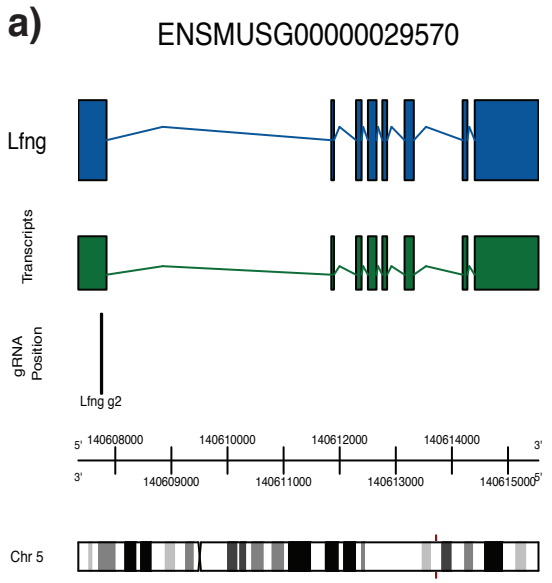

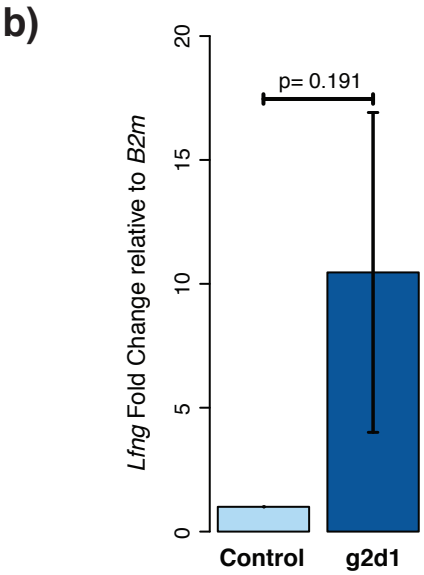

c)

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ENSMUST00000031555                 MLQRCGRRLL LALVGALLAC LLVLTADPPP TPMPAERGRR ALRSLAGSSG
Lfng_g2_g2d1_clone_protein_Tru     MLQRCGRRLL LALVGALLAC LLVLTADPPP TPMPAERGRR ALRSLAGSSG

                                  51
ENSMUST00000031555                 GAPASGSRAA VDPGVLTREV HSLSEYFSLL TRARRDADPP PGVASRQGDG
Lfng_g2_g2d1_clone_protein_Tru     GAPASGSRAA VDPGVLTREV HSLSEYFSLL TRARRDADPP PGVASRQGDG

                                 101
ENSMUST00000031555                 HPRPPAEVLS PRDVFIAVKT TRKFHRARLD LLFETWISRH KEMTFIFTDG
Lfng_g2_g2d1_clone_protein_Tru     HPRPPAEVLS PR-------- -RRLHRR--- ---------- ----------

                                 151
ENSMUST00000031555                 EDEALAKLTG NVVLTNCSSA HSRQALSCKM AVEYDRFIES GKKWFCHVDD
Lfng_g2_g2d1_clone_protein_Tru     ---------- ---------- QDHQKVSPRA A--------- ----------

                                 201
ENSMUST00000031555                 DNYVNLRALL RLLASYPHTQ DVYIGKPSLD RPIQATERIS EHKVRPVHFW
Lfng_g2_g2d1_clone_protein_Tru     ------RSAV RDLDLAPQGD DVHLH----- ---------- ----------

                                 251
ENSMUST00000031555                 FATGGAGFCI SRGLALKMGP WASGGHFMST AERIRLPDDC TIGYIVEALL
Lfng_g2_g2d1_clone_protein_Tru     ---------- ---------- ---------- ---------- ----------

                                 301
ENSMUST00000031555                 GVPLIRSGLF HSHLENLQQV PTTELHEQVT LSYGMFENKR NAVHIKGPFS
Lfng_g2_g2d1_clone_protein_Tru     ---------- ---------- ---------- ---------- ----------

                                 351
ENSMUST00000031555                 VEADPSRFRS VHCHLYPDTP WCPRSAIF
Lfng_g2_g2d1_clone_protein_Tru     ---------- ---------- --------
```