



Figure S4 FGF10 expression analysis. **(a)** Temporal analysis of the differential FGF family and its receptor transcript collection showed significant enrichment in three expression patterns ($P < 0.05$); **(b)** dynamic trends in FGF10 expression; **(c)** differential expression analysis showed that FGF10 was more highly expressed at E75 than at E125, and the differential expression was significant.