DeJong et al. Supplemental Table 7. Differential Exon Usage Characteristics

Stage 10: top 10 named genes as sorted by adjusted P-value

Gene Number if retained introns Which intron(s) retained?

Gene	Number if retained introns	Which intron(s) retained?
fgfr4	1	1st
ets2	2	3rd and 4th
tfap4	1	3rd
pygm	3	Throughout 19th
eif2s2	1	7th
gcfc1	1	6th
irx2	4	Throughout 1st and 3rd
dgka	2	16th and 17th
asxl1	1	7th
prpf6	1	16th

Stage 15: top 10 named genes as sorted by adjusted P-value

Gene	Number if retained introns	Which intron(s) retained?
ctdsp2	22	throughout whole transcript
pax8	1	5th
fgfr4	1	1st
masp1	1	4th
c5	1	12th
mapre2	1	3rd
vsig8	1	1st
mcm3ap	2	Throughout the 23rd
per2	2	1st and 2nd
gpr110	1	9th