Of genotype: 4ko



total = 15742 variables

# Nerve sheath tumor, low grade vs. Glioma, grade 4

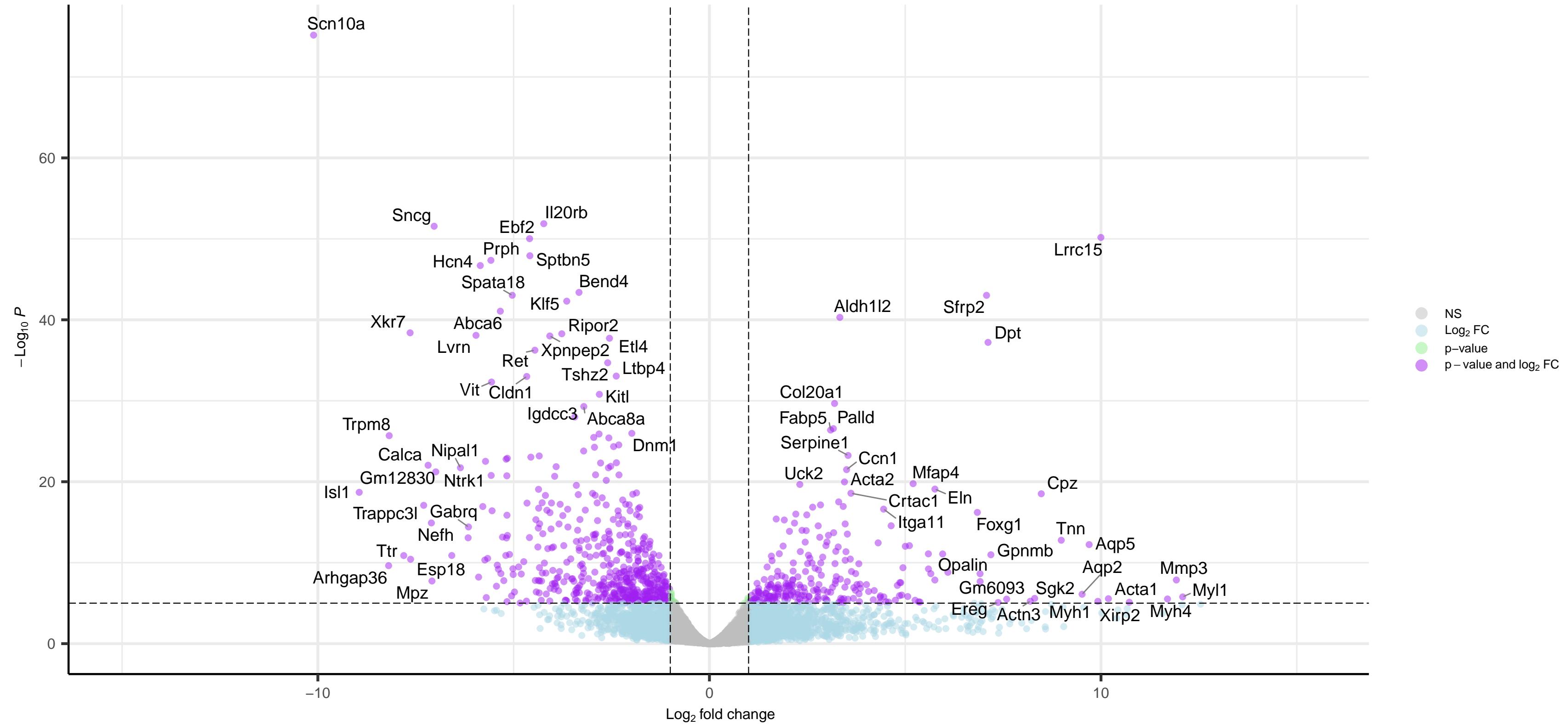
Of genotype: 4ko



total = 15415 variables

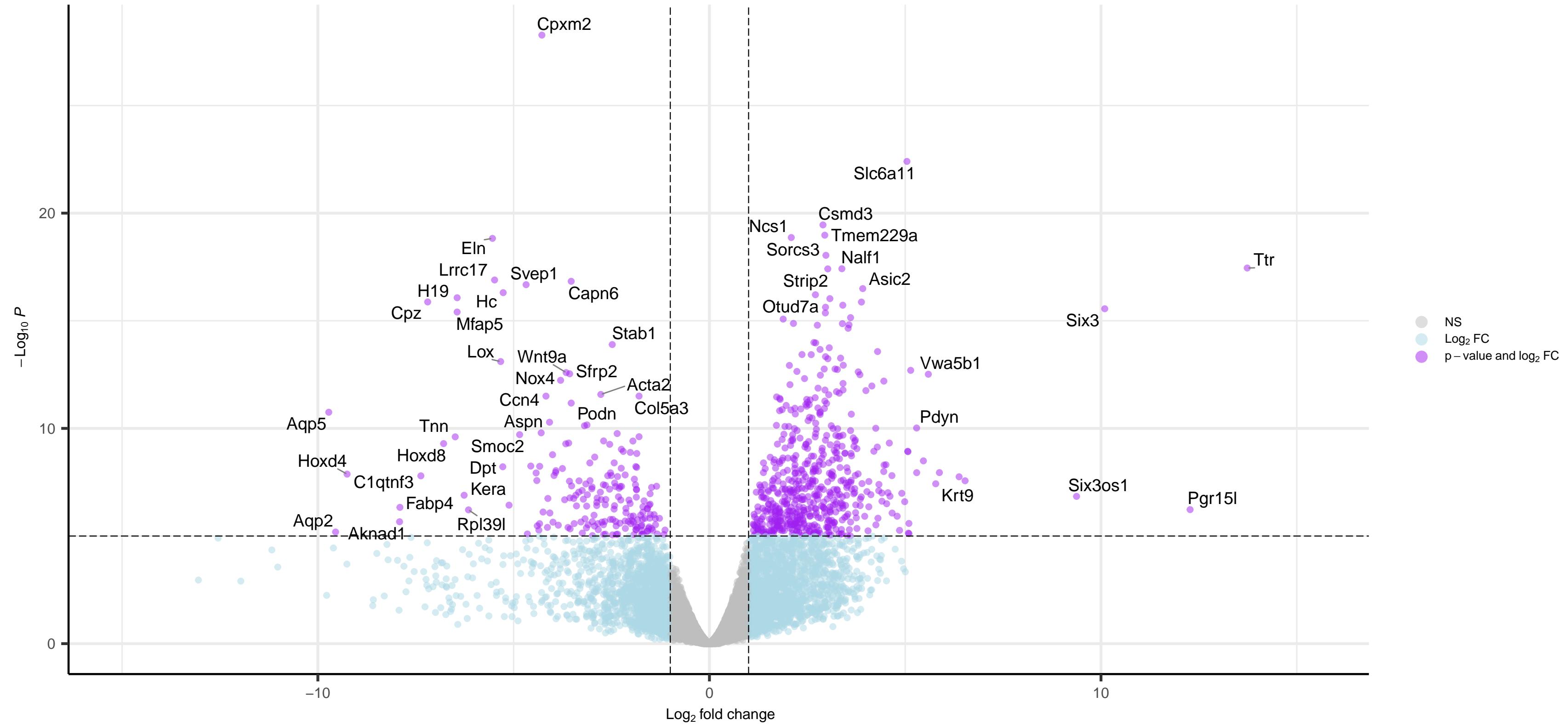
# Nerve sheath tumor, low grade vs. Spindle and epithelioid tumor, high grade

Of genotype: 4ko



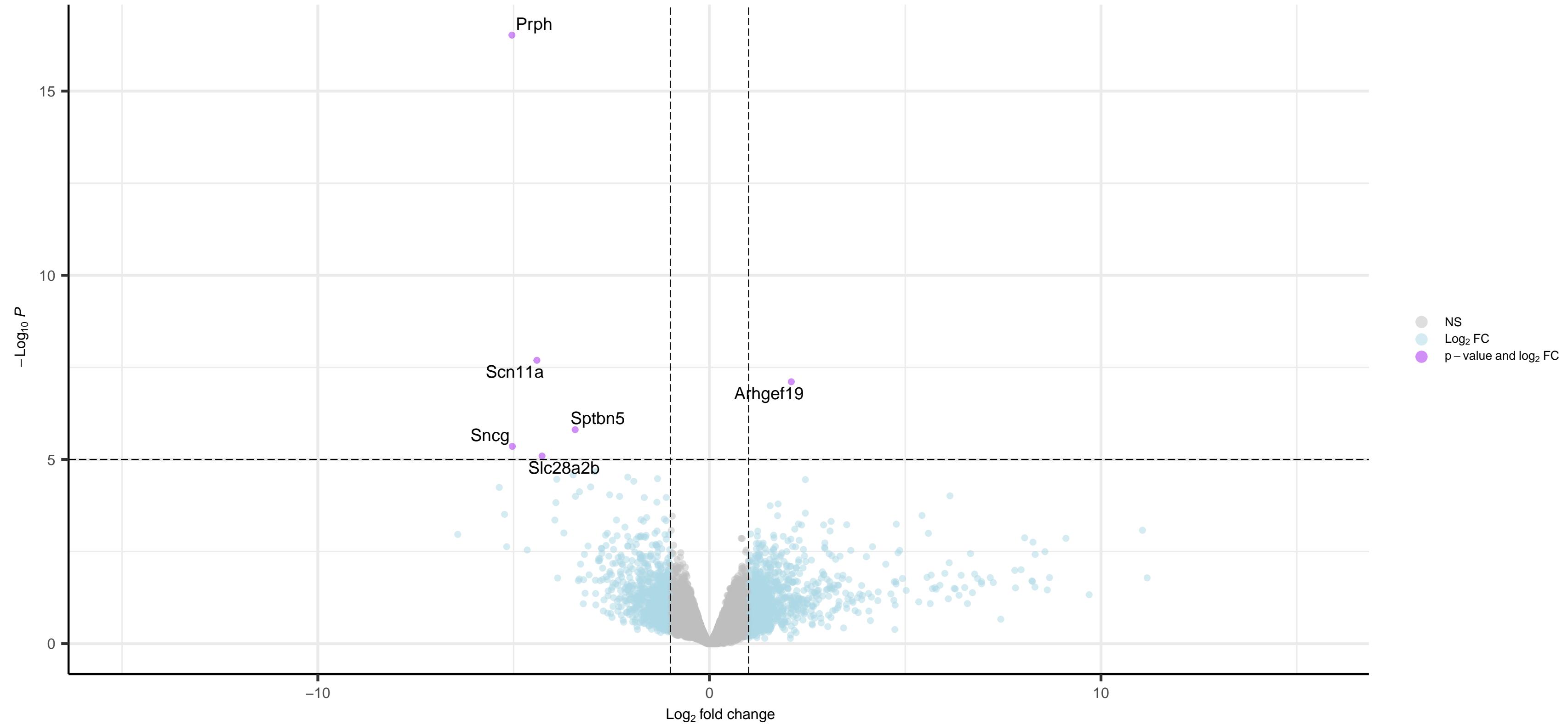
# Spindle and epithelioid tumor, high grade vs. Glioma, grade 4

Of genotype: 4ko



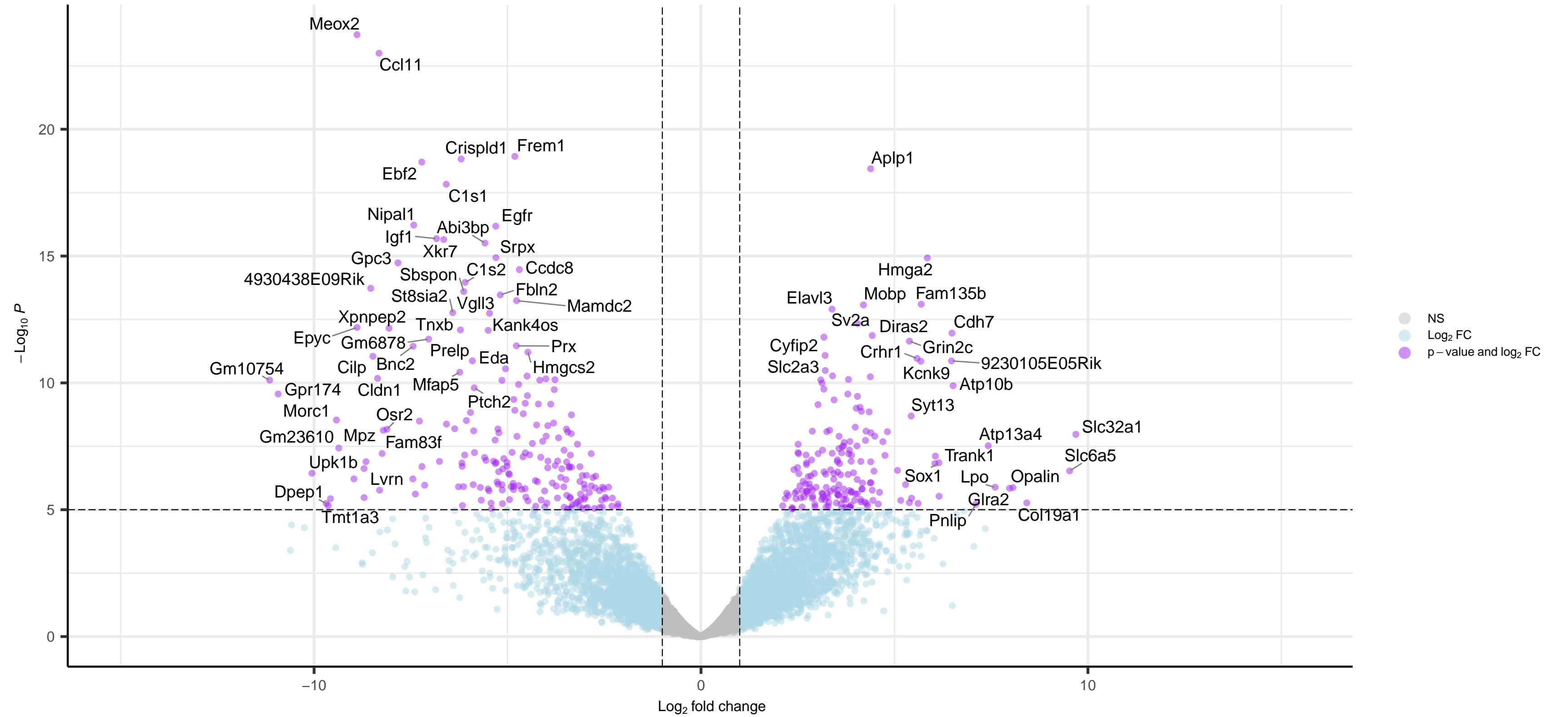
## 2 vs. 4

Of genotype: 4ko



# Nerve sheath tumor, low grade vs. Glioblastoma (GBM)

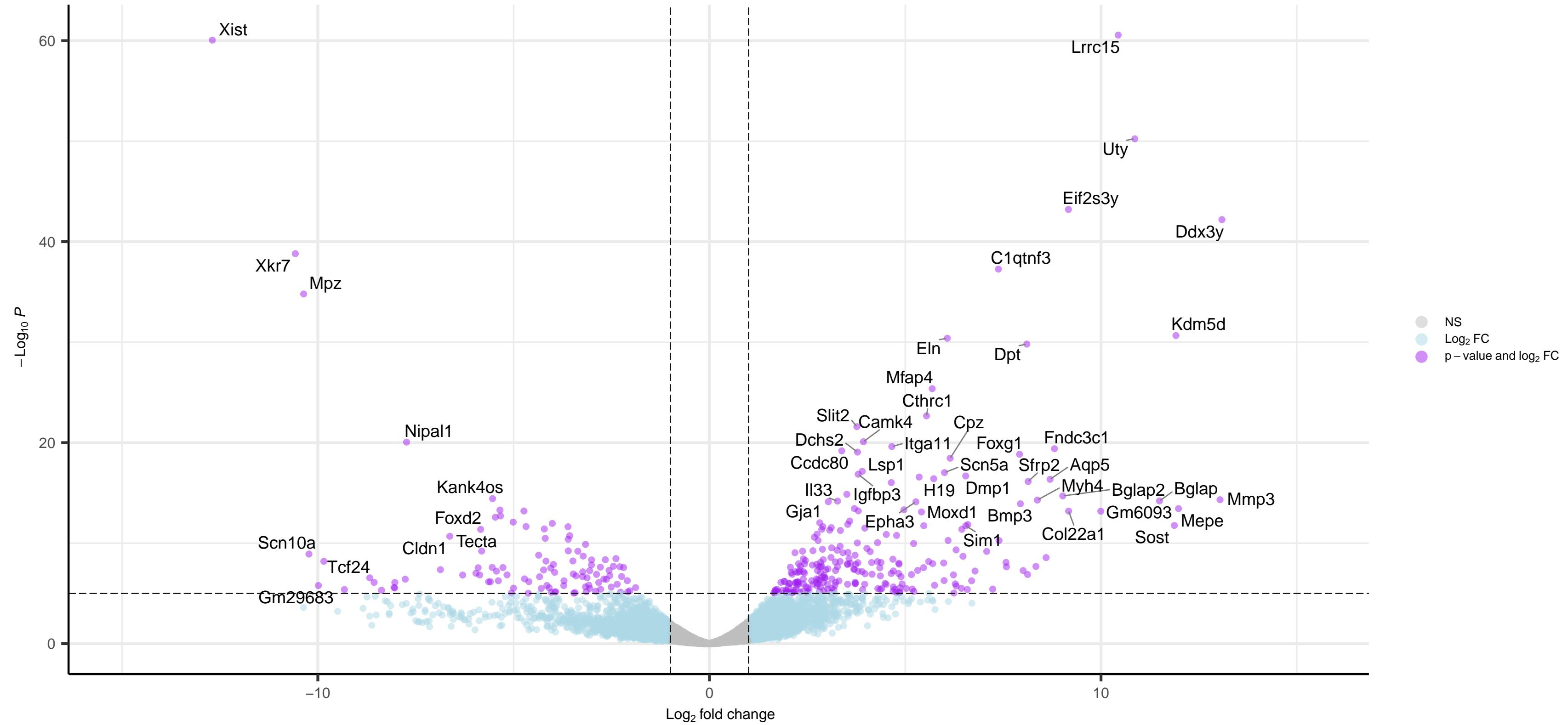
Of genotype: atrxwt



total = 15709 variables

# Nerve sheath tumor, low grade vs. Spindle and epithelioid tumor, high grade

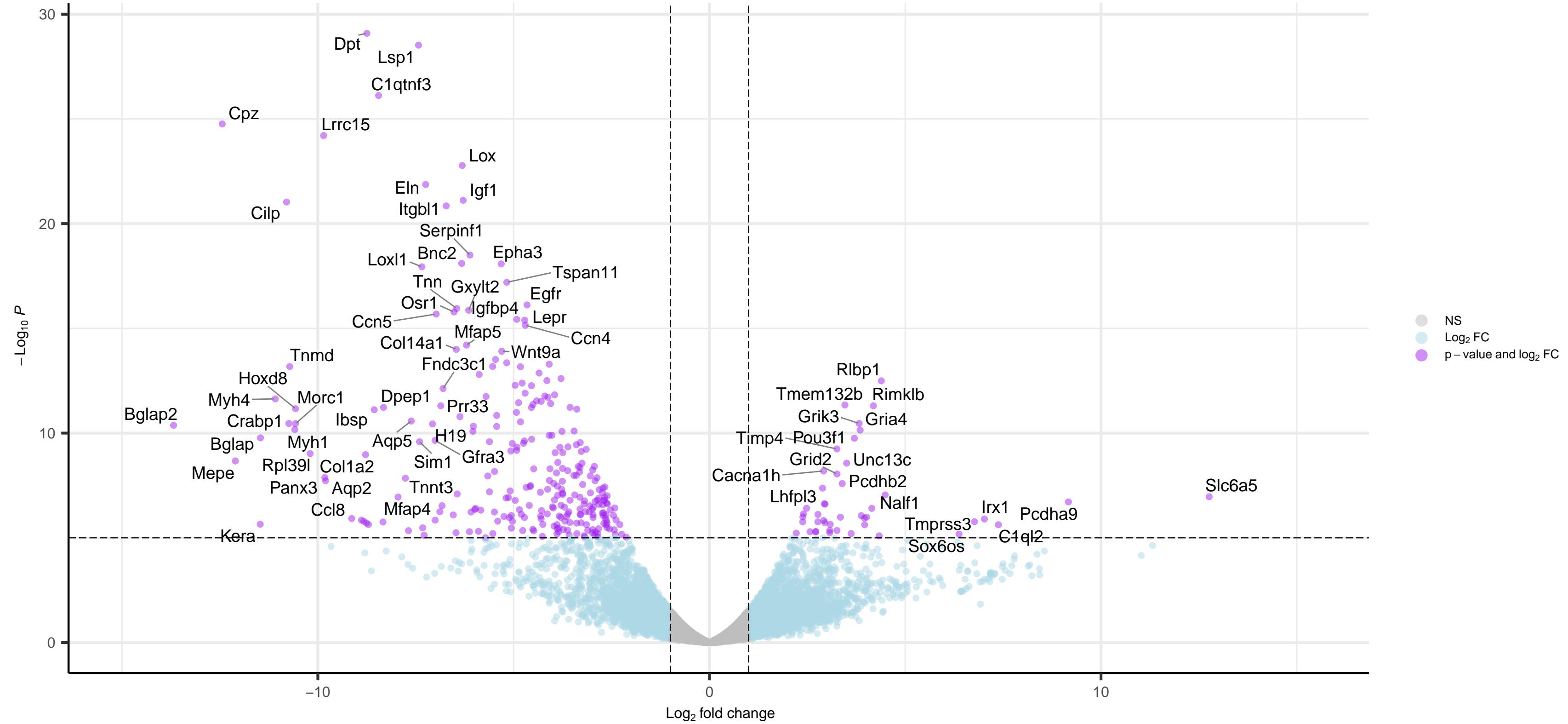
Of genotype: atrxwt



total = 15593 variables

# Spindle and epithelioid tumor, high grade vs. Glioblastoma (GBM)

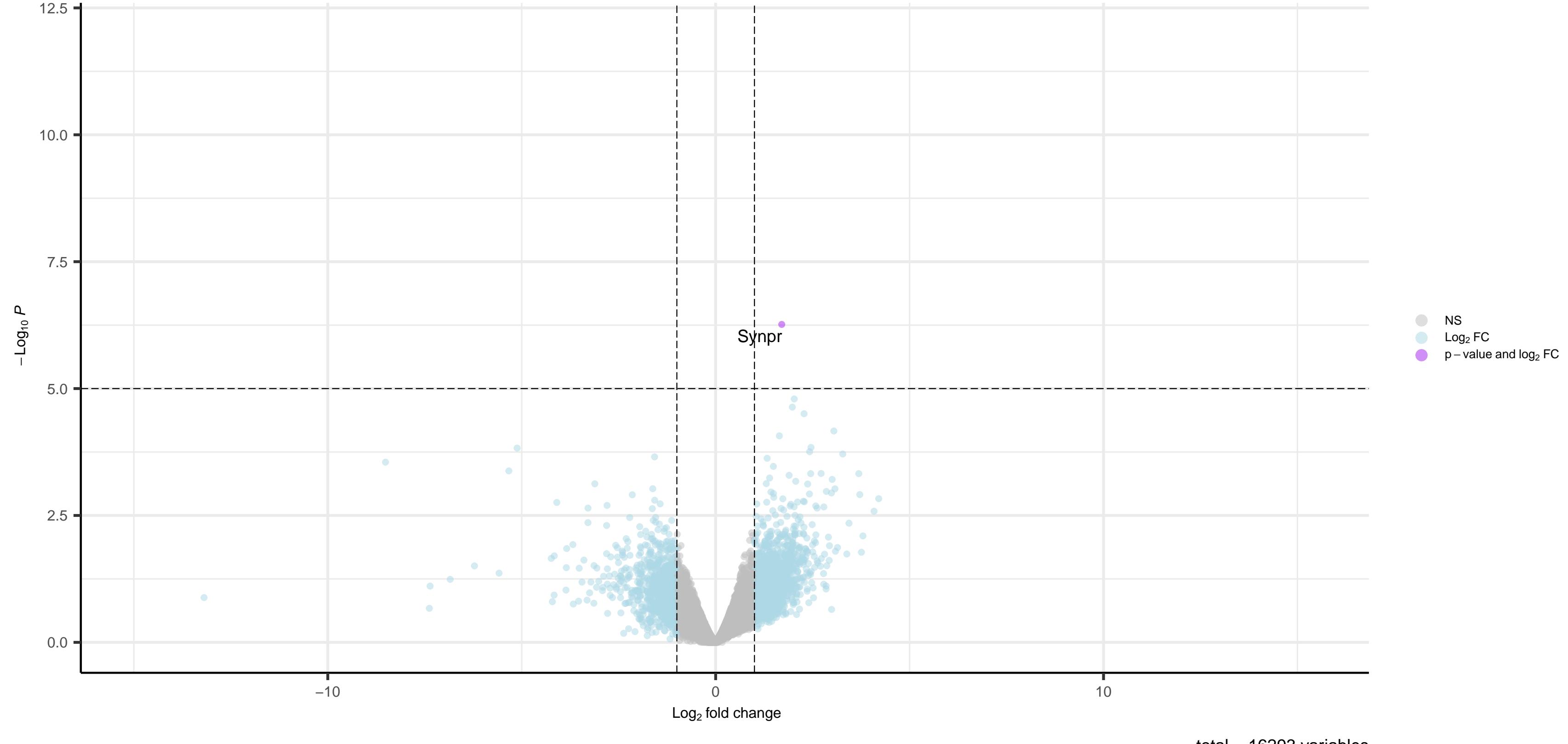
Of genotype: atrxwt



total = 15647 variables

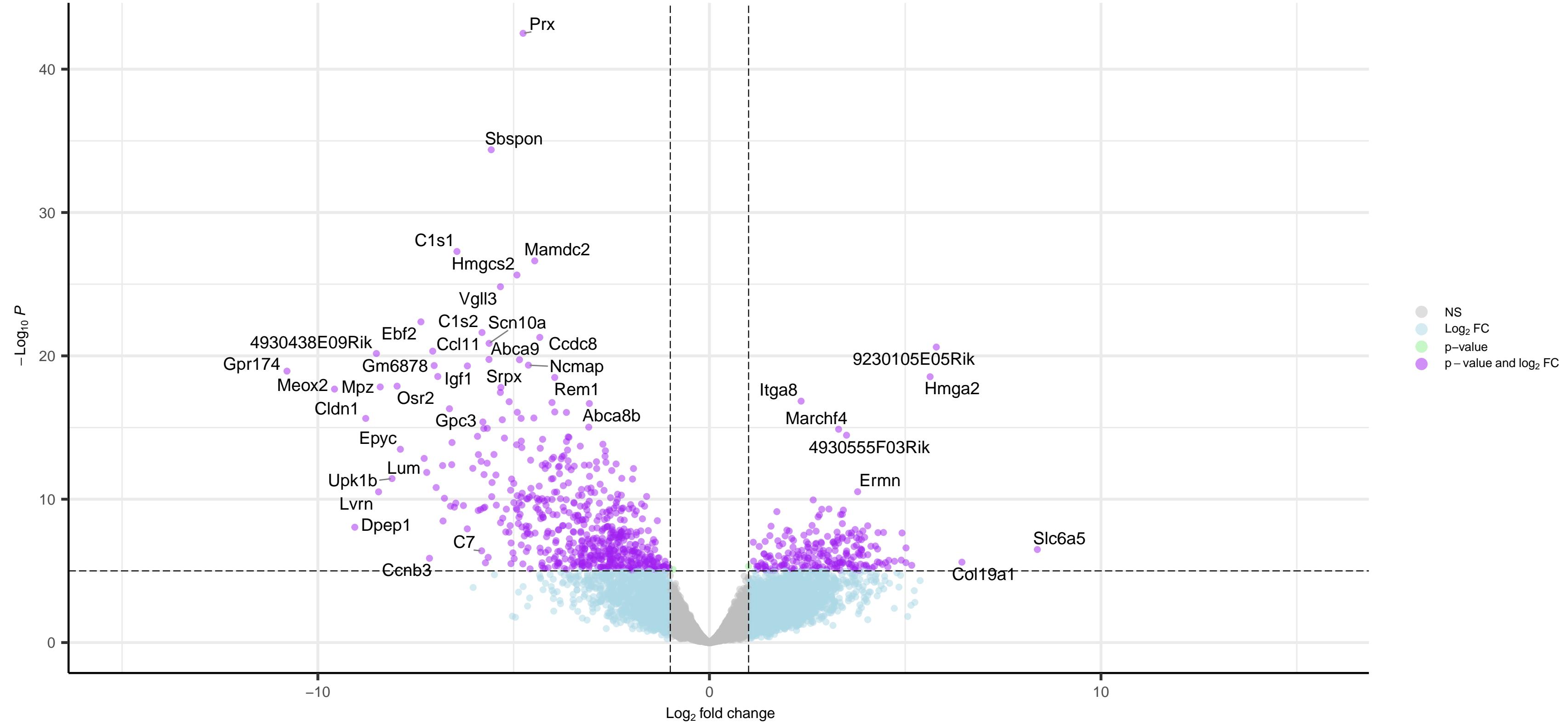
# Gliomas vs. Glioneuronal tumors

Of genotype: atrxwt



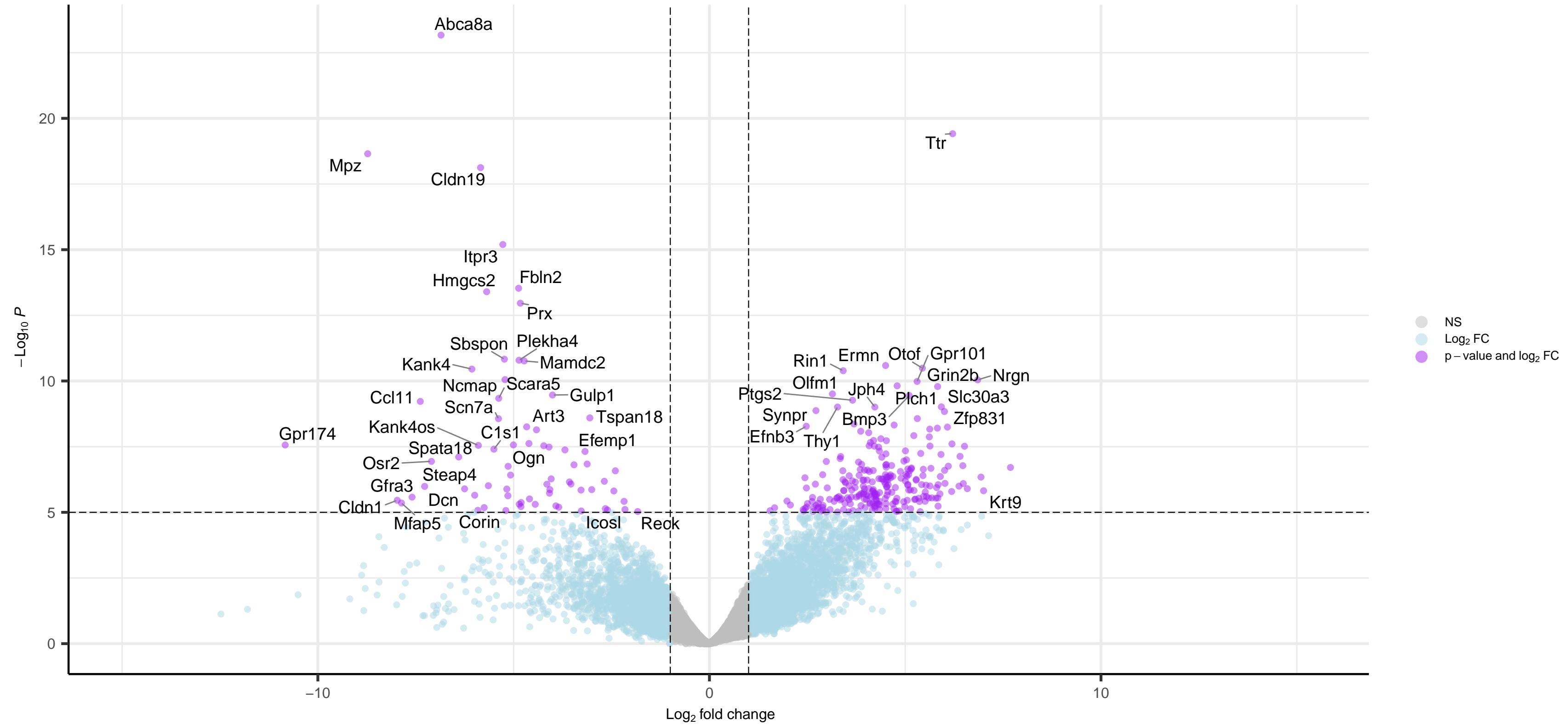
# Nerve sheath tumors vs. Gliomas

Of genotype: atrxwt



# Nerve sheath tumors vs. Glioneuronal tumors

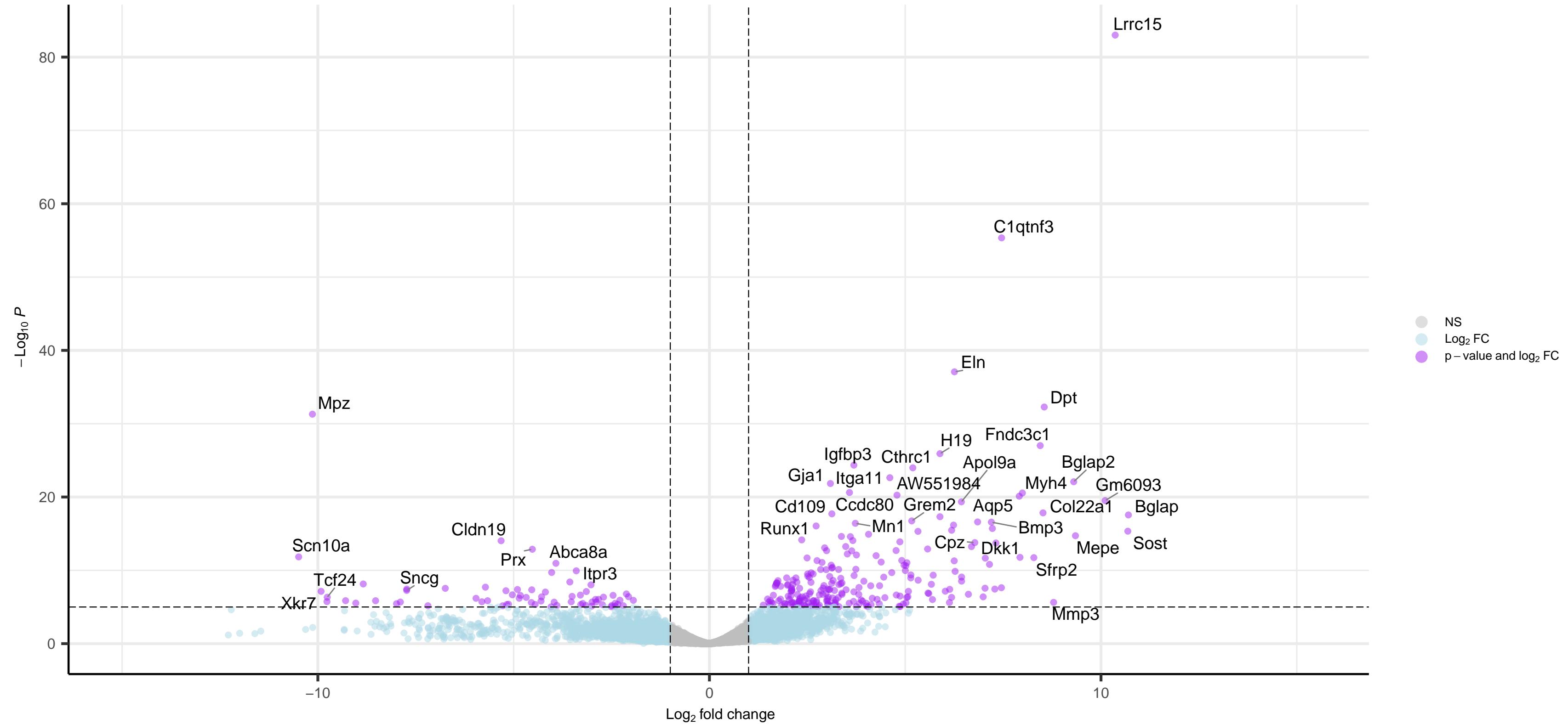
Of genotype: atrxwt



total = 16188 variables

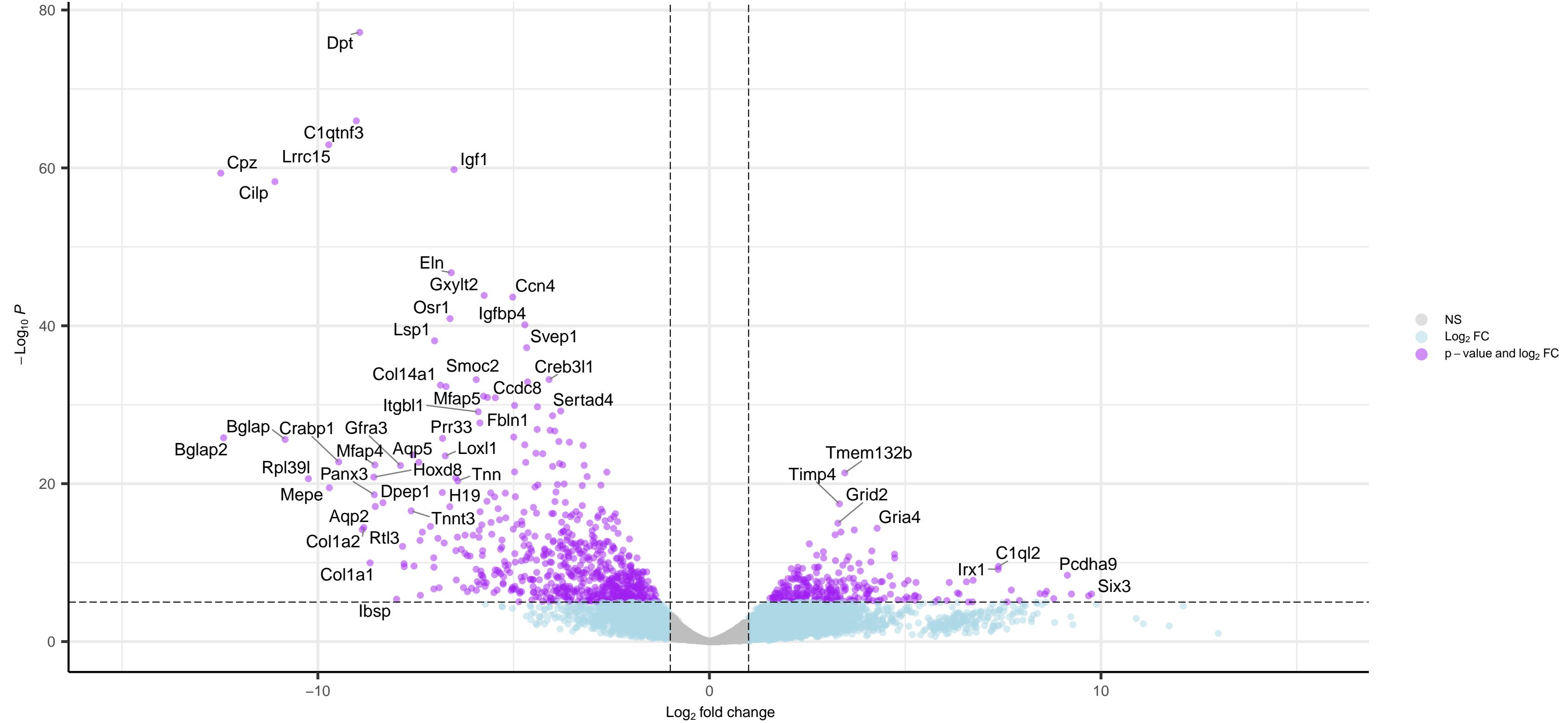
# Nerve sheath tumors vs. Spindle and epithelioid tumors

Of genotype: atrxwt



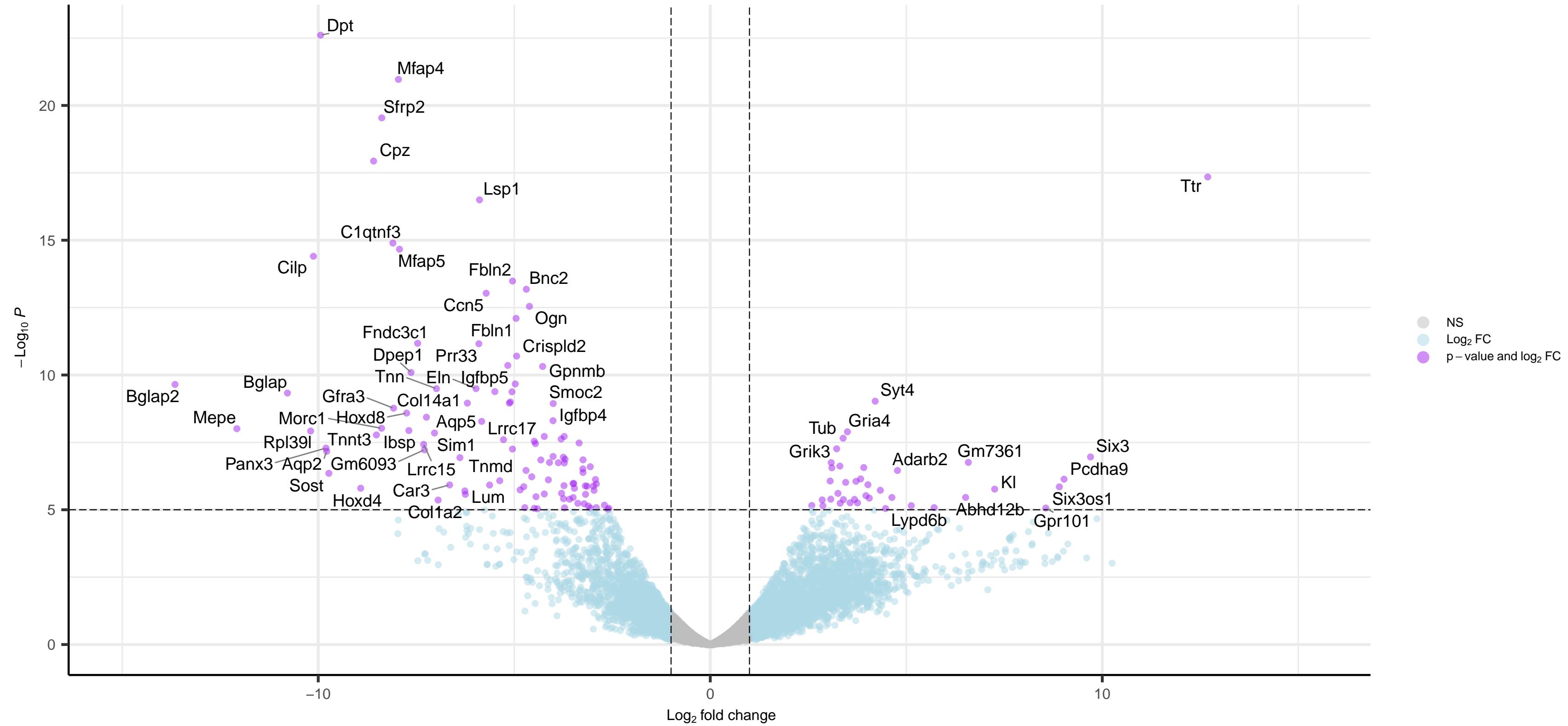
# Spindle and epithelioid tumors vs. Gliomas

Of genotype: atrxwt



## **Spindle and epithelioid tumors vs. Glioneuronal tu**

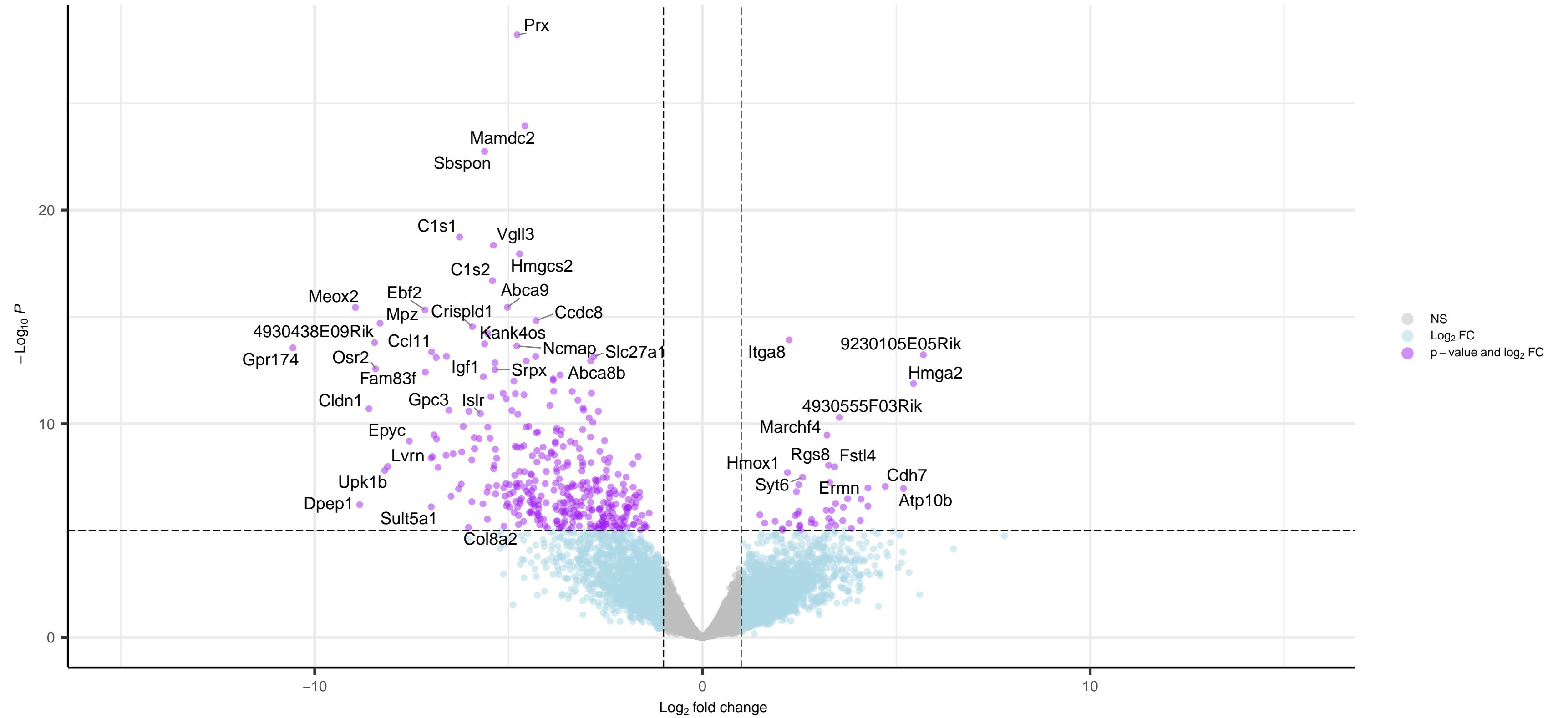
Of genotype: atrxwt



total = 15735 variables

# Nerve sheath tumor, low grade vs. Glioma, grade 4

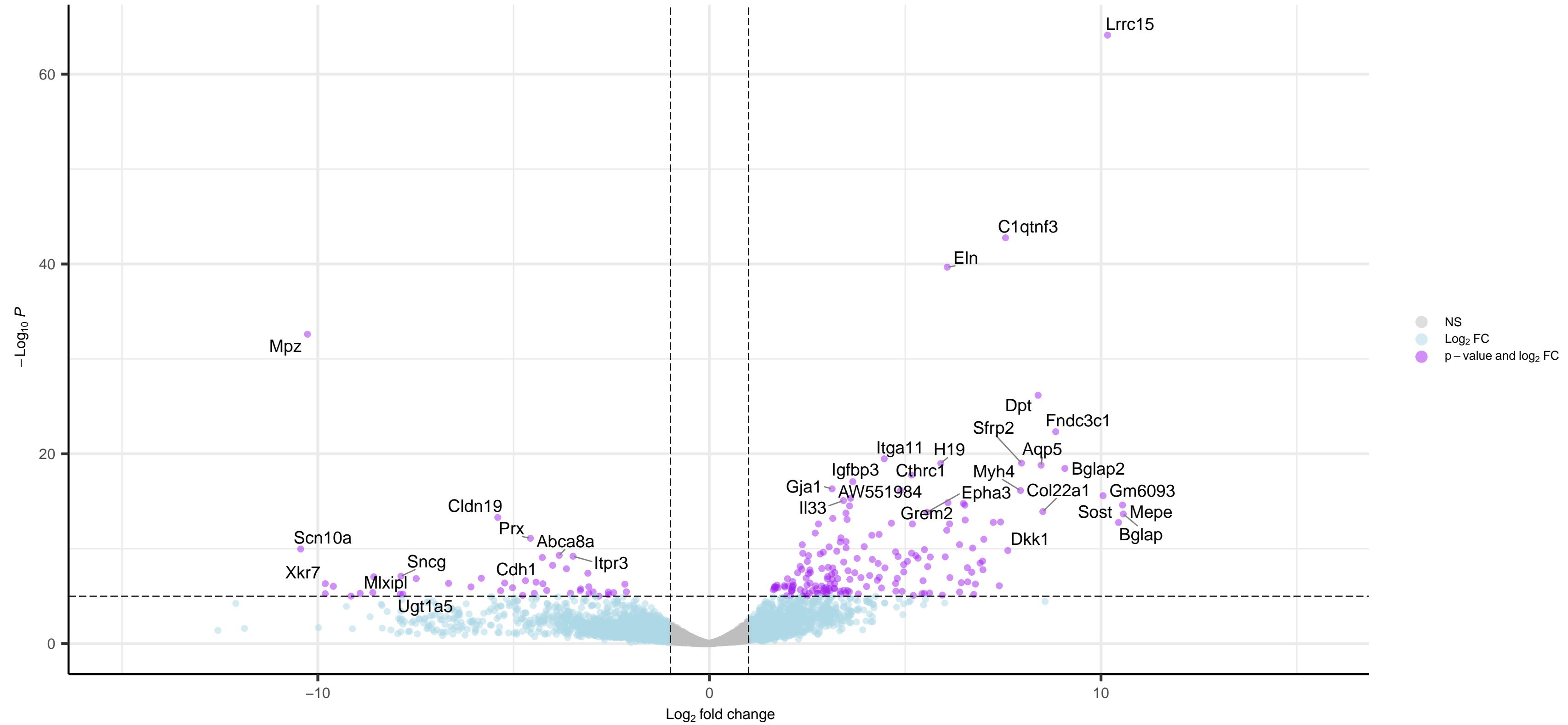
Of genotype: atrxwt



total = 15474 variables

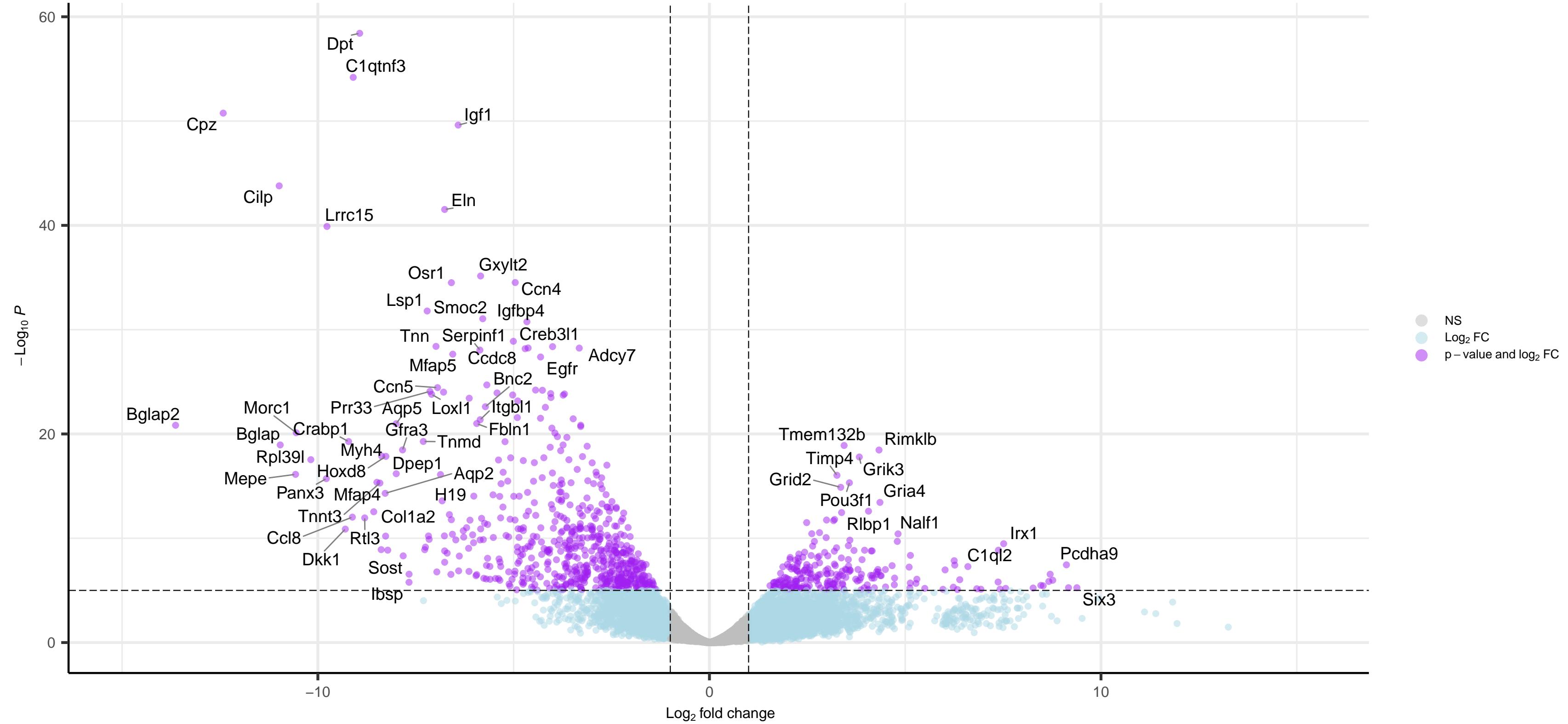
# Nerve sheath tumor, low grade vs. Spindle and epithelioid tumor, high grade

Of genotype: atrxwt



# Spindle and epithelioid tumor, high grade vs. Glioma, grade 4

Of genotype: atrxwt



total = 16018 variables

## 2 vs. 4

Of genotype: atrxwt

