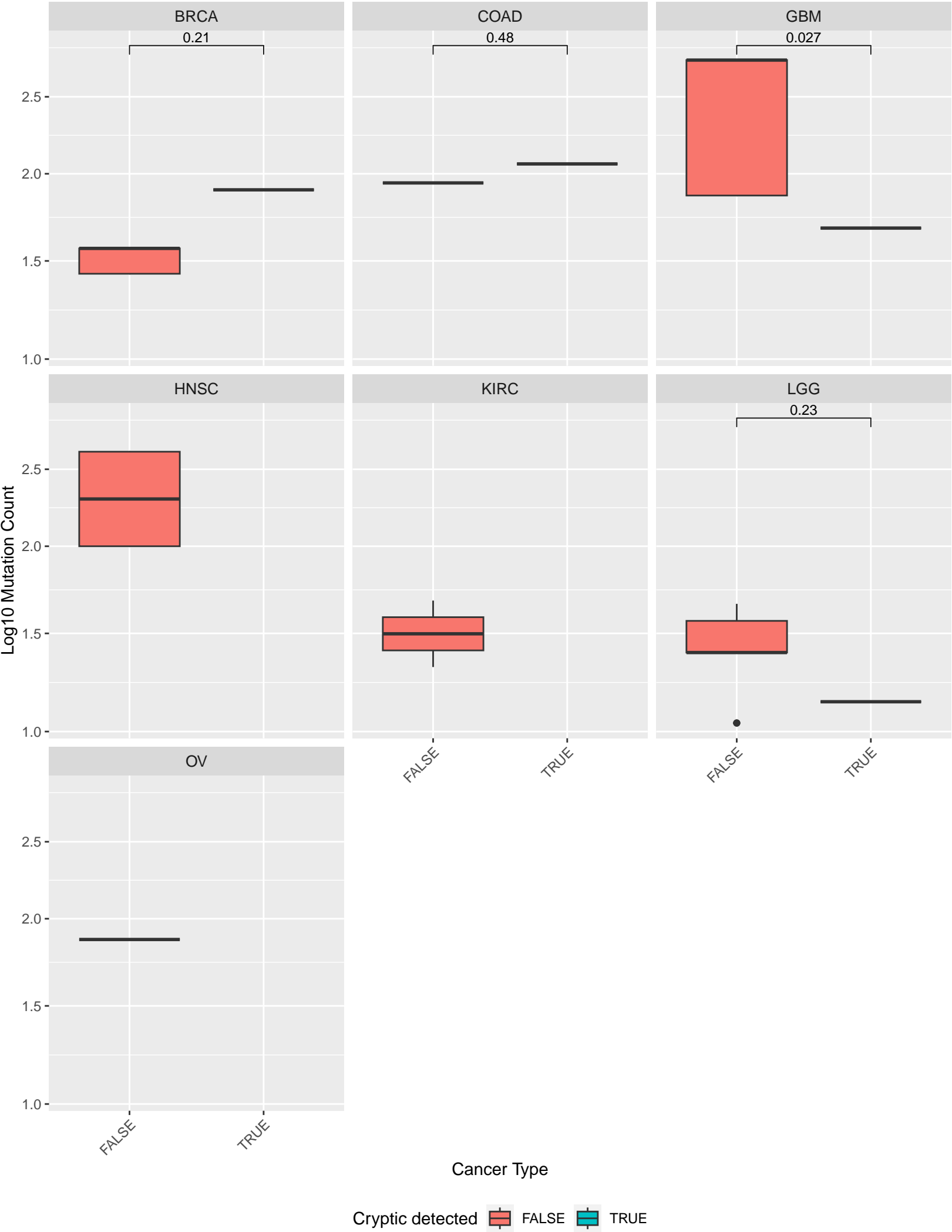


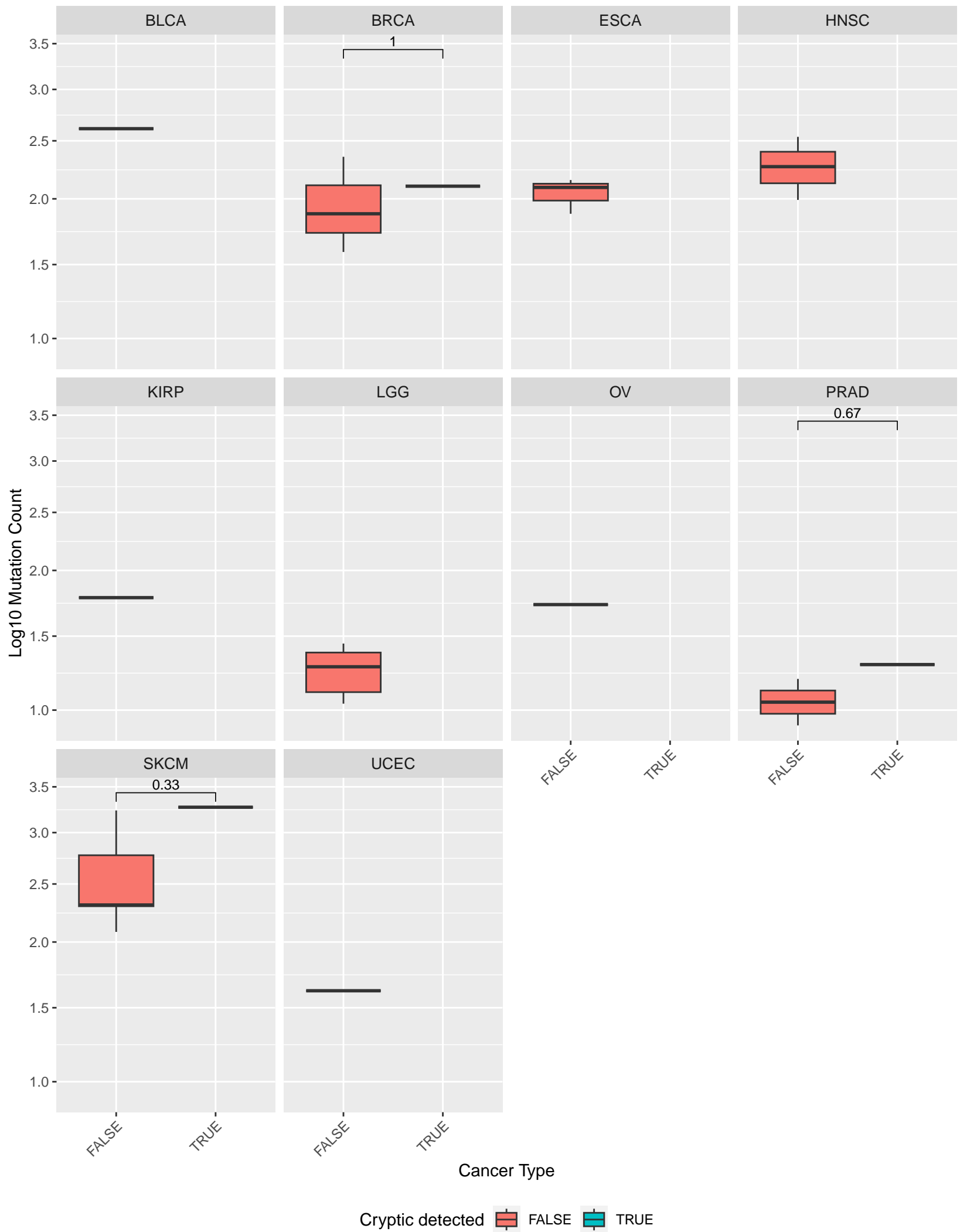
chr1:108896432-108896863 – GPSM2

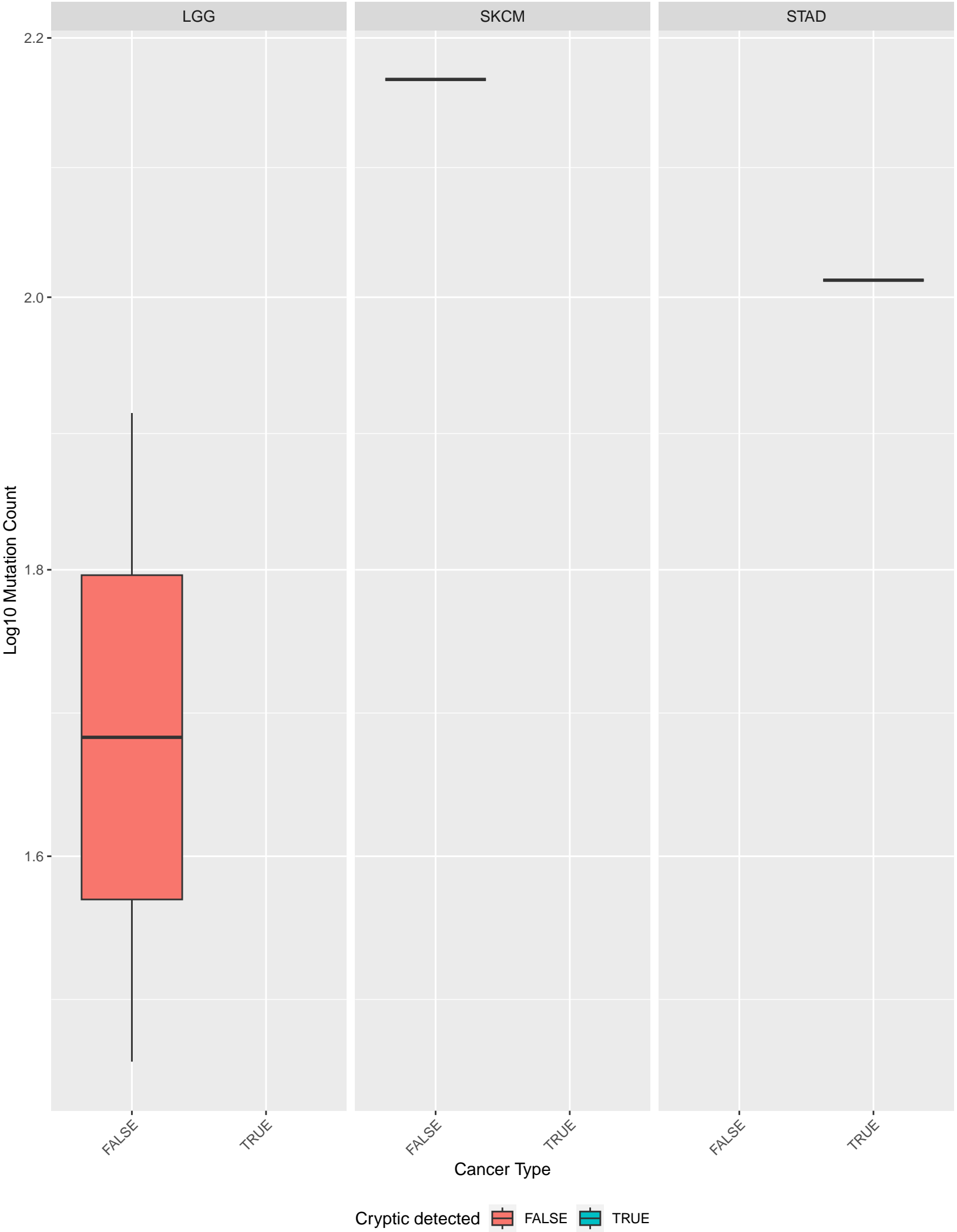


chr1:243512427-243523308 – AKT3



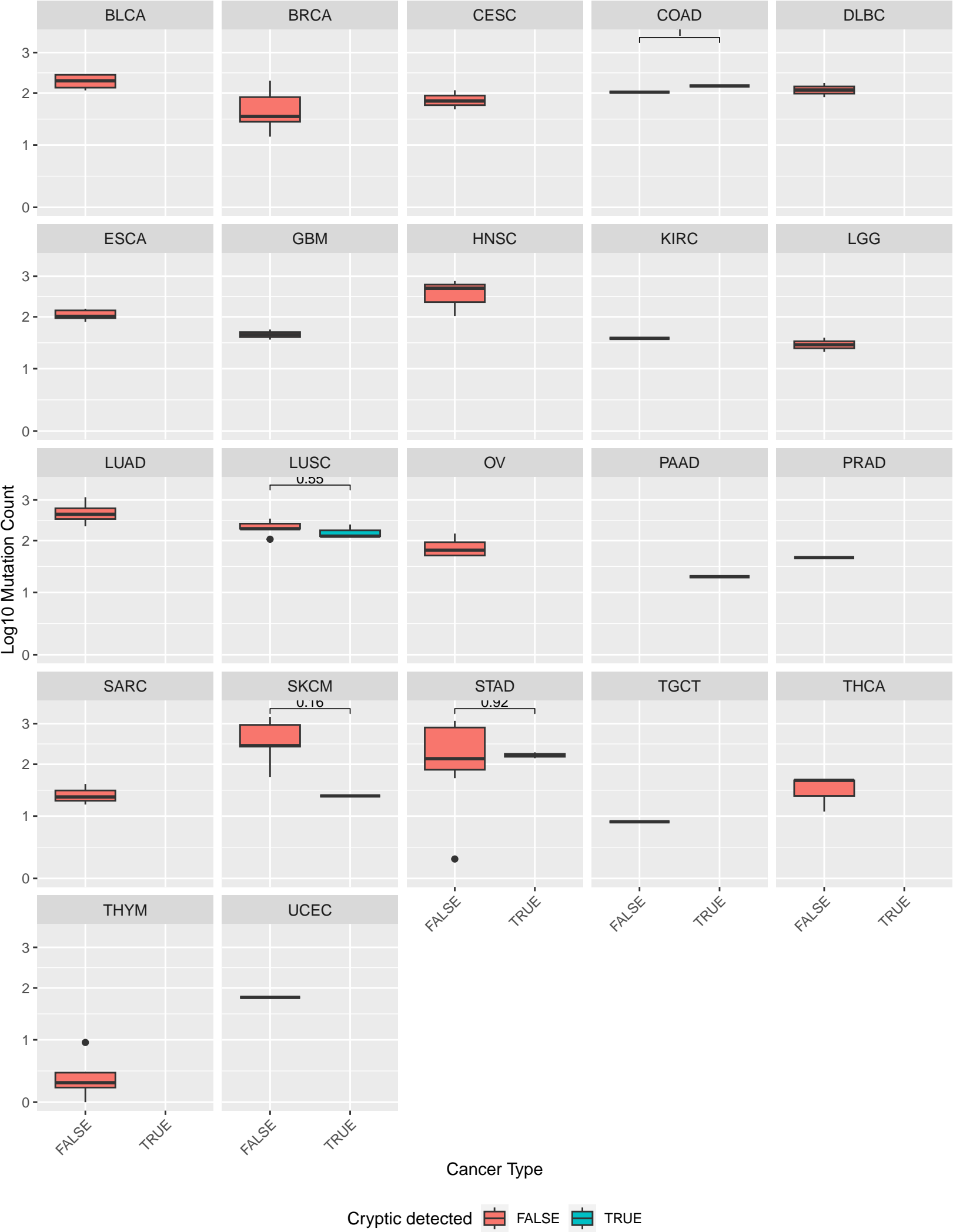
chr1:243523468-243545509 – AKT3



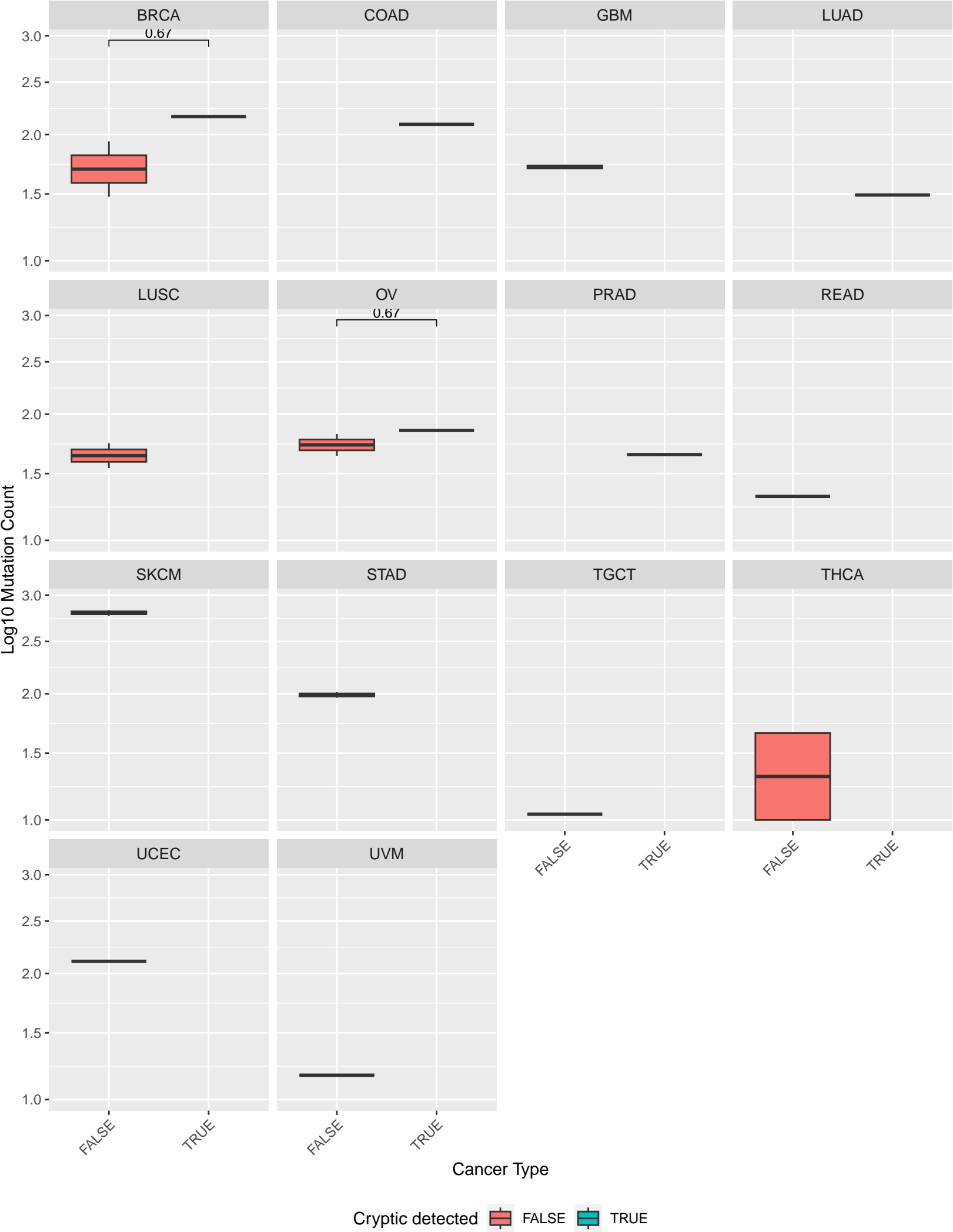


chr2:3458516–3460262 – TRAPPC12



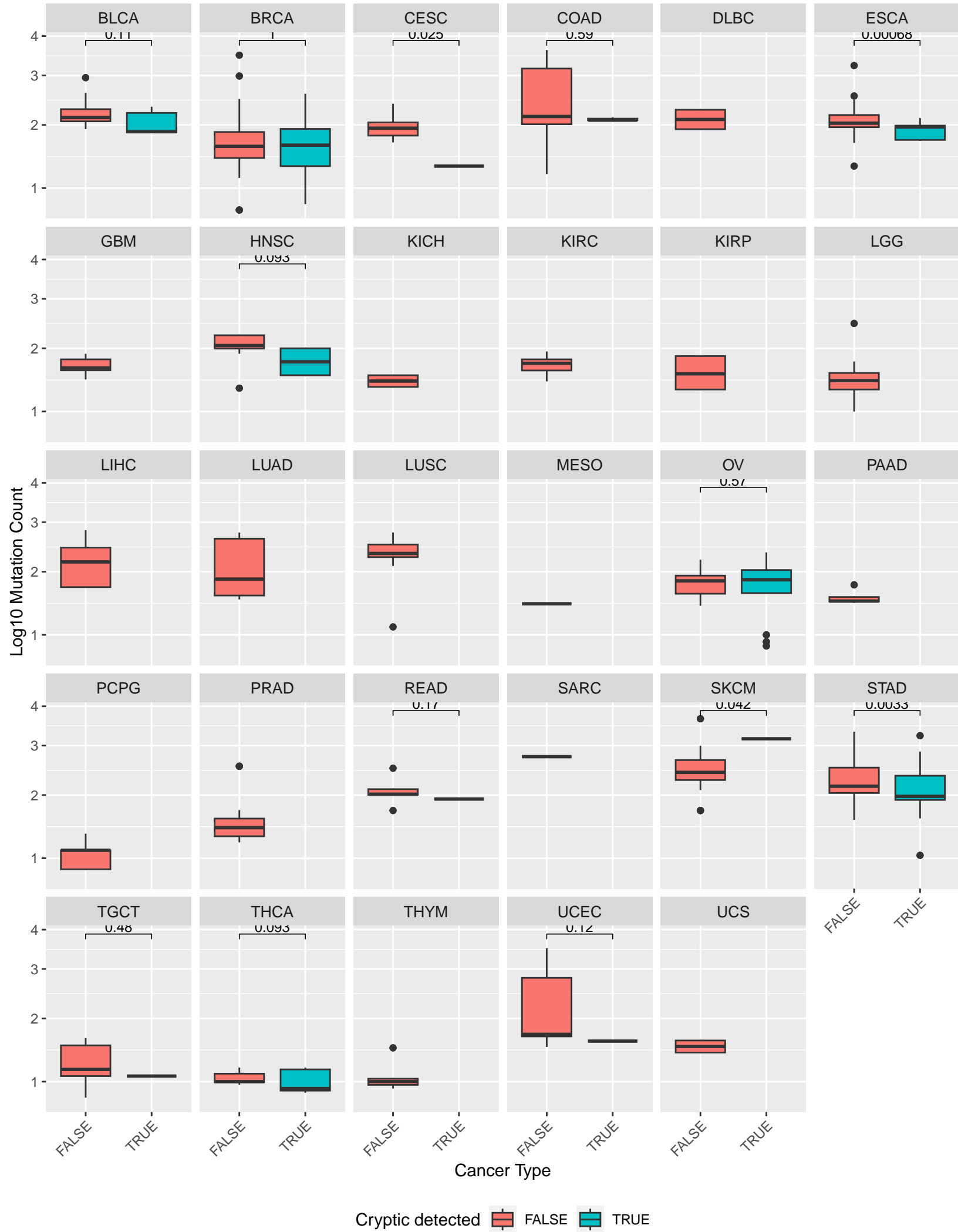


chr2:241668986–241670725 – ATG4B

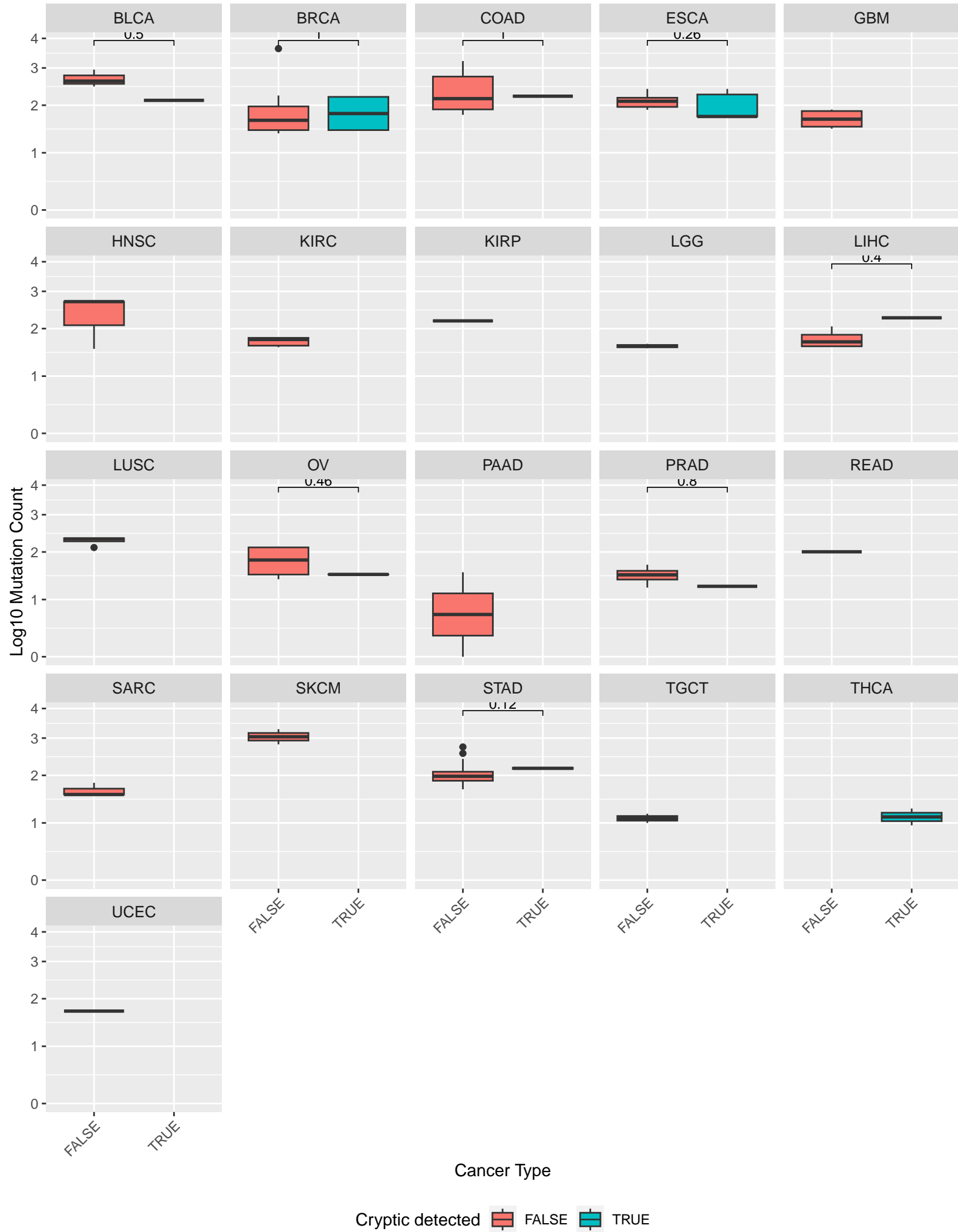




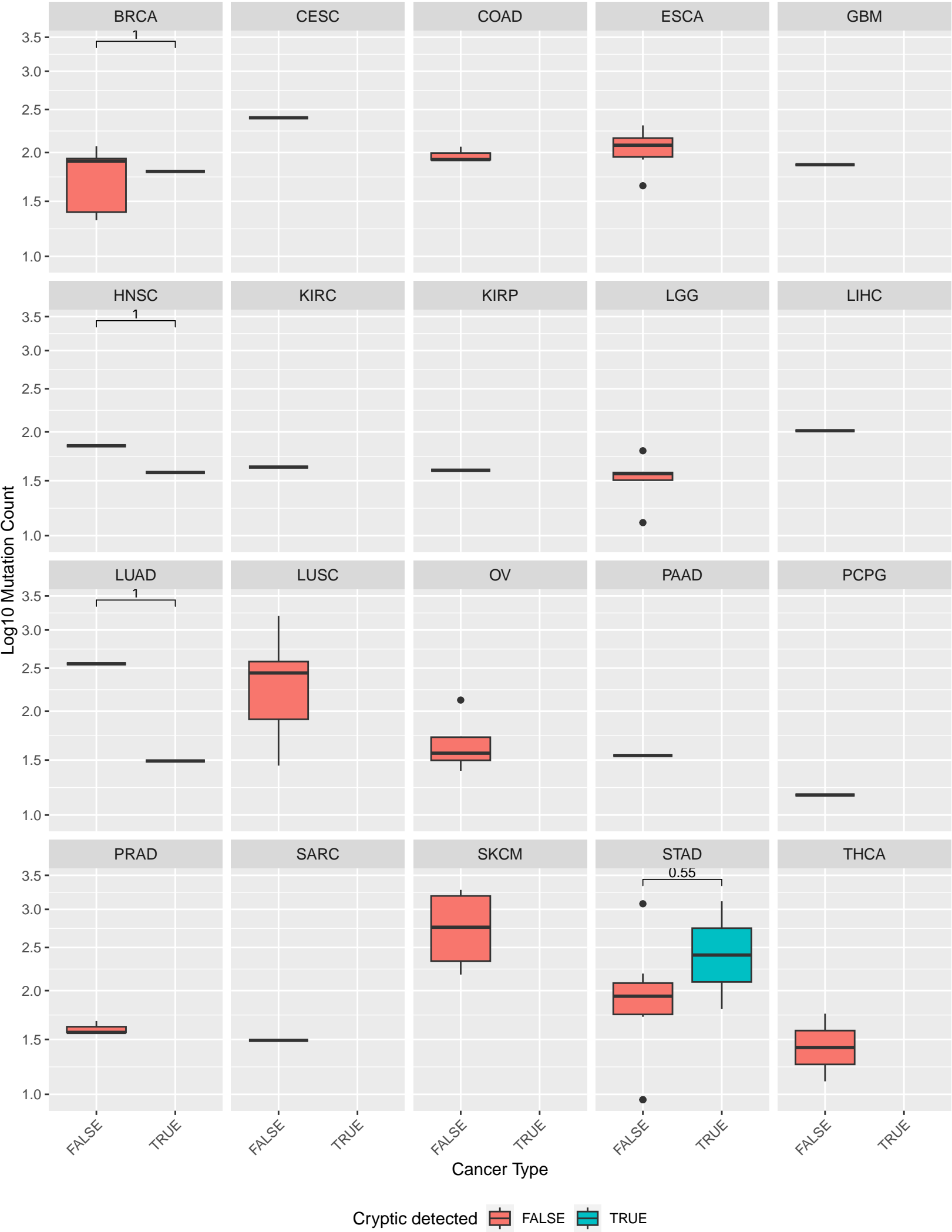
chr3:9468616–9470458 – SETD5



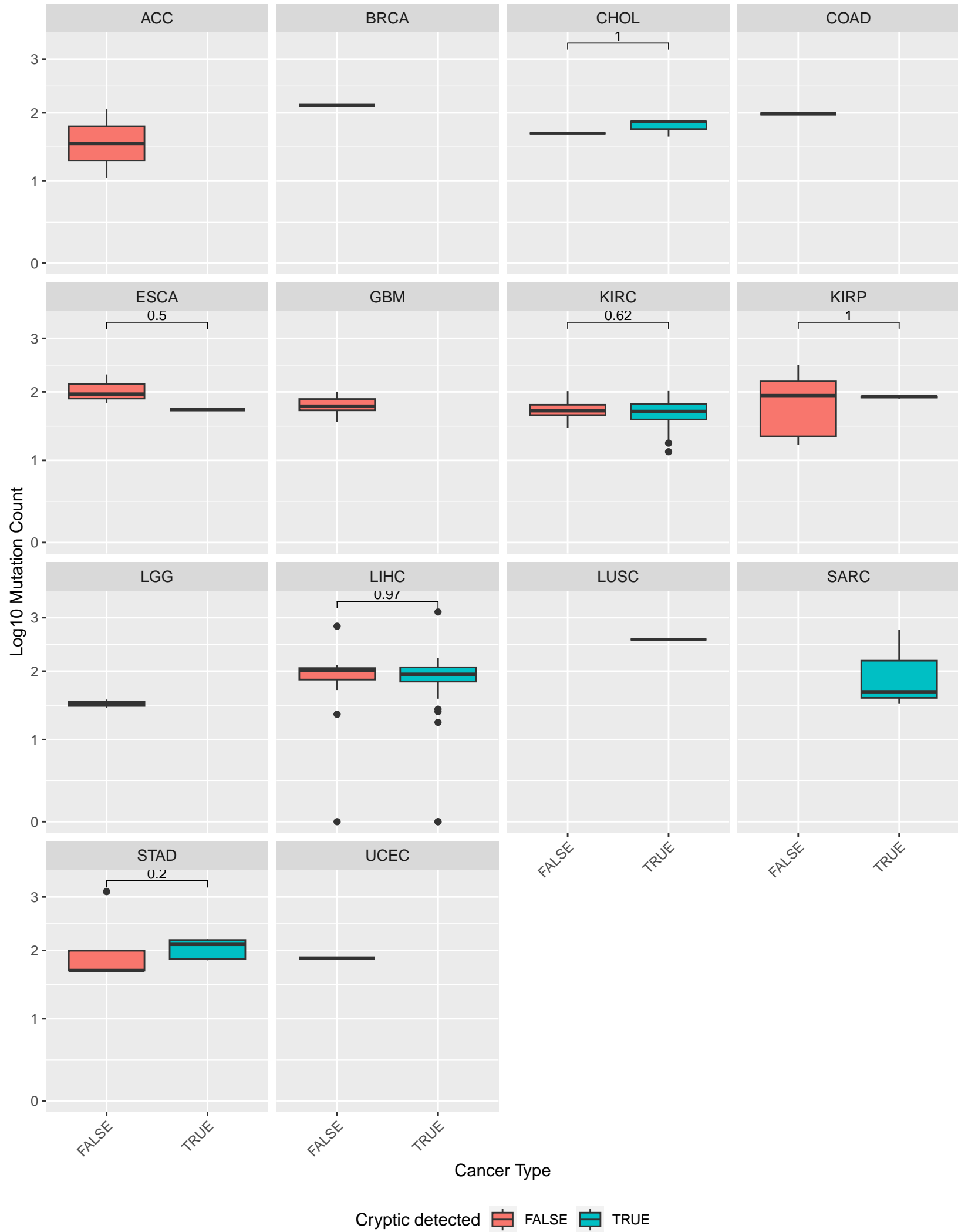
chr3:124700034–124701092 – KALRN



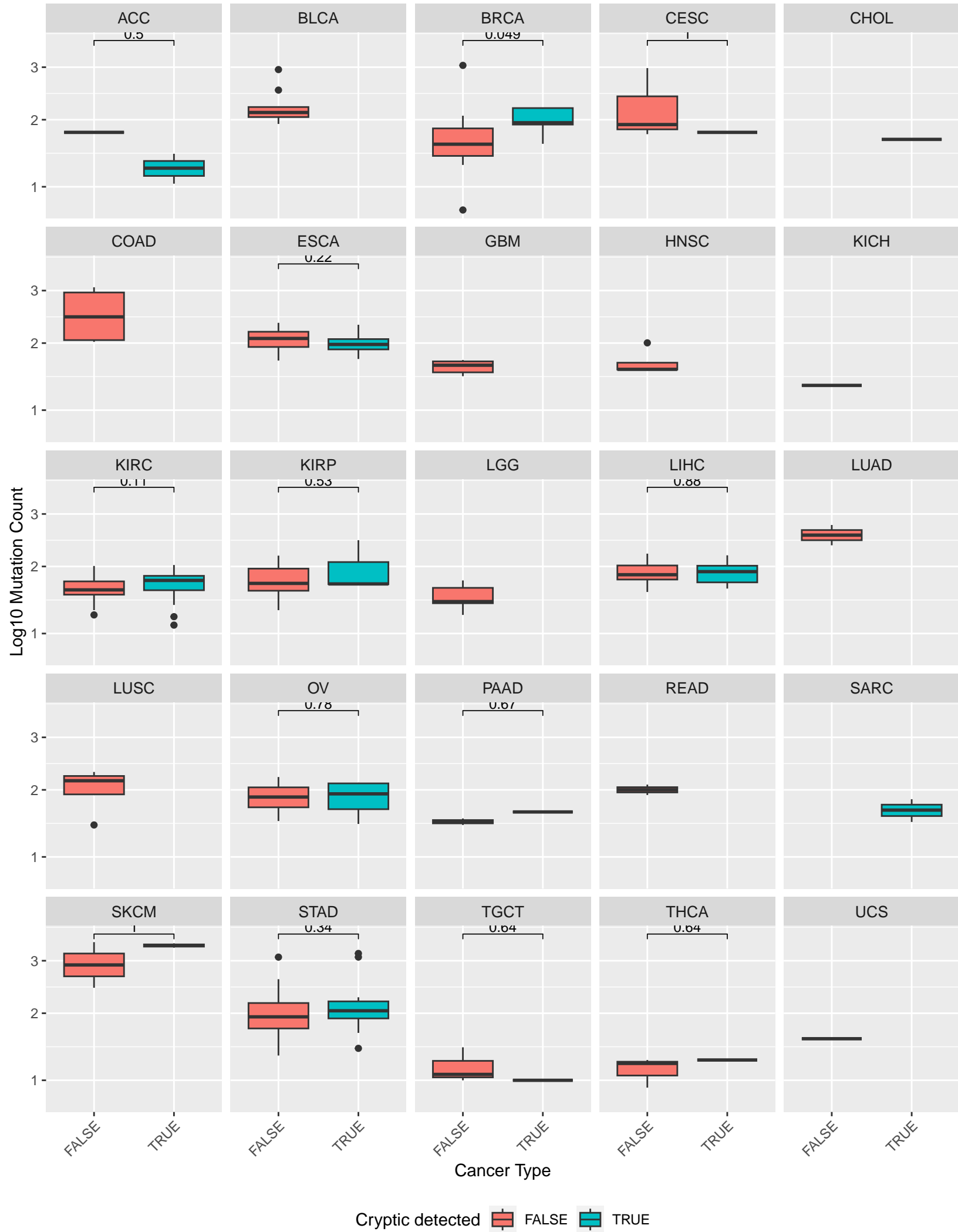
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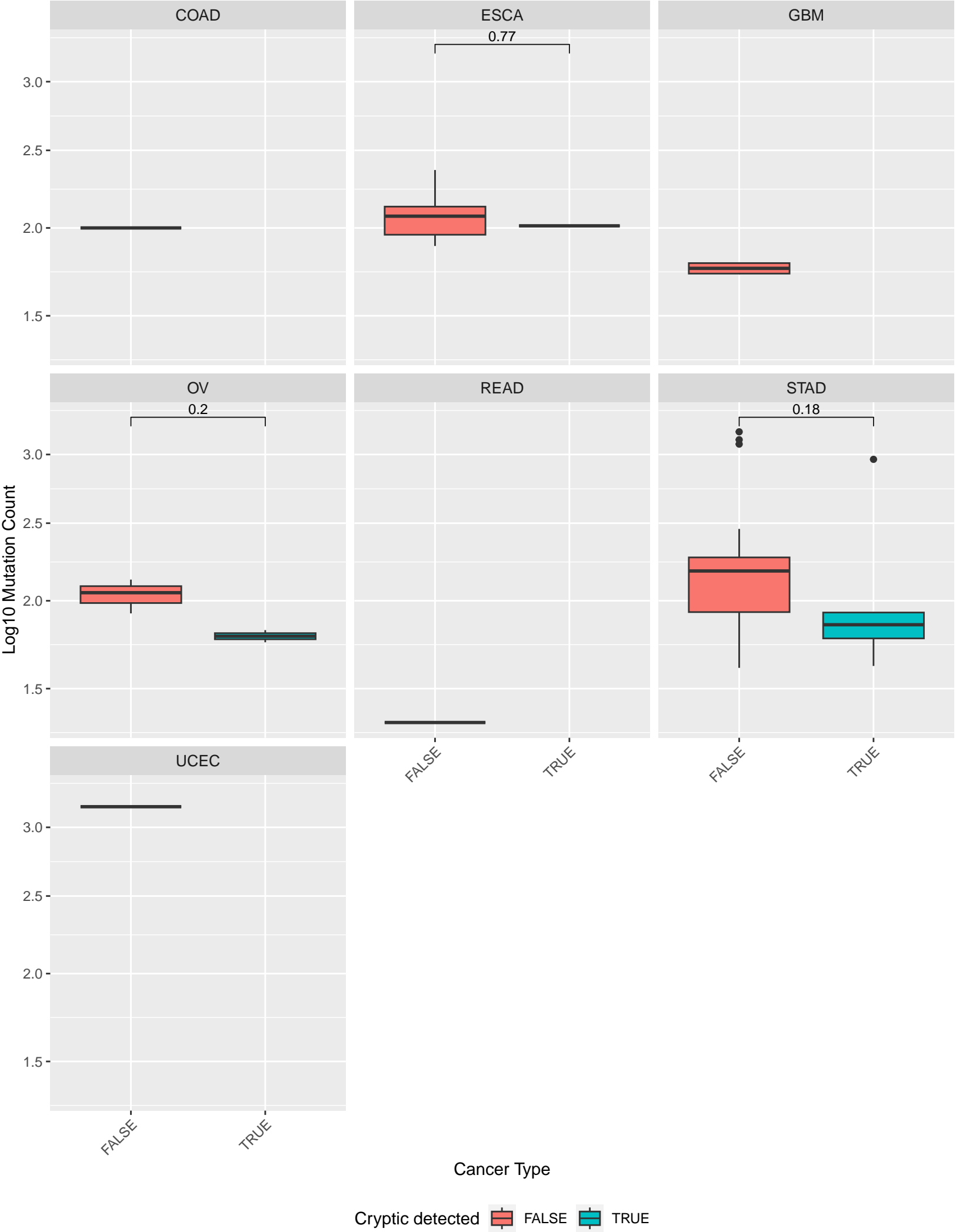


chr3:124701599–124702037 – KALRN



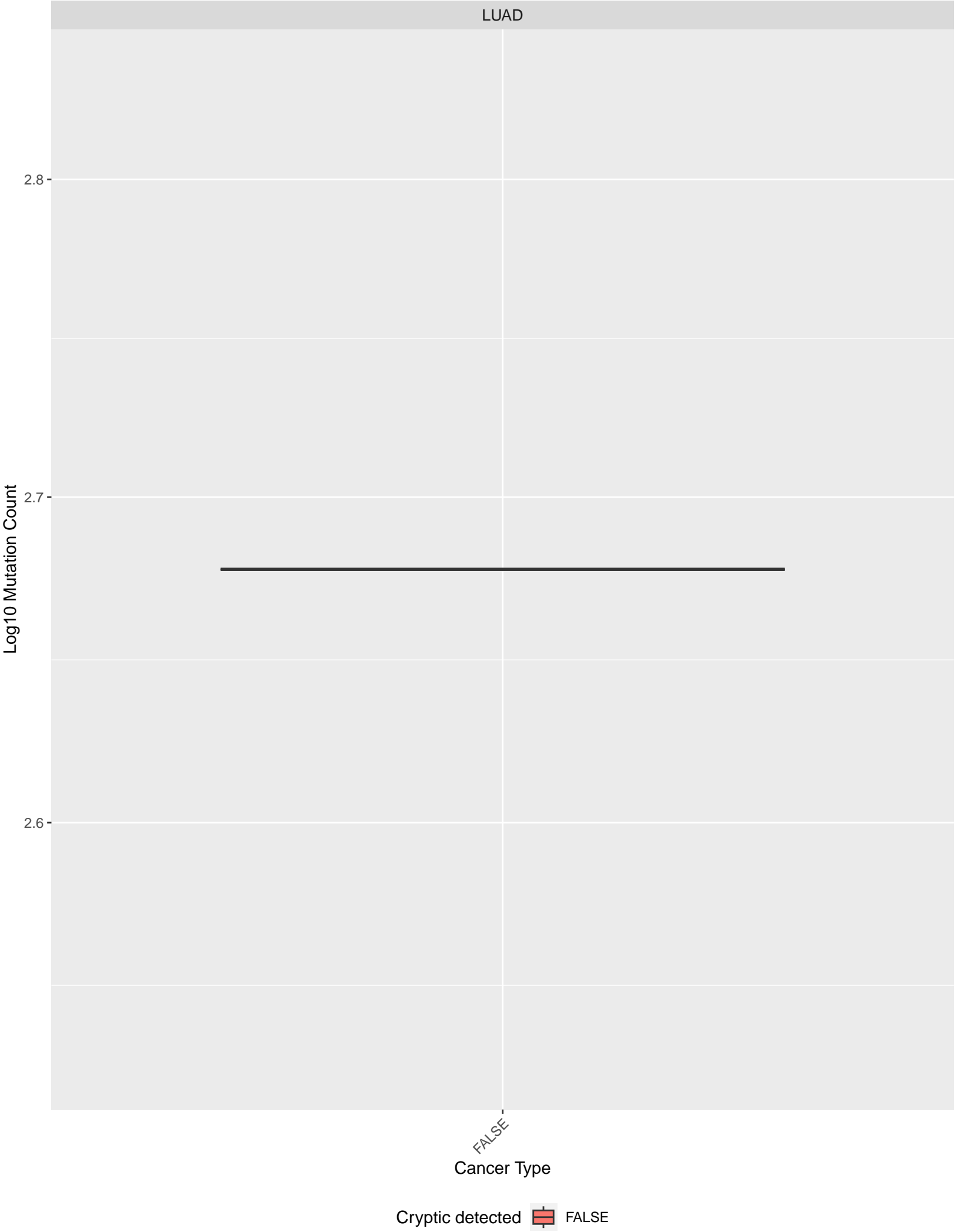
chr3:124701256–124702037 – KALRN



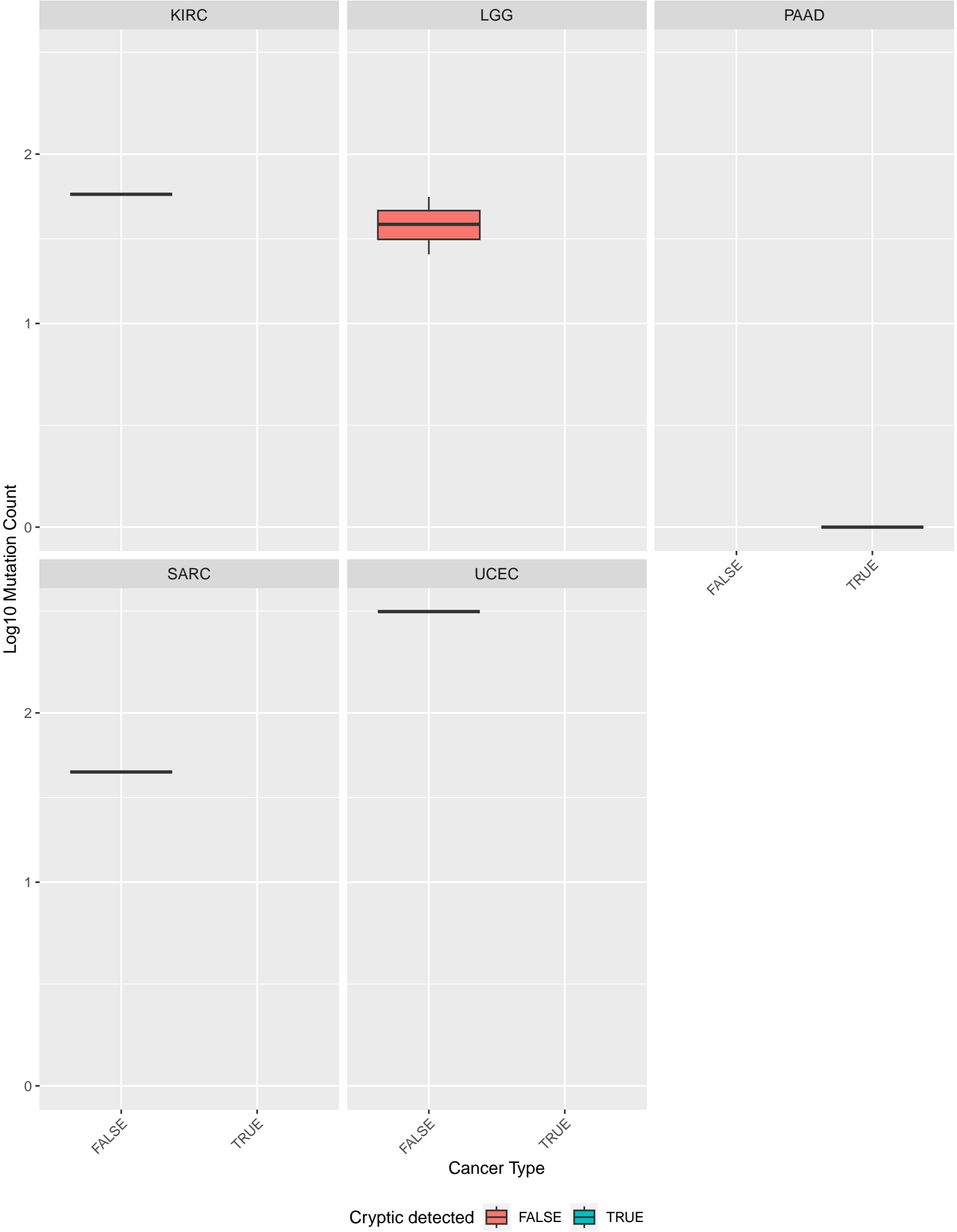


chr5:69261459–69262204 – CDK7

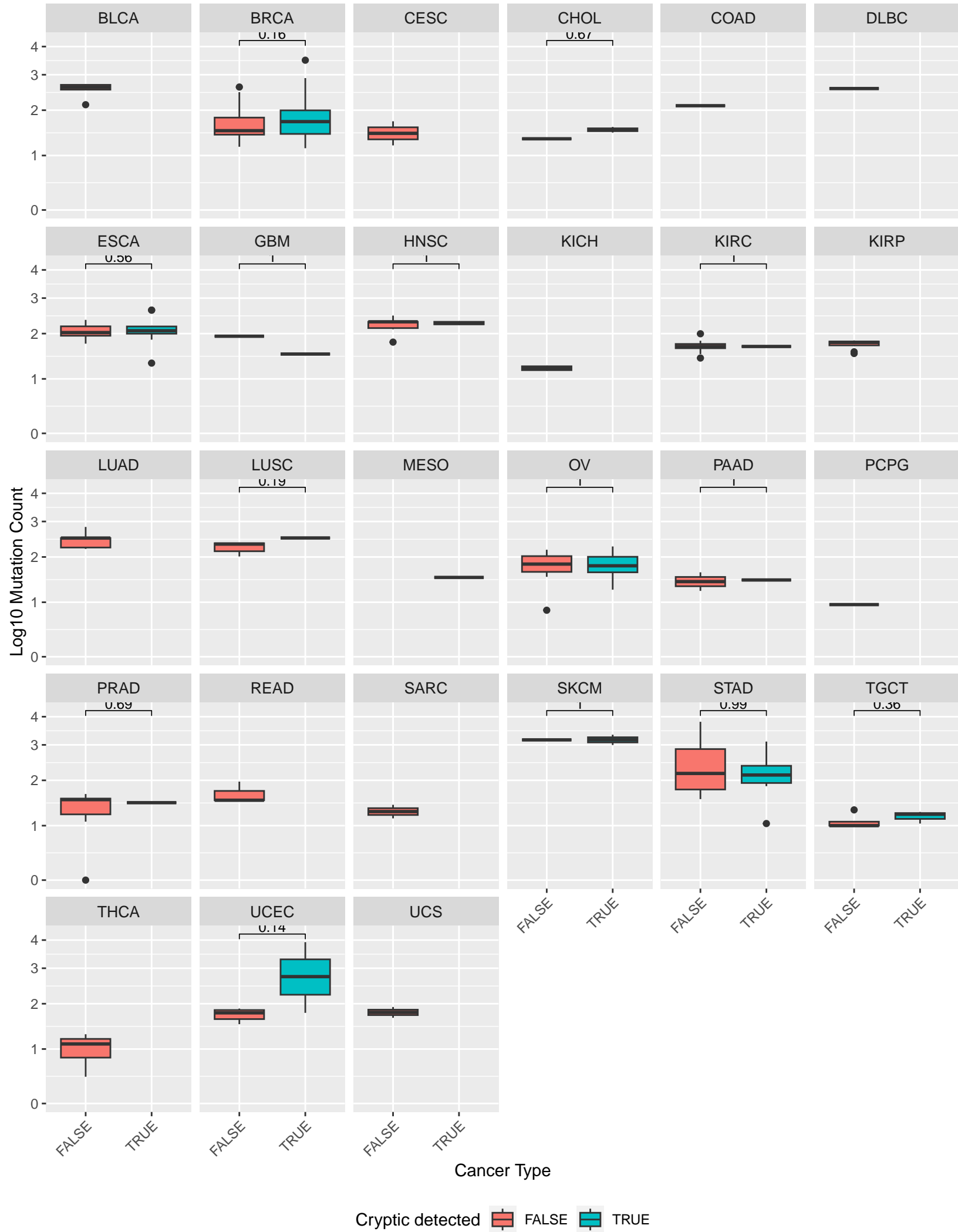






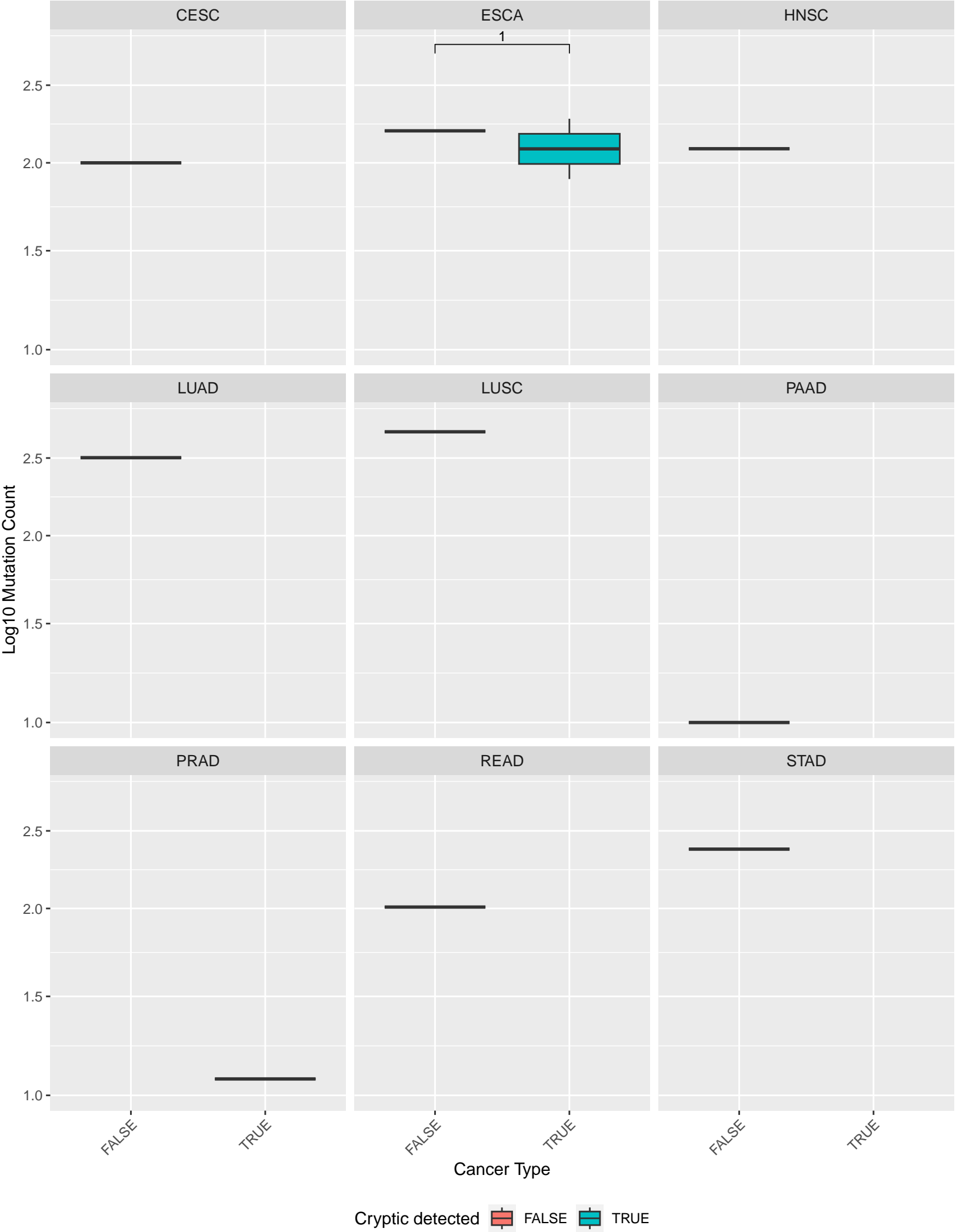


chr5:112266331-112267209 – EPB41L4A

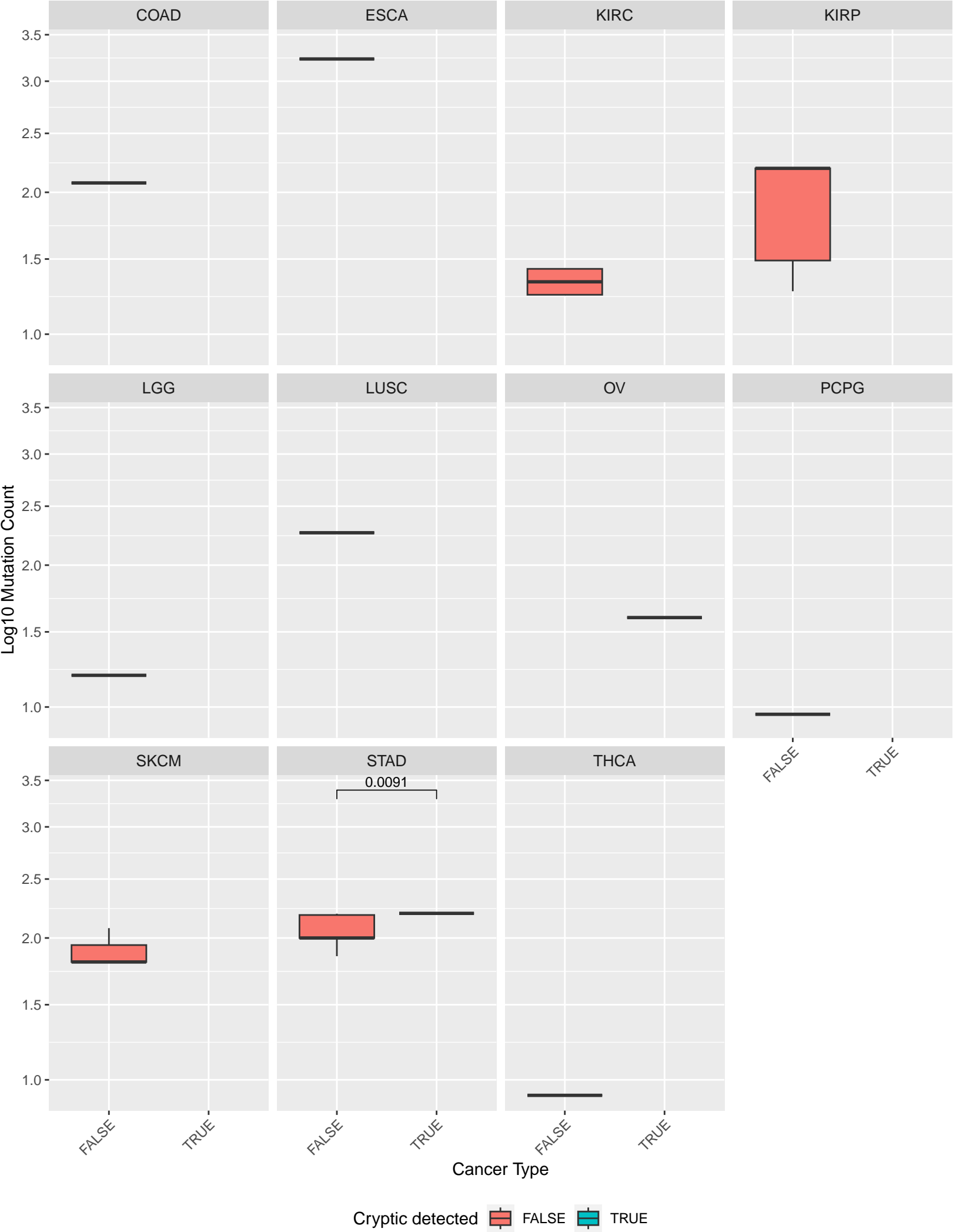


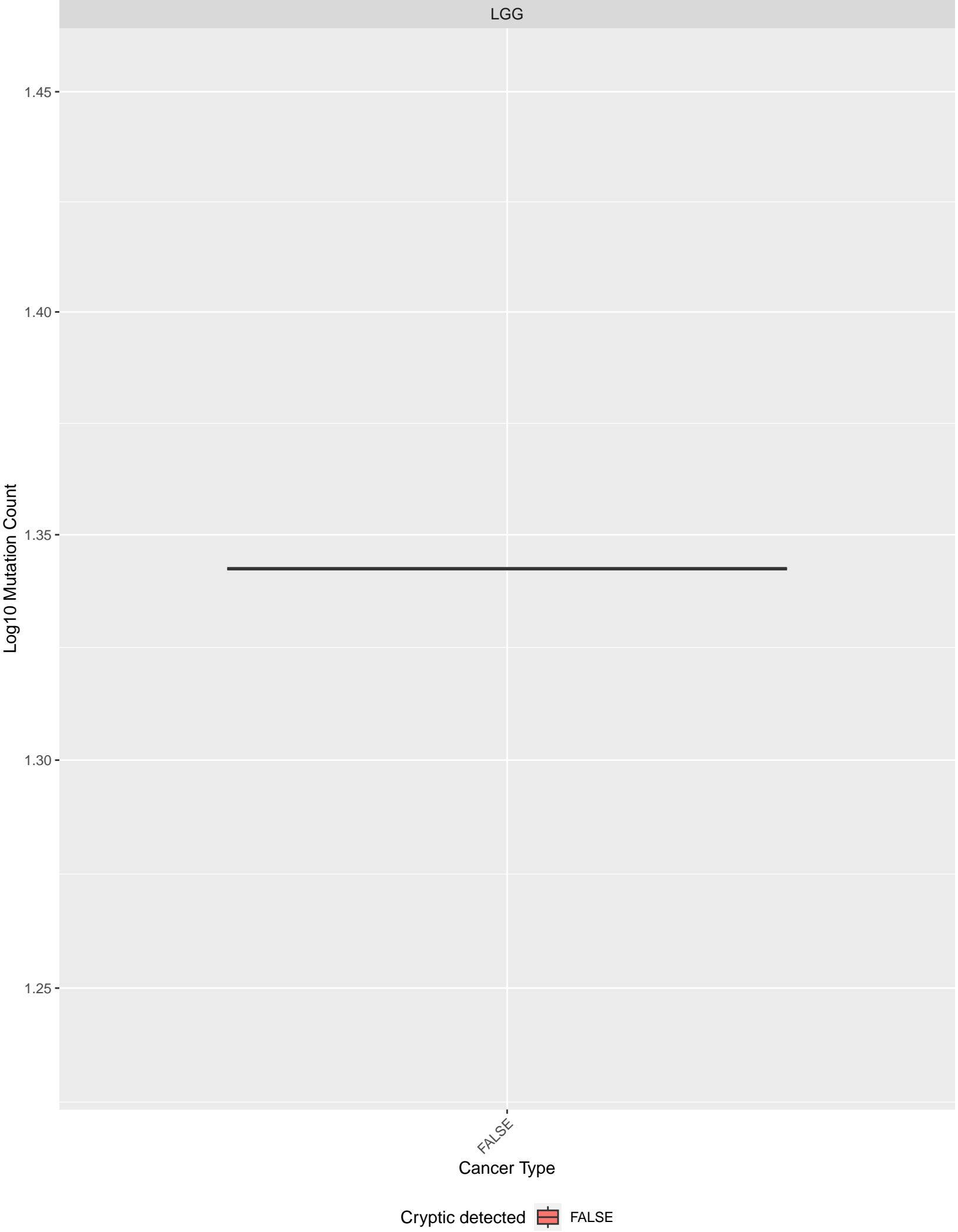
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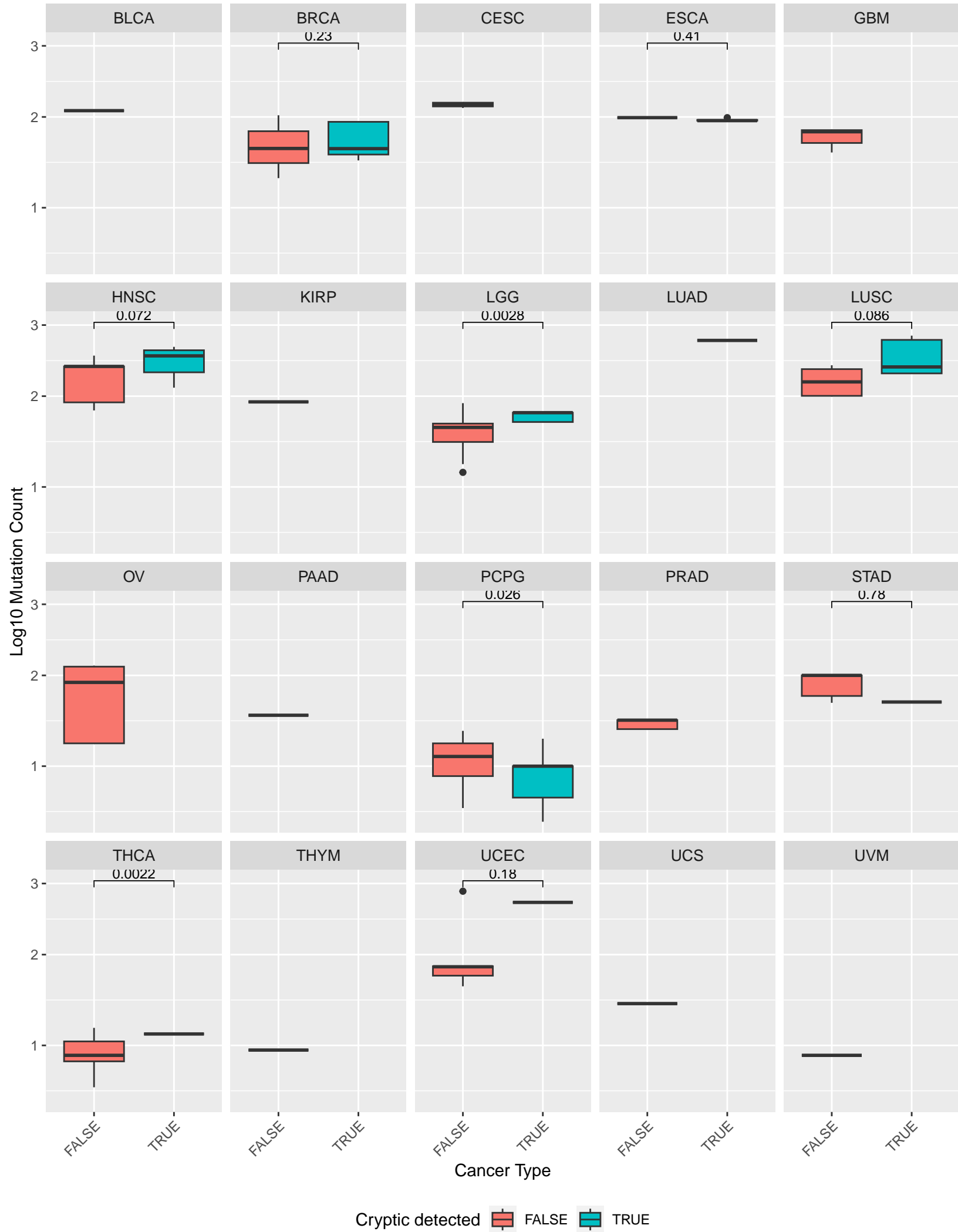




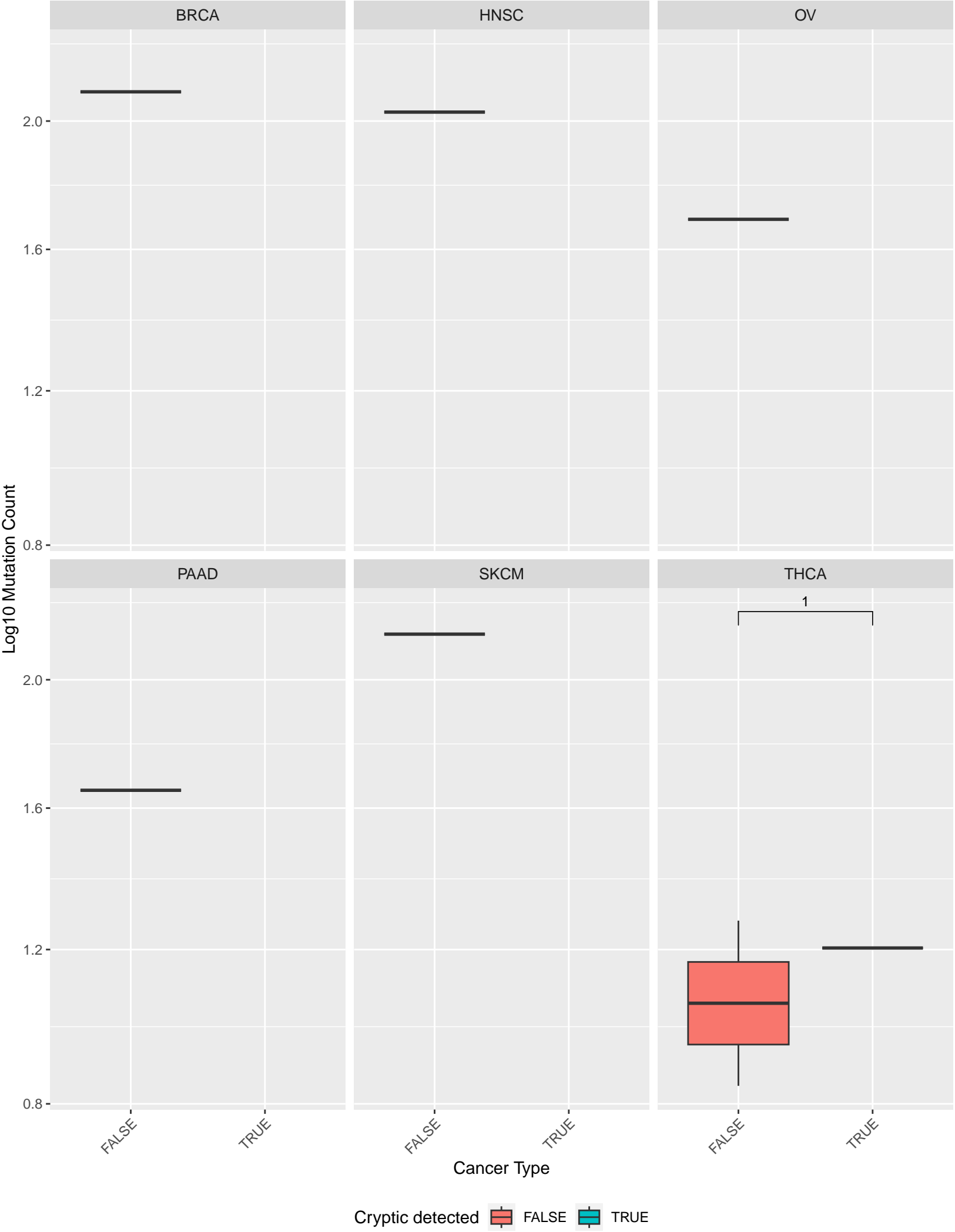
chr7:2565072–2567115 – IQCE



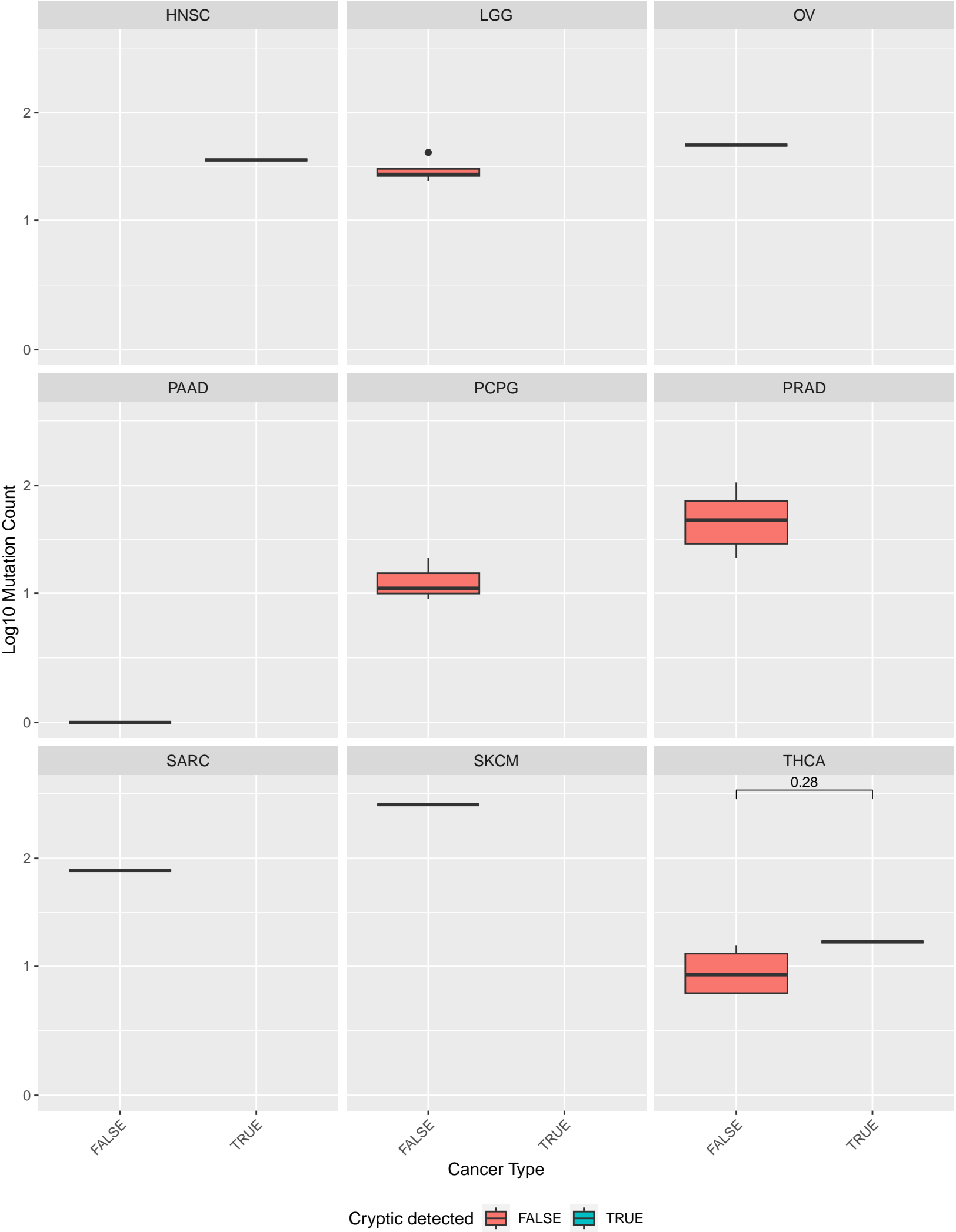




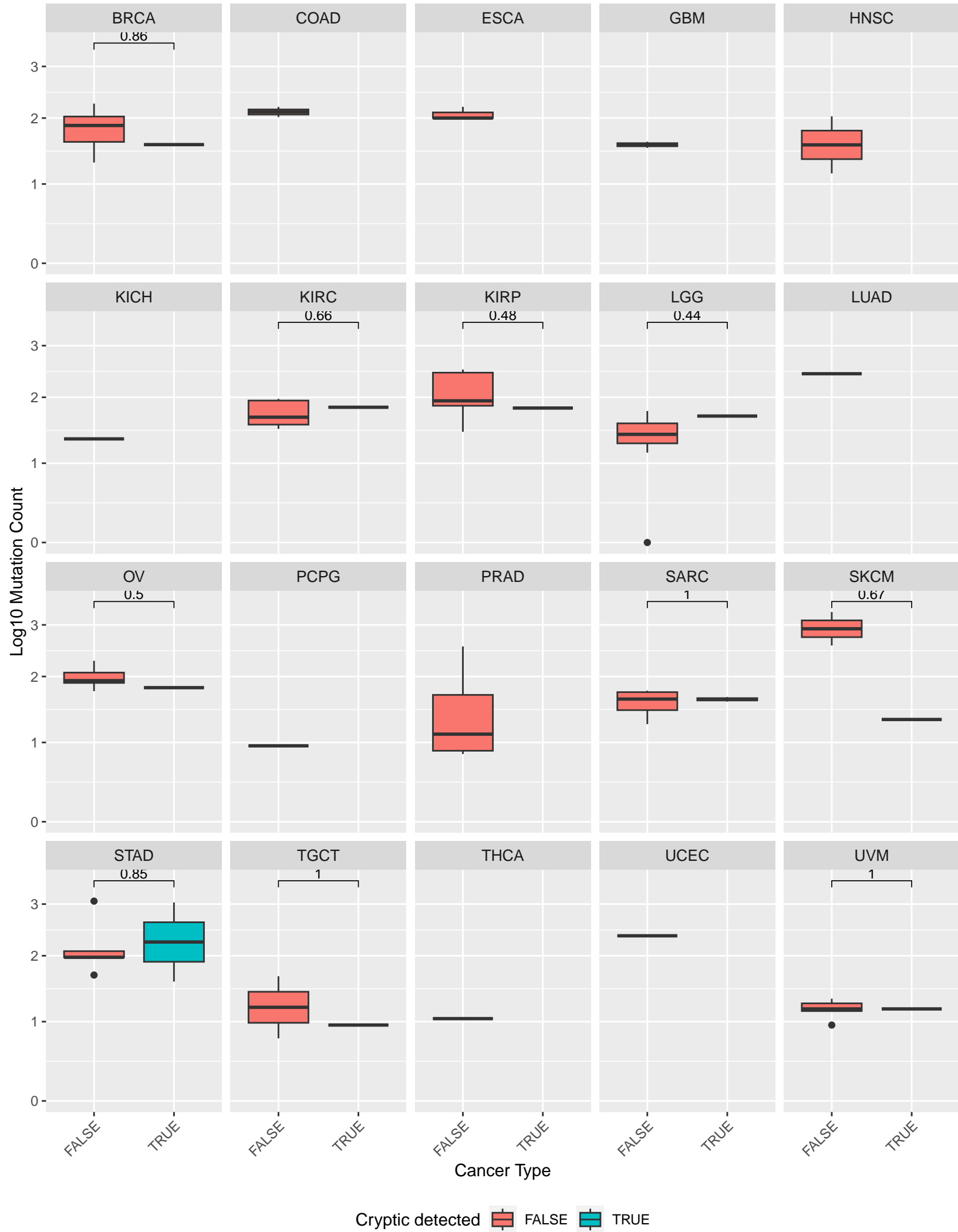
chr7:44254608–44258489 – CAMK2B



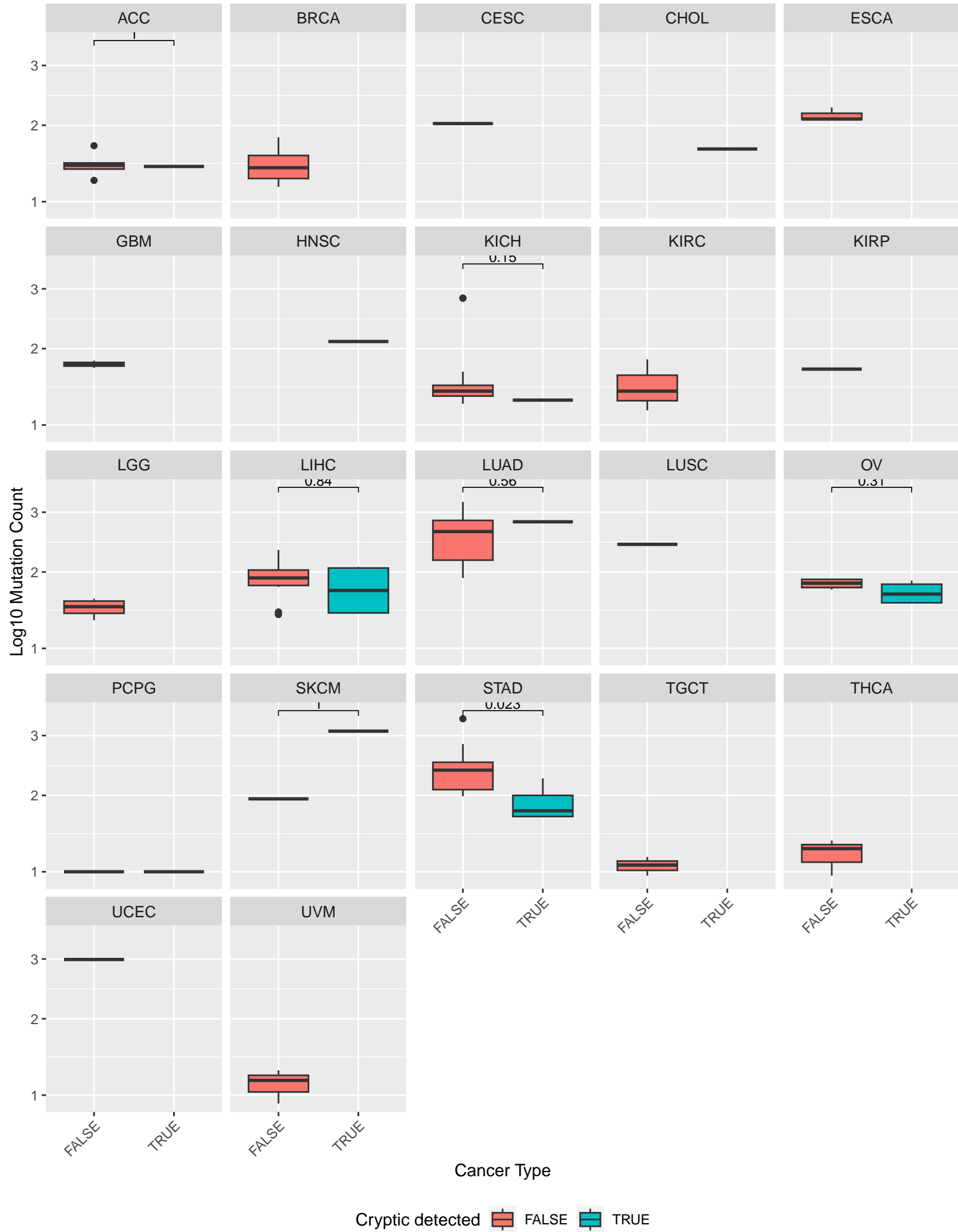




chr9:76638289–76641789 – PRUNE2

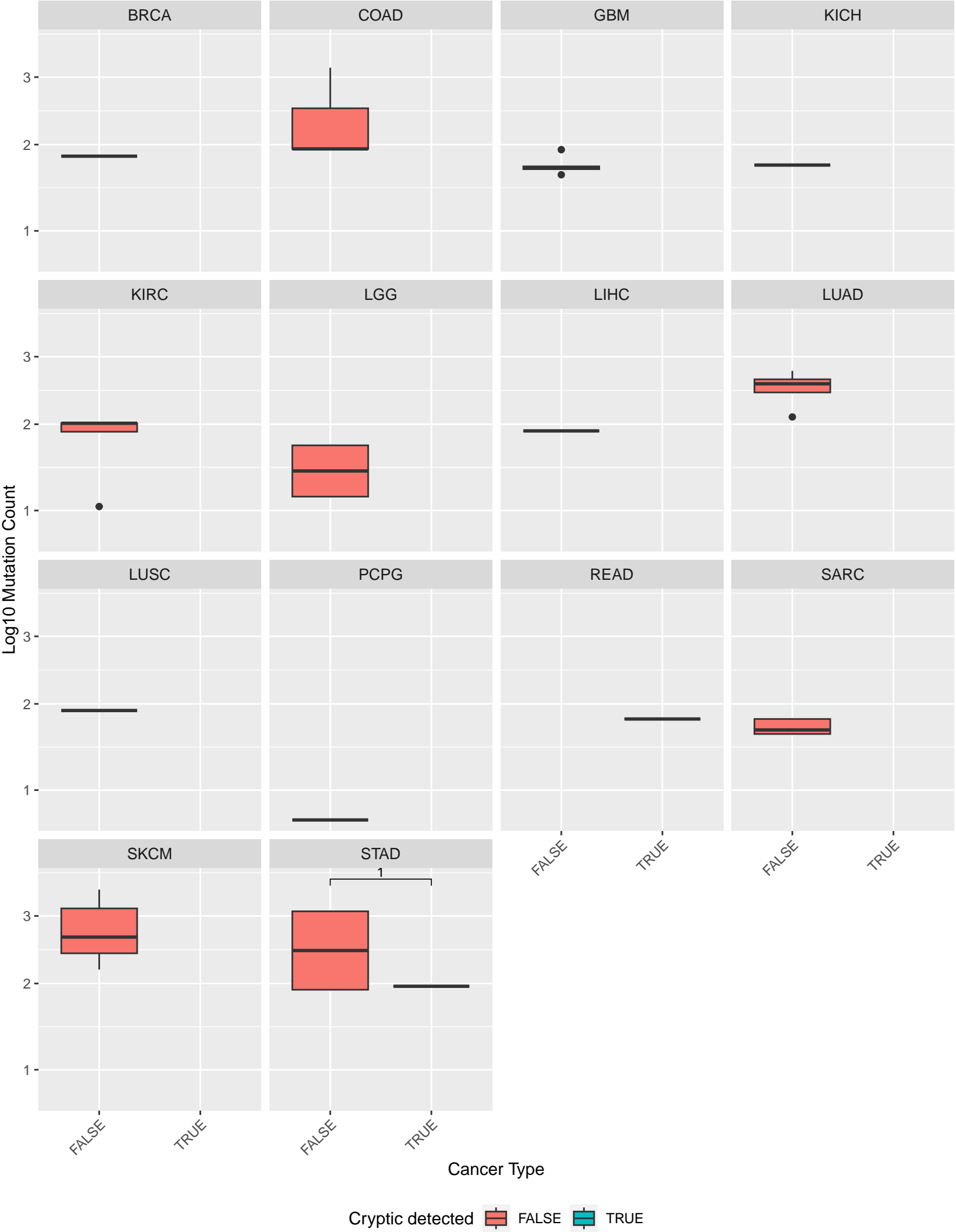


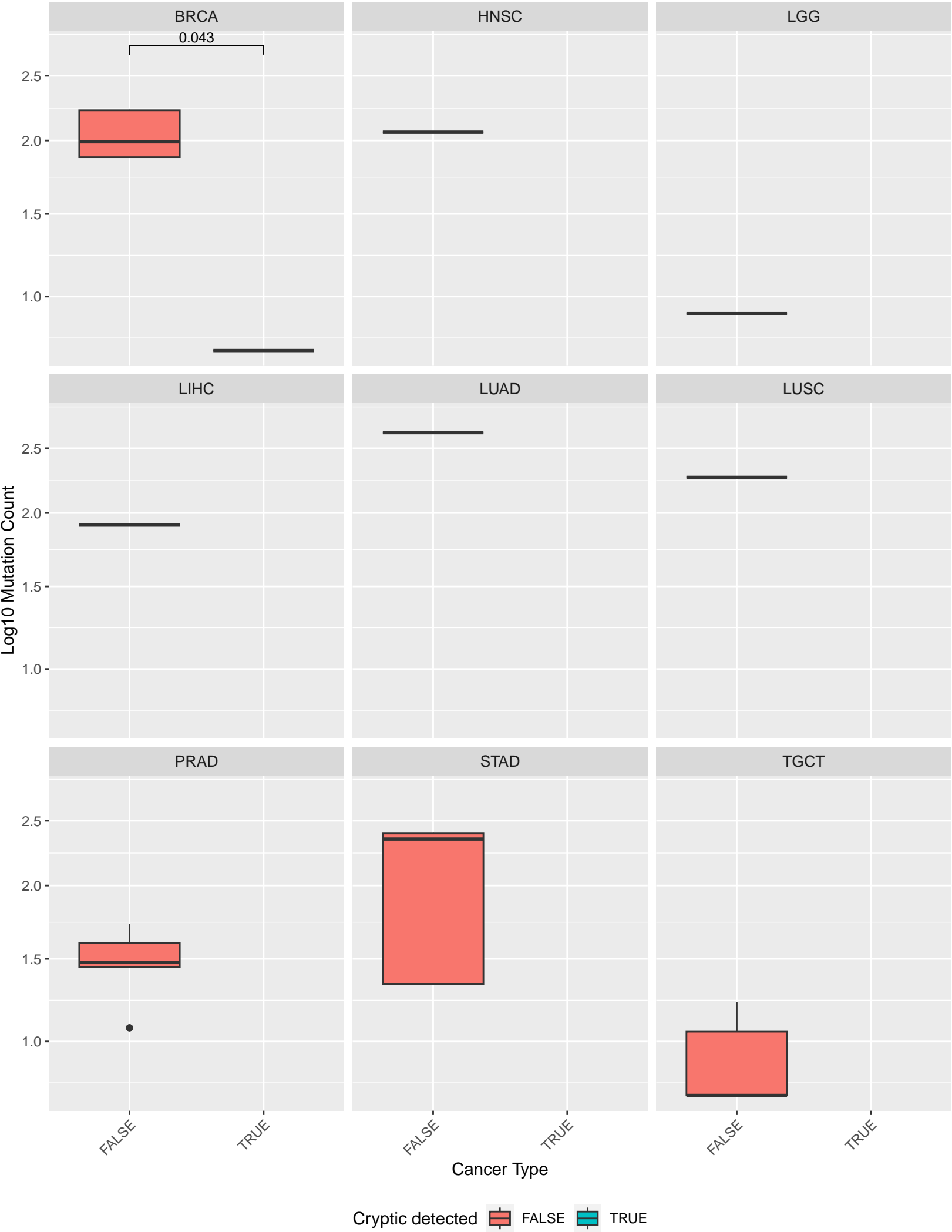
chr10:133166940–133167380 – KNDC1



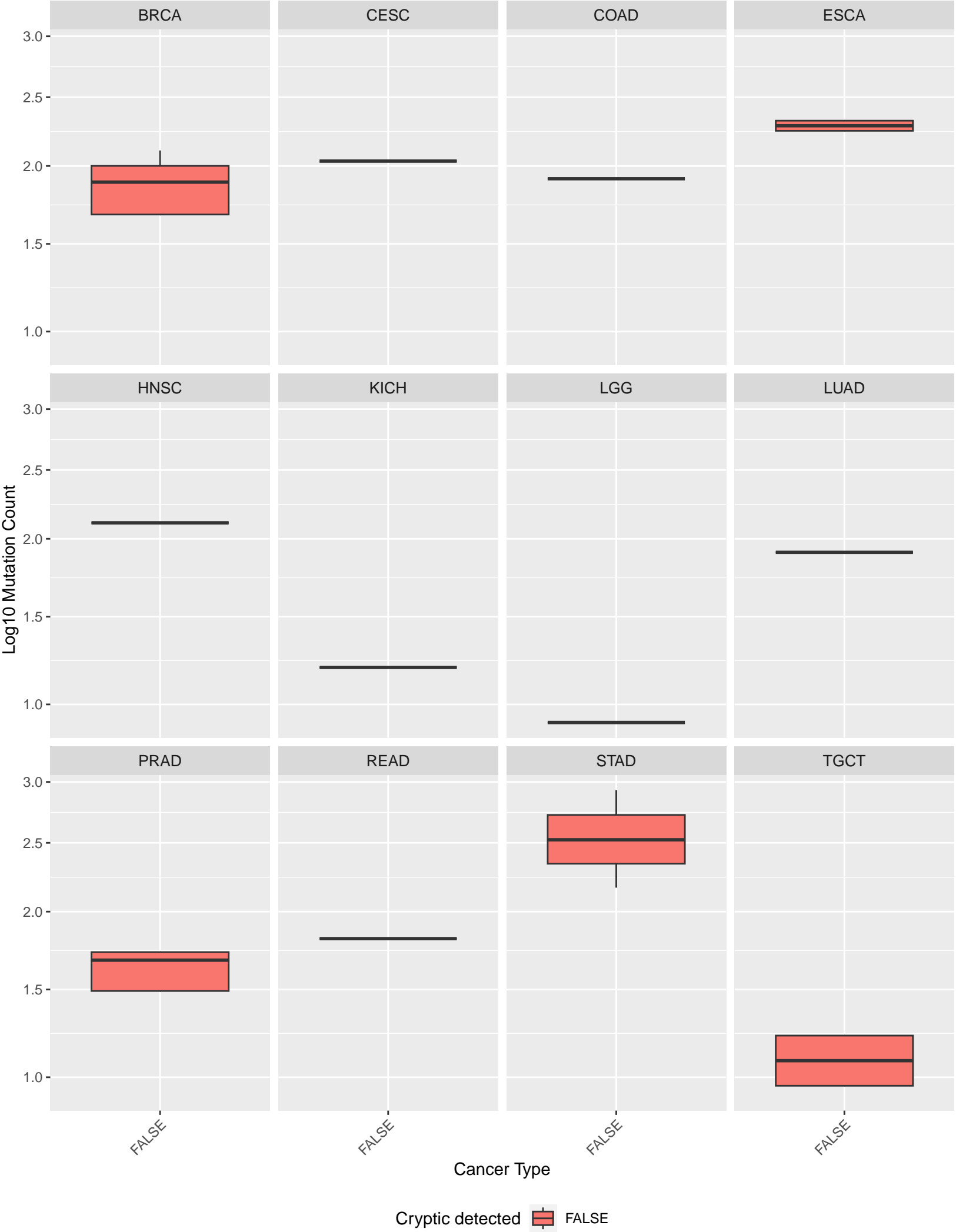
chr10:48583153–48583377 – ARHGAP22



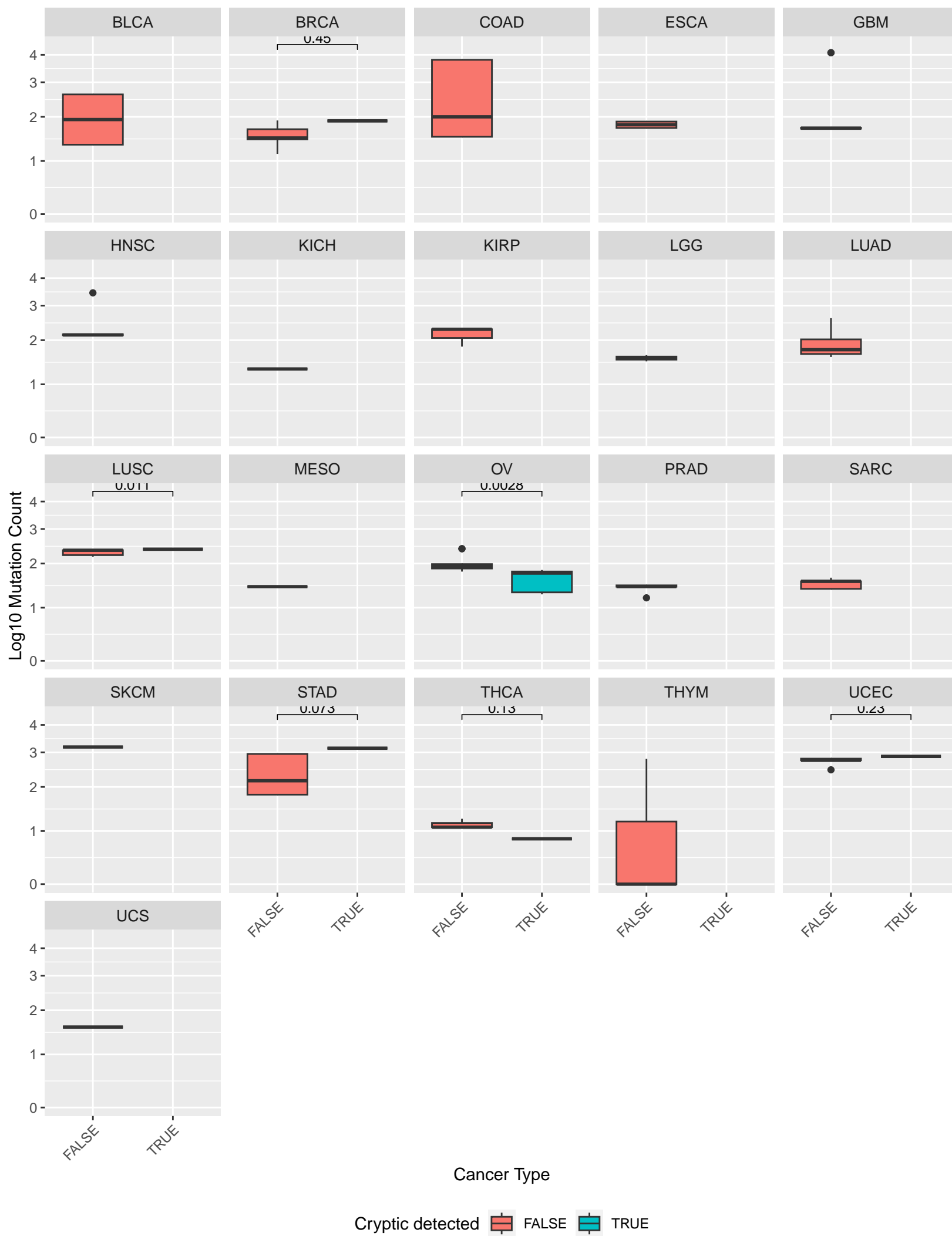




chr11:61547621-61551383 – SYT7

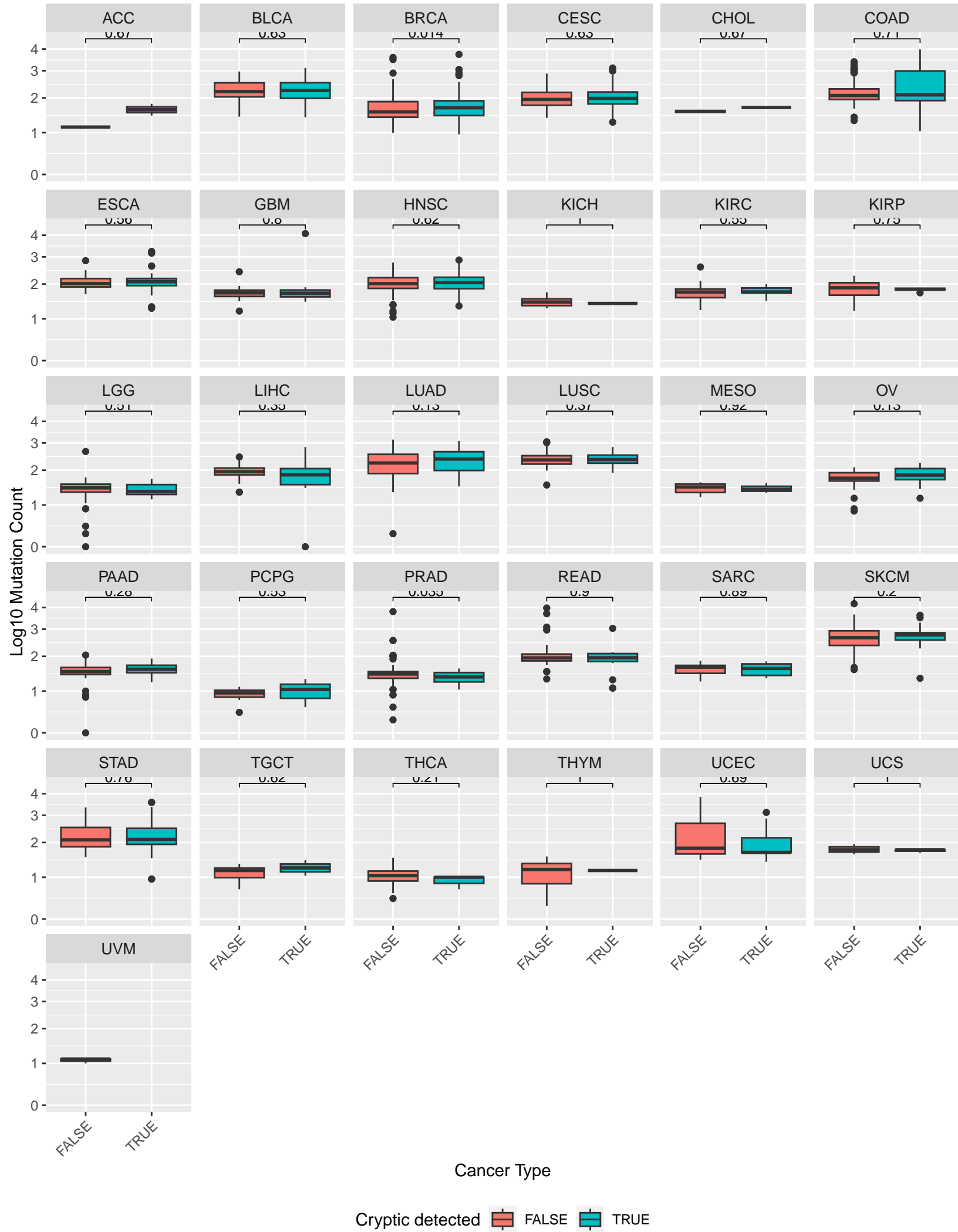


chr11:77813788–77820527 – RSF1

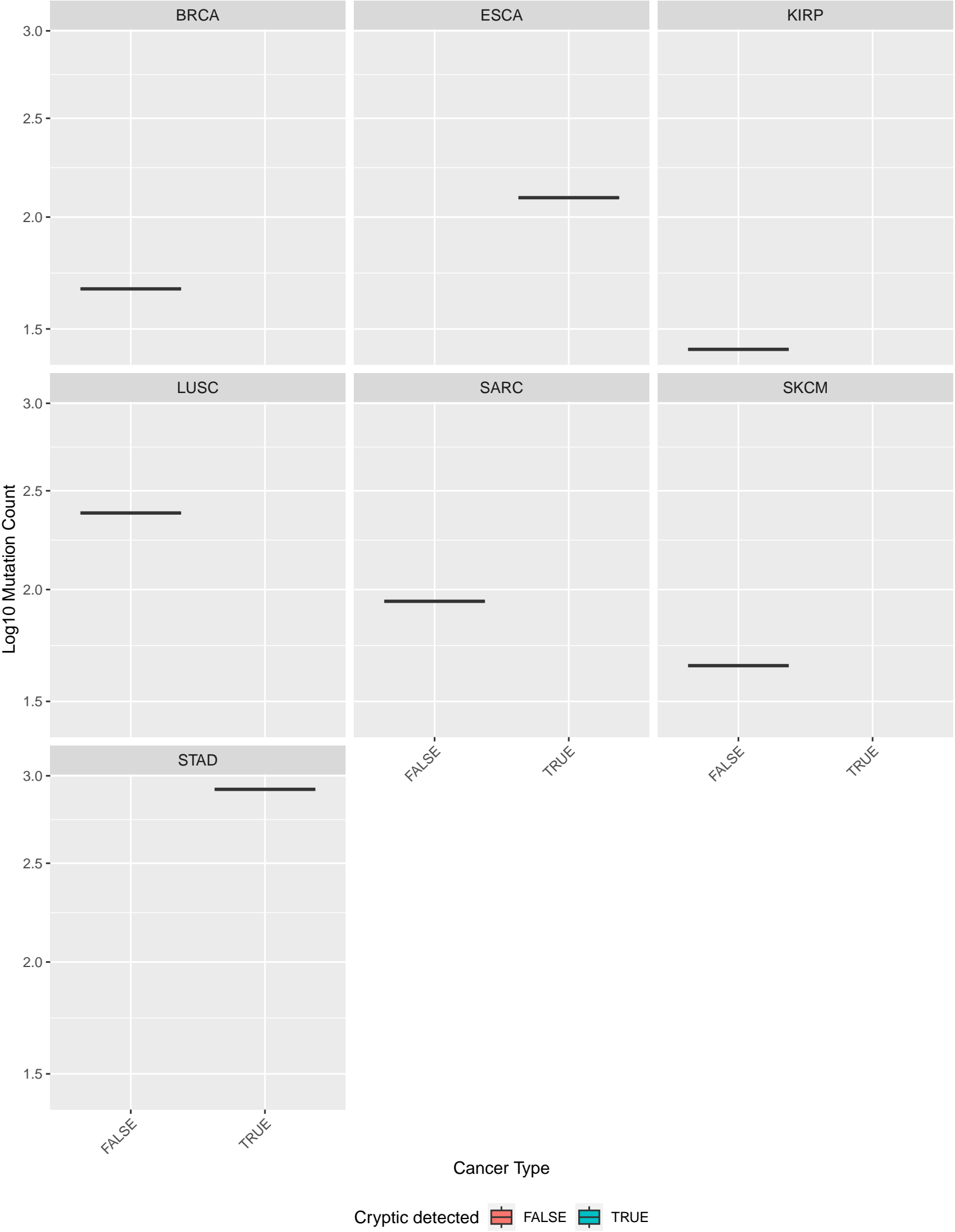




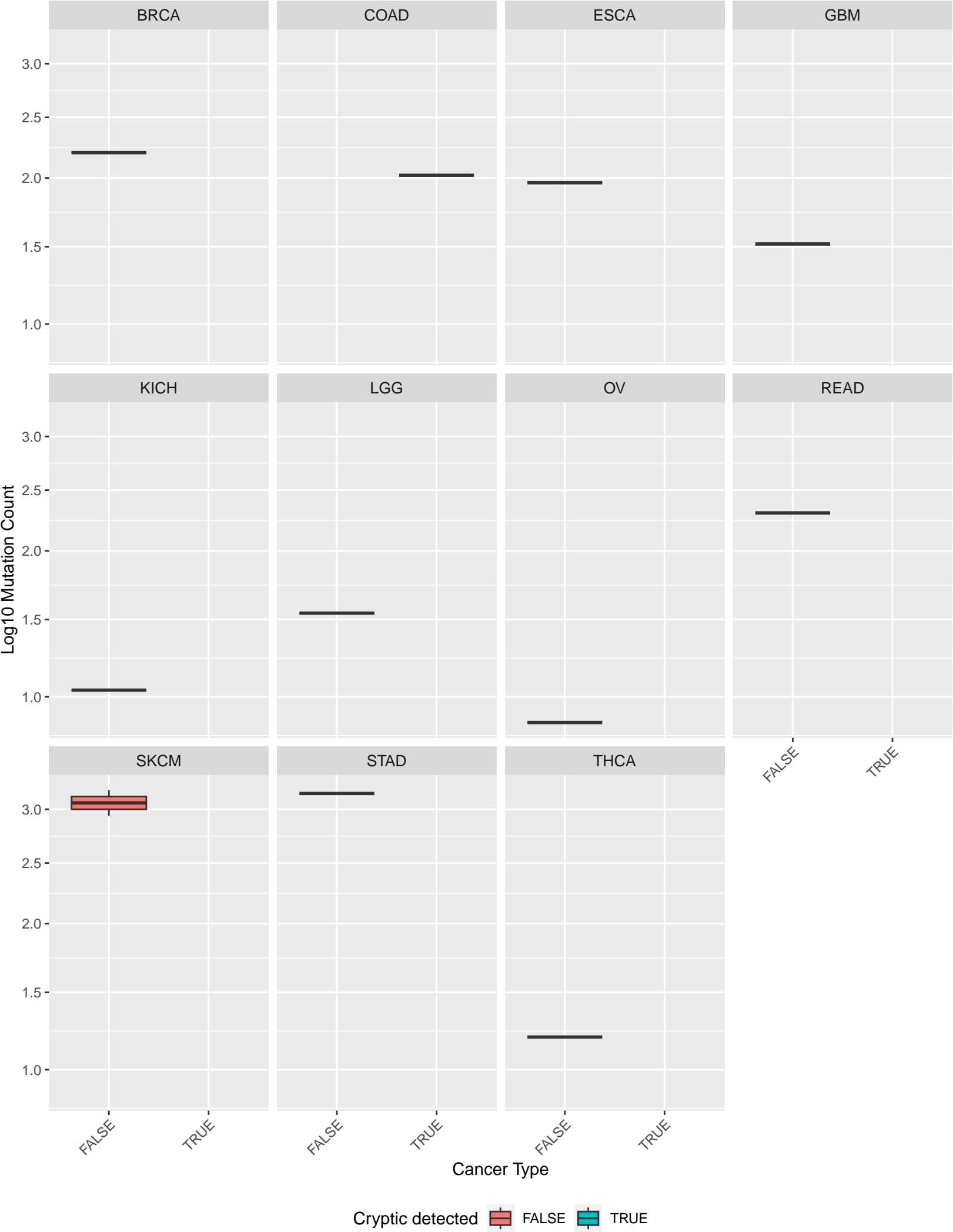
chr11:128992047-128998318 – ARHGAP32



chr12:43836197-43836743 - TMEM117

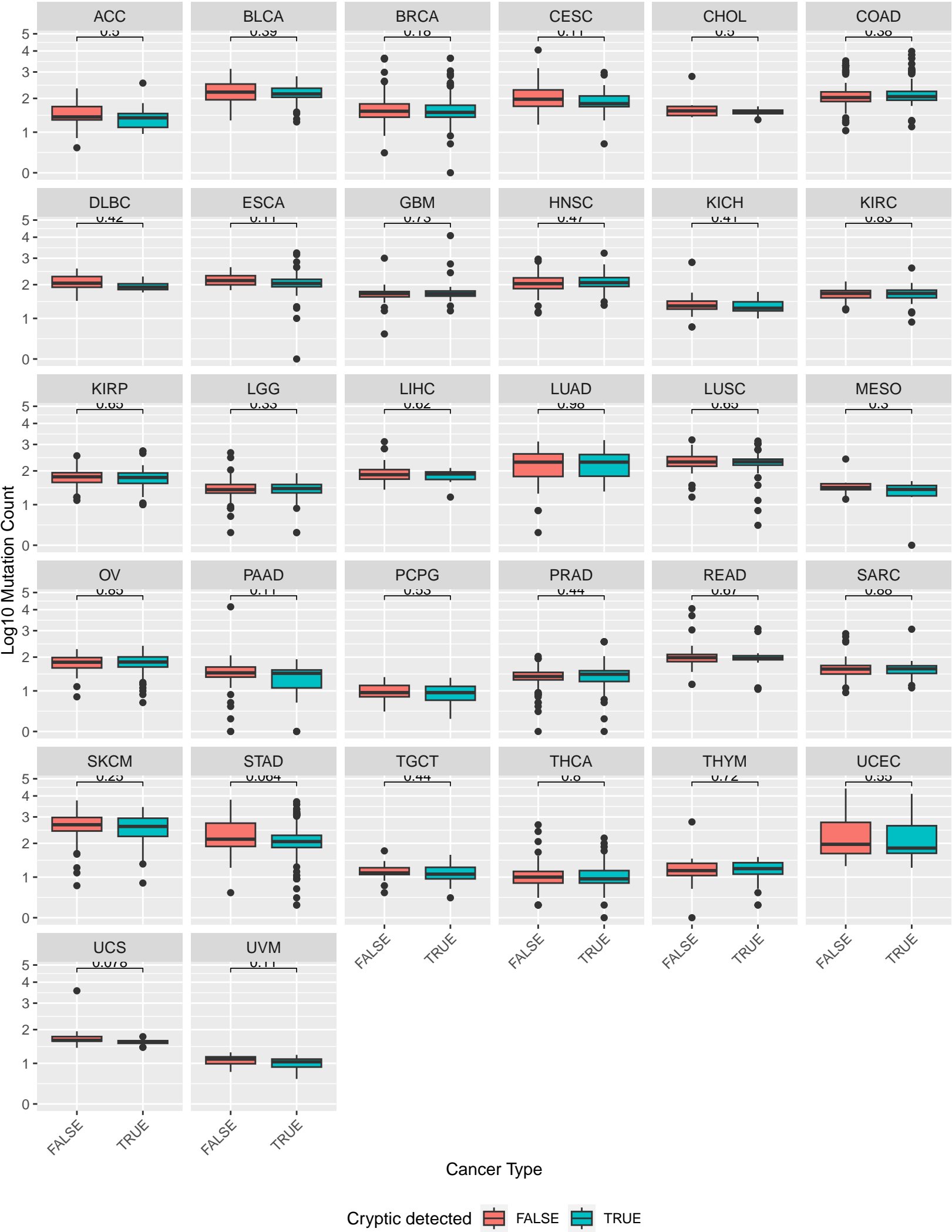


chr12:43836902–43844623 – TMEM117



chr12:88086790–88087779 – CEP290

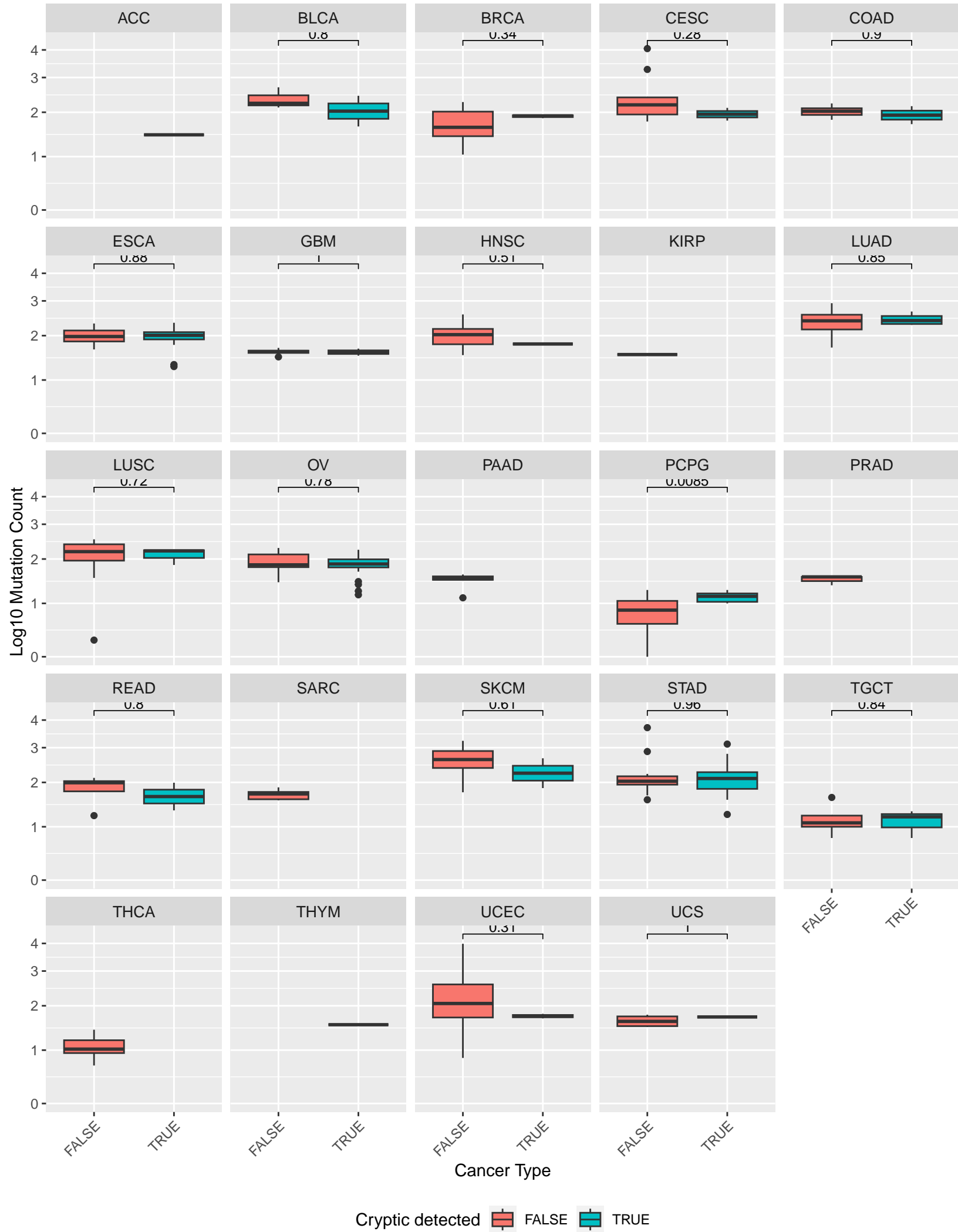






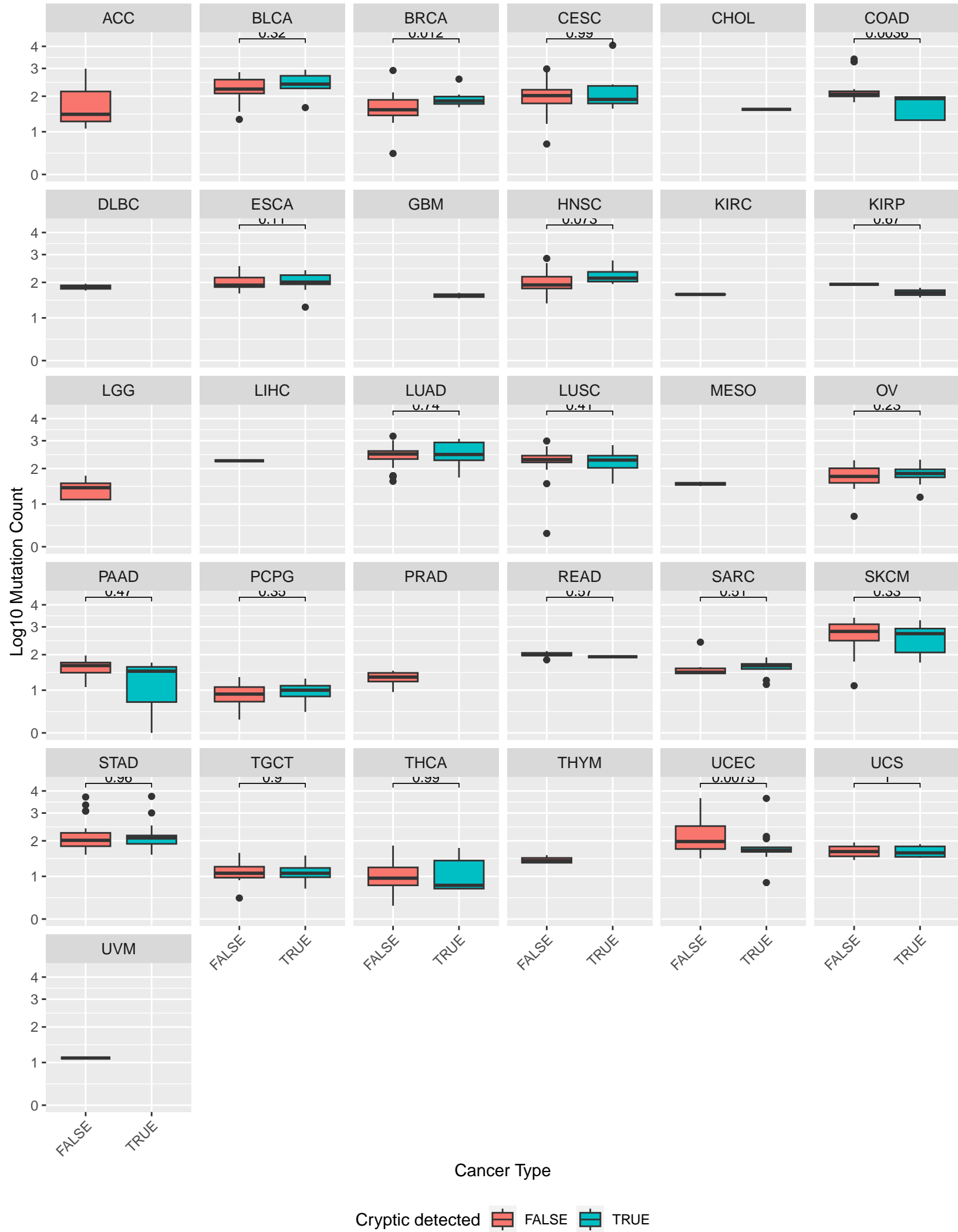
chr15:37096431-37097933 - MEIS2



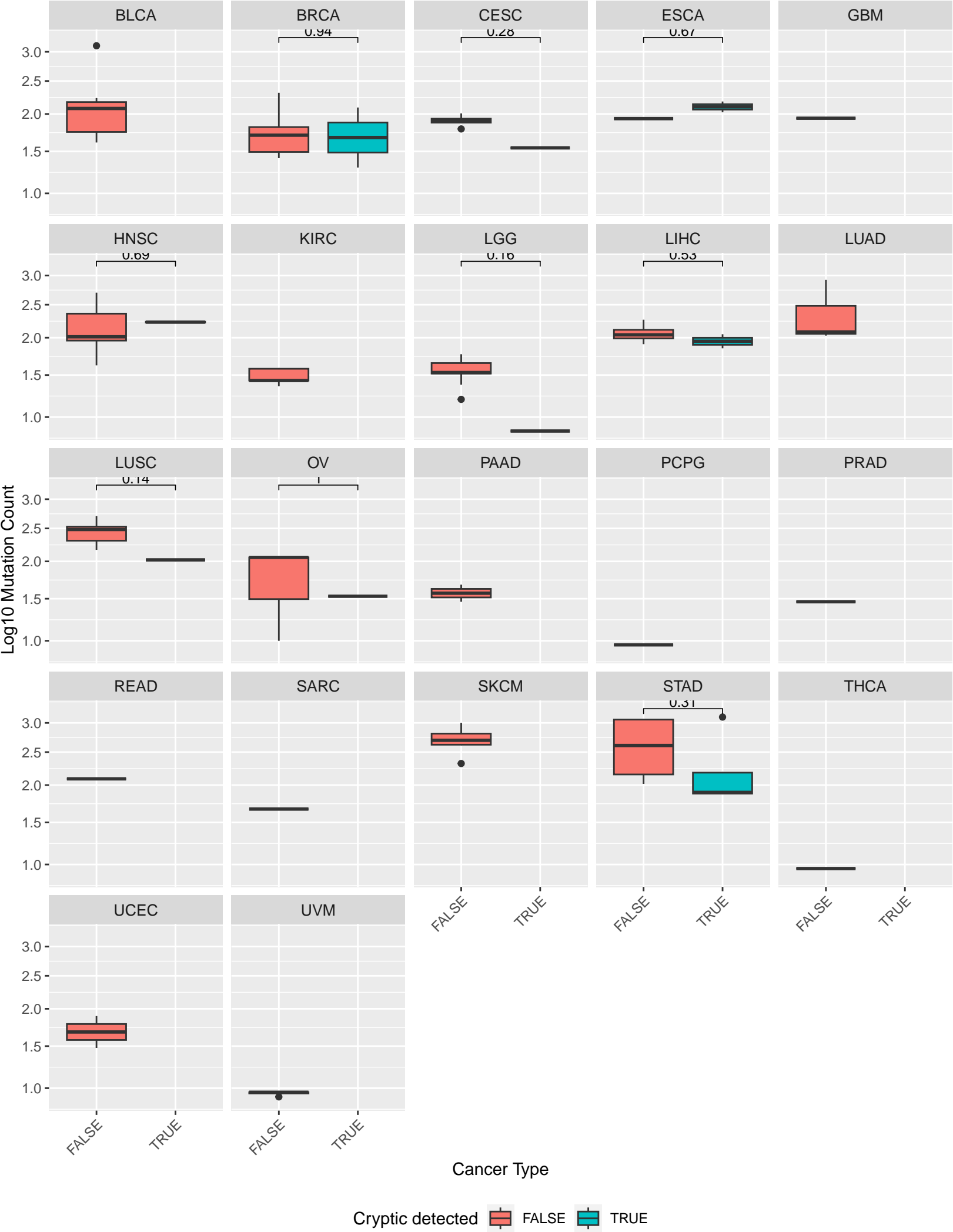




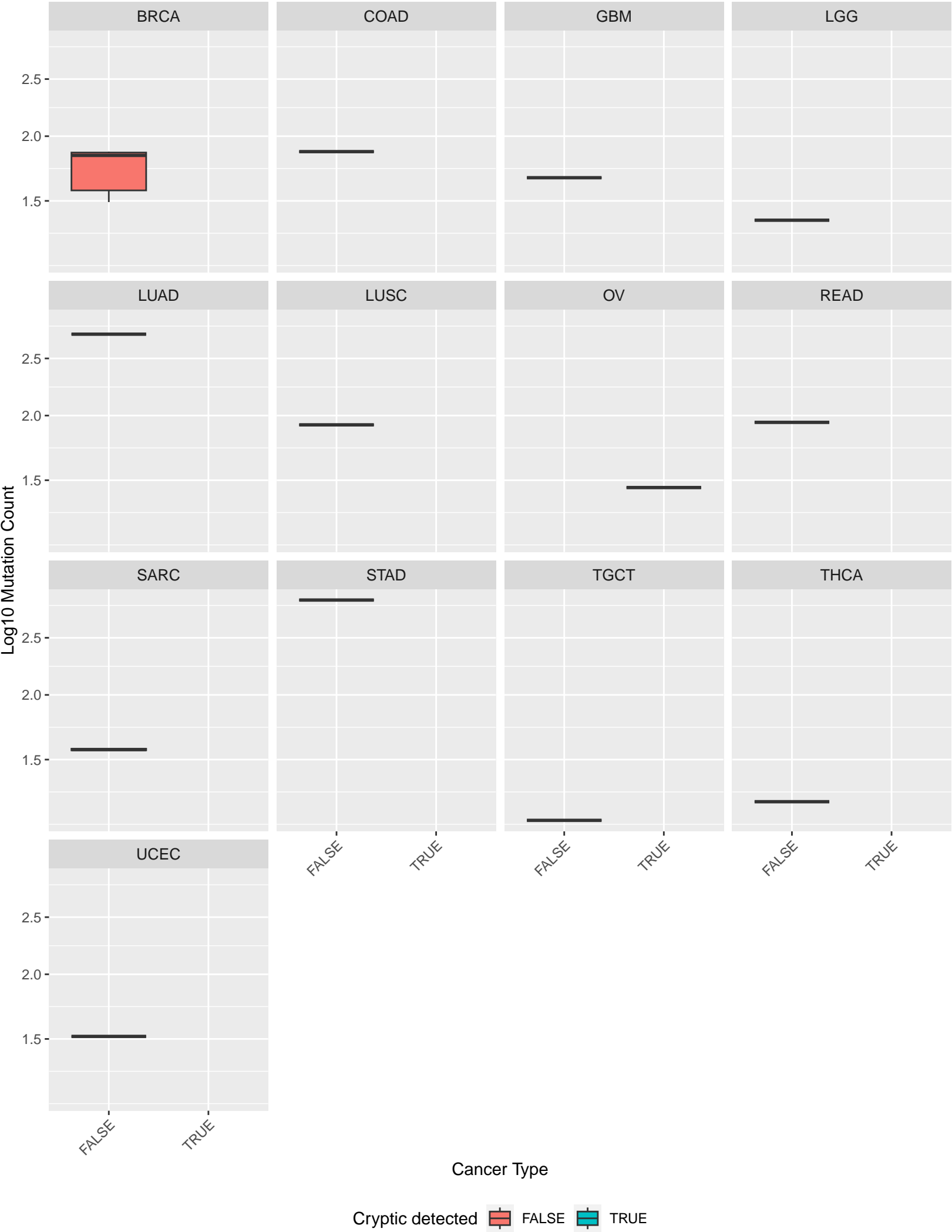
chr15:74193137-74193799 - STRA6



chr16:70271973–70272795 – AARS1

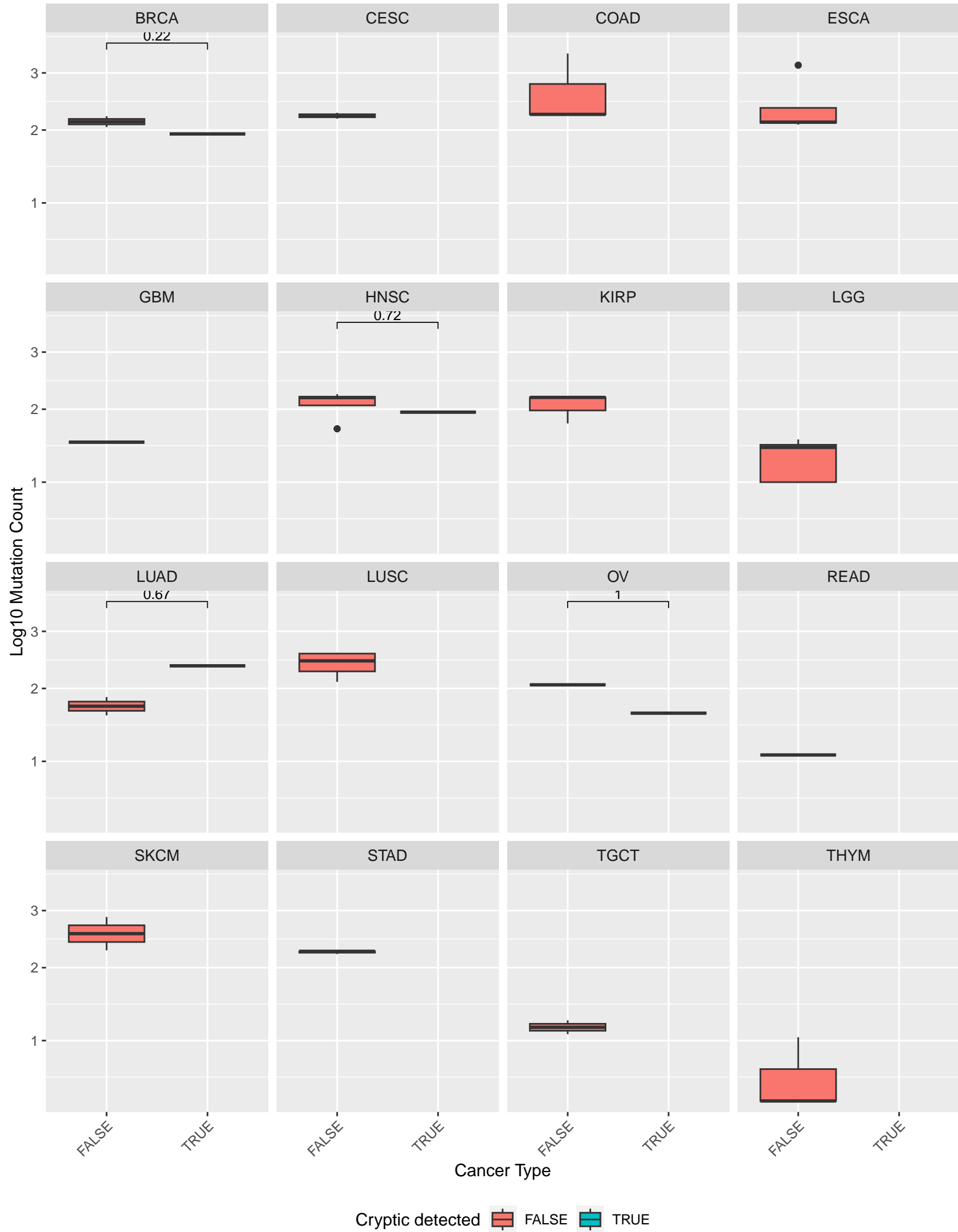


chr18:12430293–12431149 – PRELID3A

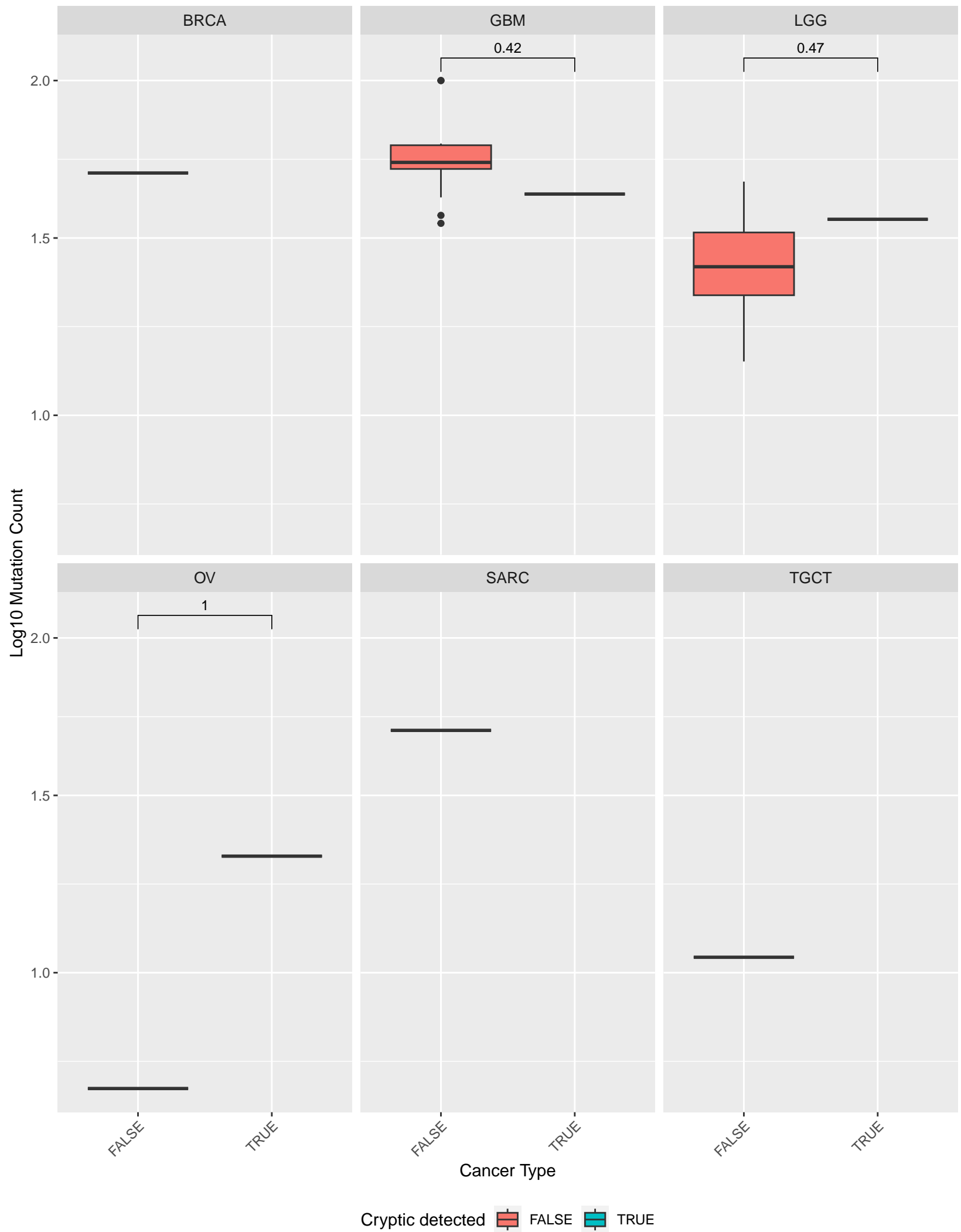


chr19:4491836–4492014 – HDGFL2

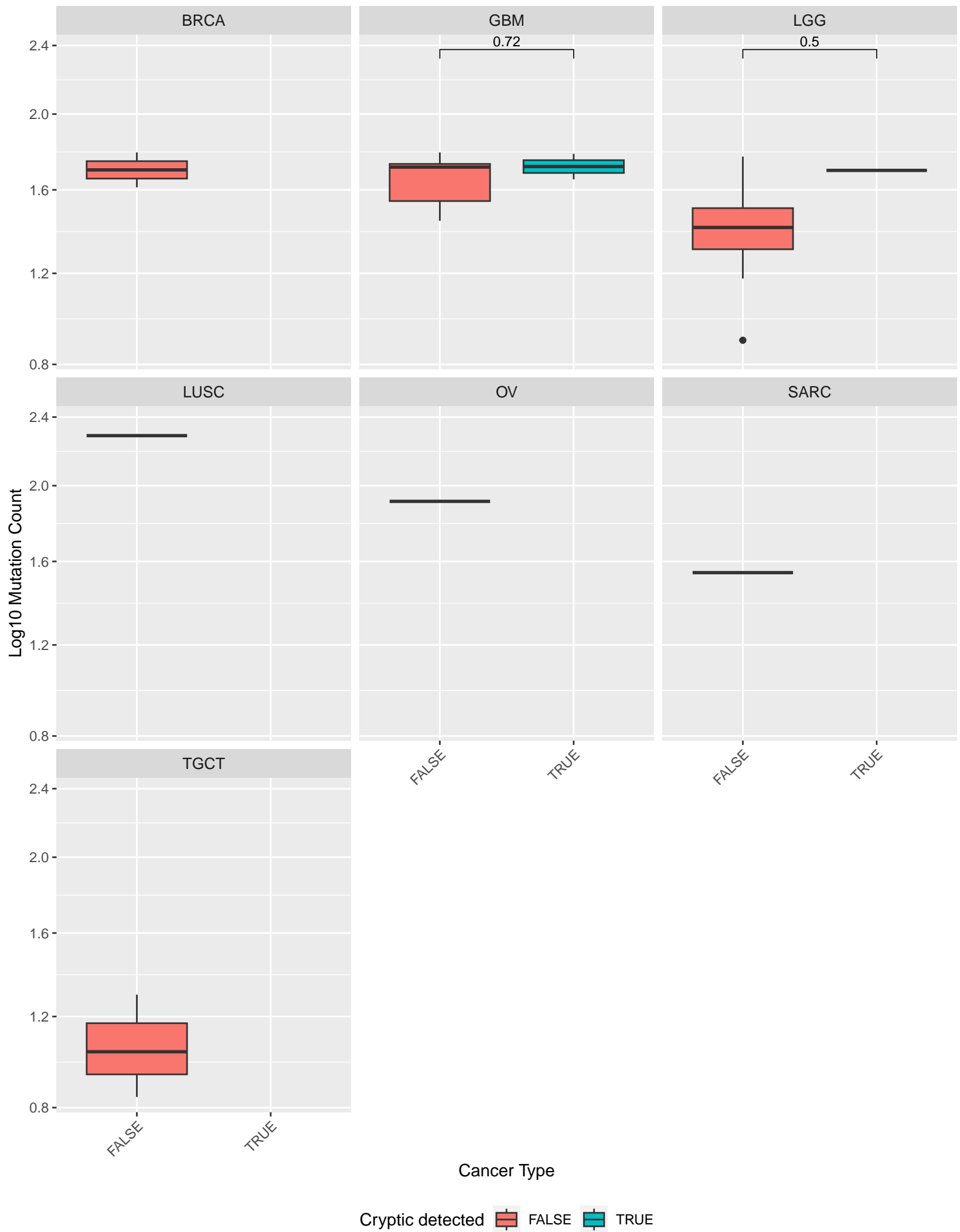




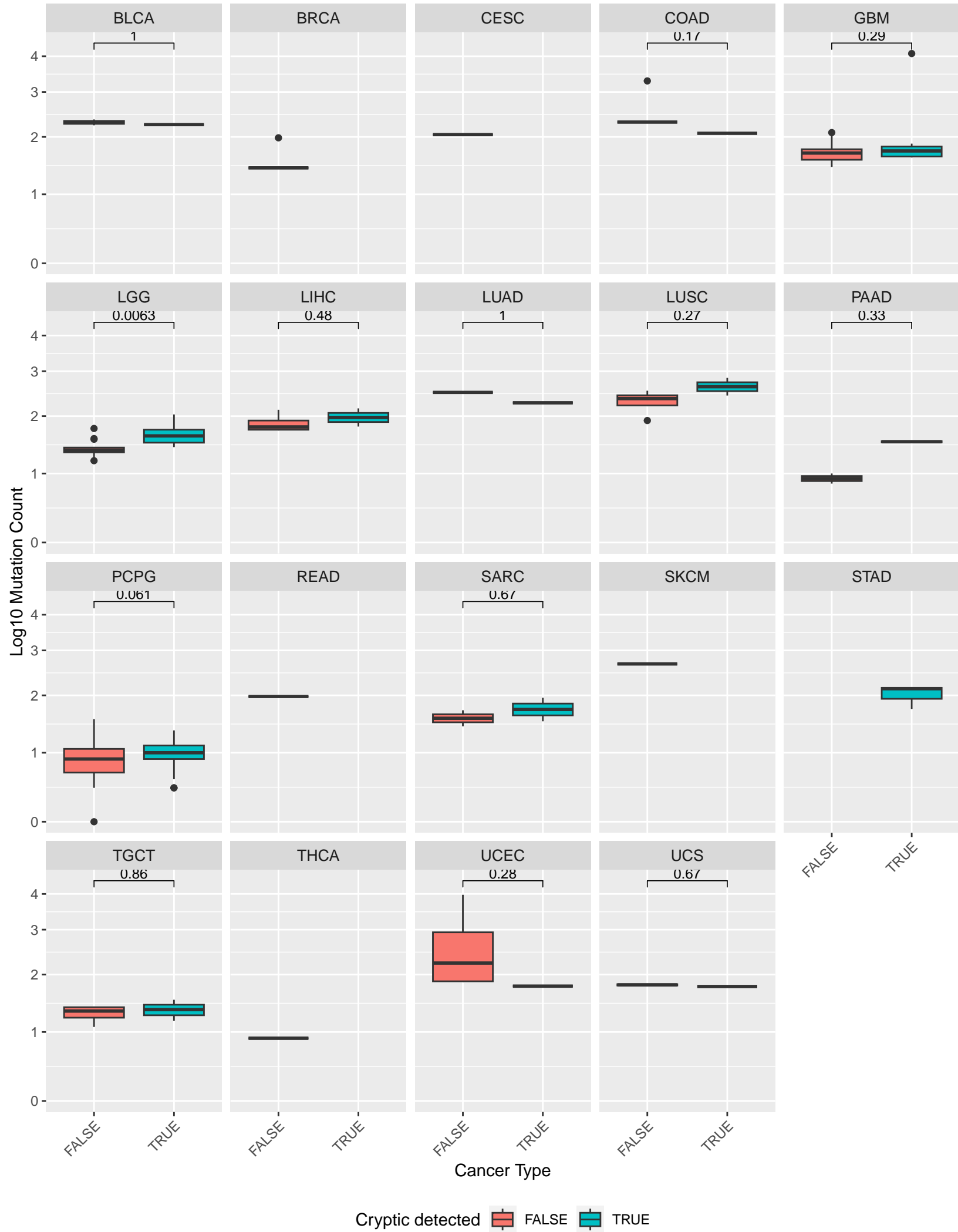
chr19:11458612–11463495 – ELAVL3



chr19:11463663–11466171 – ELAVL3



chr8:79611215–79616821 – STMN2





chr6:158017291–158019983 – SYNJ2

